





Description*Introduction and state of the art*

5 The invention relates to antibody drug conjugates (ADCs) of  
kinesin spindle protein inhibitors, to active metabolites of  
these ADCs, to processes for preparing these ADCs, to the use  
of these ADCs for the treatment and/or prophylaxis of diseases  
and to the use of these ADCs for preparing medicaments for  
10 treatment and/or prevention of diseases, in particular  
hyperproliferative and/or angiogenic disorders such as, for  
example, cancer diseases. Such treatments can be carried out as  
monotherapy or else in combination with other medicaments or  
further therapeutic measures.

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Cancer diseases are the consequence of uncontrolled cell growth  
of the most diverse tissues. In many cases, the new cells  
penetrate into existing tissue (invasive growth), or they  
metastase into remote organs. Cancer diseases occur in the most  
20 diverse organs and often have tissue-specific courses of the  
disease. The term cancer as a generic term therefore describes  
a large group of defined diseases of various organs, tissue and  
cell types.

25 Tumours in early stages can possibly be removed by surgical and  
radiotherapy measures. Metastased tumours as a rule can only be  
treated palliatively by chemotherapeutics. The aim here is to  
achieve the optimum combination of an improvement in the quality  
of life and prolonging of life.

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Conjugates of binder proteins with one or more active compound  
molecules are known, in particular in the form of antibody drug  
conjugates (ADCs) in which an internalising antibody directed  
against a tumour-associated antigen is covalently attached via  
35 a linker to a cytotoxic agent. Following introduction of the  
ADCs into the tumour cell and subsequent dissociation of the  
conjugate, either the cytotoxic agent itself or a cytotoxic  
metabolite formed therefrom is released within the tumour cell

and can unfold its action therein directly and selectively. In this manner, in contrast to conventional chemotherapy, damage to normal tissue is contained in significantly narrower limits [see, for example, J. M. Lambert, *Curr. Opin. Pharmacol.* 5, 543-549 (2005); A. M. Wu and P. D. Senter, *Nat. Biotechnol.* 23, 1137-1146 (2005); P. D. Senter, *Curr. Opin. Chem. Biol.* 13, 235-244 (2009); L. Ducry and B. Stump, *Bioconjugate Chem.* 21, 5-13 (2010)]. Thus, WO2012/171020 describes ADCs in which a plurality of toxophor molecules are attached via a polymeric linker to an antibody. As possible toxophors, WO2012/171020 mentions, among others, the substances SB 743921, SB 715992 (Ispinesib), MK-0371, AZD8477, AZ3146 and ARRY-520.

The substances mentioned last are kinesin spindle protein inhibitors. Kinesin spindle protein (KSP, also known as Eg5, HsEg5, KNSL1 or KIF11) is a kinesin-like motorprotein which is essential for the bipolar mitotic spindle to function. Inhibition of KSP leads to mitotic arrest and, over a relatively long term, to apoptosis (Tao et al., *Cancer Cell* 2005 Jul 8(1), 39-59). After the discovery of the first cell-penetrating KSP inhibitor, Monastrol, KSP inhibitors have established themselves as a class of novel chemotherapeutics (Mayer et al., *Science* 286: 971-974, 1999), and they are subject of a number of patent applications (e.g. WO2006/044825; WO2006/002236; WO2005/051922; WO2006/060737; WO03/060064; WO03/040979; and WO03/049527). However, since KSP unfolds its action only during a relatively short period of time during the mitosis phase, KSP inhibitors have to be present in a sufficiently high concentration during these initial phases.

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### ***Summary of the invention***

Against this background it is an object of the present invention to provide substances which, after administration at a relatively low concentration, unfold apoptotic action and may therefore be of benefit for cancer therapy.

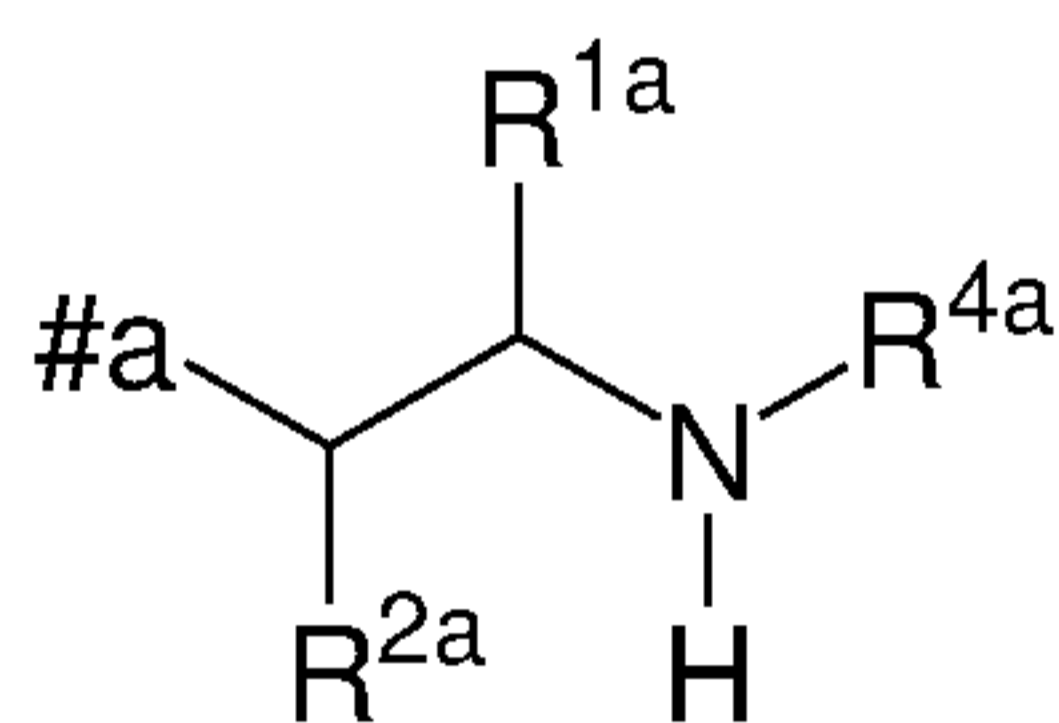
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To achieve this object, the invention provides conjugates of a

binder or derivatives thereof with one or more active compound molecules, the active compound molecule being a kinesin spindle protein inhibitor (KSP inhibitor) attached to the binder via a linker L. The binder is preferably a binder protein or peptide,  
 5 particularly preferably a human, humanized or chimeric monoclonal antibody or an antigen-binding fragment thereof, in particular an anti-TWEAKR antibody or an antigen-binding fragment thereof or an anti-EGFR antibody or an antigen-binding fragment thereof. Particular preference is given to an anti-  
 10 TWEAKR antibody which binds specifically to amino acid D in position 47 (D47) of TWEAKR (SEQ ID NO:169), in particular the anti-TWEAKR antibody TPP-2090, or the anti-EGFR antibodies cetuximab or nimotuzumab.

15 The inventors have found a number of ways to attach the binder to the KSP inhibitor in order to achieve the object mentioned above.

According to the invention, the kinesin spindle protein  
 20 inhibitors may have the substructure I(sub) below:



I (sub)

where

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#a represents a bond to the remainder of the molecule;

R<sup>1a</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

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where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' (e.g. -(CH<sub>2</sub>)<sub>0-3</sub>Z') or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub>, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where W

represents H or OH, where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ ;

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$R^{2a}$  and  $R^{4a}$  independently of one another represent H,  $-CO-CHY^4-NHY^5$  or  $-(CH_2)_{0-3}Z$ , or  $R^{2a}$  and  $R^{4a}$  together represent (with formation of a pyrrolidine ring)  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$  represents H,  $NH_2$ ,  $COOH$ ,  $SO_3H$ ,  $SH$  or  $OH$ , and where  $Z$  represents

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$-H$ ,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ;

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where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ , and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where

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$Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl.

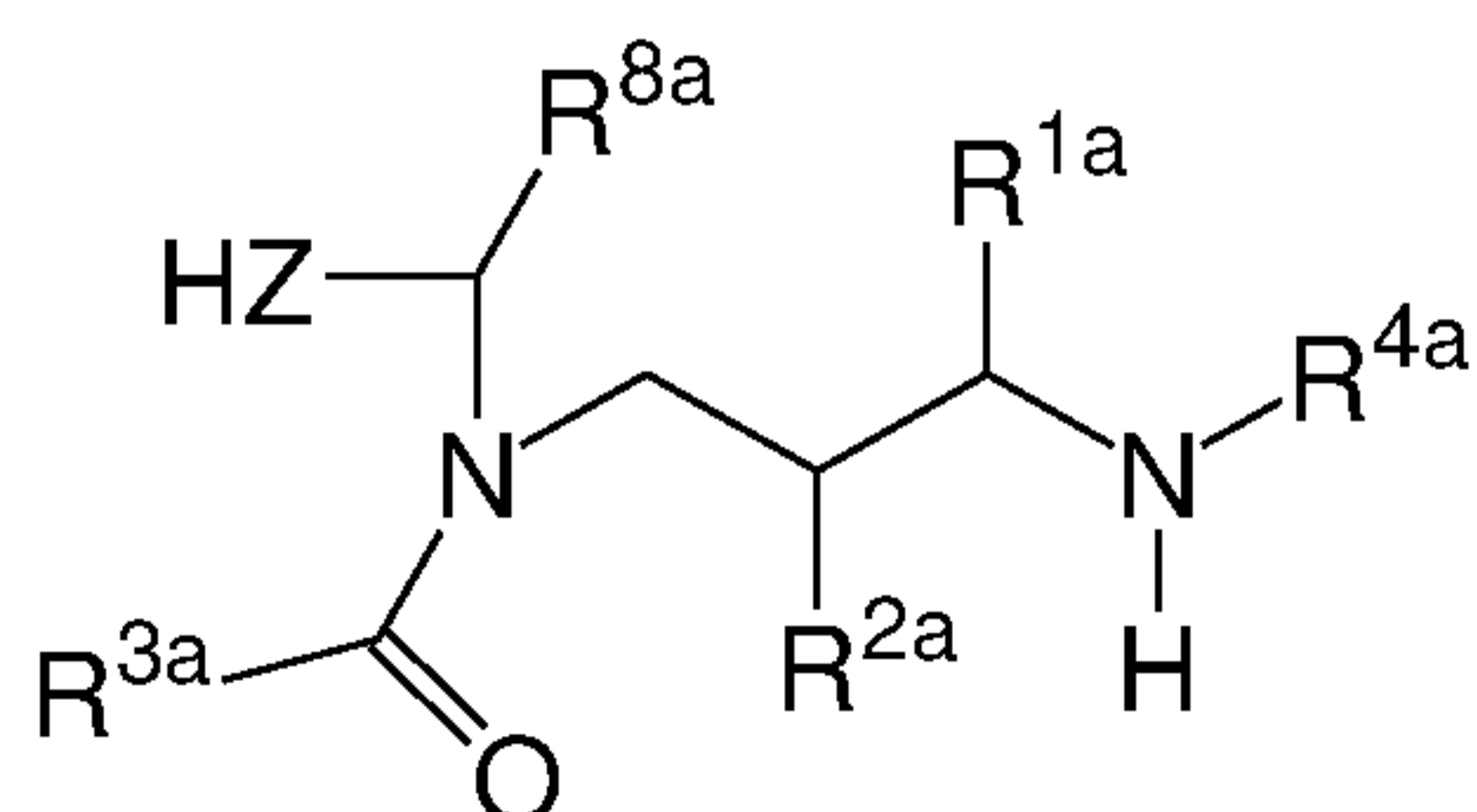
According to the invention, the kinesin spindle protein inhibitor may be attached to the binder via a linker by substitution of a hydrogen atom at  $R^{1a}$ ,  $R^{2a}$ ,  $R^{4a}$  or  $R^{10}$ .

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The KSP inhibitor which is attached to this binder (or the KSP inhibitors, since frequently more than one KSP inhibitor is attached to the binder), is preferably a compound of the formula (Ia) or (IIa) below:

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Formula (Ia):



(Ia)

where

R<sup>1a</sup> represents H, -MOD or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H,  
5 halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -  
(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' (e.g. -(CH<sub>2</sub>)<sub>0-3</sub>Z') or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup>  
represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub>,  
10 COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where W  
represents H or OH,

where Y<sup>4</sup> independently of one another represents straight-chain  
or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -  
15 NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally  
substituted by -NH<sub>2</sub>;

R<sup>2a</sup> and R<sup>4a</sup> independently of one another represent H, -CO-CHY<sup>4</sup>-  
NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or R<sup>2a</sup> and R<sup>4a</sup> together (with formation of a  
20 pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup>  
represents H, SO<sub>3</sub>H, NH<sub>2</sub>, COOH, SH or OH,

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -  
CO-OY<sup>3</sup>,

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where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  
-(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z'  
represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

30 where Y<sup>4</sup> independently of one another represents straight-chain  
or branched C<sub>1-6</sub> alkyl which is optionally substituted by -  
NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally  
substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where  
Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

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R<sup>3a</sup> represents -MOD or an optionally substituted alkyl,  
cycloalkyl, aryl, heteroaryl, heteroalkyl or heterocycloalkyl  
group,

preferably a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl or C<sub>5-10</sub>-heterocycloalkyl group which may be substituted by 1-3 -OH groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH

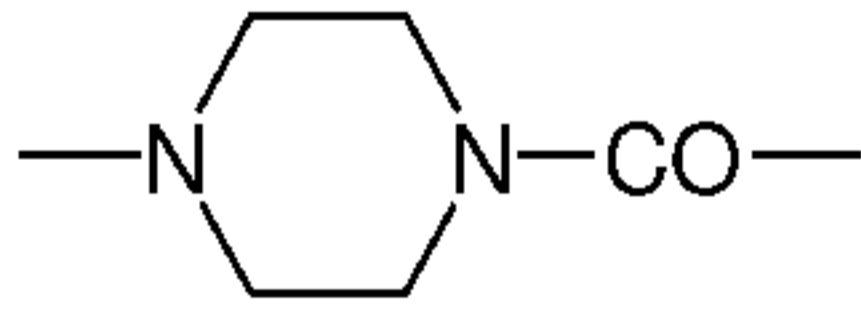
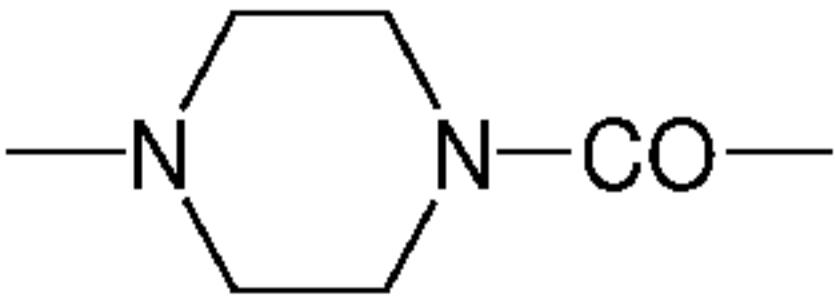
(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl);

R<sup>8a</sup> represents C<sub>1-10</sub>-alkyl or -(CH<sub>2</sub>)<sub>0-2</sub>-(HZ<sup>2</sup>), where HZ<sup>2</sup> represents a 4- to 7-membered heterocycle having up to two heteroatoms selected from the group consisting of N, O and S;

HZ represents a mono- or bicyclic heterocycle which may be substituted by one or more substituents selected from the group consisting of halogen, C<sub>1-10</sub>-alkyl groups, C<sub>6-10</sub>-aryl groups and C<sub>6-10</sub>-aralkyl groups which may optionally be substituted by halogen;

where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

R<sup>10</sup> represents H or C<sub>1-3</sub>-alkyl;

G1 represents -NHCO- , -CONH- or  (where, if G1 represents -NHCO- or  , R<sup>10</sup> does not represent NH<sub>2</sub>);

n is 0 or 1;



o is 0 or 1; and

G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>- (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl), where the hydrocarbon chain including any side chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group -MOD preferably has at least one group -COOH;

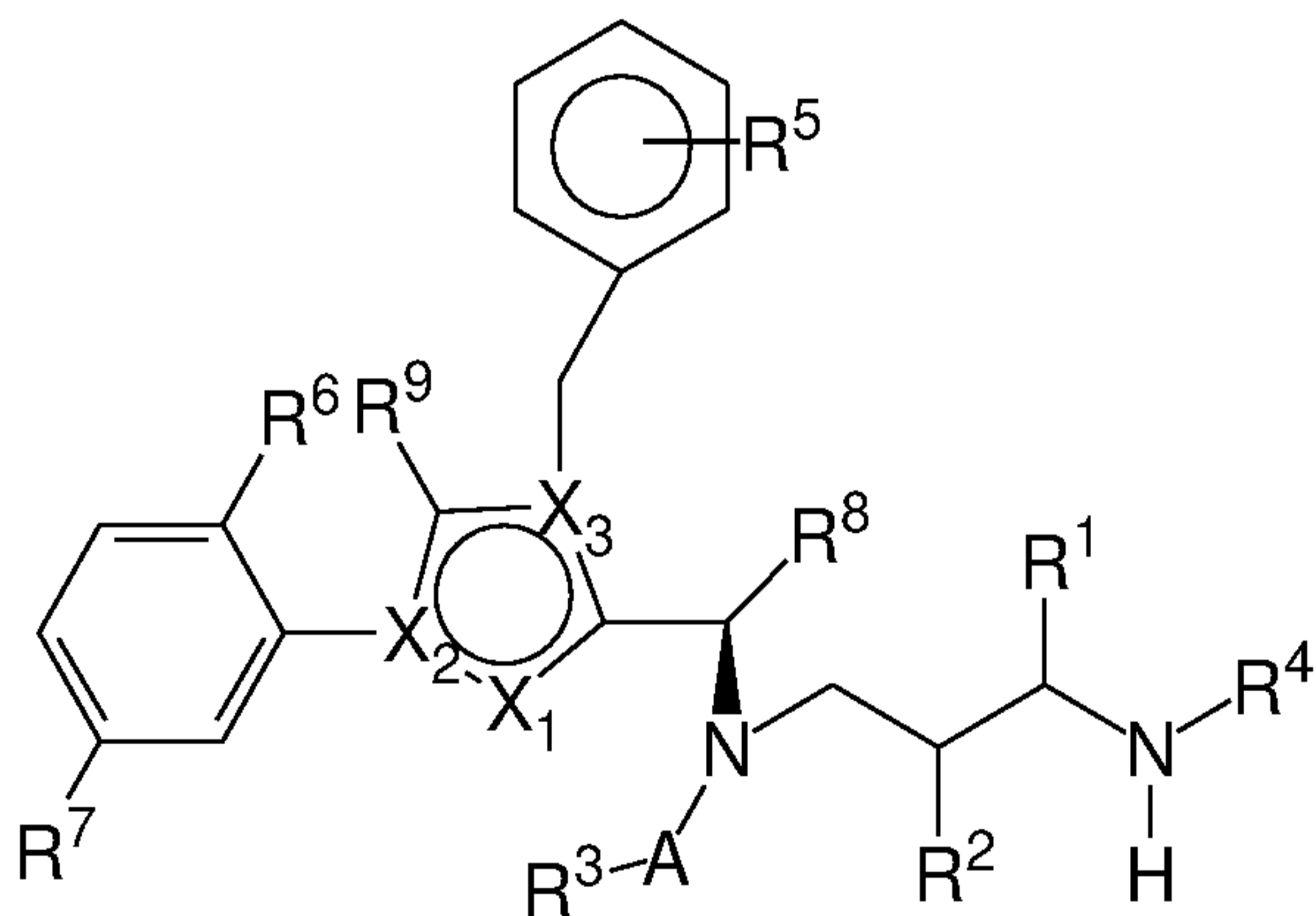
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where the kinesin spindle protein inhibitor is attached to the linker by substitution of a hydrogen atom at R<sup>1a</sup>, R<sup>2a</sup>, R<sup>3a</sup>, R<sup>4a</sup>, R<sup>8a</sup> or R<sup>10</sup> or optionally via one of the substituents of HZ, in particular via R<sup>1a</sup>, R<sup>2a</sup>, R<sup>3a</sup>, R<sup>4a</sup> or R<sup>10</sup>,

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and the salts, solvates and salts of the solvates thereof.

Formula (IIa):



25 (IIa)

where

X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C; or

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X<sub>1</sub> represents CH or CF, X<sub>2</sub> represents C and X<sub>3</sub> represents N; or

X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C; or

5 X<sub>1</sub> represents CH, X<sub>2</sub> represents N and X<sub>3</sub> represents C

(with X<sub>1</sub> representing CH, X<sub>2</sub> representing C and X<sub>3</sub> representing N being preferred);

10 R<sup>1</sup> represents H, -L-#1, -MOD or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' (e.g. -(CH<sub>2</sub>)<sub>0-3</sub>Z') or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup>

15 represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where W represents H or OH,

20 where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>;

25 R<sup>2</sup> represents -L-#1, H, -MOD, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents L-#1, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH, or OH;

30 where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

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where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally

substituted by  $-\text{NH}_2$ , and  $\text{Y}^5$  represents H or  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ , where  $\text{Y}^6$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl;

$\text{R}^4$  represents  $-\text{L}-\#1$ , H,  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  or  $-(\text{CH}_2)_{0-3}\text{Z}$ ,

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where Z represents  $-\text{H}$ , halogen,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$  or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , and  $\text{Y}^3$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$  represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$ ;

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where  $\text{Y}^4$  independently of one another represents straight-chain or branched  $\text{C}_{1-6}$  alkyl which is optionally substituted by  $-\text{NHCONH}_2$ , or represents aryl or benzyl which are optionally substituted by  $-\text{NH}_2$ , and  $\text{Y}^5$  represents H or  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ , where  $\text{Y}^6$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl;

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or  $\text{R}^2$  and  $\text{R}^4$  together (with formation of a pyrrolidine ring) represent  $-\text{CH}_2-\text{CHR}^{10}-$  or  $-\text{CHR}^{10}-\text{CH}_2-$ , where  $\text{R}^{10}$  represents  $-\text{L}-\#1$ , H,  $\text{NH}_2$ ,  $\text{SO}_3\text{H}$ ,  $\text{COOH}$ , SH or OH;

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A represents CO, SO,  $\text{SO}_2$ ,  $\text{SO}_2\text{NH}$  or CNNH;

$\text{R}^3$  represents  $-\text{L}-\#1$ ,  $-\text{MOD}$  or an optionally substituted alkyl, cycloalkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably  $-\text{L}-\#1$  or a  $\text{C}_{1-10}$ -alkyl,  $\text{C}_{6-10}$ -aryl or  $\text{C}_{6-10}$ -aralkyl,  $\text{C}_{5-10}$ -heteroalkyl,  $\text{C}_{1-10}$ -alkyl-O- $\text{C}_{6-10}$ -aryl or  $\text{C}_{5-10}$ -heterocycloalkyl group which may be substituted by 1-3  $-\text{OH}$  groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3  $-\text{SH}$  groups, 1-3  $-\text{S}$ -alkyl groups, 1-3  $-\text{O}-\text{CO}$ -alkyl groups, 1-3  $-\text{O}-\text{CO}-\text{NH}$ -alkyl groups, 1-3  $-\text{NH}-\text{CO}$ -alkyl groups, 1-3  $-\text{NH}-\text{CO}-\text{NH}$ -alkyl groups, 1-3  $-\text{S}(\text{O})_n$ -alkyl groups, 1-3  $-\text{SO}_2-\text{NH}$ -alkyl groups, 1-3  $-\text{NH}$ -alkyl groups, 1-3  $-\text{N}(\text{alkyl})_2$  groups, 1-3  $-\text{NH}_2$  groups or 1-3  $-(\text{CH}_2)_{0-3}\text{Z}$  groups, where Z represents  $-\text{H}$ , halogen,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $-\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ , where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$  or  $-(\text{CH}_2)_{0-3}\text{Z}'$  and  $\text{Y}^3$  represents H,  $-(\text{CH}_2)_{0-3}$ -

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$\text{CH}(\text{NHCOCH}_3)\text{Z}'$ ,  $-(\text{CH}_2)_{0-3}\text{CH}(\text{NH}_2)\text{Z}'$  or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$  represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$

(where "alkyl" preferably represents  $\text{C}_{1-10}$ -alkyl);

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$\text{R}^5$  represents  $-\text{L}-\#1$ , H,  $\text{NH}_2$ ,  $\text{NO}_2$ , halogen (in particular F, Cl, Br),  $-\text{CN}$ ,  $\text{CF}_3$ ,  $-\text{OCF}_3$ ,  $-\text{CH}_2\text{F}$ ,  $-\text{CH}_2\text{F}$ , SH or  $-(\text{CH}_2)_{0-3}\text{Z}$ , where Z represents  $-\text{H}$ ,  $-\text{OY}^3$ ,  $-\text{SY}^3$ , halogen,  $\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

10 where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$  or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , and  $\text{Y}^3$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$  represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$ ;

$\text{R}^6$  and  $\text{R}^7$  independently of one another represent H, cyano, 15 (optionally fluorinated)  $\text{C}_{1-10}$ -alkyl, (optionally fluorinated)  $\text{C}_{2-10}$ -alkenyl, (optionally fluorinated)  $\text{C}_{2-10}$ -alkynyl, hydroxy,  $\text{NO}_2$ ,  $\text{NH}_2$ ,  $\text{COOH}$  or halogen (in particular F, Cl, Br),

$\text{R}^8$  represents (optionally fluorinated)  $\text{C}_{1-10}$ -alkyl, (optionally 20 fluorinated)  $\text{C}_{2-10}$ -alkenyl, (optionally fluorinated)  $\text{C}_{2-10}$ -alkynyl, (optionally fluorinated)  $\text{C}_{4-10}$ -cycloalkyl or  $-(\text{CH}_2)_{0-2}(\text{HZ}^2)$ , where  $\text{HZ}^2$  represents a 4- to 7-membered heterocycle having up to two heteroatoms selected from the group consisting of N, O and S, where each of these groups may be substituted by  $-\text{OH}$ , 25  $\text{CO}_2\text{H}$  or  $\text{NH}_2$  or  $-\text{L}-\#1$ ;

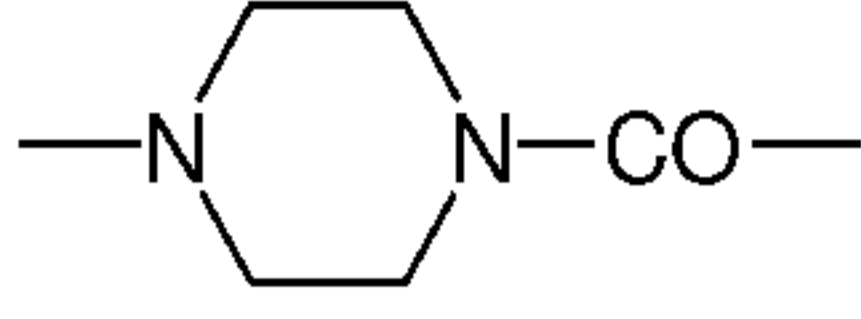
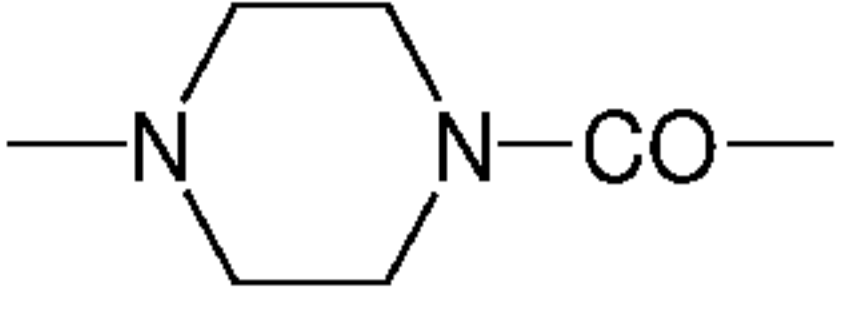
where one or none of the substituents  $\text{R}^1$ ,  $\text{R}^2$ ,  $\text{R}^3$ ,  $\text{R}^4$ ,  $\text{R}^5$ ,  $\text{R}^8$  and  $\text{R}^{10}$  represents or (in the case of  $\text{R}^8$ ) contains  $-\text{L}-\#1$ ,

30 L represents the linker and #1 represents the bond to the binder or derivative thereof,

$\text{R}^9$  represents H, F,  $\text{CH}_3$ ,  $\text{CF}_3$ ,  $\text{CH}_2\text{F}$  or  $\text{CHF}_2$ ;

35 where  $-\text{MOD}$  represents  $-(\text{NR}^{10})_n-(\text{G}1)_o-\text{G}2-\text{H}$ , where

$\text{R}^{10}$  represents H or  $\text{C}_1$ - $\text{C}_3$ -alkyl;

G1 represents  $\text{-NHCO-}$ ,  $\text{-CONH-}$  or  (where, if G1 represents  $\text{-NHCO-}$  or ,  $\text{R}^{10}$  does not represent  $\text{NH}_2$ );

n is 0 or 1;

5

o is 0 or 1; and

G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups  $\text{-O-}$ ,  $\text{-S-}$ ,  $\text{-SO-}$ ,  $\text{SO}_2$ ,  $\text{-NR}^y\text{-}$ ,  $\text{-NR}^y\text{CO-}$ ,  $\text{CONR}^y\text{-}$ ,  $\text{-NR}^y\text{NR}^y\text{-}$ ,  $\text{-SO}_2\text{NR}^y\text{NR}^y\text{-}$ ,  $\text{-CONR}^y\text{NR}^y\text{-}$  (where  $\text{R}^y$  represents H, phenyl,  $\text{C}_1\text{-C}_{10}$ -alkyl,  $\text{C}_2\text{-C}_{10}$ -alkenyl or  $\text{C}_2\text{-C}_{10}$ -alkynyl, each of which may be substituted by  $\text{-NHCONH}_2$ ,  $\text{-COOH}$ ,  $\text{-OH}$ ,  $\text{-NH}_2$ ,  $\text{NH-CNNH}_2$ , sulphonamide, sulphone, sulphoxide or sulphonic acid),  $\text{-CO-}$ ,  $\text{-CR}^x\text{=N-O-}$  (where  $\text{R}^x$  represents H,  $\text{C}_1\text{-C}_3$ -alkyl or phenyl), where the hydrocarbon chain including any side chains may be substituted by  $\text{-NHCONH}_2$ ,  $\text{-COOH}$ ,  $\text{-OH}$ ,  $\text{-NH}_2$ ,  $\text{NH-CNNH}_2$ , sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group  $\text{-MOD}$  preferably has at least one group  $\text{-COOH}$ ;

15  
20

and the salts, solvates and salts of the solvates thereof.

The conjugates according to the invention can have chemically labile linkers, enzymatically labile linkers or stable linkers. Particular preference is given to stable linkers and linkers which can be cleaved by cathepsin.

25

The invention furthermore provides processes for preparing the conjugates according to the invention, and also precursors and intermediates for the preparation.

30

The preparation of the conjugates according to the invention regularly comprises the following steps:

(i) Preparation of a linker precursor which optionally carries protective groups and has a reactive group which is capable of coupling to the binder;

35

(ii) Conjugation of the linker precursor to the derivative, which optionally carries protective groups, of a low-molecular weight KSP inhibitor (preferably a KSP inhibitor having the substructure I(sub), particularly preferably of formula (Ia) and in particular of formula (IIa), where in these formulae there is as yet no bond to a linker), giving a reactive KSP inhibitor/linker conjugate which optionally carries protective groups;

10

(iii) Removal of any protective groups present in the KSP inhibitor/linker conjugate and

(iv) Conjugation of the binder to the KSP inhibitor/linker conjugate, giving the binder/KSP inhibitor conjugate according to the invention.

Attachment of the reactive group may also take place after the construction of an optionally protected KSP inhibitor/linker precursor conjugate.

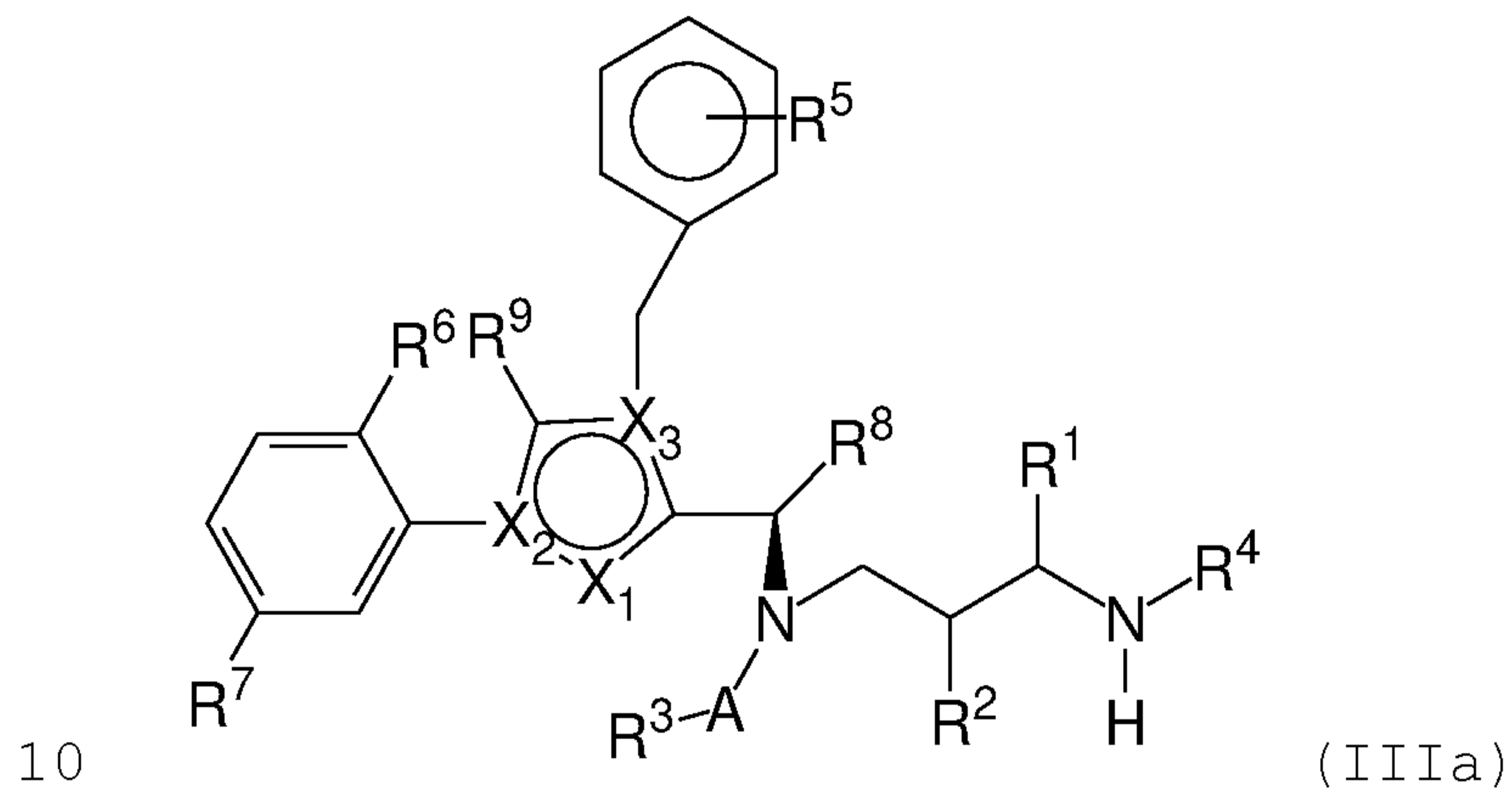
Depending on the linker, succinimide-linked ADCs may, after conjugation, be converted according to Scheme 26 into the open-chain succinamides, which have an advantageous stability profile.

As illustrated above, conjugation of the linker precursor to a low-molecular weight KSP inhibitor may take place by substitution of a hydrogen atom at  $R^{1a}$ ,  $R^{2a}$ ,  $R^{4a}$  or  $R^{10}$  in substructure I(sub),  $R^{1a}$ ,  $R^{2a}$ ,  $R^{3a}$ ,  $R^{4a}$ ,  $R^{8a}$  or  $R^{10}$  in formula (Ia), or  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$ ,  $R^8$  or  $R^{10}$  in formula (IIa) by the linker. In the synthesis steps prior to the conjugation, any functional groups present may also be present in protected form. Prior to the conjugation step, these protective groups are removed by known methods of peptide chemistry. Conjugation can take place chemically by various routes, as shown in an exemplary manner in Schemes 7 to 31 in the examples. In particular, it is optionally possible to modify the low-molecular weight KSP

35

inhibitor for conjugation to the linker, for example by introduction of protective groups or leaving groups to facilitate substitution.

- 5 In particular, the invention provides novel low-molecular weight KSP inhibitors which may optionally be conjugated to a binder. These KSP inhibitors or their binder conjugates have the following general formula (IIIa):



where

15  $X_1$  represents N,  $X_2$  represents N and  $X_3$  represents C; or

$X_1$  represents CH or CF,  $X_2$  represents C and  $X_3$  represents N; or

$X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C; or

20  $X_1$  represents CH,  $X_2$  represents N and  $X_3$  represents C

(with  $X_1$  representing CH,  $X_2$  representing C and  $X_3$  representing N being preferred);

25  $R^1$  represents H, -L-BINDER, -MOD or  $-(CH_2)_{0-3}Z$ , where Z represents -H, -NH $Y^3$ , -O $Y^3$ , -S $Y^3$ , halogen, -CO-N $Y^1Y^2$  or -CO-O $Y^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H, NH $_2$ , -  
 (CH $_2$ CH $_2$ O) $_{0-3}$ -(CH $_2$ ) $_{0-3}Z'$  (e.g. -(CH $_2$ ) $_{0-3}Z'$ ) or -CH(CH $_2$ W)Z', and  $Y^3$   
 30 represents H or  $-(CH_2)_{0-3}Z'$ , where Z' represents H, NH $_2$ , SO $_3$ H, COOH, -NH-CO-CH $_2$ -CH $_2$ -CH(NH $_2$ )COOH or -(CO-NH-CH $Y^4$ ) $_{1-3}$ COOH, where W

represents H or OH,

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>;

$R^2$  represents -L-BINDER, H, -MOD, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where  $R^{10}$  represents L-#1, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH, or OH;

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where  $Y^1$  and  $Y^2$  independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and  $Y^3$  represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and  $Y^5$  represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

$R^4$  represents -L-BINDER, H, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z,

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where  $Y^1$  and  $Y^2$  independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and  $Y^3$  represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and  $Y^5$  represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where



Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents -L-BINDER, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH or OH;

A represents CO, SO, SO<sub>2</sub>, SO<sub>2</sub>NH or CNNH;

R<sup>3</sup> represents -L-BINDER, -MOD or an optionally substituted alkyl, cycloalkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably -L-BINDER, or a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl or C<sub>5-10</sub>-heterocycloalkyl group which may be substituted by 1-3 -OH groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH (where "alkyl" preferably represents C<sub>1-10</sub>-alkyl);

R<sup>5</sup> represents -L-BINDER,, H, NH<sub>2</sub>, NO<sub>2</sub>, halogen (in particular F, Cl, Br), -CN, CF<sub>3</sub>, -OCF<sub>3</sub>, -CH<sub>2</sub>F, -CH<sub>2</sub>F, SH or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl or -(CH<sub>2</sub>)<sub>0-2</sub>-(HZ<sup>2</sup>), where HZ<sup>2</sup> represents a 4- to 7-membered heterocycle having

up to two heteroatoms selected from the group consisting of N, O and S (preferably oxetane), where each of these groups may be substituted by -OH, CO<sub>2</sub>H or NH<sub>2</sub> or -L-BINDER;

5 where L represents a linker and BINDER represents a binder or a derivative thereof, where the binder may optionally be attached to a plurality of active compound molecules,

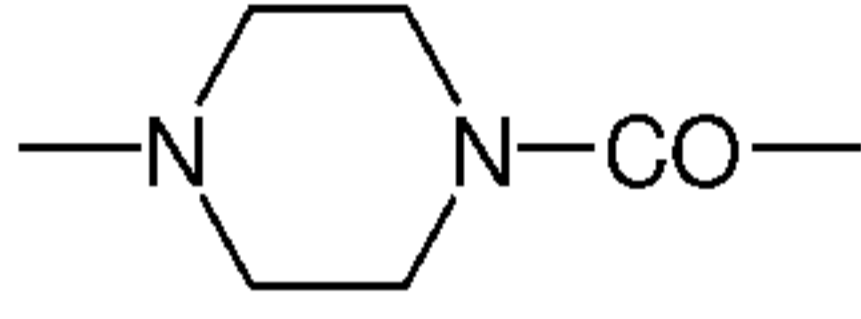
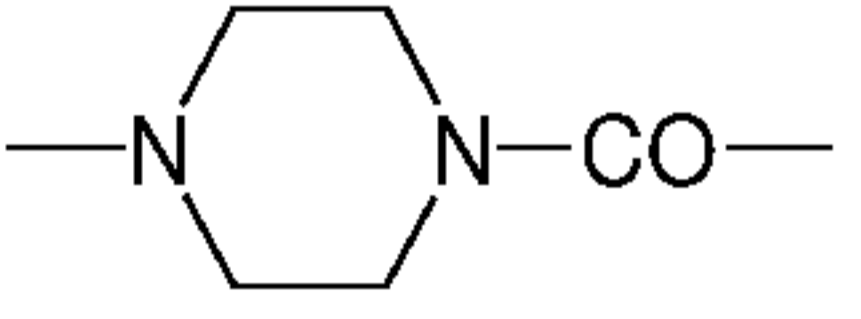
10 where at most one representative of R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>8</sup> and R<sup>10</sup> represents -L-binder;

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano, (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy, 15 NO<sub>2</sub>, NH<sub>2</sub>, COOH or halogen (in particular F, Cl, Br),

R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

20 where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

R<sup>10</sup> represents H or C<sub>1-3</sub>-alkyl;

G1 represents -NHCO- , -CONH- or  (where, if G1 represents -NHCO- or  , R<sup>10</sup> does not represent NH<sub>2</sub>);

25 n is 0 or 1;

o is 0 or 1; and

30 G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>- (where R<sup>y</sup> represents H, phenyl, C<sub>1-10</sub>-alkyl, C<sub>2-10</sub>-alkenyl or 35 C<sub>2-10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1-3</sub>-

alkyl or phenyl), where the hydrocarbon chain including any side chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group -MOD preferably has at least one group -COOH;

5

and the salts, solvates and salts of the solvates thereof.

### ***Description of the figures***

10 **Figure 1:** Alignment of the TWEAKR cysteine-rich domain (amino acids 34 to 68) of various species. (The numbers show the amino acid position in full-length constructs including the signal sequences; SEQ ID NO: 169).

15 **Figure 2:** A - Schematic diagram of the structure of TWEAKR (SEQ ID NO: 169). The diagram shows the extracellular domain (amino acids 28-80) (SEQ ID NO: 168) including the cysteine-rich domain (36-67), the transmembrane domain - TM (81-101) and the intracellular domain (102-129). TPP-2202 - the complete  
20 ectodomain (28-80), fused to the Fc domain of hIgG1. TPP-2203 - extracellular domain with N- and C-terminal truncation (34-68), fused to the Fc domain of hIgG1. Disulphide bridges Cys36-Cys49, Cys52-Cys67 and Cys55-Cys64 are indicated by black bars. N-terminally and C-terminally, TPP-2203 contains two amino acids  
25 more and one amino acid more, respectively, than the unmodified cysteine-rich domain to ensure proper folding. TPP-1984 - extracellular domain having C-terminal truncation (28-68), fused to an HIS6 tag. All three constructs show comparable binding to the antibodies according to the invention and PDL-192 (TPP-  
30 1104). P4A8 (TPP-1324) binds only to the full-length extracellular domain (TPP-2202).

B - Amino acid sequence of the extracellular domain: It has been published that the amino acid 64 is essential for TWEAK ligand  
35 binding; and the amino acid 47 is essential for binding of the antibodies according to the invention, as was determined here.

**Figure 3:** Interaction of the TWEAKR ectodomain with antibodies

and reference antibodies. What is shown is the result of an ELISA with TWEAKR-Fc fusion protein coating (TPP-2202, 1 µg/ml) and with 0.08 µg/ml (open bars) and 0.03 µg/ml (solid bars) of biotinylated IgG as soluble binding partner. Detection was carried out using streptavidin-HRP and Amplex Red substrate. Y is the "ELISA signal intensity [Rfu]"; X are the "tested antibody constructs": a is "TPP-2090"; b is "TPP-2084"; c is "PDL-192(TPP-1104)"; d is "P4A8(TPP-1324)"; e is "P3G5(TPP-2195)"; f is "136.1(TPP-2194)"; h is "ITEM1"; i is "ITEM4"; j is a mouse isotype control; k is a human isotype control. All antibodies examined show saturated binding at a concentration of 80 ng/ml.

**Figure 4:** Interaction of the cysteine-rich domain of TWEAKR with antibodies according to the invention and reference antibodies. What is shown is the result of an ELISA with TWEAKR (34-68)-Fc fusion protein coating (TPP-2203, 1 µg/ml) and 0.08 µg/ml (open bars) and 0.3 µg/ml (solid bars) of biotinylated IgG as soluble binding partner. Detection was carried out using streptavidin-HRP and Amplex Red substrate. X are the "antibody constructs tested": a is "TPP-2090"; b is "TPP-2084"; c is "PDL-192(TPP-1104)"; d is "P4A8(TPP-1324)"; e is "P3G5(TPP-2195)"; f is "136.1(TPP-2194)"; h is "ITEM1"; i is "ITEM4"; j is a mouse isotype control; k is a human isotype control. All antibodies examined show saturated binding at a concentration of 80 ng/ml.

**Figure 5:** Interaction of TWEAKR (28-68) with antibodies according to the invention and reference antibodies. What is shown is the result of an ELISA with TWEAKR (28-68)-HIS coating (TTP-1984, 1 µg/ml) and 0.08 µg/ml (open bars) and 0.3 µg/ml (solid bars) of biotinylated IgG as soluble binding partner. Detection was carried out using streptavidin-HRP and Amplex Red substrate. X are the "antibody constructs tested": a is "TPP-2090"; b is "TPP-2084"; c is "PDL-192(TPP-1104)"; d is "P4A8(TPP-1324)"; e is "P3G5(TPP-2195)"; f is "136.1(TPP-2194)"; h is "ITEM1"; i is "ITEM4"; j is a mouse isotype control; k is a human isotype control. All antibodies examined show saturated binding at a concentration of 80 ng/ml.

**Figure 6:** A - Alanine scan of the cysteine-rich domain. Muteins of TWEAKR(34-68)-Fc were analysed for PDL-192 (TPP-1104) (X)- and TPP-2090 (Y)-binding. S37A, R38A, S40A, W42A, S43A, D45A, D47A, K48A, D51A, S54A, R56A, R58A, P59A, H60A, S61A, D62A, F63A and L65A muteins were expressed in HEK293 cells (black diamonds). PFL192 (TPP-1104) and TPP-2090 were coated (1 µg/ml) and an 8-fold diluted supernatant of the HEK293 fermentation broth was added for TWEAKR protein binding. X is the "ELISA intensity of the PDL-192/TPP-1104) interaction [Rfu]", Y is the "ELISA intensity of the TPP-2090 interaction [Rfu]". TPP-2090 (Y) shows reduced binding for the D74A-TWEAKR mutein (closed box), and PDL-192 (TPP-1104) (X) shows reduced binding to R56A (spotted box).

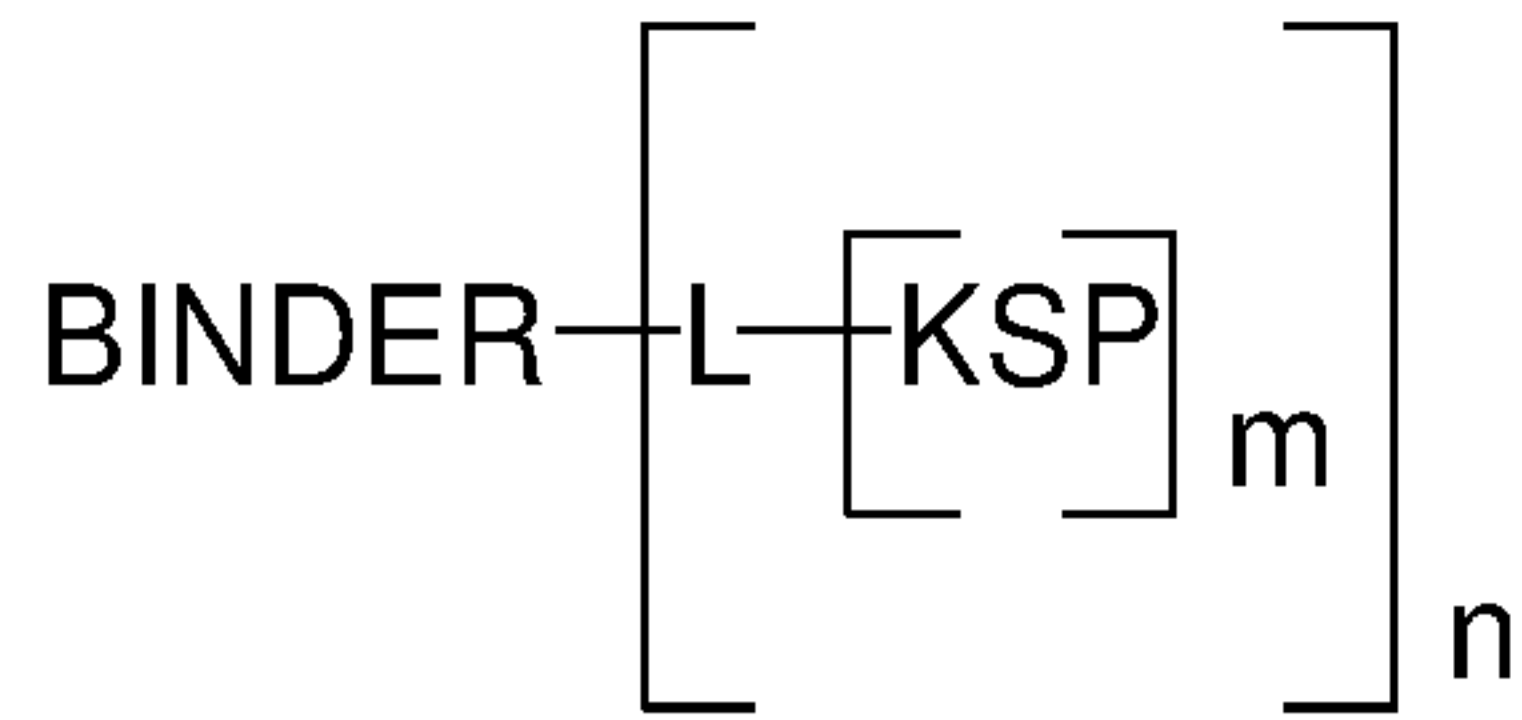
B - Y is the "binding in % normalized to the wild-type binding signal [%]", 1 is "TPP-2090"; 2 is "PDL-192 (TPP-1104)"; 3 is "P4A8 (TPP-1324)". (1 µg/ml), the TWEAKR variant was added at 250 ng/ml, detection was via anti-HIS HRP. Compared to the wild-type construct, TTP-2090 shows less than 5% binding.

**Figure 7:** NMR structure of the TWEAKR ectodomain as published by Pellegrini et al. (FEBS 280:1818-1829). TWEAK binding depends on L46 (Pellegrini et al.), TTP-2090 binding depends on D47 and PDL-192 binds to R56. PDL-192 binds opposite the TWEAK ligand binding site, TPP-2090 binds directly to the TWEAK ligand site.

#### ***Detailed description of the invention***

The invention provides conjugates of a binder or derivative thereof with one or more active compound molecules, the active compound molecule being a kinesin spindle protein inhibitor (KSP inhibitor) attached to the binder via a linker L.

The conjugate according to the invention can be represented by the general formula



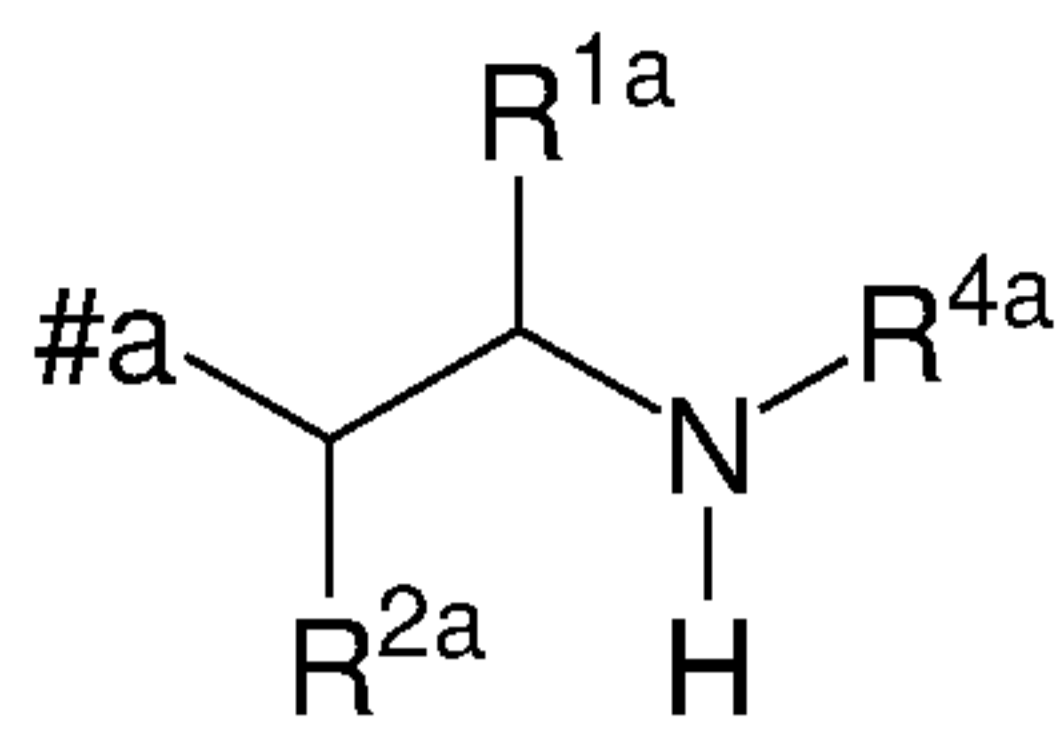
where BINDER represents the binder, preferably an antibody, L represents the linker, KSP represents the KSP inhibitor, m represents a number from 1 to 2, preferably 1, and n represents a number from 1 to 50, preferably from 1.2 to 20 and particularly preferably from 2 to 8. Here, m is the number of KSP inhibitors per linker and n a mean of the number of KSP inhibitor/linker conjugates per BINDER. The sum of all KSP present in the conjugate is thus the product of m and n. KSP-L preferably has the formula (I) or (II) shown above. The binder is preferably a binder peptide or protein such as, for example, an antibody. Furthermore, the linker is preferably attached to different amino acids of the binder peptide or protein or derivative thereof. Particular preference is given to binding to different cysteine residues of the binder.

Binders which can be used according to the invention, KSP inhibitors which can be used according to the invention and linkers which can be used according to the invention which can be used in combination without any limitation are described below. In particular, the binders represented in each case as preferred or particularly preferred can be employed in combination with the KSP inhibitors represented in each case as preferred or particularly preferred, optionally in combination with the linkers represented in each case as preferred or particularly preferred.

### ***KSP inhibitors and their binder conjugates***

Low-molecular weight KSP inhibitors are known, for example, from WO2006/044825; WO2006/002236; WO2005/051922; WO2006/060737; WO03/060064; WO03/040979; and WO03/049527.

As a rule, KSP inhibitors have the following substructure I(sub):



I (sub)

where

5

#a represents a bond to the remainder of the molecule;

$R^{1a}$  represents H or  $-(CH_2)_{0-3}Z$ , where Z represents -H, halogen,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

10

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  (e.g.  $-(CH_2)_{0-3}Z'$ ) or  $-CH(CH_2W)Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$ ,  $COOH$ ,  $-NH-CO-CH_2-CH_2-CH(NH_2)COOH$  or  $-(CO-NH-CHY^4)_{1-3}COOH$ , where W

15

represents H or OH,

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ ;

20

$R^{2a}$  and  $R^{4a}$  independently of one another represent H,  $-CO-CHY^4-NHY^5$  or  $-(CH_2)_{0-3}Z$ , or  $R^{2a}$  and  $R^{4a}$  together (with formation of a pyrrolidine ring) represent  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$

25

represents H,  $NH_2$ ,  $COOH$ ,  $SO_3H$ , SH or OH,

where Z represents -H,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

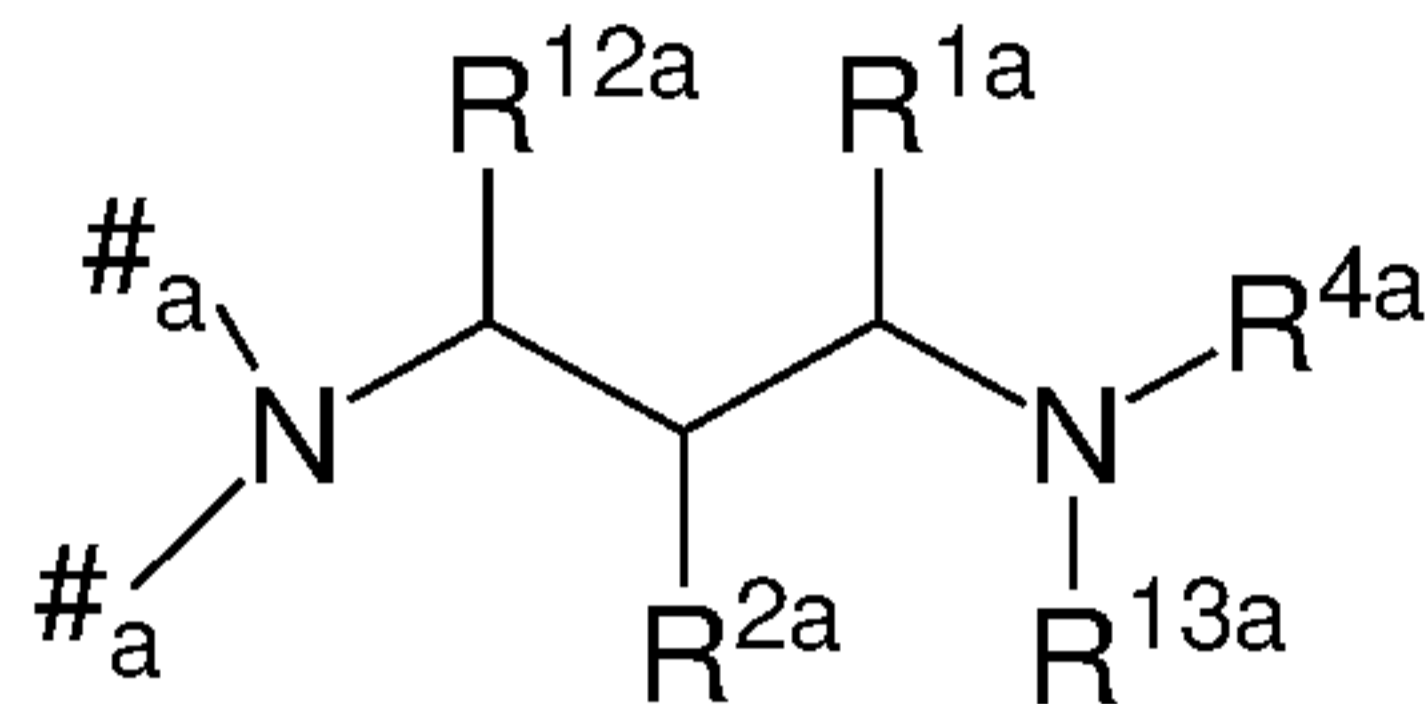
where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ,

30

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by -

NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl.

5 Particularly frequently encountered is the following substructure II(sub)



II(sub)

10 where #a, R<sup>1a</sup>, R<sup>2a</sup>, R<sup>4a</sup> have the same meaning as in I(sub) and R<sup>12a</sup> and R<sup>13a</sup> represent H or R<sup>12a</sup> and R<sup>13a</sup> together (with formation of a piperidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents H, NH<sub>2</sub>, COOH, SO<sub>3</sub>H, SH or OH;

15 where Z represents -H, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' (e.g. -(CH<sub>2</sub>)<sub>0-3</sub>Z'), and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub>, COOH or -(CO-NH-  
 20 CHY<sup>4</sup>)<sub>1-3</sub>COOH, where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>.

25 In particular, a number of KSP inhibitors have the substructure II(sub) where R<sup>1a</sup>, R<sup>2a</sup>, R<sup>4a</sup>, R<sup>12a</sup> and R<sup>13a</sup> represent H.

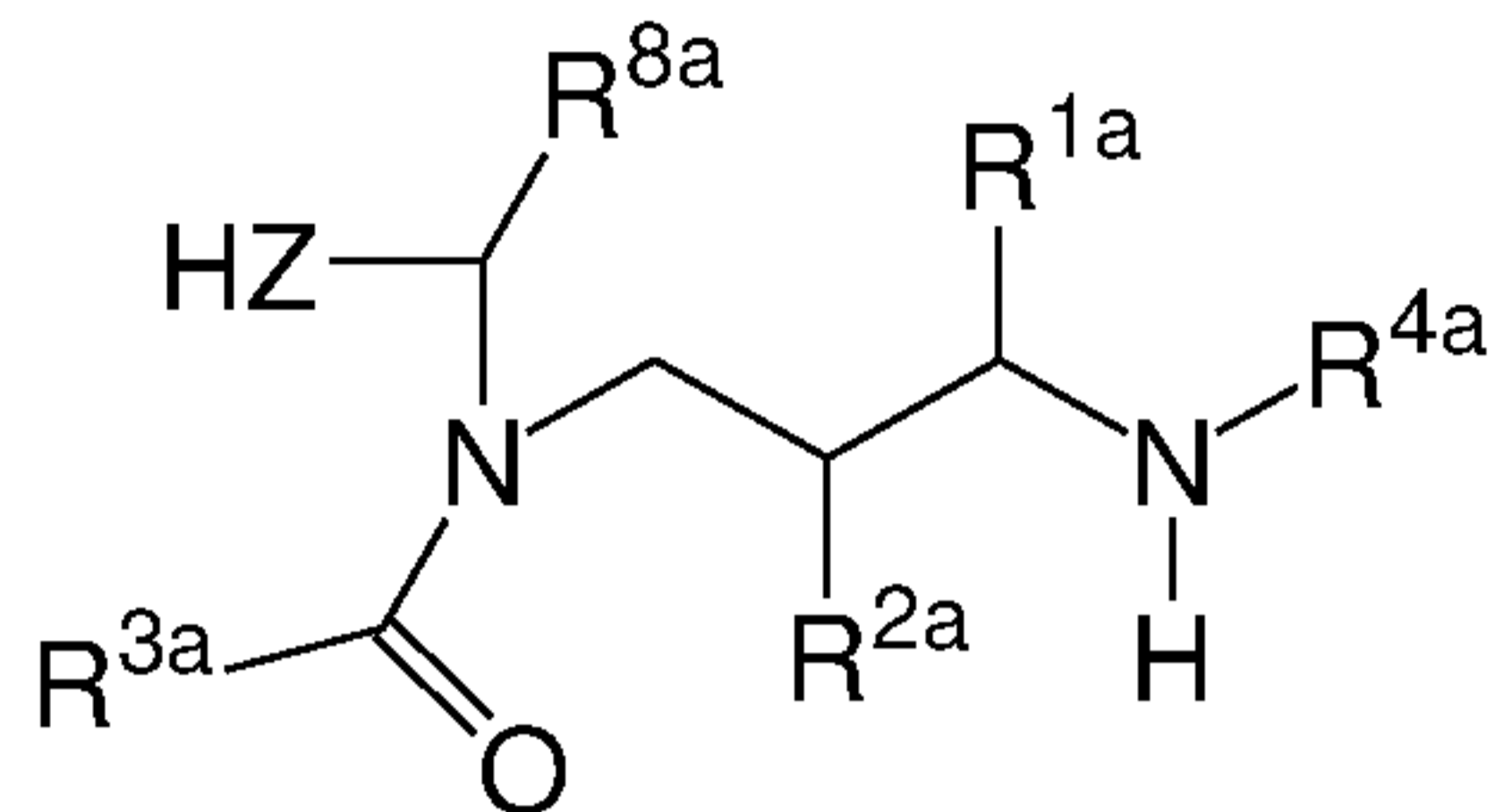
According to the invention, use may be made of KSP inhibitors of the substructure I(sub) or the substructure II(sub). The KSP  
 30 inhibitors which are used in accordance with the invention also include, for example, ispinesib (Cytokinetics/GSK), MK-0731 (Merck), AZD4877 (AstraZeneca), ARRY-520 (Array BioPharma) and ARQ 621 (ArQule).



KSP inhibitors which are preferred in accordance with the invention have the following basic structure:

Formula (I):

5



(I)

where

10  $R^{1a}$  represents H, -MOD or  $-(CH_2)_{0-3}Z$ , where Z represents -H, halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  (e.g.  $-(CH_2)_{0-3}Z'$ ) or  $-CH(CH_2W)Z'$ , and  $Y^3$   
 15 represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$ ,  $COOH$ ,  $-NH-CO-CH_2-CH_2-CH(NH_2)COOH$  or  $-(CO-NH-CHY^4)_{1-3}COOH$ , where W represents H or OH,

where  $Y^4$  independently of one another represents straight-chain  
 20 or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ ;

$R^{2a}$  and  $R^{4a}$  independently of one another represent H,  $-CO-CHY^4-$   
 25  $NHY^5$  or  $-(CH_2)_{0-3}Z$ , or  $R^{2a}$  and  $R^{4a}$  together (with formation of a pyrrolidine ring) represent  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$  represents H,  $SO_3H$ ,  $NH_2$ ,  $COOH$ , SH or OH,

where Z represents -H, halogen,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  
 30  $CO-OY^3$ ,

where  $Y_1$  and  $Y_2$  independently of one another represent H,  $NH_2$  or  
 $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$

represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl.

R<sup>3a</sup> represents -MOD or an optionally substituted alkyl, cycloalkyl, aryl, heteroaryl, heteroalkyl or heterocycloalkyl group,

preferably a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl or C<sub>5-10</sub>-heterocycloalkyl group which may be substituted by 1-3 -OH groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH

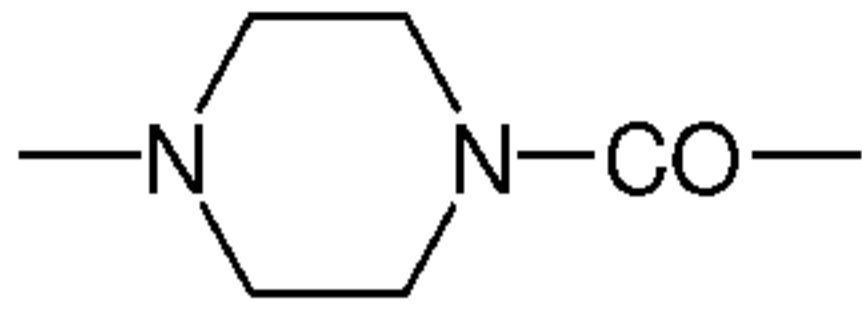
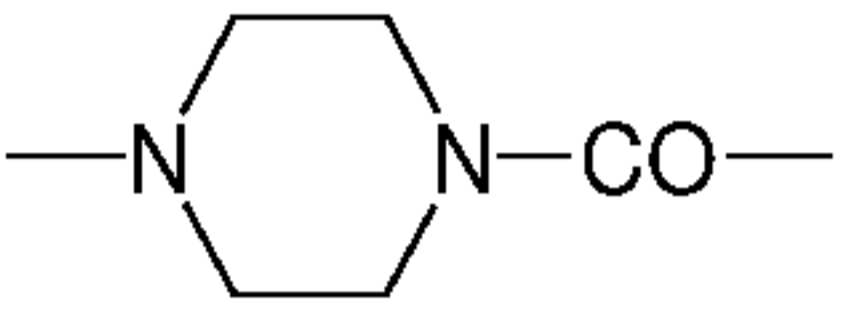
(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl);

R<sup>8a</sup> represents C<sub>1-10</sub>-alkyl;

HZ represents a mono- or bicyclic heterocycle which may be substituted by one or more substituents selected from the group consisting of halogen, C<sub>1-10</sub>-alkyl groups, C<sub>6-10</sub>-aryl groups and C<sub>6-10</sub>-aralkyl groups which may optionally be substituted by halogen;

where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

R<sup>10</sup> represents H or C<sub>1</sub>-C<sub>3</sub>-alkyl;

G1 represents -NHCO- , -CONH- or  (where, if G1  
 5 represents -NHCO- or  , R<sup>10</sup> does not represent NH<sub>2</sub>);

n is 0 or 1;

o is 0 or 1; and

10

G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>-  
 15 (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl), where the hydrocarbon chain including any side  
 20 chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group -MOD preferably has at least one group -COOH;

and the salts, solvates and salts of the solvates thereof.

25

According to the invention, such a kinesin spindle protein inhibitor can be attached to the linker by substitution of a hydrogen atom at R<sup>1a</sup>, R<sup>2a</sup>, R<sup>3a</sup>, R<sup>4a</sup>, R<sup>8a</sup> or R<sup>10</sup> or optionally via one of the substituents of HZ, in particular via R<sup>1a</sup>, R<sup>2a</sup>, R<sup>3a</sup>,  
 30 R<sup>4a</sup> or R<sup>10</sup>.

The substituents of the formula (I) preferably have the following meanings, where these preferred meanings are preferably combined with one another:

35

R<sup>1a</sup> preferably represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  (e.g.  $-(CH_2)_{0-3}Z'$ ) or  $-CH(CH_2W)Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$ ,  
 5  $COOH$ ,  $-NH-CO-CH_2-CH_2-CH(NH_2)COOH$  or  $-(CO-NH-CHY^4)_{1-3}COOH$ , where  $W$  represents H or OH,

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally  
 10 substituted by  $-NH_2$ ;

$R^{2a}$  and  $R^{4a}$  independently of one another preferably represent H,  $-COCHY^4-NHY^5$  or  $-(CH_2)_{0-3}Z$ , or  $R^{2a}$  and  $R^{4a}$  together (with formation  
 15 of a pyrrolidine ring) represent  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$  represents H,  $SO_3H$ ,  $NH_2$ ,  $COOH$ ,  $SH$  or  $OH$ ,

where  $Z$  represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,  
 20

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ , where  $Y^4$  independently of one  
 25 another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$  and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl.

$R^{3a}$  preferably represents an optionally substituted alkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably  $-L-$   
 30  $\#1$  or a  $C_{1-10}$ -alkyl,  $C_{6-10}$ -aryl or  $C_{6-10}$ -aralkyl,  $C_{5-10}$ -heteroalkyl,  $C_{1-10}$ -alkyl- $O-C_{6-10}$ -aryl or  $C_{5-10}$ -heterocycloalkyl group which may be substituted by 1-3  $-OH$  groups, 1-3 halogen atoms, 1-3  
 35 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3  $O$ -alkyl groups, 1-3  $-SH$  groups, 1-3  $-S$ -alkyl groups, 1-3  $-O-CO$ -alkyl groups, 1-3  $-O-CO-NH$ -alkyl groups, 1-3  $-NH-CO$ -alkyl groups, 1-3  $-NH-CO-NH$ -alkyl groups, 1-3  $-S(O)_n$ -alkyl groups, 1-

3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub>  
groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z  
represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,  
where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  
5 -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-  
3-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or  
COOH,

(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl).

10

R<sup>8a</sup> preferably represents C<sub>1-10</sub>-alkyl.

HZ preferably represents a mono- or bicyclic heterocycle which  
may be substituted by one or more substituents selected from the  
15 group consisting of halogen, C<sub>1-10</sub>-alkyl groups, C<sub>6-10</sub>-aryl groups  
and C<sub>6-10</sub>-aralkyl groups which may optionally be substituted by  
halogen.

C<sub>1-10</sub>-Alkyl in the context of the invention (i.e. in the formula  
20 above and also in the formulae that follow) represents a  
straight-chain or branched alkyl radical having 1 to 10 carbon  
atoms. Examples which may be mentioned as being preferred are:  
methyl, ethyl, n-propyl, isopropyl, n-butyl, isobutyl, 1-  
methylpropyl and *tert*-butyl.

25

C<sub>6-10</sub>-Aryl- in the context of the invention represents a mono- or  
bicyclic aromatic homocycle, for example phenyl and naphthyl.

C<sub>6-10</sub>-Aralkyl group in the context of the invention represents a  
30 monocyclic aromatic homocycle, by way of example phenyl, to  
which a C<sub>1-4</sub>-alkyl group is attached. An exemplary C<sub>6-10</sub>-aralkyl  
group is benzyl.

C<sub>5-10</sub>-Heteroaryl in the context of the invention represents a  
35 mono- or bicyclic aromatic heterocycle having a total of 6 to  
10 ring atoms, where the ring(s) contains/contain one or two  
ring heteroatoms from the group consisting of N, O, S, SO and  
SO<sub>2</sub> and which is attached via a ring carbon atom or optionally

a ring nitrogen atom. Examples which may be mentioned are pyridyl, furanyl, pyrimidyl, imidazolyl, thienyl, thiophenyl, isoxazolyl, isothiazoyl, 1,2,3-oxadiazoyl, furazanyl, 1,2,3-triazoyl, 1,2,4-triazoyl, pyridazyl, pyrrolyl, triazinyl, 5 indolyl, quinolinyl, quinazolinyl, 1,3-benzodioxol, isoindolyl, indazolyl, 1H-pyrazolo[3,4-d]pyrimidyl, benzotriazolyl, isoquinolinyl, cinolinyl, phthalazinyl, pteridinyl, naphthyridinyl, benzimidazolyl, benzothiazolinyl, benzoxazolyl, 3,4-methylenedioxyphenyl and benzo[6]furanyl.

10

Mono- or bicyclic heterocycle in the context of the invention represents a mono- or bicyclic heterocycle having a total of 5 to 10 ring carbon atoms, where the ring(s) contains/contain one to three ring heteroatoms from the group consisting of N, O, S, 15 SO and SO<sub>2</sub> and which is attached via a ring carbon atom or optionally a ring nitrogen atom. Examples which may be mentioned are piperidyl, pyrrolinyl, morpholinyl, 3,4-methylenedioxyphenyl and tetrahydrofuranyl.

20 Halogen atom in the context of the invention represents F, Cl, Br or I.

By substitution of a hydrogen atom at R<sup>1a</sup>, R<sup>2a</sup>, R<sup>4a</sup> or R<sup>10</sup> in substructure I(sub) or substructure II(sub), or R<sup>1a</sup>, R<sup>2a</sup>, R<sup>3a</sup>, 25 R<sup>4a</sup>, R<sup>8a</sup> or R<sup>10</sup> at HZ in formula (I), the compound of the formula (I) may be attached to a linker in a manner known to the person of average skill. Particularly preferably, the substitution of the hydrogen atom takes place at R<sup>1a</sup>, R<sup>2a</sup>, R<sup>3a</sup>, R<sup>4a</sup> or at the pyrrolidine ring formed by R<sup>2a</sup> and R<sup>4a</sup>. This conjugation can take 30 place chemically by various routes, as shown in an exemplary manner in Schemes 7 to 31 in the examples. In particular, it is optionally possible to modify the low-molecular weight KSP inhibitor for the conjugation to the linker, for example by introducing protective groups or leaving groups to facilitate 35 substitution (such that in the reaction said leaving group, and not a hydrogen atom, is substituted by the linker). The KSP inhibitor - linker molecules obtained in this manner (where the linker has a reactive group for coupling to the binder) can then

be reacted with the binder to give a binder conjugate according to the invention. In the experimental section, this procedure is illustrated in an exemplary manner by a large number of examples.

5

Preferred for  $R^{1a}$  are H, -COOH, -CONH<sub>2</sub>, -(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub>, -CONZ''(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub> and -CONZ''CH<sub>2</sub>COOH, where Z'' represents H or NH<sub>2</sub>.

10 Preferred for  $R^{2a}$  and  $R^{4a}$  are H, or  $R^{2a}$  and  $R^{4a}$  together (with formation of a pyrrolidine ring) represent CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where  $R^{10}$  represents H.

15 Preferred for  $R^{3a}$  is C<sub>1-10</sub>-alkyl-, which may be substituted by -OH, O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-NH-alkyl, NH-CO-alkyl, NH-CO-NH-alkyl, S(O)<sub>n</sub>-alkyl, SO<sub>2</sub>-NH-alkyl, NH-alkyl, N(alkyl)<sub>2</sub> or NH<sub>2</sub> (where alkyl is preferably C<sub>1-3</sub>-alkyl).

20 Preferred for  $R^{8a}$  is a branched C<sub>1-5</sub>-alkyl group, preferably methyl, ethyl, n-propyl, isopropyl, n-butyl, isobutyl, 1-methylpropyl and *tert*-butyl.

25 Preferred for HZ is a mono- or bicyclic heterocycle which may be substituted by one or more substituents selected from the group consisting of halogen, C<sub>1-10</sub>-alkyl groups, C<sub>6-10</sub>-aryl groups and C<sub>6-10</sub>-aralkyl groups which may optionally be substituted by halogen.

30 Particularly preferably, HZ is a substituted pyrrole or pyrazole which is substituted in the ortho-position (with respect to the substituents with R<sub>1a</sub> etc.) by an optionally substituted benzyl group. Furthermore, the substituted pyrrole or pyrazole can preferably be substituted by oxo (in the case of dihydroquinazoline) or a phenyl group substituted by 1 or 2  
35 halogen atoms. Particularly preferably, HZ is a substituted pyrrole.

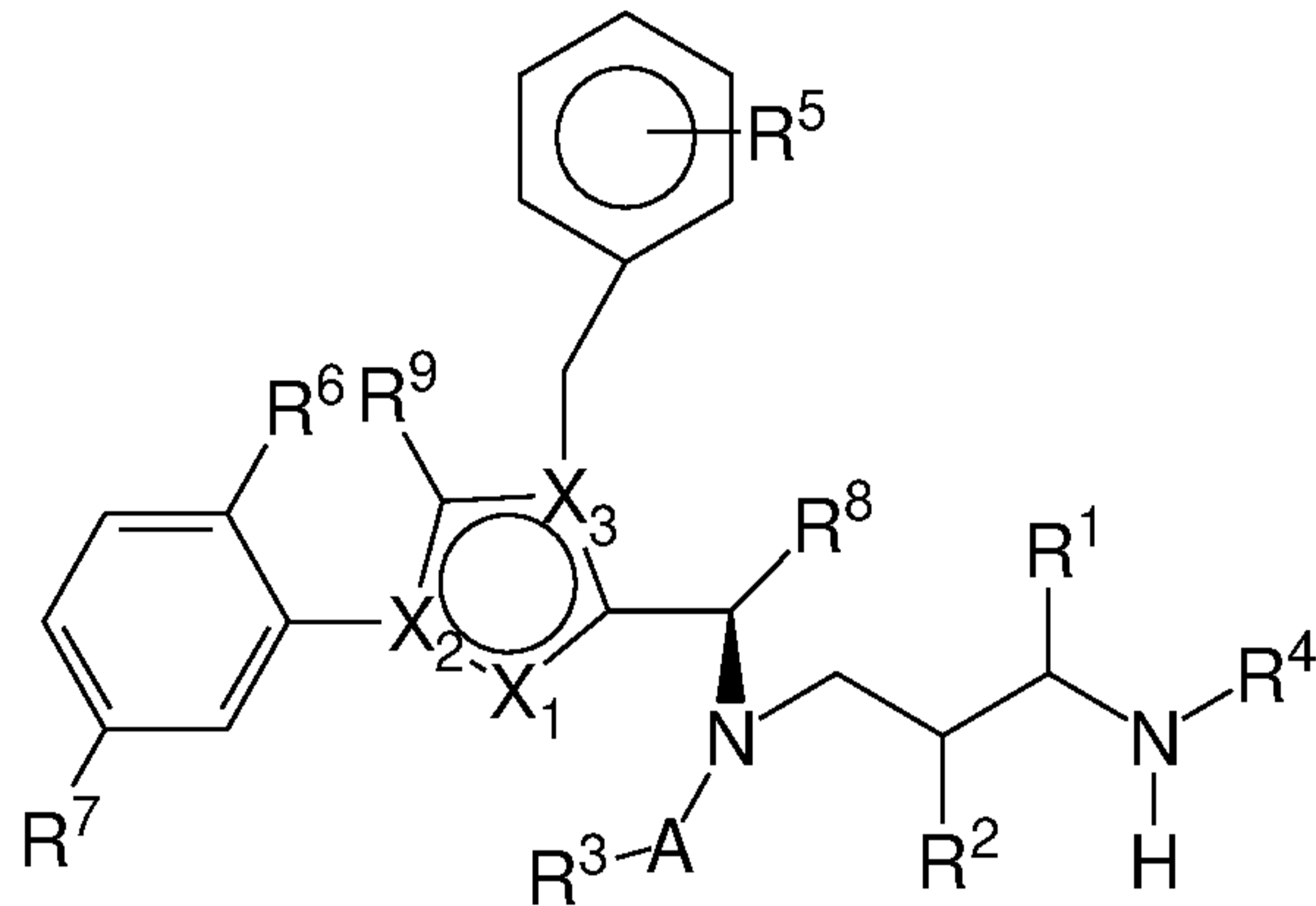
A KSP inhibitor which is preferably used is ispinesib. A further

preferred KSP inhibitor is Arry-520.

Other particularly preferred compounds have the formula (IIa) or (II) below:

5

Formula (IIa):



(IIa)

10 where

$X_1$  represents N,  $X_2$  represents N and  $X_3$  represents C; or

$X_1$  represents CH or CF,  $X_2$  represents C and  $X_3$  represents N; or

15

$X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C; or

$X_1$  represents CH or CF,  $X_2$  represents N and  $X_3$  represents C;

20 (with  $X_1$  representing CH,  $X_2$  representing C and  $X_3$  representing N being preferred);

$R^1$  represents H, -L-#1, -MOD or  $-(CH_2)_{0-3}Z$ , where Z represents -H, -NH $Y^3$ , -O $Y^3$ , -S $Y^3$ , halogen, -CO-N $Y^1Y^2$  or -CO-O $Y^3$ ,

25

where  $Y^1$  and  $Y^2$  independently of one another represent H, NH $_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  (e.g.  $-(CH_2)_{0-3}Z'$ ) or  $-CH(CH_2W)Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H, NH $_2$ , SO $_3H$ , COOH, -NH-CO-CH $_2$ -CH $_2$ -CH(NH $_2$ )COOH or  $-(CO-NH-CHY^4)_{1-3}COOH$ , where W

30 represents H or OH,



where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ ;

$R^2$  represents  $-L-#1$ , H,  $-MOD$ ,  $-CO-CHY^4-NHY^5$  or  $-(CH_2)_{0-3}Z$ ,

where Z represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $CO-OY^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ;

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ , and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

$R^4$  represents  $-L-#1$ , H,  $-CO-CHY^4-NHY^5$  or  $-(CH_2)_{0-3}Z$ ,

where Z represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $CO-OY^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ;

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ , and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring) represent  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$  represents  $L-#1$ ,

H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH or OH;

A represents CO, SO, SO<sub>2</sub>, SO<sub>2</sub>NH or CNNH;

5 R<sup>3</sup> represents -L-#1, -MOD or an optionally substituted alkyl, cycloalkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl or C<sub>5-10</sub>-heterocycloalkyl group which may be substituted by 1-3 -OH groups, 1-3 halogen  
 10 atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -  
 15 N(alkyl)<sub>2</sub> groups, 1-3 -NH((CH<sub>2</sub>CH<sub>2</sub>O)<sub>1-20</sub>H) groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where  
 20 Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH (where "alkyl" is preferably C<sub>1-10</sub>-alkyl);

R<sup>5</sup> represents -L-#1, H, -MOD, NH<sub>2</sub>, NO<sub>2</sub>, halogen (in particular F, Cl, Br), -CN, CF<sub>3</sub>, -OCF<sub>3</sub>, -CH<sub>2</sub>F, -CF<sub>3</sub>, SH or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where  
 25 Z represents -H, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

30

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano, (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy, NO<sub>2</sub>, NH<sub>2</sub>, COOH or halogen (in particular F, Cl, Br),

35

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl or -(CH<sub>2</sub>)<sub>0-2</sub>-

(HZ<sup>2</sup>), where HZ<sup>2</sup> represents a 4- to 7-membered heterocycle having up to two heteroatoms selected from the group consisting of N, O and S (preferably oxetane), where each of these groups may be substituted by -OH, CO<sub>2</sub>H or NH<sub>2</sub> or L-#1;

5

where one or none of the substituents R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>8</sup> and R<sup>10</sup> represents (or in the case of R<sup>8</sup> contains) -L-#1,

L represents the linker and #1 represents the bond to the binder or derivative thereof,

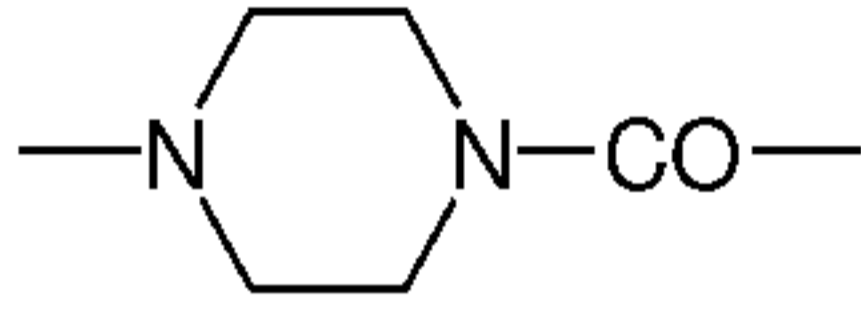
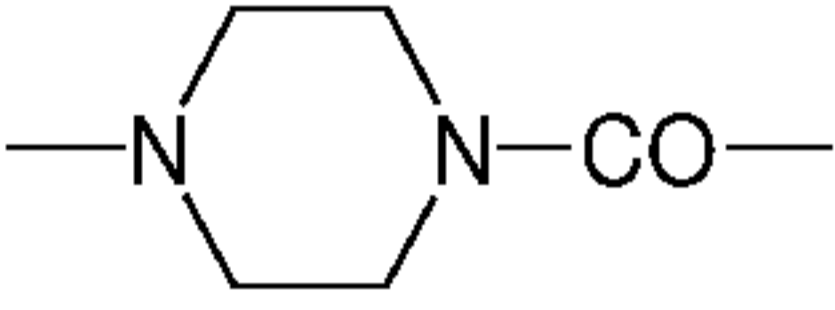
10

R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

15

R<sup>10</sup> represents H or C<sub>1</sub>-C<sub>3</sub>-alkyl;

G1 represents -NHCO-, -CONH- or  (where, if G1 represents -NHCO- or , R<sup>10</sup> does not represent NH<sub>2</sub>);

20

n is 0 or 1;

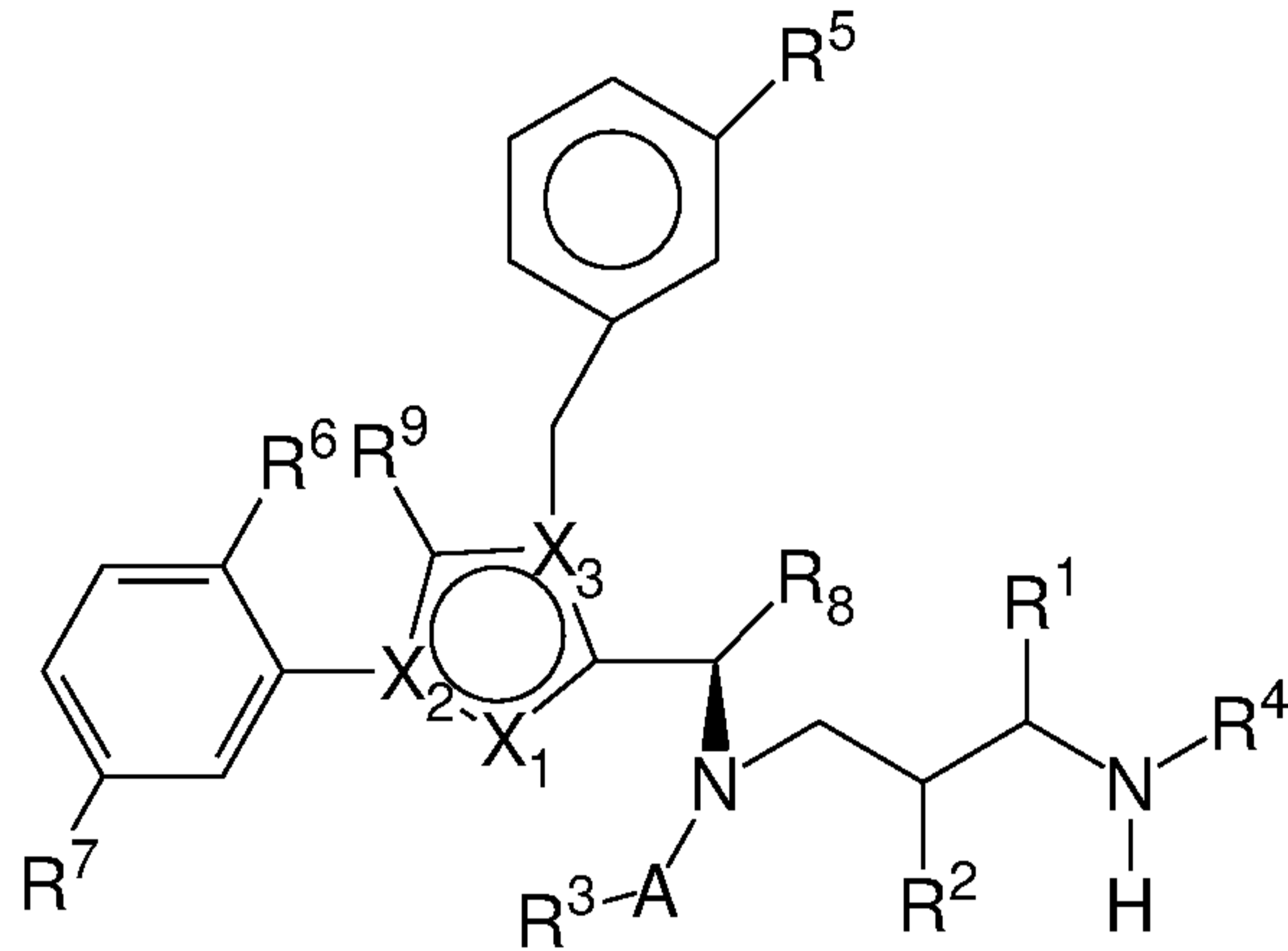
o is 0 or 1; and

G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>- (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl), where the hydrocarbon chain including any side chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group -MOD preferably has at least one group -COOH;

35

and the salts, solvates and salts of the solvates thereof.

Formula (II):



5 (II)

where

$X_1$  represents N,  $X_2$  represents N and  $X_3$  represents C; or

10

$X_1$  represents CH or CF,  $X_2$  represents C and  $X_3$  represents N; or

$X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C; or

15

$X_1$  represents CH,  $X_2$  represents N and  $X_3$  represents C

$R^1$  represents H,  $-L-\#1$  or  $-(CH_2)_{0-3}Z$ , where Z represents  $-H$ ,  $-NHY^3$ ,  $-OY^3$ ,  $-SY^3$ , halogen,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

20

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  (e.g.  $-(CH_2)_{0-3}Z'$ ) or  $-CH(CH_2W)Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $NH_2$ ,  $SO_3H$ ,  $COOH$ ,  $-NH-CO-CH_2-CH_2-CH(NH_2)COOH$  or  $-(CO-NH-CHY^4)_{1-3}COOH$ , where W represents H or OH;

25

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ ;

R<sup>2</sup> and R<sup>4</sup> independently of one another represent H, -L-#1, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where  
 5 R<sup>10</sup> represents H, -L-#1, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH or OH,

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

10 where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where Y<sup>4</sup> independently of one another represents straight-chain  
 15 or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub> or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub> or represents aryl or benzyl which are  
 20 optionally substituted by -NH<sub>2</sub> and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

A represents CO, SO, SO<sub>2</sub>, SO<sub>2</sub>NH or CNNH;

25 R<sup>3</sup> represents an optionally substituted alkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably -L-#1 or a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl or C<sub>5-10</sub>-heterocycloalkyl group which may be substituted by 1-3 -OH groups, 1-3 halogen atoms, 1-3  
 30 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub>  
 35 groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-</sub>

${}^3\text{-CH}(\text{NH}_2)\text{Z}'$  or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$  represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$ ,

(where "alkyl" preferably represents  $\text{C}_{1-10}$ -alkyl);

5

$\text{R}^5$  represents H, F,  $\text{NH}_2$ ,  $\text{NO}_2$ , halogen, SH or  $-(\text{CH}_2)_{0-3}\text{Z}$ , where Z represents -H, halogen,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$  or  
10  $-(\text{CH}_2)_{0-3}\text{Z}'$ , and  $\text{Y}^3$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$  represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$ ;

$\text{R}^6$  and  $\text{R}^7$  independently of one another represent H, cyano, (optionally fluorinated)  $\text{C}_{1-10}$ -alkyl, (optionally fluorinated)  
15  $\text{C}_{2-10}$ -alkenyl, (optionally fluorinated)  $\text{C}_{2-10}$ -alkynyl, hydroxy or halogen,

$\text{R}^8$  represents (optionally fluorinated)  $\text{C}_{1-10}$ -alkyl, (optionally fluorinated)  $\text{C}_{4-10}$ -cycloalkyl or optionally substituted oxetane;  
20 and

$\text{R}^9$  represents H, F,  $\text{CH}_3$ ,  $\text{CF}_3$ ,  $\text{CH}_2\text{F}$  or  $\text{CHF}_2$ ;

and the salts, solvates and salts of the solvates thereof.

25

By substitution of a hydrogen atom at  $\text{R}^1$ ,  $\text{R}^2$ ,  $\text{R}^3$ ,  $\text{R}^4$ ,  $\text{R}^5$  or  $\text{R}^8$  or at the pyrrolidine ring ( $\text{R}^{10}$ ) formed by  $\text{R}^2$  and  $\text{R}^4$ , in a manner known to the person of average skill the compound of the formula (IIa) or (II) in which none of the substituents  $\text{R}^1$ ,  $\text{R}^2$ ,  $\text{R}^3$ ,  $\text{R}^4$ ,  
30  $\text{R}^5$ ,  $\text{R}^8$  and  $\text{R}^{10}$  represents  $-\text{L}-\#1$  may be attached to a linker. This gives conjugates of the formula (IIa) or (II) where one of the substituents  $\text{R}^1$ ,  $\text{R}^2$ ,  $\text{R}^3$ ,  $\text{R}^4$ ,  $\text{R}^5$ ,  $\text{R}^8$  or  $\text{R}^{10}$  represents  $-\text{L}-\#1$ , L represents the linker and #1 represents the bond to the binder or the derivative thereof. If the KSP inhibitor according to  
35 formula (IIa) or (II) is conjugated with a binder, one of the substituents  $\text{R}^1$ ,  $\text{R}^2$ ,  $\text{R}^3$ ,  $\text{R}^4$ ,  $\text{R}^5$ ,  $\text{R}^8$  or  $\text{R}^{10}$  thus represents  $-\text{L}-\#1$ , where L represents the linker and #1 represents the bond to the binder or the derivative thereof. That is in the case of the

conjugates one of the substituents  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$ ,  $R^8$  and  $R^{10}$  represents  $-L-#1$ , where  $-L-#1$  is attached to the binder, for example an antibody. With particular preference, one of the substituents  $R^1$  and  $R^3$  represents  $-L-#1$ . The binder is preferably  
5 a human, humanized or chimeric monoclonal antibody or an antigen-binding fragment thereof, in particular an anti-TWEAKR antibody or an antigen-binding fragment thereof or an anti-EGFR antibody or an antigen-binding fragment thereof. Particular preference is given to an anti-TWEAKR antibody which binds  
10 specifically to amino acid D in position 47 (D47) of TWEAKR (SEQ ID NO:169), in particular the anti-TWEAKR antibody TPP-2090, or the anti-EGFR antibodies cetuximab or nimotuzumab.

Instead of  $-L-#1$ , it is also possible for the group  $-L-#3$  to be  
15 present in the compound, where L represents the linker and #3 represents the reactive group for binding to the binder or the derivative thereof. Compounds comprising  $-L-#3$  are reactive compounds which react with the binder or the derivative thereof. #3 is preferably a group which reacts with an amino or thiol  
20 group with formation of a covalent bond, preferably with the cysteine residue in a protein. The cysteine residue in a protein may of course be present naturally in the protein, may be introduced by biochemical methods or, preferably, may be generated by prior reduction of disulphides of the binder.

25 The compounds of the formula (IIa) or (II) in which one of the substituents  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$  and  $R^{10}$  represents  $-L-#1$  and in which

30  $X^1$  represents N,  $X_2$  represents N and  $X_3$  represents C;

$X_1$  represents CH or CF,  $X_2$  represents C and  $X_3$  represents N;

$X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C; or

35  $X_1$  represents CH,  $X_2$  represents N and  $X_3$  represents C

are particularly preferred,

in particular those in which

X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C; or X<sub>1</sub>  
 5 represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represents N. Particular  
 preference is given to compounds in which X<sub>1</sub> represents CH, X<sub>2</sub>  
 represents C and X<sub>3</sub> represents N.

For A, preference is given to CO (carbonyl).

10

Preferred for R<sup>1</sup> are -L-#1, H, -COOH, -CONHNH<sub>2</sub>, -(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub>, -  
 CONZ''(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub> and -CONZ''CH<sub>2</sub>COOH, where Z'' represents H or  
 NH<sub>2</sub>.

15 Preferred for R<sup>2</sup> and R<sup>4</sup> are H, -L-#1, or R<sup>2</sup> and R<sup>4</sup> together (with  
 formation of a pyrrolidine ring) represent CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-  
 CH<sub>2</sub>-, where R<sup>10</sup> represents H or -L-#1.

Preferred for R<sup>3</sup> is -L-#1 or C<sub>1-10</sub>-alkyl-, which may optionally  
 20 be substituted by -OH, O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-  
 NH-alkyl, NH-CO-alkyl, NH-CO-NH-alkyl, S(O)<sub>n</sub>-alkyl, SO<sub>2</sub>-NH-  
 alkyl, NH-alkyl, N(alkyl)<sub>2</sub> or NH<sub>2</sub> (where alkyl is preferably C<sub>1</sub>-  
 3-alkyl).

25 Preferred for R<sup>5</sup> is -L-#1, H or F.

Preferred for R<sup>6</sup> and R<sup>7</sup>, independently of one another, are H,  
 (optionally fluorinated) C<sub>1-3</sub>-alkyl, (optionally fluorinated) C<sub>2</sub>-  
 4-alkenyl, (optionally fluorinated) C<sub>2-4</sub>-alkynyl, hydroxy or  
 30 halogen,

Preferred for R<sup>8</sup> is a branched C<sub>1-5</sub>-alkyl group, in particular a  
 group of the formula -C(CH<sub>3</sub>)<sub>2</sub>-(CH<sub>2</sub>)<sub>0-2</sub>-R<sub>y</sub>, where R<sub>y</sub> represents -  
 H, -OH, CO<sub>2</sub>H, NH<sub>2</sub> or -L-#1. Particular preference is given to  
 35 the group of the formula -C(CH<sub>3</sub>)<sub>2</sub>-(CH<sub>2</sub>)<sub>0-2</sub>-R<sub>y</sub>, where R<sub>y</sub> represents  
 -H or -L-#1.

Preferred for R<sup>9</sup> is H or F.



Particular preference is given to compounds of the formula (IIa) or (II) in which none or one of the substituents  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$ ,  $R^8$  and  $R^{10}$  represents  $-L-#1$ , and

5

in which

$X_1$  represents N,  $X_2$  represents N and  $X_3$  represents C;

10  $X_1$  represents CH or CF,  $X_2$  represents C and  $X_3$  represents N;

$X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C; or

$X_1$  represents CH,  $X_2$  represents N and  $X_3$  represents C

15

A represents CO (carbonyl);

$R^1$  represents H,  $-COOH$ ,  $-CONHNH_2$ ,  $-(CH_2)_{1-3}NH_2$ ,  $-CONZ''(CH_2)_{1-3}NH_2$  and  $-CONZ''CH_2COOH$ , where  $Z''$  represents H or  $NH_2$ ;

20

$R^2$  and  $R^4$  represent H or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring) represent  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$  represents H or  $-L-#1$ ;

25  $R^3$  represents a phenyl group which may be mono- or polysubstituted by halogen (in particular F) or optionally fluorinated  $C_{1-3}$ -alkyl, or represents an optionally fluorinated  $C_{1-10}$ -alkyl group which may optionally be substituted by  $-OY^4$ ,  $-SY^4$ ,  $-O-CO-Y^4$ ,  $-O-CO-NH-Y^4$ ,  $NH-CO-Y^4$ ,  $-NH-CO-NH-Y^4$ ,  $S(O)_n-Y^4$   
30 (where n represents 0, 1 or 2),  $-SO_2-NH-Y^4$ ,  $NH-Y^4$  or  $N(Y^4)_2$ , where  $Y^4$  represents H, phenyl (optionally mono- or polysubstituted by halogen (in particular F) or optionally fluorinated  $C_{1-3}$ -alkyl), or alkyl (where the alkyl group may be substituted by  $-OH$ ,  $-COOH$ , and/or  $-NHCO-C_{1-3}$ -alkyl and where alkyl preferably  
35 represents  $C_{1-3}$ -alkyl);

where particularly preferably  $R^3$  may be substituted by  $-OH$ , O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-NH-alkyl, NH-CO-alkyl, NH-

CO-NH-alkyl, S(O)<sub>n</sub>-alkyl, SO<sub>2</sub>-NH-alkyl, NH-alkyl, N(alkyl)<sub>2</sub> or NH<sub>2</sub> (where alkyl preferably means C<sub>1-3</sub>-alkyl)

R<sup>5</sup> represents H or F;

5

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, (optionally fluorinated) C<sub>1-3</sub>-alkyl, (optionally fluorinated) C<sub>2-4</sub>-alkenyl, (optionally fluorinated) C<sub>2-4</sub>-alkynyl, hydroxy or halogen;

10 R<sup>8</sup> represents a branched C<sub>1-5</sub>-alkyl group; and

R<sup>9</sup> represents H or F.

Furthermore, it is preferred when (alone or in combination)

15

- R<sup>1</sup> represents -L-#1, COOH or H,

- R<sup>2</sup> and R<sup>4</sup> independently of one another represent -L-#1 or H or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents H or -L-#1,

20

- A represents CO,

25

- R<sup>3</sup> represents -(CH<sub>2</sub>)OH, -CH(CH<sub>3</sub>)OH, -CH<sub>2</sub>SCH<sub>2</sub>CH(COOH)NHCOCH<sub>3</sub>, -CH(CH<sub>3</sub>)OCH<sub>3</sub>, a phenyl group which may be substituted by 1-3 halogen atoms, 1-3 amino groups or 1-3 alkyl groups (which may optionally be halogenated), or represents -L-#1,

30

- R<sup>5</sup> represents -L-#1 or H,

- R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, C<sub>1-3</sub>-alkyl or halogen; in particular, R<sup>6</sup> and R<sup>7</sup> represent F;

35

- R<sup>8</sup> represents C<sub>1-4</sub>-alkyl (preferably tert-butyl); and/or

- R<sup>9</sup> represents H,

- where one of the substituents  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$  and  $R^{10}$  represents -L-#1.

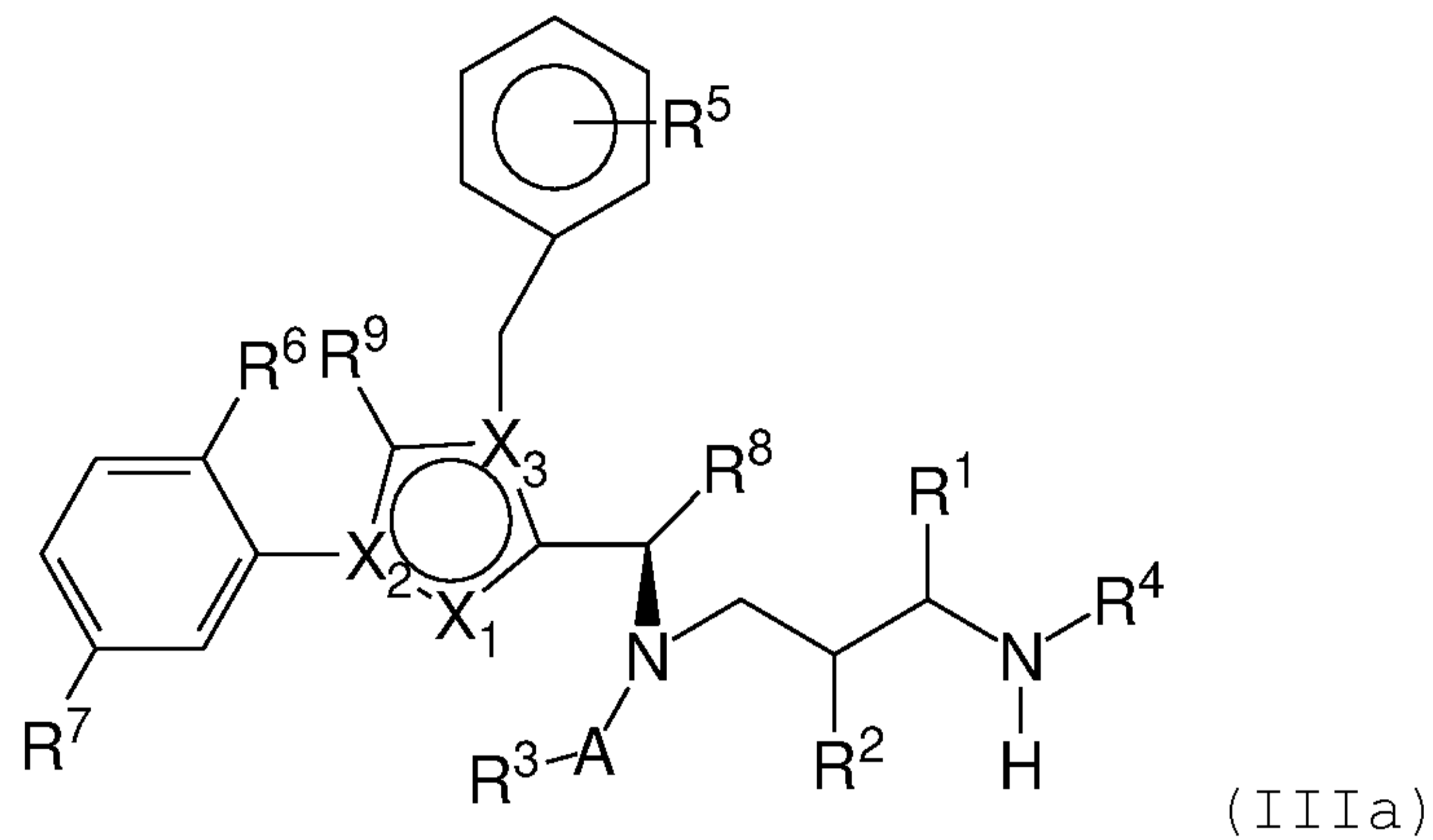
Additionally, in accordance with the invention it is preferred  
5 when

- $R^1$  represents -L-#1, COOH or H,
- $R^2$  and  $R^4$  independently of one another represent -L-#1 or H  
10 or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring)  
represent  $-\text{CH}_2\text{-CHR}^{10}\text{-}$  or  $-\text{CHR}^{10}\text{-CH}_2\text{-}$ , where  $R^{10}$  represents H or -  
L-#1,
- A represents CO,  
15
- $R^3$  represents  $-(\text{CH}_2)\text{OH}$ ,  $-\text{CH}(\text{CH}_3)\text{OH}$ ,  $-\text{CH}_2\text{SCH}_2\text{CH}(\text{COOH})\text{NHCOCH}_3$ ,  
 $-\text{CH}(\text{CH}_3)\text{OCH}_3$ , a phenyl group which may be substituted by 1-3  
halogen atoms, 1-3 amino groups or 1-3 alkyl groups (which may  
optionally be halogenated), or represents -L-#1,  
20
- $R^5$  represents -L-#1 or H,
- $R^6$  and  $R^7$  independently of one another represent H,  $\text{C}_{1-3}$ -  
alkyl or halogen; in particular,  $R^6$  and  $R^7$  represent F;  
25
- $R^8$  represents  $\text{C}_{1-4}$ -alkyl (preferably tert-butyl); and
- $R^9$  represents H,
- where one of the substituents  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$  and  $R^{10}$   
30 represents -L-#1.

Other particularly preferred compounds have the formula (IIIa)  
or (III) below:

35

Formula (IIIa):



where

5  $X_1$  represents N,  $X_2$  represents N and  $X_3$  represents C; or

$X_1$  represents CH or CF,  $X_2$  represents C and  $X_3$  represents N; or

$X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C; or

10

$X_1$  represents CH,  $X_2$  represents N and  $X_3$  represents C

(with  $X_1$  representing CH,  $X_2$  representing C and  $X_3$  representing N being preferred);

15

$R^1$  represents H, -L-BINDER, -MOD or  $-(CH_2)_{0-3}Z$ , where Z represents -H, -NH $Y^3$ , -O $Y^3$ , -S $Y^3$ , halogen, -CO-N $Y^1Y^2$  or -CO-O $Y^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H, NH $_2$ , -  
 20  $(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  (e.g.  $-(CH_2)_{0-3}Z'$ ) or -CH(CH $_2W$ )Z', and  $Y^3$  represents H or  $-(CH_3)_{0-3}Z'$ , where Z' represents H, NH $_2$ , SO $_3H$ , -COOH, -NH-CO-CH $_2$ -CH $_2$ -CH(NH $_2$ )COOH or  $-(CO-NH-CHY^4)_{1-3}COOH$ , where W represents H or OH,

25 where  $Y^4$  independently of one another represents straight-chain or branched C $_{1-6}$ -alkyl which is optionally substituted by -NHCONH $_2$ , or represents aryl or benzyl which are optionally substituted by -NH $_2$ ;

30  $R^2$  represents H, -L-BINDER, -MOD, -CO-CH $Y^4$ -NH $Y^5$  or  $-(CH_2)_{0-3}Z$ , where  $Y^4$  independently of one another represents straight-chain

or branched C<sub>1-6</sub> alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

5

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  
10 -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

R<sup>4</sup> represents H, -L-BINDER, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z,

15 where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  
20 -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally  
25 substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents -L-  
30 BINDER, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH or OH;

A represents CO, SO, SO<sub>2</sub>, SO<sub>2</sub>NH or CNNH;

R<sup>3</sup> represents -L-BINDER, -MOD or an optionally substituted alkyl,  
35 cycloalkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably -L-#1 or a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl or C<sub>5-10</sub>-heterocycloalkyl group which may be substituted by 1-3 -OH

groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3  
 5 3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-  
 10 CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH

(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl);

15 R<sup>5</sup> represents -L-BINDER, H, NH<sub>2</sub>, NO<sub>2</sub>, halogen (in particular F, Cl, Br), -CN, CF<sub>3</sub>, -OCF<sub>3</sub>, -CH<sub>2</sub>F, -CH<sub>2</sub>F, SH or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  
 20 -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano, (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated)  
 25 C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy, NO<sub>2</sub>, NH<sub>2</sub>, COOH or halogen (in particular F, Cl, Br),

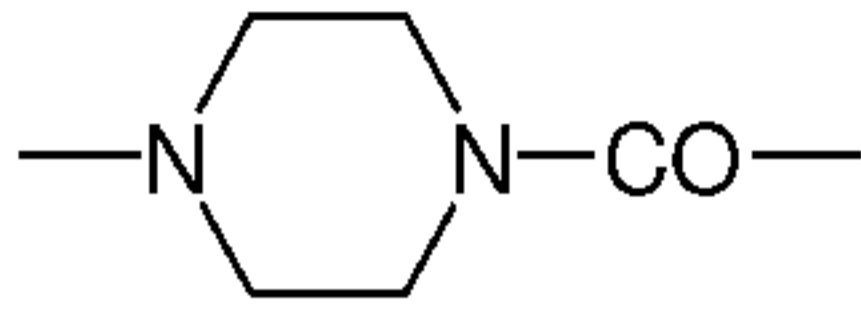
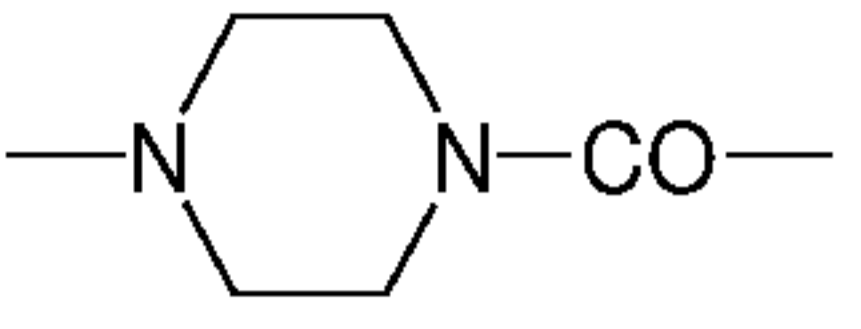
R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl or -(CH<sub>2</sub>)<sub>0-2</sub>-(HZ<sup>2</sup>), where HZ<sup>2</sup> represents a 4- to 7-membered heterocycle having up to two heteroatoms selected from the group consisting of N, O and S, where each of these groups may be substituted by -OH, CO<sub>2</sub>H or NH<sub>2</sub> or -L-BINDER;

35

R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

R<sup>10</sup> represents H or C<sub>1</sub>-C<sub>3</sub>-alkyl;

G1 represents -NHCO- , -CONH- or  (where, if G1  
 5 represents -NHCO- or  , R<sup>10</sup> does not represent NH<sub>2</sub>);

n is 0 or 1;

o is 0 or 1; and

10

G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>-  
 15 (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl), where the hydrocarbon chain including any side  
 20 chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group -MOD preferably has at least one group -COOH;

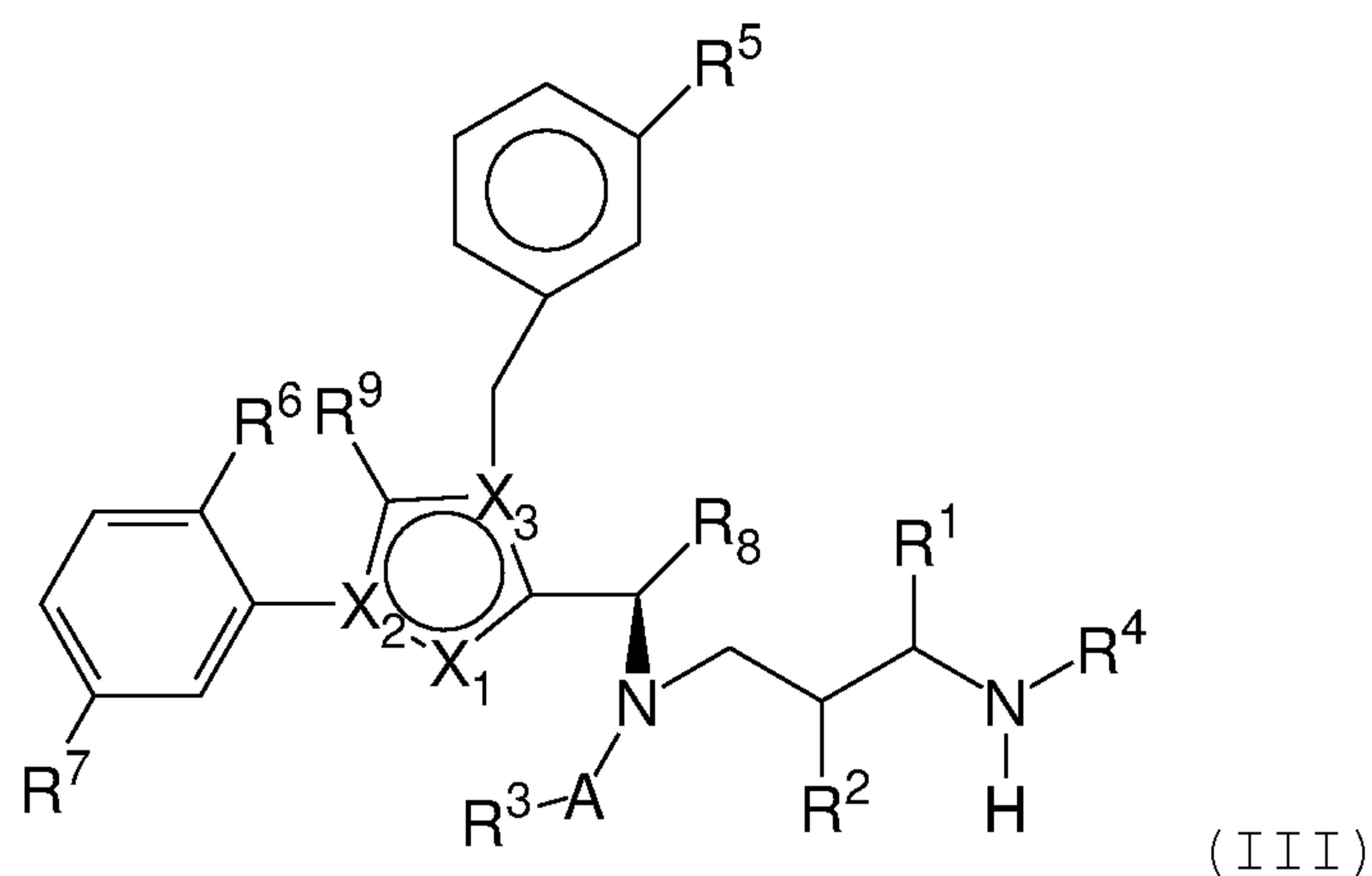
and the salts, solvates and salts of the solvates thereof.

25

In the case of binder conjugates of the KSP inhibitors of the formula (IIIa), at most one representative of R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>8</sup> and R<sup>10</sup> (alternatively to one of the conditions given above) may represent -L-BINDER, where L represents a linker and BINDER  
 30 represents a binder or a derivative thereof, where the binder may optionally be attached to a plurality of active compound molecules.

Formula (III):

35



where

- 5  $X_1$  represent N,  $X_2$  represents N and  $X_3$  represents C, or  $X_1$  represents CH,  $X_2$  represents C and  $X_3$  represents N or  $X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C, or  $X_1$  represents CH,  $X_2$  represents N and  $X_3$  represents C;
- 10  $R_1$  represents -L-BINDER, H or  $-(CH_2)_{0-3}Z$ , where Z represents -H,  $-NHY^3$ ,  $-OY^3$ ,  $-SY^3$ , halogen,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  or  $-CH(CH_2W)Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $NH_2$ ,  $SO_3H$ ,  $COOH$ ,  $-NH-CO-CH_2-CH_2-CH(NH_2)COOH$  or  $-(CO-NH-CHY^4)_{1-3}COOH$ ; where W represents H or OH;

15

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ ;

20

$R^2$  and  $R^4$  independently of one another represent -L-BINDER, H,  $-CO-CHY^4-NHY^5$  or  $-(CH_2)_{0-3}Z$ , or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring) represent  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$  represents L-#1, H,  $NH_2$ ,  $SO_3H$ ,  $COOH$ , SH or OH,

25

where Z represents -H, halogen,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

30



where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ;

5

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ , and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where  
10  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

A represents CO, SO,  $SO_2$ ,  $SO_2NH$  or  $CNNH$ ;

$R^3$  represents  $-L$ -BINDER or an optionally substituted alkyl, aryl,  
15 heteroaryl, heteroalkyl, heterocycloalkyl group, preferably  $-L$ -  
#1, or a  $C_{1-10}$ -alkyl,  $C_{6-10}$ -aryl or  $C_{6-10}$ -aralkyl,  $C_{5-10}$ -heteroalkyl,  
 $C_{1-10}$ -alkyl- $O$ - $C_{6-10}$ -aryl or  $C_{5-10}$ -heterocycloalkyl group which may  
be substituted by 1-3  $-OH$  groups, 1-3 halogen atoms, 1-3  
halogenated alkyl groups (each having 1-3 halogen atoms), 1-3  
20  $O$ -alkyl groups, 1-3  $-SH$  groups, 1-3  $-S$ -alkyl groups, 1-3  $-O-CO$ -  
alkyl groups, 1-3  $-O-CO-NH$ -alkyl groups, 1-3  $-NH-CO$ -alkyl  
groups, 1-3  $-NH-CO-NH$ -alkyl groups, 1-3  $-S(O)_n$ -alkyl groups, 1-  
3  $-SO_2-NH$ -alkyl groups, 1-3  $-NH$ -alkyl groups, 1-3  $-N(alkyl)_2$   
groups, 1-3  $-NH_2$  groups or 1-3  $-(CH_2)_{0-3}Z$  groups, where  $Z$   
25 represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,  
where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  
 $-(CH_2)_{0-3}Z'$  and  $Y^3$  represents H,  $-(CH_2)_{0-3}-CH(NHCOCH_3)Z'$ ,  $-(CH_2)_{0-3}-$   
 $CH(NH_2)Z'$  or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  
 $COOH$ ,

30

(where "alkyl" preferably represents  $C_{1-10}$ -alkyl);

$R^5$  represents  $-L$ -BINDER, H, F,  $NH_2$ ,  $NO_2$ , halogen, SH or  $-(CH_2)_{0-3}Z$ , where  $Z$  represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$   
35 or  $-CO-OY^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$

represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where L represents a linker and BINDER represents a binder or a derivative thereof, where the binder may optionally be attached  
5 to a plurality of active compound molecules,

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano, (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy or  
10 halogen,

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl or optionally substituted oxetane;  
and

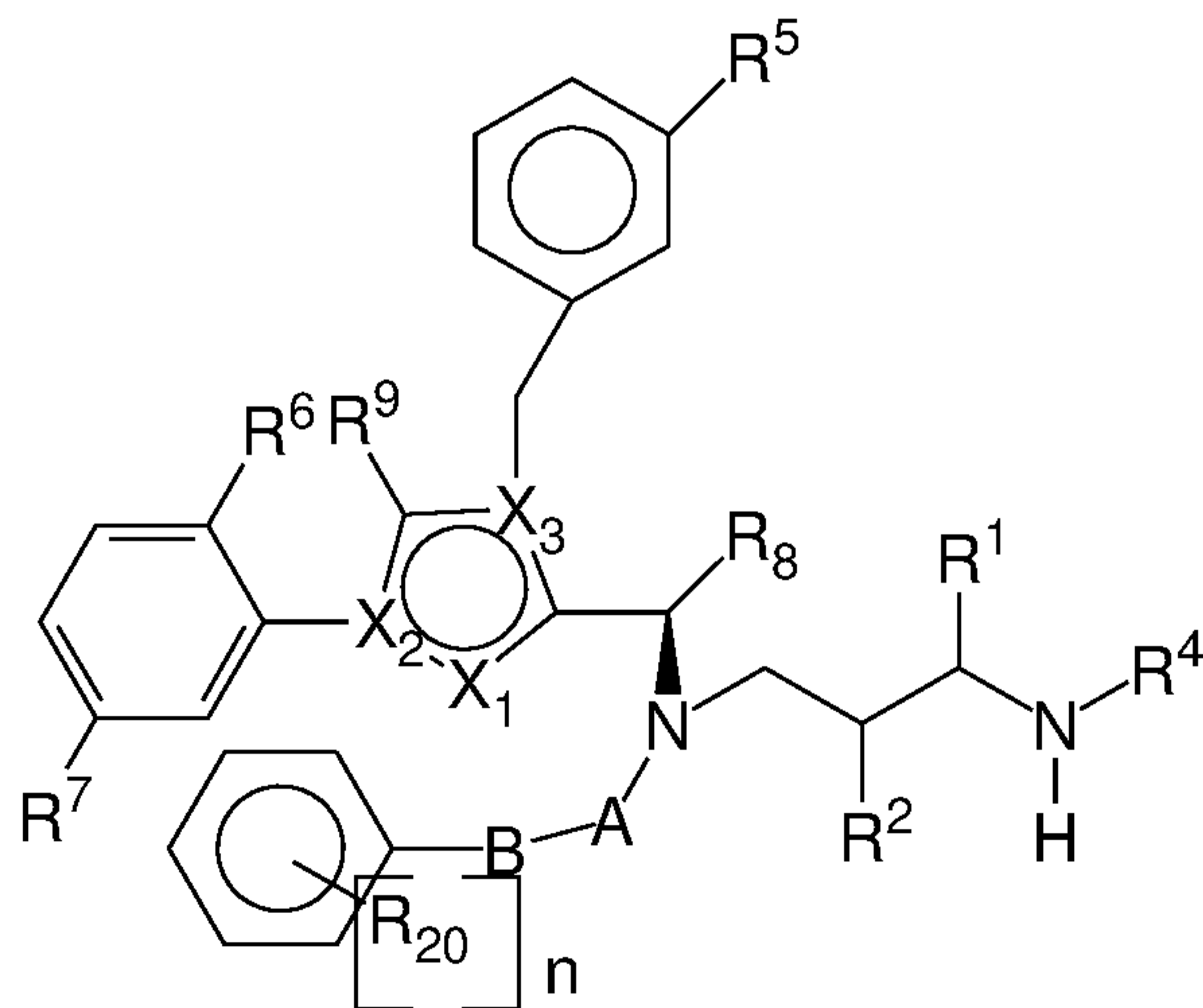
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R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

and the salts, solvates and salts of the solvates thereof.

20 Furthermore, preference according to the invention is given to the following KSP inhibitors and their binder conjugates:

Formula (IIIb):

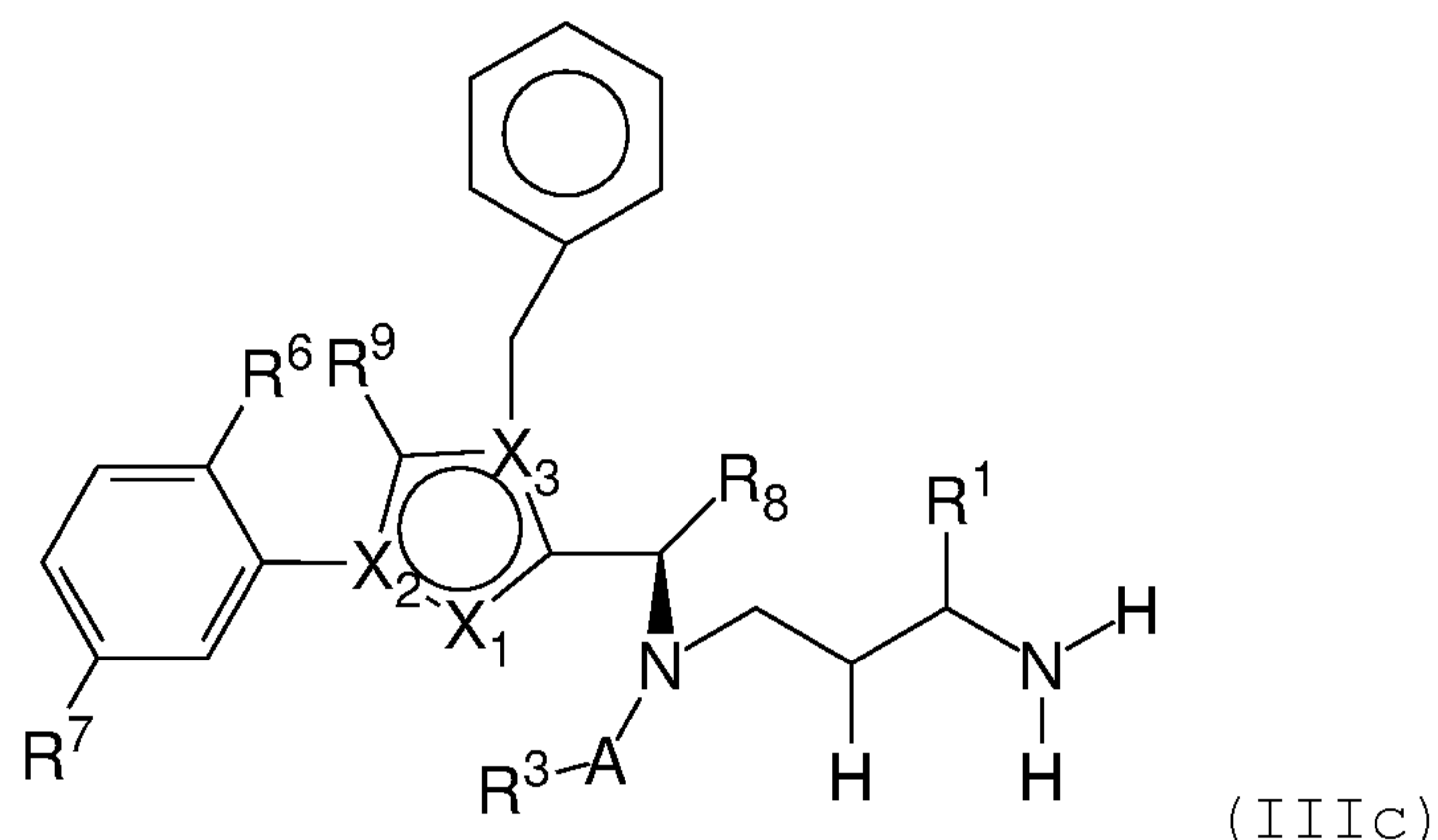


(IIIb)

where X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub> have the same meaning as in formula (IIIa) or (III) (where preferably X<sub>1</sub> represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represent N), R<sup>1</sup>, R<sup>2</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>6</sup>, R<sup>7</sup>, R<sup>8</sup> and R<sup>9</sup> have the same

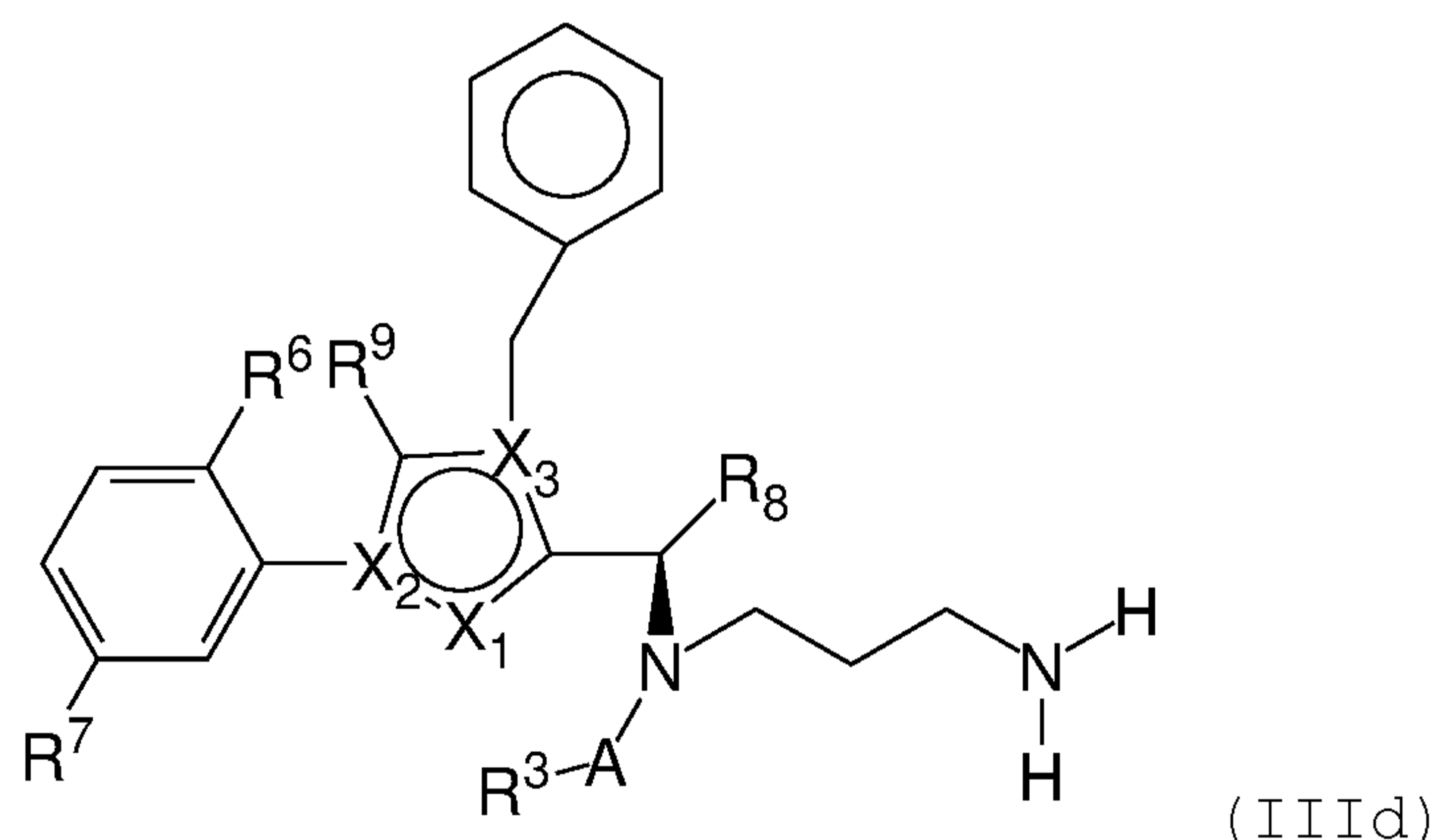
meaning as in formula (IIIa) or (III), A represents CO, B represents a single bond,  $-O-CH_2-$  or  $-CH_2-O-$  and  $R^{20}$  represents  $NH_2$ , F,  $CF_3$  or  $CH_3$  and n represents 0, 1 or 2.

5 Formula (IIIc):



10 where  $X_1$ ,  $X_2$ ,  $X_3$  have the same meaning as in formula (IIIa) or (III) (where preferably  $X_1$  represents CH,  $X_2$  represents C and  $X_3$  represents N), A,  $R^1$ ,  $R^3$ ,  $R^6$ ,  $R^7$ ,  $R^8$  and  $R^9$  have the same meaning as in formula (IIIa) or (III), A preferably represents CO and  $R^3$  represents  $-CH_2OH$ ,  $-CH_2OCH_3$ ,  $CH(CH_3)OH$  or  $CH(CH_3)OCH_3$ .

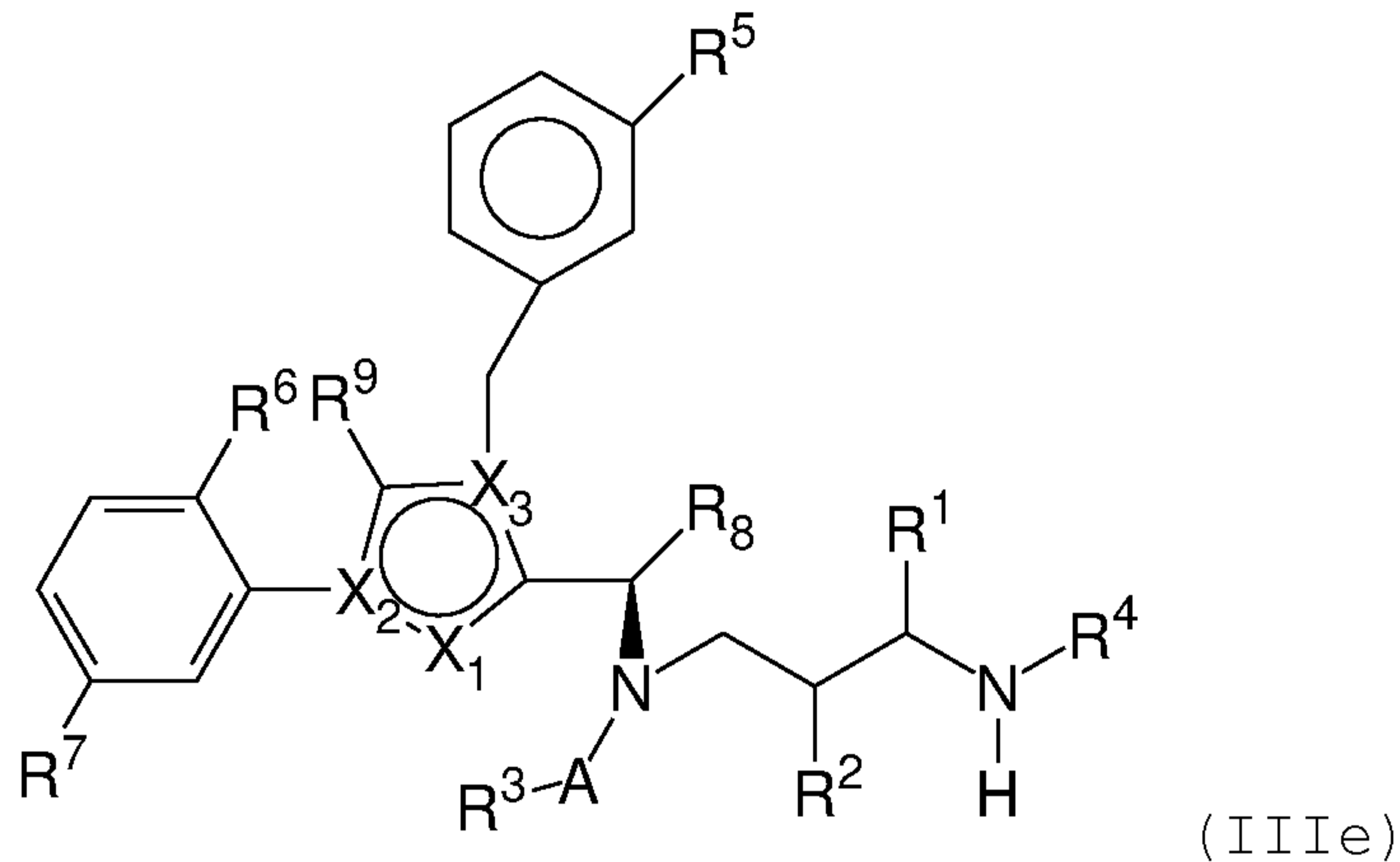
15 Formula (IIIId):



20 where  $X_1$ ,  $X_2$ ,  $X_3$  have the same meaning as in formula (IIIa) or (III) (where preferably  $X_1$  represents CH,  $X_2$  represents C and  $X_3$  represents N), A,  $R^3$ ,  $R^6$ ,  $R^7$ ,  $R^8$  and  $R^9$  have the same meaning as in formula (IIIa) or (III), where A preferably represents CO and  $R^3$  represents  $-CH_2-S_x-(CH_2)_{0-4}-CHY^5-COOH$ , where x is 0 or 1 and  $Y^5$

represents H or  $\text{NH}Y^6$ , where  $Y^6$  represents H or  $-\text{COCH}_3$ .

Formula (IIIe):



where  $X_1$  represents CH,  $X_2$  represents C and  $X_3$  represents N, A,  $R^3$ ,  $R^6$ ,  $R^7$ ,  $R^8$  and  $R^9$  have the same meaning as in formula (IIIa) or (III) and  $R^1$  represents -L-BINDER.

10

Furthermore, it is preferred when in the compounds of the formulae (III), (IIIa), (IIIb), (IIIc), (IIId) and (IIIe) (alone or in combination):

15

- Z represents Cl or Br;
- $R^1$  represents  $-(\text{CH}_2)_{0-3}Z$ , where Z represents  $-\text{CO}-\text{NY}^1\text{Y}^2$ , where  $Y^2$  represents  $-(\text{CH}_2\text{CH}_2\text{O})_{0-3}-\text{CH}_2\text{CH}_2Z'$  and  $Y^1$  represents H,  $\text{NH}_2$  or  $-(\text{CH}_2\text{CH}_2\text{O})_{0-3}-\text{CH}_2\text{CH}_2Z'$ ;

20

- $Y^1$  represents H,  $Y^2$  represents  $-(\text{CH}_2\text{CH}_2\text{O})_3-\text{CH}_2\text{CH}_2Z'$  and  $Z'$  represents  $-\text{COOH}$ ;

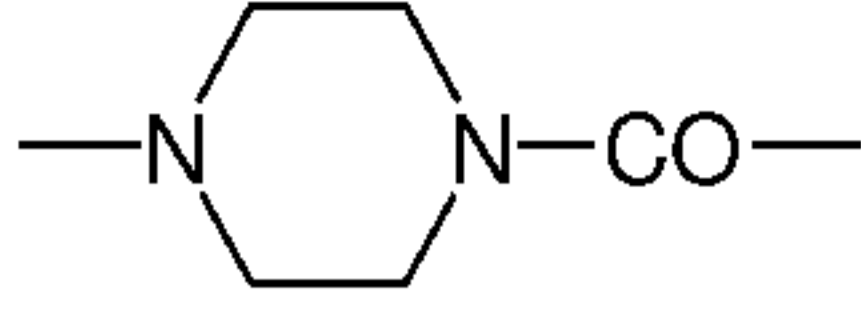
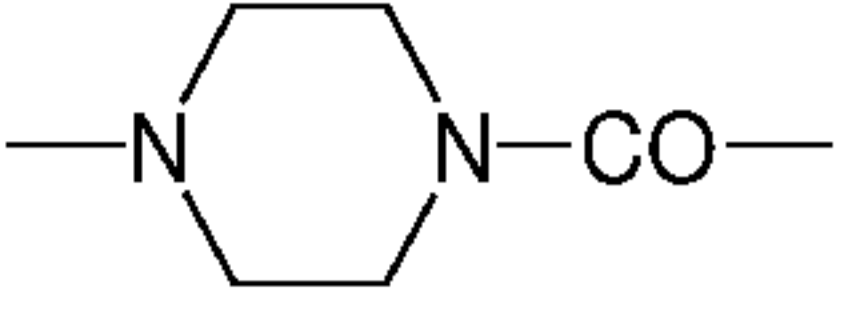
25

- $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2Z'$  and  $Z'$  represents  $-(\text{CONHCHY}^4)_2\text{COOH}$ ;

30

- $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2Z'$ ,  $Z'$  represents  $-(\text{CONHCHY}^4)_2\text{COOH}$  and one of the  $Y^4$  radicals represents *i*-propyl and the other  $-(\text{CH}_2)_3-\text{NHCONH}_2$ ;

- $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2\text{Z}'$ ,  $\text{Z}'$  represents  $-(\text{CONHCHY}^4)_2\text{COOH}$  and one of the  $Y^4$  radicals represents  $-\text{CH}_3$  and the other  $-(\text{CH}_2)_3-\text{NHCONH}_2$ ;
- 5 ●  $Y^4$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl which is optionally substituted by  $-\text{NHCONH}_2$ ;
- at least one  $Y^4$  representative is selected from the group consisting of *i*-propyl and  $-\text{CH}_3$ ;
- 10 ●  $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2\text{Z}'$ ,  $\text{Z}'$  represents  $-\text{CONHCHY}^4\text{COOH}$  and  $Y^4$  represents aryl or benzyl which are optionally substituted by  $-\text{NH}_2$ ;
- 15 ●  $Y^4$  represents aminobenzyl;
- $\text{R}^2$  represents  $-(\text{CH}_2)_{0-3}\text{Z}$  and Z represents  $-\text{SY}^3$ ;
  - $\text{R}^4$  represents  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  and  $Y^5$  represents H;
- 20 ●  $\text{R}^4$  represents  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  and  $Y^5$  represents  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ ;
- $Y^4$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl which is optionally substituted by  $-\text{NHCONH}_2$ .
- 25 Furthermore, it is preferred when  $\text{R}^1$ ,  $\text{R}^2$  or  $\text{R}^3$  in formula (IIa) or (IIIa) represents  $-\text{MOD}$ , in particular when  $\text{R}^4$  represents  $-\text{L}-\#1$  or  $-\text{L}-\text{BINDER}$  (in particular when  $-\text{L}$  is a cleavable linker which cleaves directly at  $-\text{N}-\text{R}^4$  or  $-\text{N}-\text{L}-\#1$  or  $-\text{L}-\text{BINDER}$ , such
- 30 that  $\text{R}^4$  or L is replaced by H).
- Particularly preferably,  $\text{R}^3$  represents  $-\text{MOD}$  and  $\text{R}^1$  or  $\text{R}^4$  represents  $-\text{L}-\#1$  or  $-\text{L}-\text{BINDER}$ ,
- 35 where  $-\text{MOD}$  represents  $-(\text{NR}^{10})_n-(\text{G1})_o-\text{G2}-\text{H}$ , where
- $\text{R}^{10}$  represents H or  $\text{C}_1-\text{C}_3$ -alkyl;

G1 represents  $\text{-NHCO-}$ ,  $\text{-CONH-}$  or  (where, if G1 represents  $\text{-NHCO-}$  or ,  $\text{R}^{10}$  does not represent  $\text{NH}_2$ );

n is 0 or 1;

5

o is 0 or 1; and

G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups  $\text{-O-}$ ,  $\text{-S-}$ ,  $\text{-SO-}$ ,  $\text{SO}_2$ ,  $\text{-NR}^y\text{-}$ ,  $\text{-NR}^y\text{CO-}$ ,  $\text{CONR}^y\text{-}$ ,  $\text{-NR}^y\text{NR}^y\text{-}$ ,  $\text{-SO}_2\text{NR}^y\text{NR}^y\text{-}$ ,  $\text{-CONR}^y\text{NR}^y\text{-}$  (where  $\text{R}^y$  represents H, phenyl,  $\text{C}_1\text{-C}_{10}$ -alkyl,  $\text{C}_2\text{-C}_{10}$ -alkenyl or  $\text{C}_2\text{-C}_{10}$ -alkynyl, each of which may be substituted by  $\text{-NHCONH}_2$ ,  $\text{-COOH}$ ,  $\text{-OH}$ ,  $\text{-NH}_2$ ,  $\text{NH-CNNH}_2$ , sulphonamide, sulphone, sulphoxide or sulphonic acid),  $\text{-CO-}$ ,  $\text{-CR}^x\text{=N-O-}$  (where  $\text{R}^x$  represents H,  $\text{C}_1\text{-C}_3$ -alkyl or phenyl), where the hydrocarbon chain including any side chains may be substituted by  $\text{-NHCONH}_2$ ,  $\text{-COOH}$ ,  $\text{-OH}$ ,  $\text{-NH}_2$ ,  $\text{NH-CNNH}_2$ , sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group  $\text{-MOD}$  preferably has at least one group  $\text{-COOH}$ ;

20

Particularly preferably, the group  $\text{-MOD}$  has a (preferably terminal)  $\text{-COOH}$  group, for example in a betaine group. Preferably, the group  $\text{-MOD}$  has the formula  $\text{-CH}_2\text{-S}_x\text{-(CH}_2\text{)}_{0-4}\text{-CHY}^5\text{-COOH}$  where x is 0 or 1, and  $\text{Y}^5$  represents H or  $\text{NHY}^6$ , where  $\text{Y}^6$  represents H or  $\text{-COCH}_3$ .

25

Furthermore, it is preferred when in the formula (IIa), (II), (III), (IIIa), (IIIb), (IIIc), (IIId) or (IIIe) (alone or in combination):

30

- Z represents Cl or Br;
- $\text{R}^1$  represents  $\text{-(CH}_2\text{)}_{0-3}\text{Z}$ , where Z represents  $\text{-CO-NY}^1\text{Y}^2$ , where  $\text{Y}^2$  represents  $\text{-(CH}_2\text{CH}_2\text{O)}_{0-3}\text{-(CH}_2\text{)}_{0-3}\text{Z}'$  and  $\text{Y}^1$  represents H,  $\text{NH}_2$  or  $\text{-(CH}_2\text{CH}_2\text{O)}_{0-3}\text{-(CH}_2\text{)}_{0-3}\text{Z}'$ ;
- $\text{Y}^1$  represents H,  $\text{Y}^2$  represents  $\text{-(CH}_2\text{CH}_2\text{O)}_3\text{-CH}_2\text{CH}_2\text{Z}'$  and  $\text{Z}'$

35

represents  $-\text{COOH}$ ;

- $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2\text{Z}'$  and  $\text{Z}'$  represents  $-(\text{CONHCHY}^4)_2\text{COOH}$ ;

5

- $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2\text{Z}'$ ,  $\text{Z}'$  represents  $-(\text{CONHCHY}^4)_2\text{COOH}$  and one  $Y^4$  representative represents *i*-propyl and the other represents  $-(\text{CH}_2)_3-\text{NHCONH}_2$ ;

10

- $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2\text{Z}'$ ,  $\text{Z}'$  represents  $-(\text{CONHCHY}^4)_2\text{COOH}$  and one  $Y^4$  representative represents  $-\text{CH}_3$  and the other represents  $-(\text{CH}_2)_3-\text{NHCONH}_2$ ;

15

- $Y^4$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl which is optionally substituted by  $-\text{NHCONH}_2$ ;

- at least one  $Y^4$  representative is selected from the group consisting of *i*-propyl and  $-\text{CH}_3$ ;

20

- $Y^1$  represents H,  $Y^2$  represents  $-\text{CH}_2\text{CH}_2\text{Z}'$ ,  $\text{Z}'$  represents  $-\text{CONHCHY}^4\text{COOH}$  and  $Y^4$  represents aryl or benzyl which are optionally substituted by  $-\text{NH}_2$ ;

- $Y^4$  represents aminobenzyl;

25

- $\text{R}^2$  represents  $-(\text{CH}_2)_{0-3}\text{Z}$  and  $\text{Z}$  represents  $-\text{SY}^3$ ;

- $\text{R}^4$  represents  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  and  $Y^5$  represents H;

30

- $\text{R}^4$  represents  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  and  $Y^5$  represents  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ ;

- $Y^4$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl which is optionally substituted by  $-\text{NHCONH}_2$ .

35

Preference is furthermore given to compounds of the formula (IIa), (II), (III) or (IIIa)

where

X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C; or

X<sub>1</sub> represents CH or CF, X<sub>2</sub> represents C and X<sub>3</sub> represents N; or

5

X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C; or

X<sub>1</sub> represents CH or CF, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

10 (with X<sub>1</sub> representing CH, X<sub>2</sub> representing C and X<sub>3</sub> representing N being preferred);

R<sup>1</sup> represents H, -L-#1 or -L-BINDER, -MOD or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

15

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' (e.g. -(CH<sub>2</sub>)<sub>0-3</sub>Z') or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup> represents H or -(CH<sub>3</sub>)<sub>0-3</sub>Z', where Z' represents H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where W

20 represents H or OH,

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>;

25

R<sup>2</sup> represents H, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z,

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

30

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

35

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally



substituted by  $-\text{NH}_2$ , and  $\text{Y}^5$  represents H or  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ , where  $\text{Y}^6$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl;

$\text{R}^4$  represents H;

5

A represents CO, SO,  $\text{SO}_2$ ,  $\text{SO}_2\text{NH}$  or  $\text{CNNH}$ ;

$\text{R}^3$  represents -L-#1 or -L-BINDER, -MOD or an optionally substituted alkyl, cycloalkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably a  $\text{C}_{1-10}$ -alkyl,  $\text{C}_{6-10}$ -aryl or  $\text{C}_{6-10}$ -aralkyl,  $\text{C}_{5-10}$ -heteroalkyl,  $\text{C}_{1-10}$ -alkyl-O- $\text{C}_{6-10}$ -aryl or  $\text{C}_{5-10}$ -heterocycloalkyl group which may be substituted by 1-3 -OH groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH((CH<sub>2</sub>CH<sub>2</sub>O)<sub>1-20</sub>H) groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH (where "alkyl" is preferably  $\text{C}_{1-10}$ -alkyl);

25

$\text{R}^5$  represents H, -MOD, NH<sub>2</sub>, NO<sub>2</sub>, halogen (in particular F, Cl, Br), -CN, CF<sub>3</sub>, -OCF<sub>3</sub>, -CH<sub>2</sub>F, -CH<sub>2</sub>F, SH or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

30 where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

$\text{R}^6$  and  $\text{R}^7$  independently of one another represent H, cyano, 35 (optionally fluorinated)  $\text{C}_{1-10}$ -alkyl, (optionally fluorinated)  $\text{C}_{2-10}$ -alkenyl, (optionally fluorinated)  $\text{C}_{2-10}$ -alkynyl, hydroxy, NO<sub>2</sub>, NH<sub>2</sub>, COOH or halogen (in particular F, Cl, Br),

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl or (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl;

5 where one or none of the substituents R<sup>1</sup> and R<sup>3</sup> represents -L-#1 or -L-BINDER,

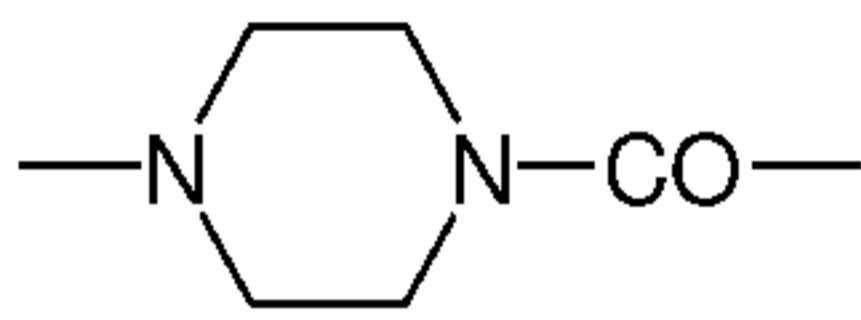
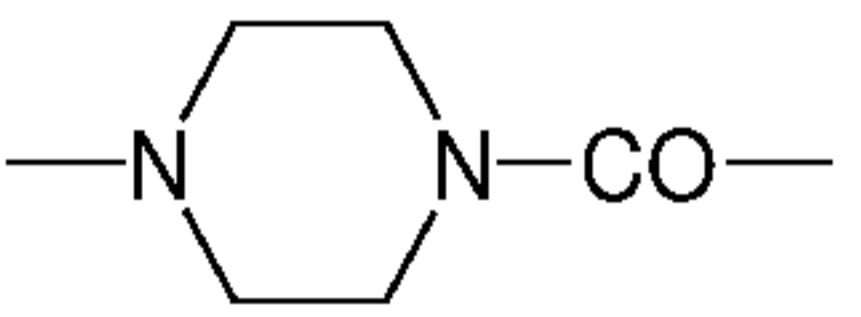
L represents the linker and #1 represents the bond to the binder or derivative thereof and BINDER represents the binder,

10

R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

15 R<sup>10</sup> represents H or C<sub>1-3</sub>-alkyl;

G1 represents -NHCO- , -CONH- or  (where, if G1 represents -NHCO- or , R<sup>10</sup> does not represent NH<sub>2</sub>);

20 n is 0 or 1;

o is 0 or 1; and

G2 represents a straight-chain and/or branched hydrocarbon group  
 25 which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>- (where R<sup>y</sup> represents H, phenyl, C<sub>1-10</sub>-alkyl, C<sub>2-10</sub>-alkenyl or C<sub>2-10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or  
 30 sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1-3</sub>-alkyl or phenyl), where the hydrocarbon chain including any side chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid,  
 35 where the group -MOD preferably has at least one group -COOH;

and the salts, solvates and salts of the solvates thereof.

Preference is furthermore given to compounds of the formula (IIa), (II), (III) or (IIIa) in which

5 X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C; or

X<sub>1</sub> represents CH or CF, X<sub>2</sub> represents C and X<sub>3</sub> represents N; or

X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C; or

10

X<sub>1</sub> represents CH or CF, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

(with X<sub>1</sub> representing CH, X<sub>2</sub> representing C and X<sub>3</sub> representing N being preferred);

15

R<sup>1</sup> represents H, -L-#1 or -L-BINDER, -MOD or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

20 where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' (e.g. -(CH<sub>2</sub>)<sub>0-3</sub>Z') or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup> represents H or -(CH<sub>3</sub>)<sub>0-3</sub>Z', where Z' represents H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where W represents H or OH,

25 where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>;

30 R<sup>2</sup> represents H, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z,

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

35 where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ , and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where  
 5  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

$R^4$  represents H;

A represents CO, SO,  $SO_2$ ,  $SO_2NH$  or CNNH;

10

$R^3$  represents  $-L-#1$  or  $-L-BINDER$ ,  $-MOD$  or an optionally substituted alkyl, cycloalkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably a  $C_{1-10}$ -alkyl,  $C_{6-10}$ -aryl or  $C_{6-10}$ -aralkyl,  $C_{5-10}$ -heteroalkyl,  $C_{1-10}$ -alkyl- $O$ - $C_{6-10}$ -aryl or  $C_{5-10}$ -heterocycloalkyl group which may be substituted by 1-3  $-OH$  groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3  $O$ -alkyl groups, 1-3  $-SH$  groups, 1-3  $-S$ -alkyl groups, 1-3  $-O-CO$ -alkyl groups, 1-3  $-O-CO-NH$ -alkyl groups, 1-3  $-NH-CO$ -alkyl groups, 1-3  $-NH-CO-NH$ -alkyl groups, 1-3  $-S(O)_n$ -alkyl groups, 1-3  $-SO_2-NH$ -alkyl groups, 1-3  $-NH$ -alkyl groups, 1-3  $-N(alkyl)_2$  groups, 1-3  $-NH((CH_2CH_2O)_{1-20}H)$  groups, 1-3  $-NH_2$  groups or 1-3  $-(CH_2)_{0-3}Z$  groups, where Z represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ , where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$  and  
 15  
 20  
 25  $Y^3$  represents H,  $-(CH_2)_{0-3}-CH(NHCOCH_3)Z'$ ,  $-(CH_2)_{0-3}-CH(NH_2)Z'$  or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$  (where "alkyl" is preferably  $C_{1-10}$ -alkyl);

$R^5$  represents H,  $-MOD$ ,  $NH_2$ ,  $NO_2$ , halogen (in particular F, Cl, Br),  $-CN$ ,  $CF_3$ ,  $-OCF_3$ ,  $-CH_2F$ ,  $-CH_2F$ , SH or  $-(CH_2)_{0-3}Z$ , where Z represents  $-H$ ,  $-OY^3$ ,  $-SY^3$ , halogen,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,  
 30

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$   
 35 represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ;

$R^6$  and  $R^7$  independently of one another represent H or halogen (in particular F, Cl, Br),

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl;

where one or none of the substituents R<sup>1</sup> and R<sup>3</sup> represents -L-#1  
5 or -L-BINDER,

L represents the linker and #1 represents the bond to the binder  
or derivative thereof and BINDER represents the binder,

10 R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

where -MOD represents -CH<sub>2</sub>-S<sub>x</sub>-(CH<sub>2</sub>)<sub>0-4</sub>-CHY<sup>5</sup>-COOH where x is 0 or  
1, and Y<sup>5</sup> represents H or NHY<sup>6</sup>, where Y<sup>6</sup> represents H or -COCH<sub>3</sub>,

15 and the salts, solvates and salts of the solvates thereof.

Preference is furthermore given to the following compounds which  
may optionally be present together with an acid such as, for  
example, trifluoroacetic acid. These compounds may be attached  
20 via the positions corresponding to the positions R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>,  
R<sup>5</sup>, R<sup>8</sup> and R<sup>10</sup>, in particular R<sup>1</sup> and R<sup>3</sup>, via a linker to a binder  
(where a hydrogen atom is substituted by the linker):

N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
25 1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide;

N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide;

30 (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-  
pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]-N-  
methylbutanamide (1:1);

N-(3-aminopropyl)-N-{(1S)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
35 1H-pyrrol-2-yl]-2,2-dimethylpropyl}acetamide;

N-(3-aminopropyl)-N-{(1S)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide;

S-[1-(2-{[2-({(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-oxoethyl)-2,5-dioxopyrrolidin-3-yl]-L-cysteine;

N-{(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl-L-alanyl-N-[4-(3-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-2,5-dioxopyrrolidin-1-yl)phenyl]-N<sup>5</sup>-carbamoyl-L-ornithinamide;

S-(1-{2-[(N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)amino]ethyl}-2,5-dioxopyrrolidin-3-yl)-L-cysteine;

S-[1-(2-{[2-({(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-oxoethyl)-2,5-dioxopyrrolidin-3-yl]-L-cysteine;

N-{(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl-L-alanyl-N-[4-(3-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-2,5-dioxopyrrolidin-1-yl)phenyl]-N<sup>5</sup>-carbamoyl-L-ornithinamide;

S-(1-{2-[(N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)amino]ethyl}-2,5-dioxopyrrolidin-3-yl)-L-cysteine;

N-[6-(3-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-2,5-dioxopyrrolidin-1-yl)hexanoyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithyl-N<sup>6</sup>-{(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-lysine;

S-[1-(2-{[2-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl)amino}-2-oxoethyl)-2,5-dioxopyrrolidin-3-yl]-L-cysteine;

S-(2-{[2-({(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl)amino}-2-oxoethyl)-L-cysteine;

S-{1-[6-(2-{(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}hydrazino)-6-oxohexyl]-2,5-dioxopyrrolidin-3-yl}-L-cysteine;

N-[19-(3(R/S)-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-2,5-dioxopyrrolidin-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-R/S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}homocysteine;

S-{(3R/S)-1-[2-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]-2,5-dioxopyrrolidin-3-yl}-L-cysteine;

N-[19-(3(R/S)-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-2,5-dioxopyrrolidin-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-R/S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-imidazol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}homocysteine;

S-[(3R/S)-1-(2-{[6-({2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulphonyl)hexanoyl]amino)ethyl)-2,5-dioxopyrrolidin-3-yl]-L-cysteine;

S-{1-[2-({[(1R,3S)-3-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)cyclopentyl]carbonyl}amino)ethyl]-2,5-dioxopyrrolidin-3-yl}-L-cysteine;

S-(2-{[2-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-oxoethyl)-L-cysteine;

N<sup>6</sup>-(N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)-N<sup>2</sup>-{N-[6-(3-[(2R)-2-amino-2-carboxyethyl]sulphonyl)-2,5-dioxopyrrolidin-1-yl]hexanoyl}-L-valyl-L-alanyl}-L-lysine;

N-[2-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]-L-glutamine;

N<sup>6</sup>-(N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)-L-lysine;

N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}acetamide;

N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-methoxyacetamide;

N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2,4-difluorobenzamide;

N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-4-methylbenzamide;



N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamide;

5 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-3,3,3-trifluoropropanamide;

10 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-fluorobenzamide;

N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}acetamide;

15 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-(trifluoromethyl)benzamide;

20 N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamide;

N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamide;

25 (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl] (glycoloyl) amino]butanoic acid;

30 (2S)-2-amino-N-(2-aminoethyl)-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl] (glycoloyl) amino]butanamide;

35 4-[(2-[[2-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl] (glycoloyl) amino]butanoyl] amino) ethyl] amino]-2-oxoethyl) amino]-3-[(2R)-2-amino-2-carboxyethyl]sulphonyl]-4-oxobutanoic acid;

4-[(2-{{2-({(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-oxoethyl)amino]-2-{{(2R)-2-amino-2-carboxyethyl}sulphanyl}-4-oxobutanoic acid;

N-{{(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanine;

10

N-{{(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-serine;

15

N-{{(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-alanine;

20

N-{{(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}glycine;

25

N-(3-aminopropyl)-N-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-methylbenzamide;

N-(3-aminopropyl)-N-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-(methylsulphanyl)benzamide;

30

(2S)-N-(3-aminopropyl)-N-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxypropanamide;

35

N-(3-aminopropyl)-N-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-(methylsulphanyl)acetamide;

(2S)-N-(3-aminopropyl)-N-{{(1R)-1-[4-benzyl-1-(2,5-

difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-hydroxypropanamide;

5 methyl 4-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-4-oxobutanoate;

10 4-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-4-oxobutanoic acid;

15 (2R)-22-[(3R/S)-3-[[2-[(2R)-2-amino-2-carboxyethyl]sulphanyl]-2,5-dioxopyrrolidin-1-yl]-2-[(2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl)sulphanyl)methyl]-4,20-dioxo-7,10,13,16-tetraoxa-3,19-diazadocosan-1-oic acid;

4-amino-N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}benzamide;

20 N-acetyl-S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-L-cysteine;

25 N-acetyl-S-[2-([3-(L-alanyl-amino)propyl]{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)-2-oxoethyl]-L-cysteine;

30 (2S)-N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}tetrahydrofuran-2-carboxamide;

3-[(2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl)sulphanyl]propanoic acid;

35 S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}homocysteine;

4-amino-N-(3-aminopropyl)-N-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]benzamide;

5 4-[(2-[[[(2R)-2-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]amino)-2-carboxyethyl]amino]-2-oxoethyl)amino]-3-[[[(2R)-2-amino-2-carboxyethyl]sulphonyl]-4-oxobutanoic acid];

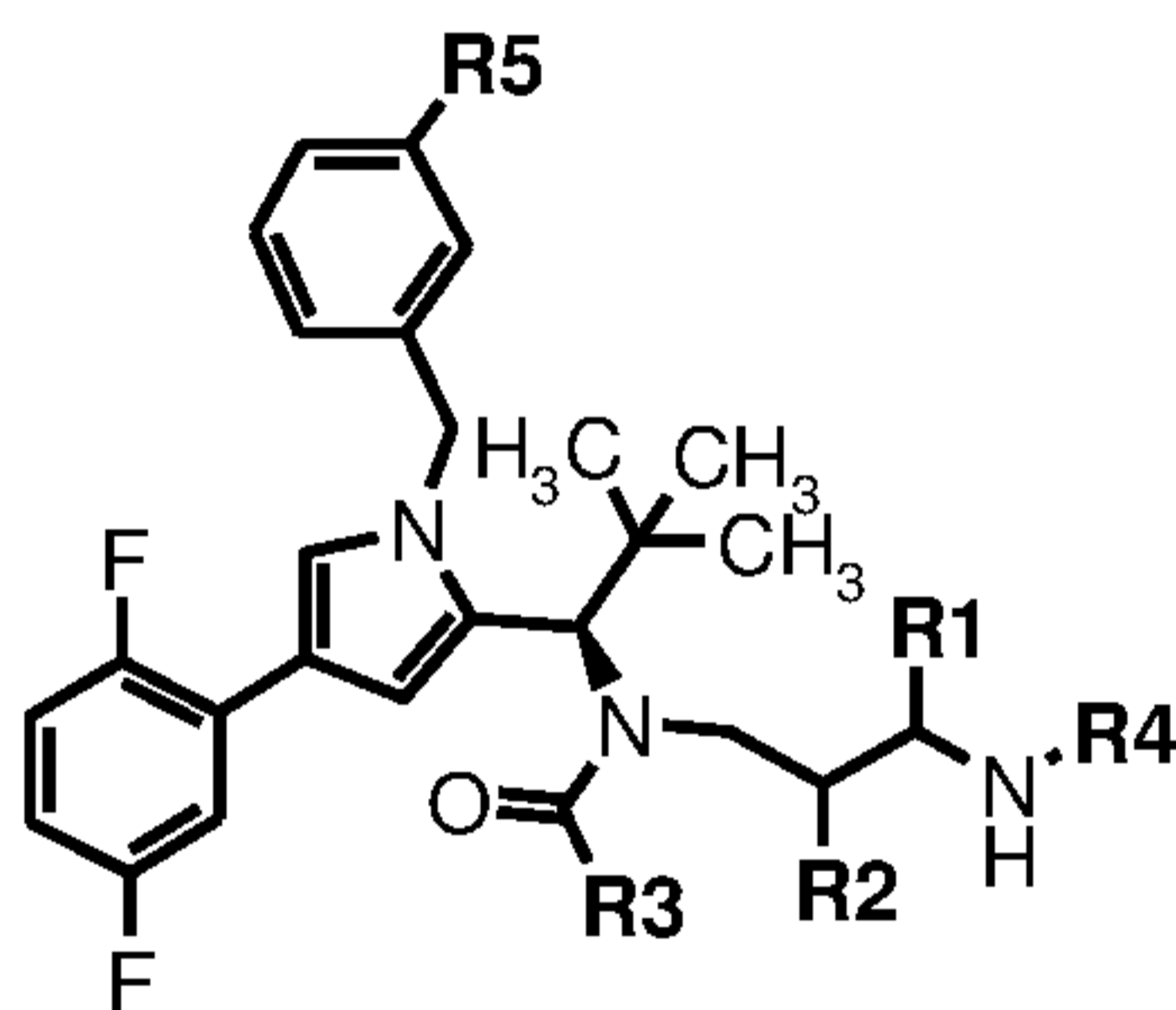
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4-[(2-[[[(2R)-2-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]amino)-2-carboxyethyl]amino]-2-oxoethyl)amino]-2-[[[(2R)-2-amino-2-carboxyethyl]sulphonyl]-4-oxobutanoic acid].

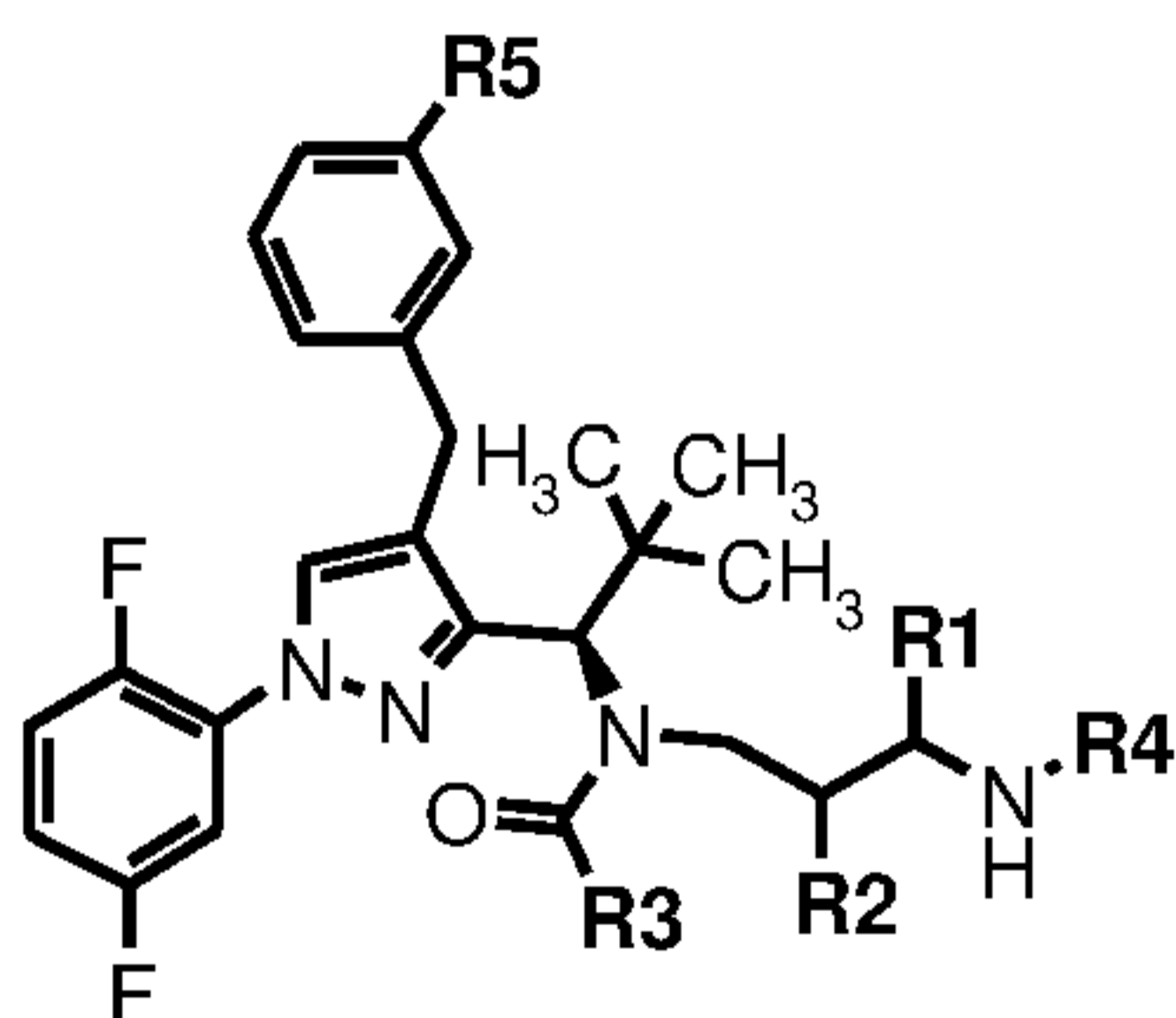
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Particular preference according to the invention is given to the following compounds of the formulae V, VI and VII, where R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup> and R<sup>5</sup> have the meanings mentioned above (as mentioned, for example for formula (IIa) or (IIIa)):

20



Formula VI



Formula VII

25

Particular preference is given to the compounds of the formulae

V, VI, VII where  $R^1$  and  $R^5$  represent H or -L-#1;  $R^2$  and  $R^4$  independently of one another represent -L-#1 or H or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-,  $R^{10}$  represents H or -L-#1; and  $R^3$  represents  
5 CH<sub>2</sub>OH, CH(CH<sub>3</sub>)OH or -L-#1, where one of the substituents  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$  and  $R^{10}$  represents -L-#1. Especially preferred are the corresponding compounds of the formula VI.

### Linkers

10

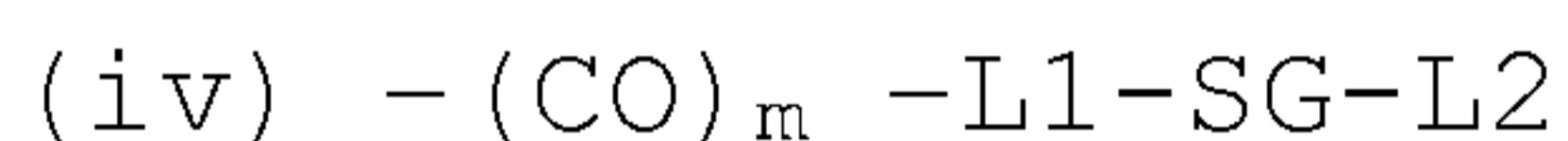
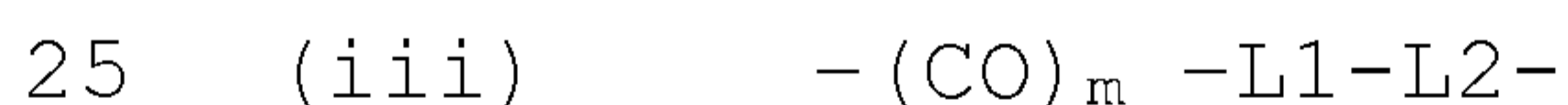
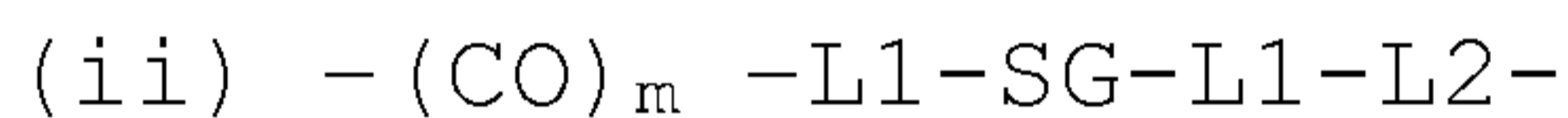
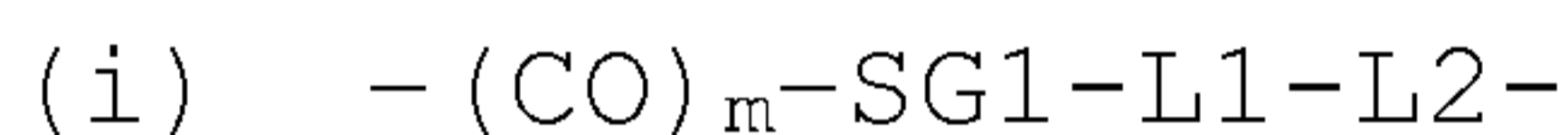
The literature discloses various options for covalently coupling (conjugating) organic molecules to binders such as, for example antibodies (see, for example, K. Lang and J. W. Chin. *Chem. Rev.* **2014**, *114*, 4764-4806, M. Rashidian et al. *Bioconjugate Chem.* **2013**, *24*, 1277-1294). Preference according to the invention is  
15 given to conjugation of the KSP inhibitors to an antibody via one or more sulphur atoms of cysteine residues of the antibody which are either already present as free thiols or generated by reduction of disulphide bridges, and/or via one or more NH groups  
20 of lysine residues of the antibody. However, it is also possible to attach the KSP inhibitor to the antibody via tyrosine residues, via glutamine residues, via residues of unnatural amino acids, via free carboxyl groups or via sugar residues of the antibody. For coupling, use is made of linkers. Linkers can  
25 be categorized into the group of the linkers which can be cleaved *in vivo* and the group of the linkers which are stable *in vivo* (see L. Ducry and B. Stump, *Bioconjugate Chem.* 21, 5-13 (2010)). The linkers which can be cleaved *in vivo* have a group which can be cleaved *in vivo*, where, in turn, a distinction may be made  
30 between groups which are chemically cleavable *in vivo* and groups which are enzymatically cleavable *in vivo*. "Chemically cleavable *in vivo*" and "enzymatically cleavable *in vivo*" means that the linkers or groups are stable in circulation and are cleaved only at or in the target cell by the chemically or enzymatically  
35 different environment therein (lower pH; elevated glutathione concentration; presence of lysosomal enzymes such as cathepsin or plasmin, or glycosidases such as, for example,  $\beta$ -glucuronidases), thus releasing the low-molecular weight KSP

inhibitor or a derivative thereof. Groups which can be cleaved chemically *in vivo* are in particular disulphide, hydrazone, acetal and aminal; groups which can be cleaved enzymatically *in vivo* are in particular the 2-8-oligopeptide group, especially a dipeptide group or glycoside. Peptide cleavage sites are disclosed in *Bioconjugate Chem.* **2002**, *13*, 855-869, and *Bioorganic & Medicinal Chemistry Letters* **8** (**1998**) 3341-3346 and also *Bioconjugate Chem.* **1998**, *9*, 618-626. These include, for example, valine-alanine, valine-lysine, valine-citrulline, alanine-lysine and phenylalanine-lysine (optionally with additional amide group).

Linkers which are stable *in vivo* are distinguished by a high stability (less than 5% metabolites after 24 hours in plasma) and do not have the chemically or enzymatically *in vivo* cleavable groups mentioned above.

The linker -L- preferably has one of the basic structures (i) to (iv) below:

20



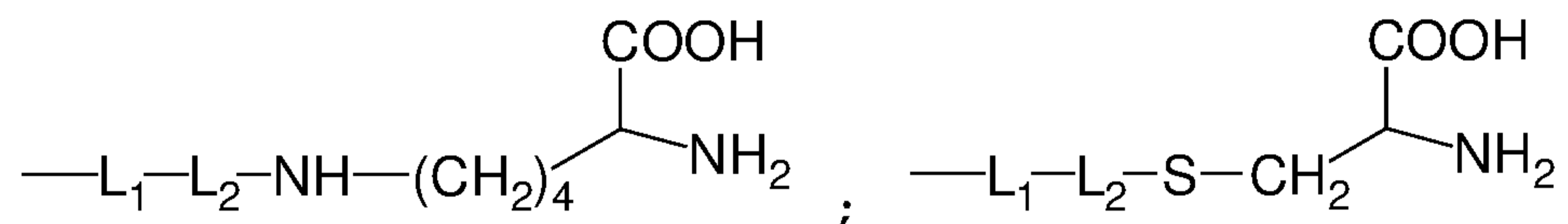
where m is 0 or 1; SG is a (chemically or enzymatically) *in vivo* cleavable group (in particular disulphide, hydrazone, acetal and aminal; or a 2-8-oligopeptide group which can be cleaved by cathepsin or plasmin), SG1 is an oligopeptide group or preferably a dipeptide group, L1 independently of one another represent *in vivo* stable organic groups, and L2 represents a coupling group to the binder or a single bond. Here, coupling is preferably to a cysteine residue or a lysine residue of the binder. Alternatively, coupling can be to a tyrosine residue, glutamine residue or to an unnatural amino acid of the binder.

35

The unnatural amino acids may contain, for example, aldehyde or keto groups (such as, for example, formylglycine) or azide or alkyne groups (see Lan & Chin, Cellular Incorporation of Unnatural Amino Acids and Bioorthogonal Labeling of Proteins, 5 Chem.Rev. 2014, 114, 4764-4806).

Particular preference according to the invention is given to the basic linker structure (iii), in particular when the binder is an anti-TWEAKR antibody or an anti-EGFR antibody. Via 10 metabolization, the administration of a conjugate according to the invention having a basic linker structure (iii) and coupling of the linker to a cysteine or lysine residue of the binder protein or peptide leads to cysteine or lysine derivatives of the formulae below:

15



where L1 is in each case attached to the low-molecular weight KSP inhibitor, for example a compound of the formula (I), (IIa), 20 (II), (III), (IIIa), (IIIb), (IIIc), (IIId), (IIIe) or (IV).

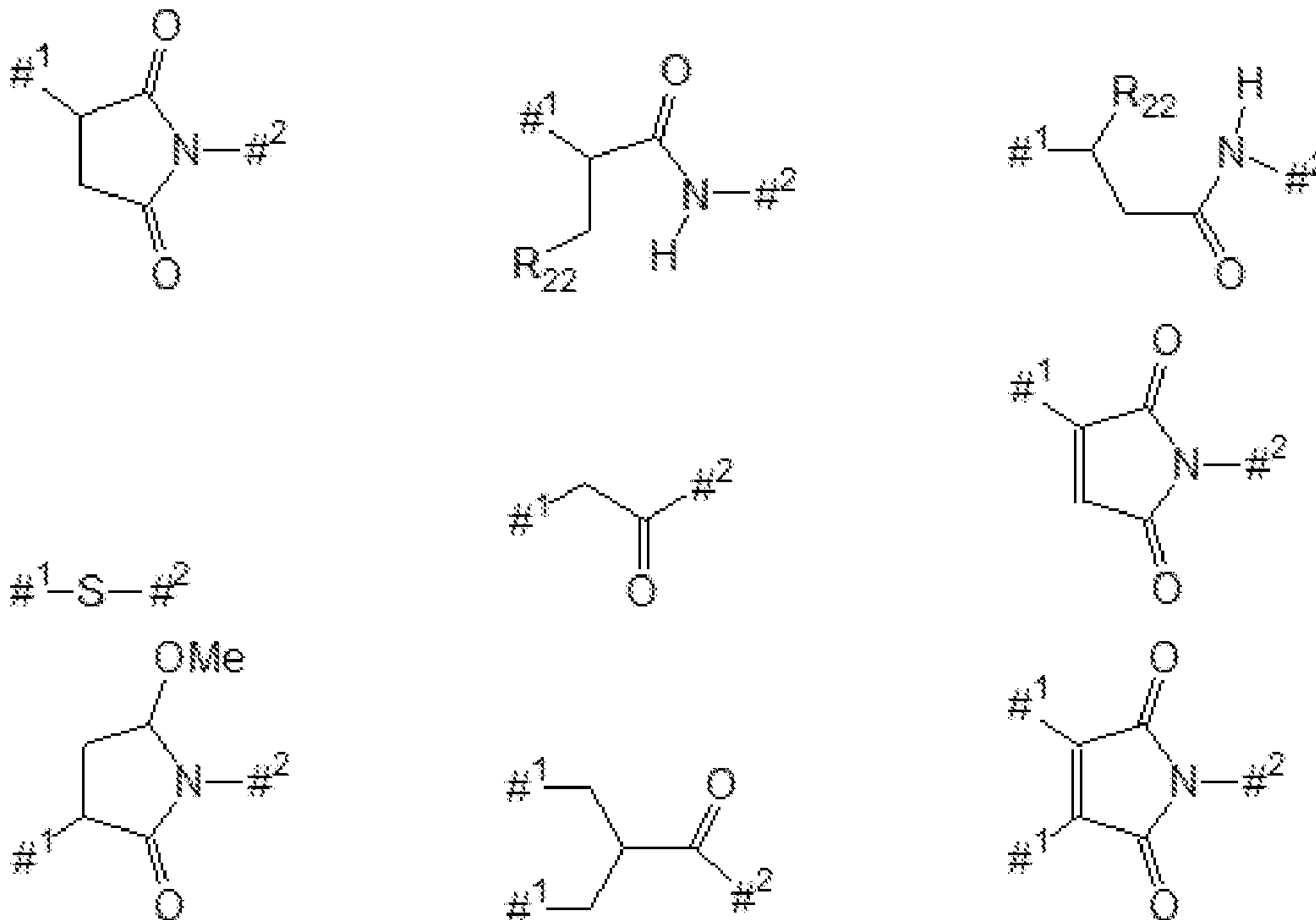
Preference according to the invention is also given to the basic linker structures (ii) and (iv), in particular when attachment is at position (iii), in particular when group L1 has one of the 25 following structures:

(a)  $\text{—NH—(CH}_2\text{)}_{0-4}\text{—(CHCH}_3\text{)}_{0-4}\text{—CHY}^5\text{—CO—Y}^7$  , where  $\text{Y}^5$  represents H or  $\text{NHY}^6$ , where  $\text{Y}^6$  represents H or  $\text{—COCH}_3$ , and  $\text{Y}^7$  represents a single bond or  $\text{—NH—(CH}_2\text{)}_{0-4}\text{—CHNH}_2\text{—CO—}$ , so that after cleavage the 30 corresponding structure  $\text{—NH—(CH}_2\text{)}_{0-4}\text{—(CHCH}_3\text{)}_{0-4}\text{—CHY}^5\text{—COOH}$  or  $\text{—NH—(CH}_2\text{)}_{0-4}\text{—(CHCH}_3\text{)}_{0-4}\text{—CHY}^5\text{—CO—NH—(CH}_2\text{)}_{0-4}\text{—CHNH}_2\text{—COOH}$  is obtained.

(b)  $\text{—CH}_2\text{—S}_x\text{—(CH}_2\text{)}_{0-4}\text{—CHY}^5\text{—CO—}$ , where x is 0 or 1, and  $\text{Y}^5$  represents H or  $\text{NHY}^6$ , where  $\text{Y}^6$  represents H or  $\text{—COCH}_3$ , such that after 35 cleavage the corresponding structure  $\text{—CH}_2\text{—S}_x\text{—(CH}_2\text{)}_{0-4}\text{—CHY}^5\text{—COOH}$  is obtained.

This embodiment is preferred when L1 is attached in each case to the low-molecular weight KSP inhibitor, for example a compound of the formula (I), (IIa), (II), (III), (IIIa), (IIIb), (IIIc), (IIId), (IIIe) or (IV), in particular at position R<sub>4</sub>.  
 5 The binder is preferably an anti-TWEAKR antibody or an anti-EGFR antibody.

If the linker is attached to a cysteine side chain or a cysteine residue, L2 is preferably derived from a group which reacts with the sulphhydryl group of the cysteine. These include haloacetyls, maleimides, aziridines, acryloyls, arylating compounds, vinylsulphones, pyridyl disulphides, TNB thiols and disulphide-reducing agents. These groups generally react in an  
 10 electrophilic manner with the sulphhydryl bond, forming a sulphide (e.g. thioether) or disulphide bridge. Preference is given to stable sulphide bridges. L2 is preferably  
 15



20

where

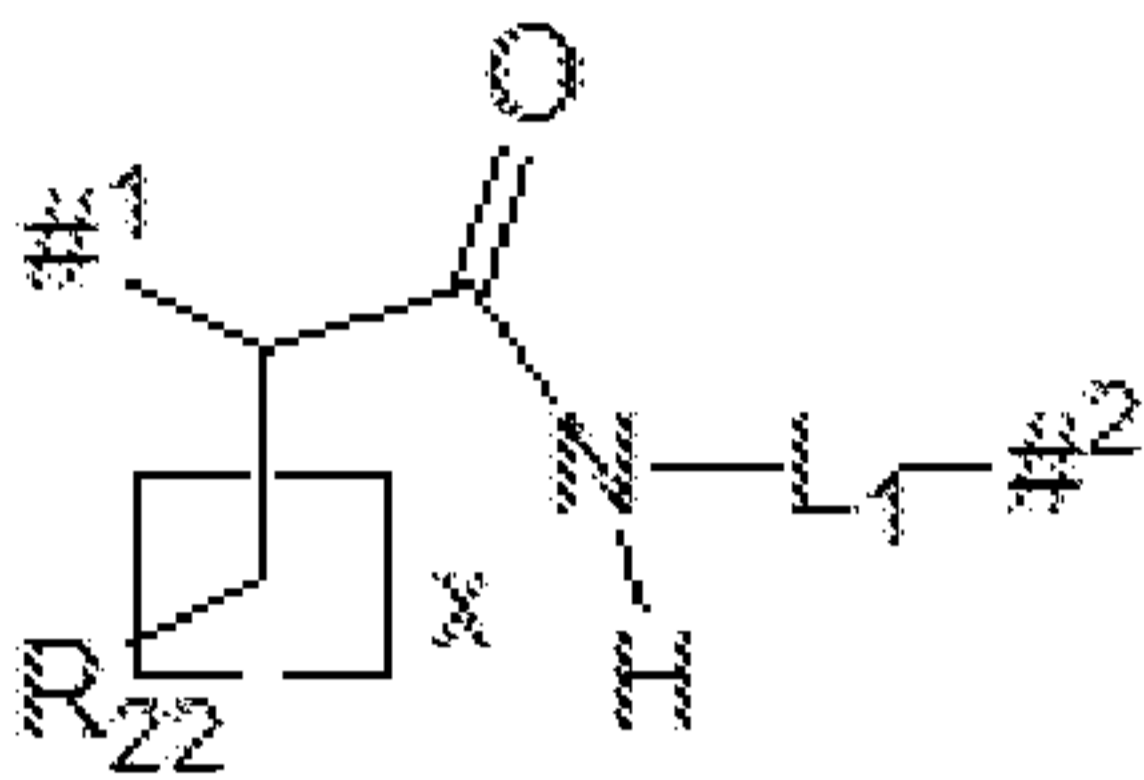
#<sup>1</sup> denotes the point of attachment to the sulphur atom of the binder,



#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>, and

R<sub>22</sub> represents COOH, COOR, COR, CONHR, CONR<sub>2</sub> (where R in each  
5 case represents C<sub>1-3</sub>-alkyl), CONH<sub>2</sub>, preferably COOH.

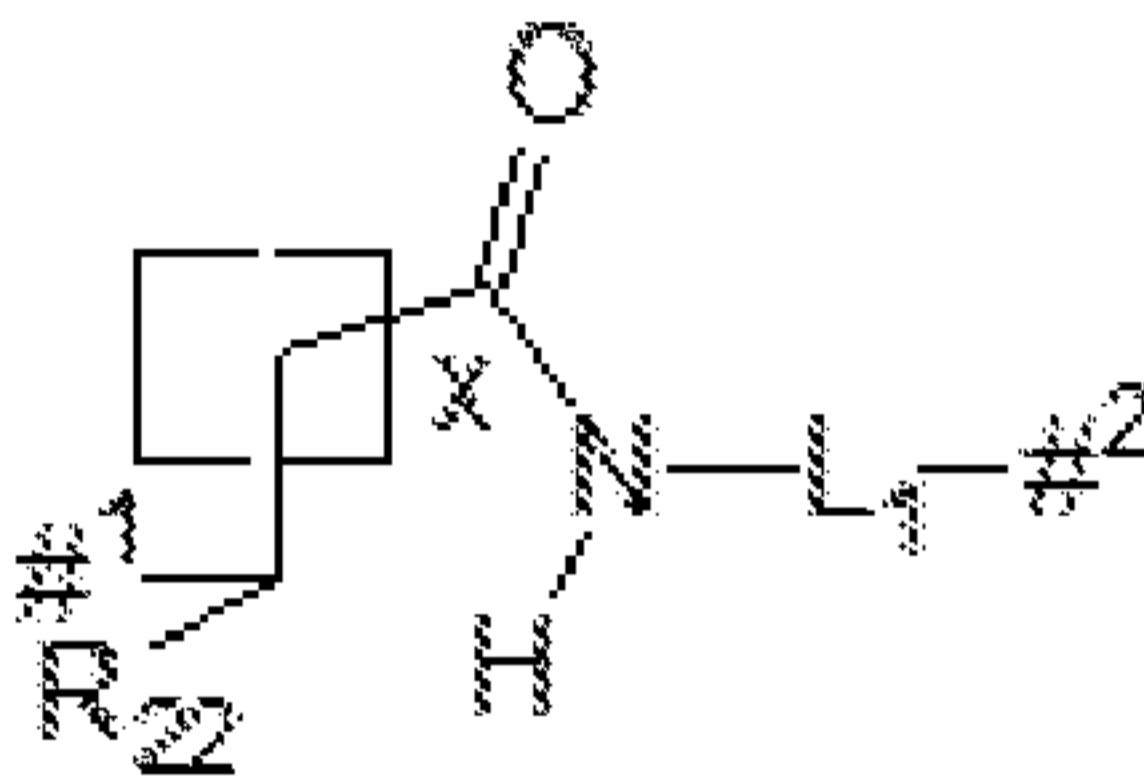
Particularly preferred for L<sub>2</sub> is:



Formula A3

10

or

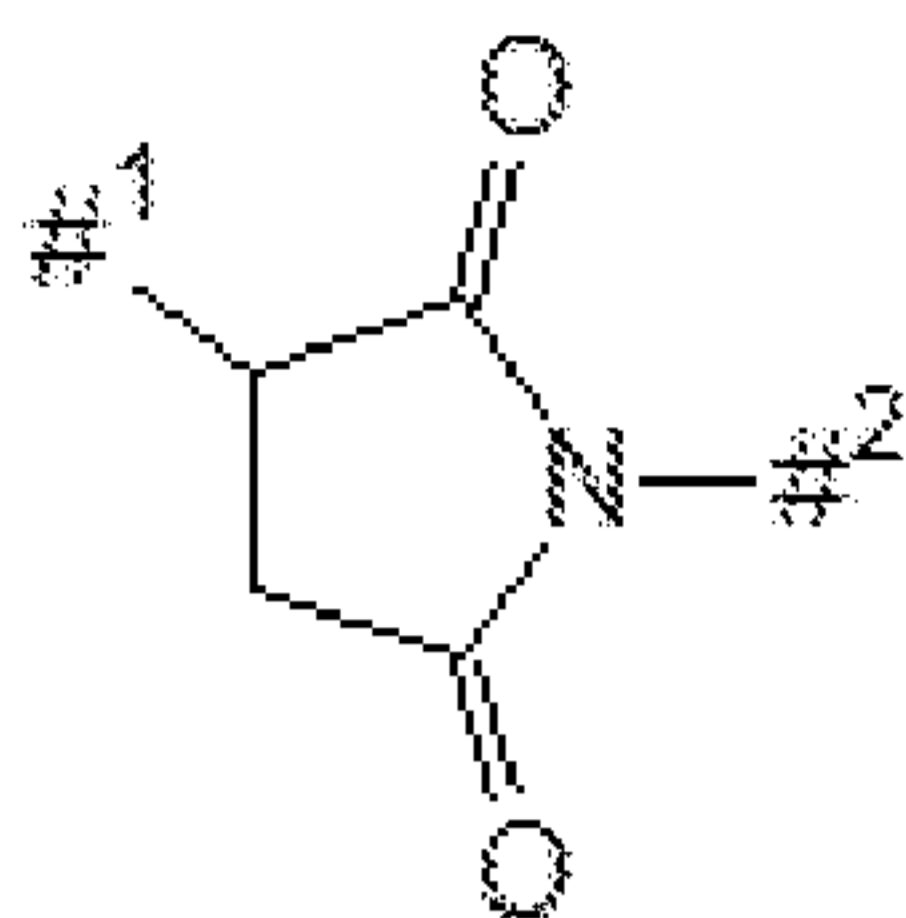


Formula A4

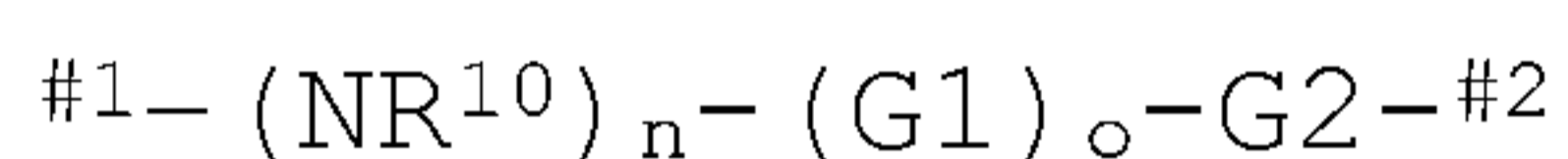
15 where #<sup>1</sup> denotes the point of attachment to the sulphur atom of  
the binder, #<sup>2</sup> denotes the point of attachment to the active  
compound, x represents 1 or 2, and R<sup>22</sup> represents COOH, COOR,  
COR, CONHR (where R in each case represents C<sub>1-3</sub>-alkyl), CONH<sub>2</sub>,  
preferably COOH. It is preferred when x=1 and R<sup>22</sup> represents  
20 COOH.

In a conjugate according to the invention or in a mixture of the  
conjugates according to the invention, the bonds to a cysteine  
residue of the binder are present, to an extent of preferably  
25 more than 80%, particularly preferably more than 90% (in each  
case based on the total number of bonds of the linker to the  
binder), particularly preferably as one of the two structures  
of the formula A3 or A4. Here, the structures of the formula A3  
or A4 are generally present together, preferably in a ratio of  
30 from 60:40 to 40:60, based on the number of bonds to the binder.

The remaining bonds are then present as the structure

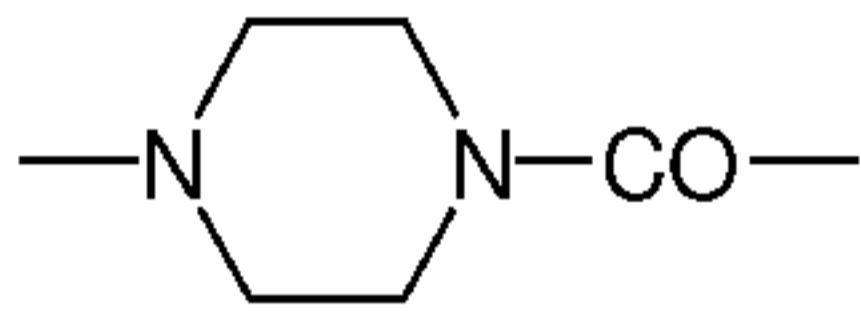
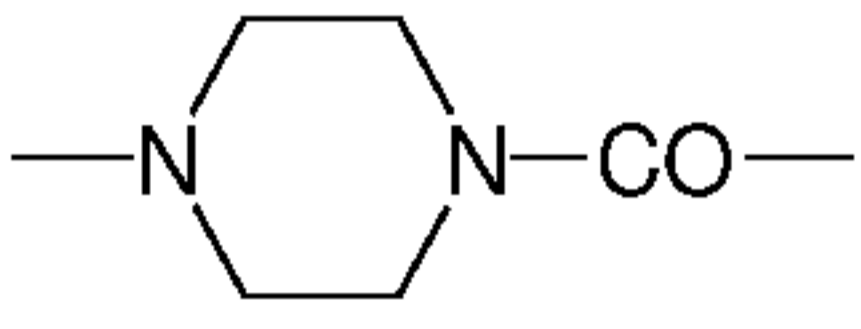


5 According to the invention, L1 is preferably represented by the formula



10 where

R<sup>10</sup> represents H, NH<sub>2</sub> or C<sub>1</sub>-C<sub>3</sub>-alkyl;

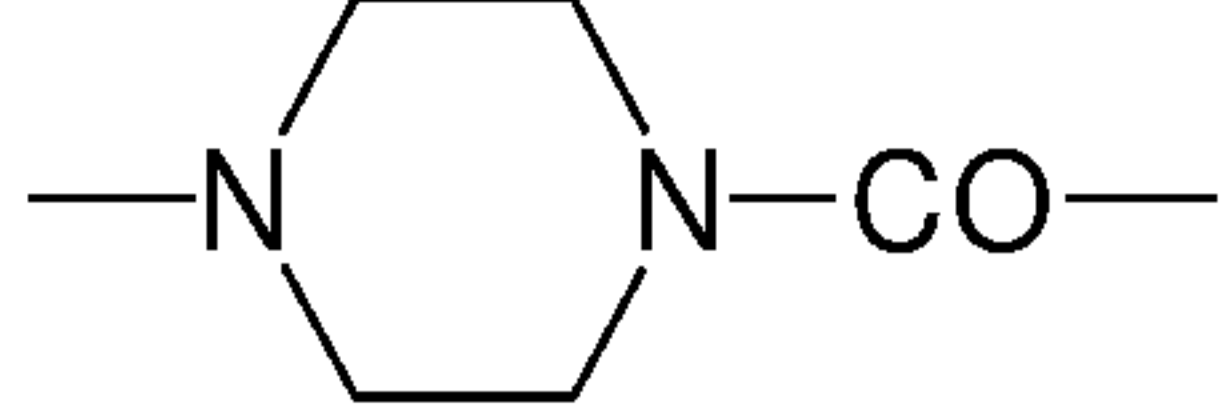
G1 represents -NHCO- , -CONH- or  ; (R<sup>10</sup> is preferably  
 15 not NH<sub>2</sub>, if G1 represents NHCO or  ).

n is 0 or 1;

o is 0 or 1; and

20

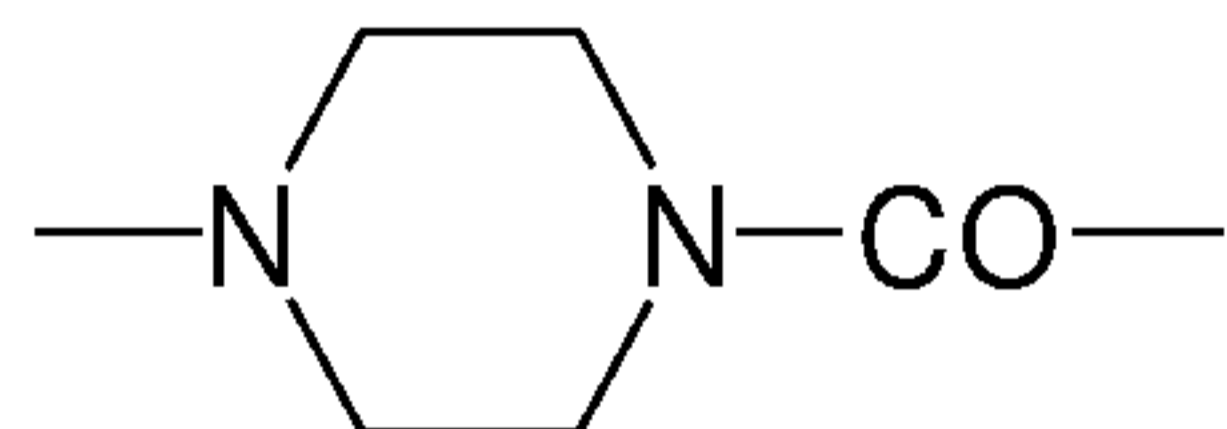
G2 represents a straight-chain or branched hydrocarbon chain which has 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more  
 25 of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, -C(NH)NR<sup>y</sup>-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>- (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -  
 30 CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl) and/or a 3- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O

and S, -SO- or -SO<sub>2</sub>- (preferably ) , where the hydrocarbon chain including the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid.

5

G2 represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the

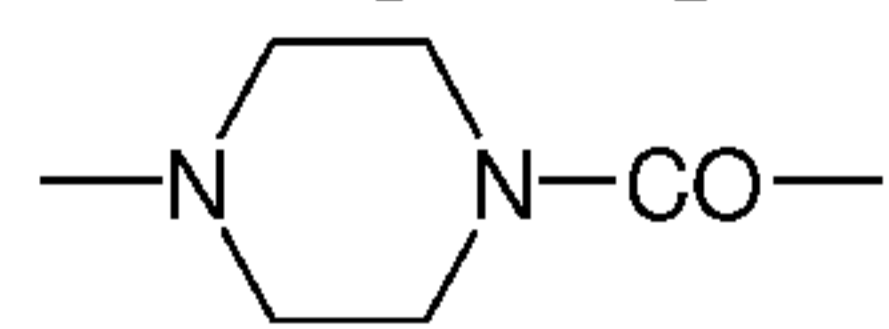
10 groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, or -SO- (preferably



15 substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid.

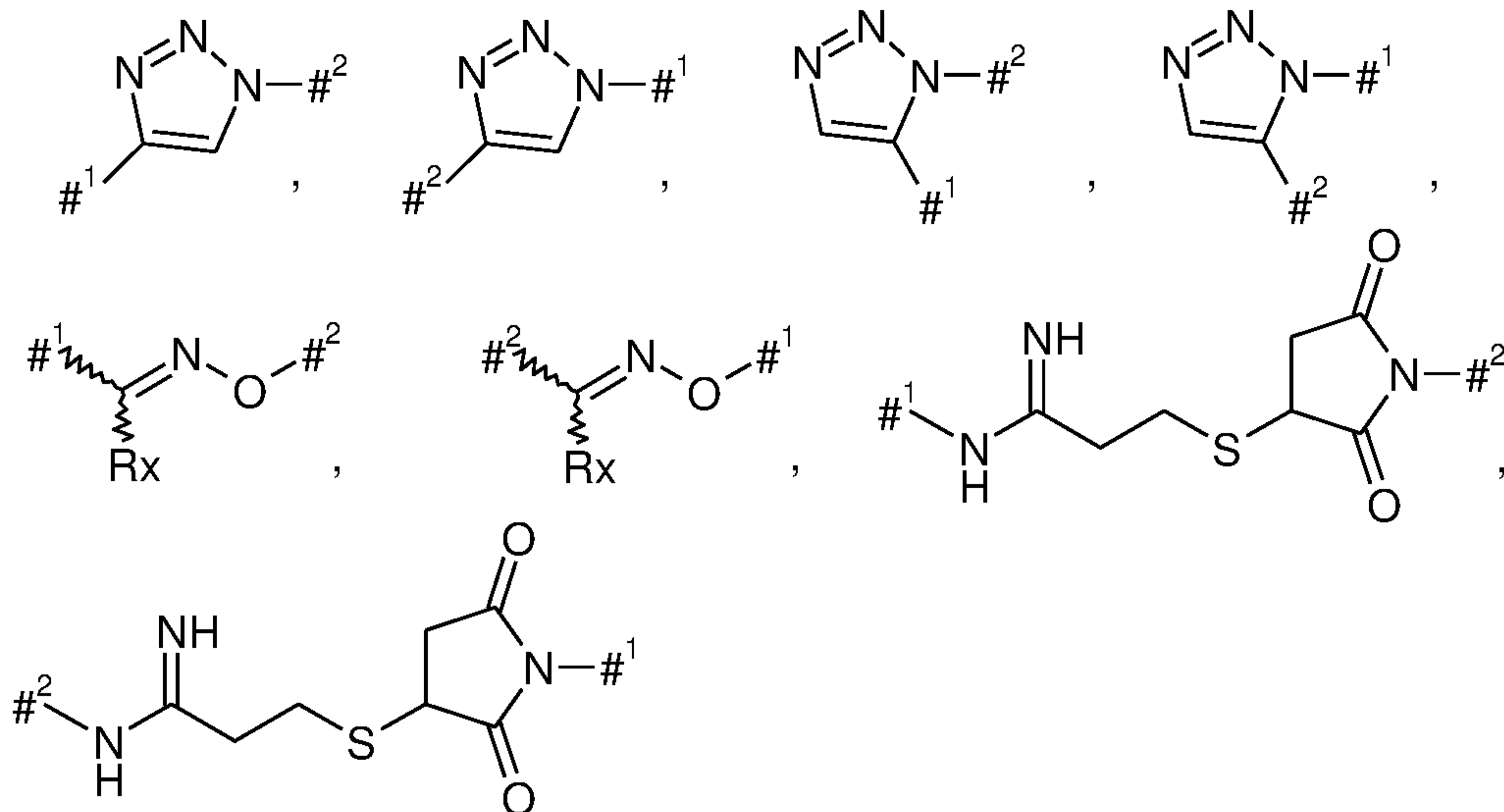
G2 preferably represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl) and a 3- to 10-

25 membered, for example 5- to 10-membered, aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, -SO- or -SO<sub>2</sub>- (preferably



30 substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid.

Further interrupting groups in G2 are preferably



where Rx represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl.

- 5 Here, #<sup>1</sup> is the bond to the KSP inhibitor and #<sup>2</sup> is the bond to the coupling group to the binder (e.g. L2).

A straight-chain or branched hydrocarbon chain of arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups generally comprises a  $\alpha,\omega$ -divalent alkyl radical having the respective number of carbon atoms stated. Examples which may be mentioned as being preferred are: methylene, ethane-1,2-diyl (1,2-ethylene), propane-1,3-diyl (1,3-propylene), butane-1,4-diyl (1,4-butylene), pentane-1,5-diyl (1,5-pentylene), hexane-1,6-diyl (1,6-hexylene), heptane-1,7-diyl (1,7-hexylene), octane-1,8-diyl (1,8-octylene), nonane-1,9-diyl (1,9-nonylene), decane-1,10-diyl (1,10-decylene). However, the alkylene groups in the hydrocarbon chain may also be branched, i.e. one or more hydrogen atoms of the straight-chain alkylene groups mentioned above may optionally be substituted by C<sub>1-10</sub>-alkyl groups, thus forming side chains. The hydrocarbon chain may furthermore contain cyclic alkylene groups (cycloalkanediyl), for example 1,4-cyclohexanediyl or 1,3-cyclopentanediy. These cyclic groups may be unsaturated. In particular, aromatic groups (arylene groups), for example phenylene, may be present in the hydrocarbon group. In turn, in the cyclic alkylene groups and the arylene groups, too, one or more hydrogen atoms may optionally be substituted by C<sub>1-10</sub>-alkyl groups. In this way, an

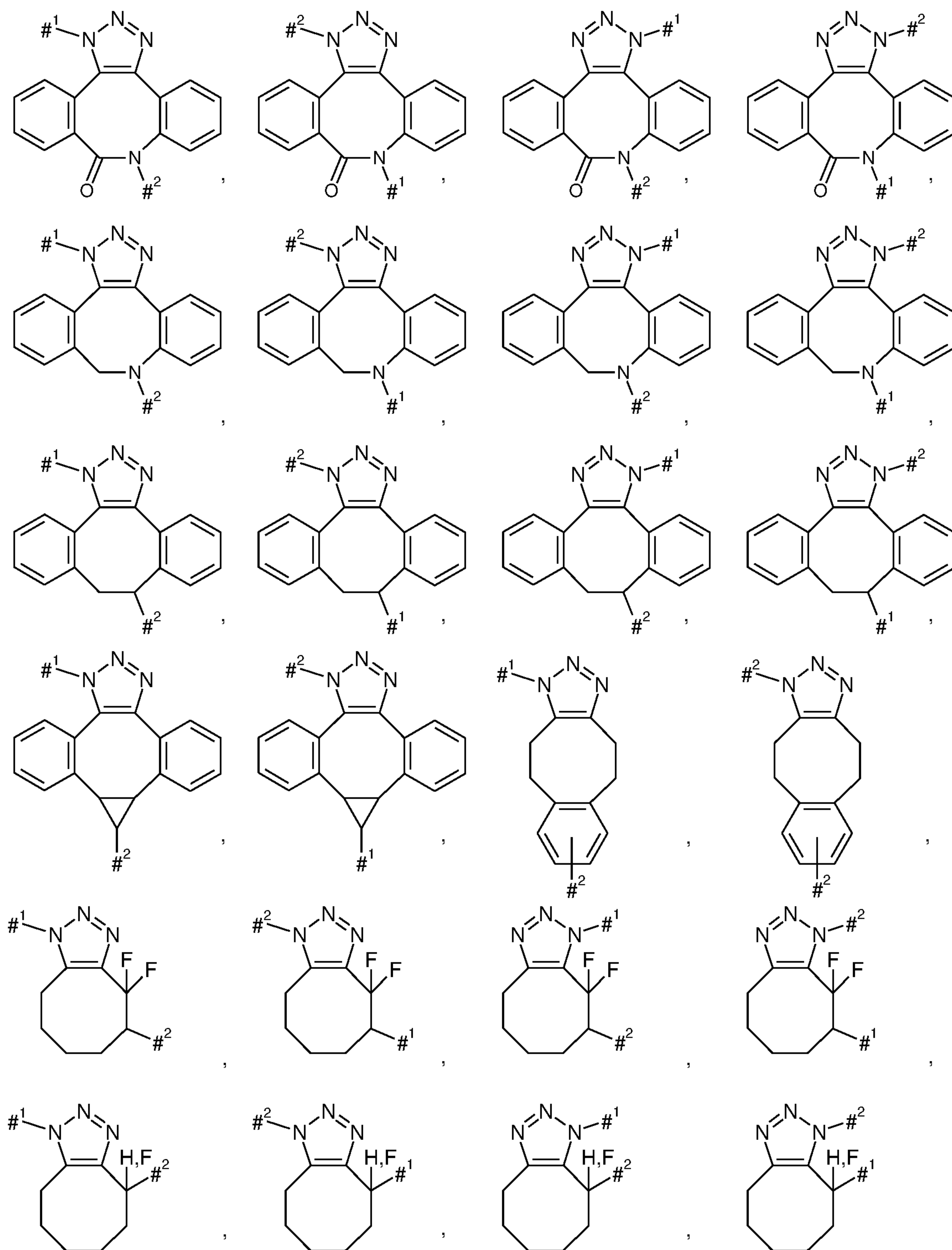
optionally branched hydrocarbon chain is formed. This hydrocarbon chain has a total of 0 to 100 carbon atoms, preferably 1 to 50, particularly preferably 2 to 25 carbon atoms.

- 5 The side chains, if present, may be substituted by  $-\text{NHCONH}_2$ ,  $-\text{COOH}$ ,  $-\text{OH}$ ,  $-\text{NH}_2$ ,  $\text{NH-CNNH}_2$ , sulphonamide, sulphone, sulphoxide or sulphonic acid.

10 The hydrocarbon chain may be interrupted once or more than once by one or more of the groups  $-\text{O}-$ ,  $-\text{S}-$ ,  $-\text{SO}-$ ,  $\text{SO}_2$ ,  $-\text{NH}-$ ,  $-\text{CO}-$ ,  $-\text{NHCO}-$ ,  $-\text{CONH}-$ ,  $-\text{NMe}-$ ,  $-\text{NHNH}-$ ,  $-\text{SO}_2\text{NHNH}-$ ,  $-\text{CONHNH}-$  and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, -

15  $\text{SO}-$  or  $-\text{SO}_2-$  (preferably  ).

Further interrupting groups in G2 are preferably



Preferably, the linker corresponds to the formula below:

5  $\text{S}-(\text{CO})_m\text{-L1-L2-SS}$

where

$m$  is 0 or 1;

§ represents the bond to the active compound molecule and

§§ represents the bond to the binder peptide or protein, and

5

L1 and L2 have the meaning given above.

Particularly preferably, L1 has the formula  $-NR^{11}B-$ , where

10  $R^{11}$  represents H or  $NH_2$ ;

B represents  $-[(CH_2)_x-(X^4)_y]_w-(CH_2)_z-$ ,

w = 0 to 20;

15

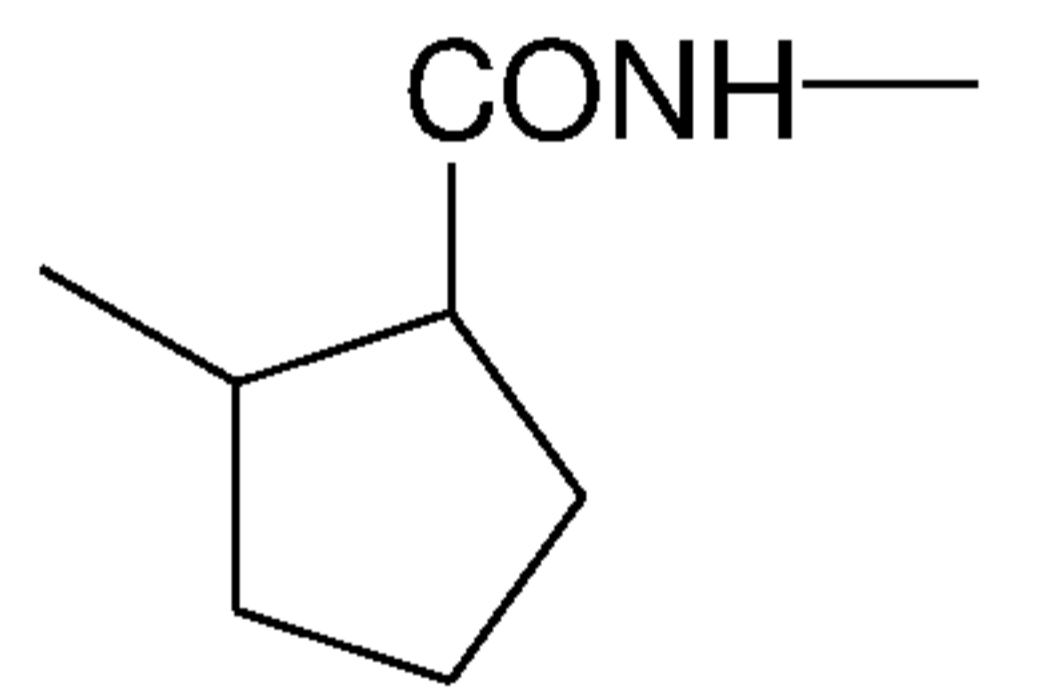
x = 0 to 5;

x = 0 to 5;

20 y = 0 or 1;

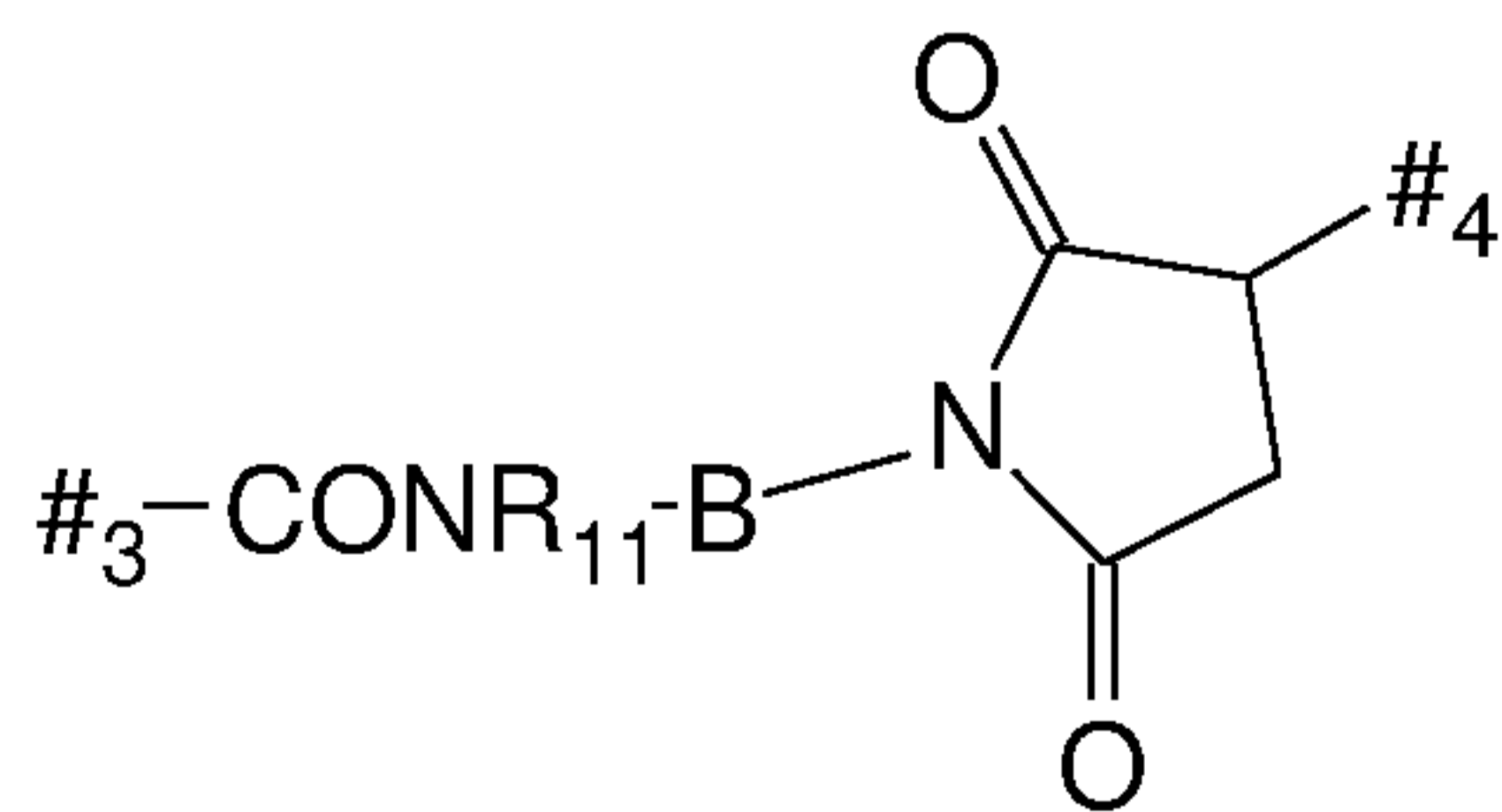
z = 0 to 5; and

$X^4$  represents  $-O-$ ,  $-CONH-$ ,  $-NHCO-$  or



25

Linkers L which are preferred in accordance with the invention have the formula below:



30

where

#3 represents the bond to the active compound molecule,

#4 represents the bond to the binder peptide or protein,

5  $R^{11}$  represents H or  $NH_2$ ;

B represents  $-[(CH_2)_x-(X^4)_y]_w-(CH_2)_z-$ ,

w = 0 to 20;

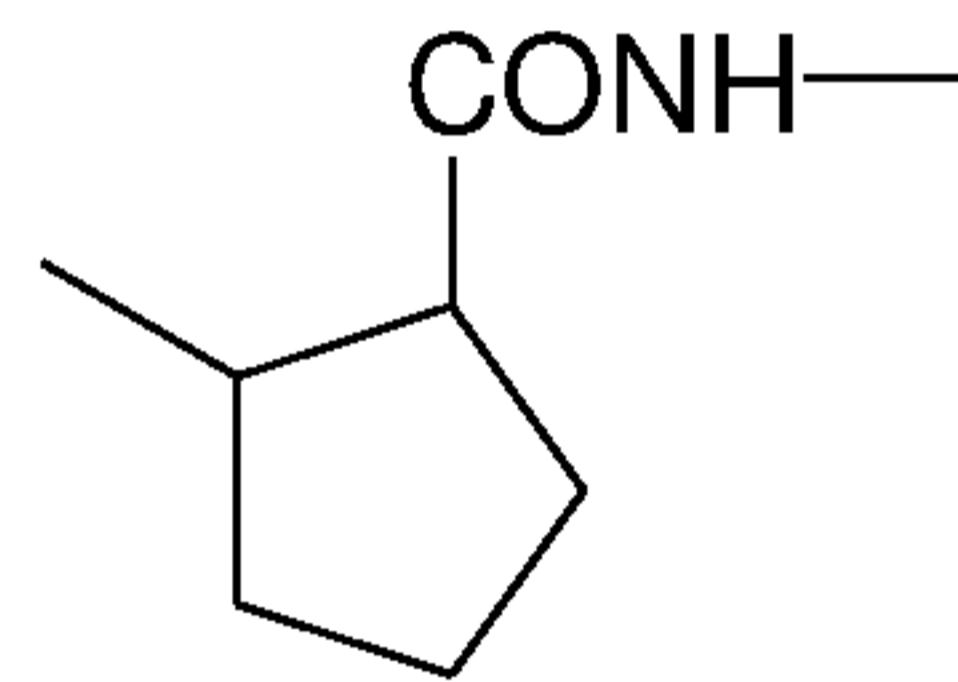
10

x = 0 to 5;

y = 0 or 1;

15 z = 1 to 5; and

$X^4$  represents  $-O-$ ,  $-CONH-$ ,  $-NHCO-$  or

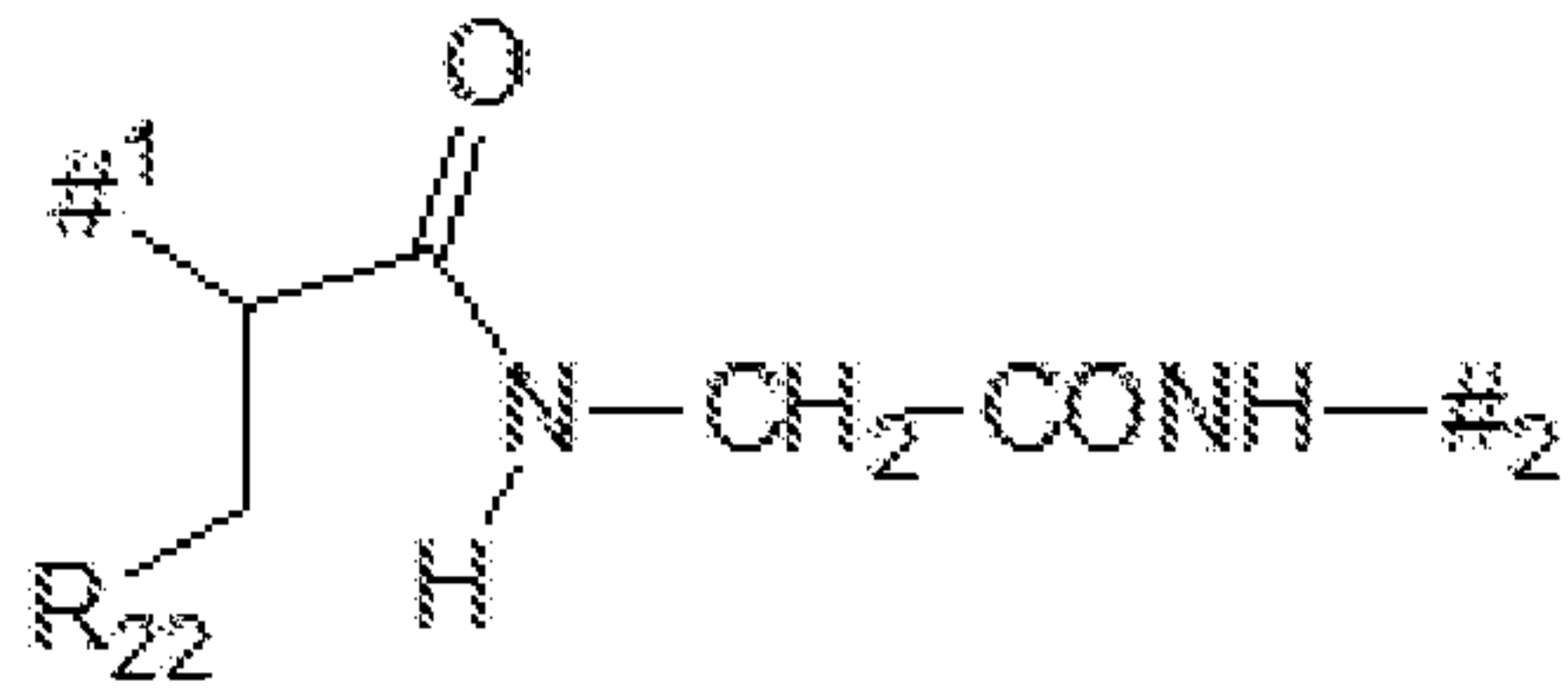


The linkers mentioned above are especially preferred in  
 20 conjugates of the formula (I) or (II) in which the linker couples  
 by substitution of a hydrogen atom at  $R^1$  or in combination with  
 a cleavable linker SG1 at  $R^4$ , i.e.  $R^1$  represents  $-L-#1$  or  $R^4$   
 represents  $-SG1-L-#1$ , where #1 represents the bond to the  
 binder.

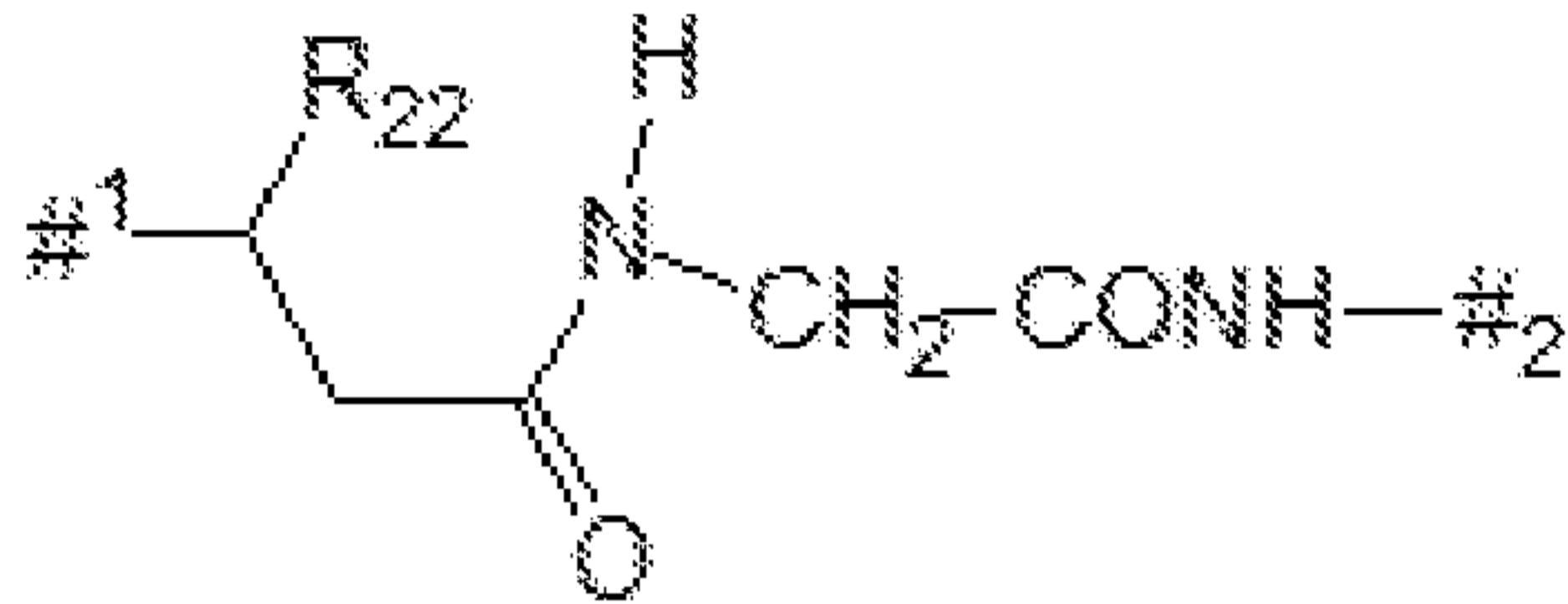
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Preference in accordance with the invention is furthermore given  
 to the linkers below: In a conjugate according to the invention  
 or in a mixture of the conjugates according to the invention,  
 the bonds to a cysteine residue of the binder are present, to  
 30 an extent of preferably more than 80%, particularly preferably  
 more than 90% (in each case based on the total number of bonds  
 of the linker to the binder), particularly preferably as one of  
 the two structures of the formula A5 or A6:





Formula A5



Formula A6

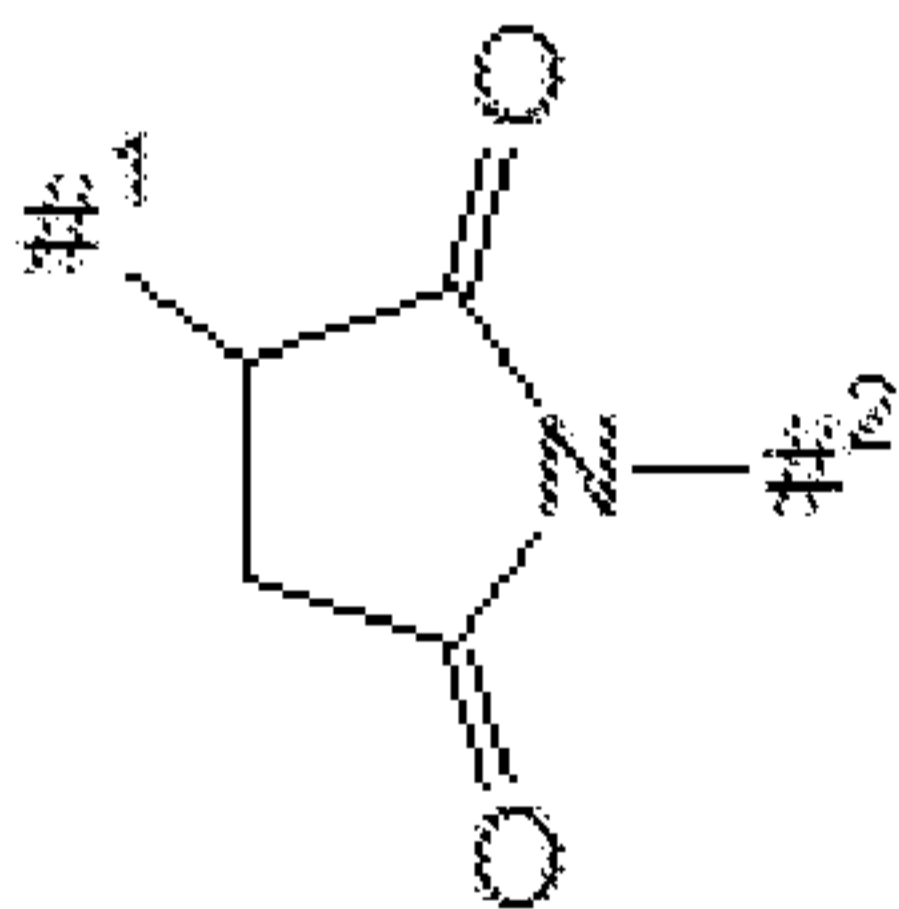
5 where

#<sup>1</sup> denotes the point of attachment to the sulphur atom of the binder,

10 #<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>, and

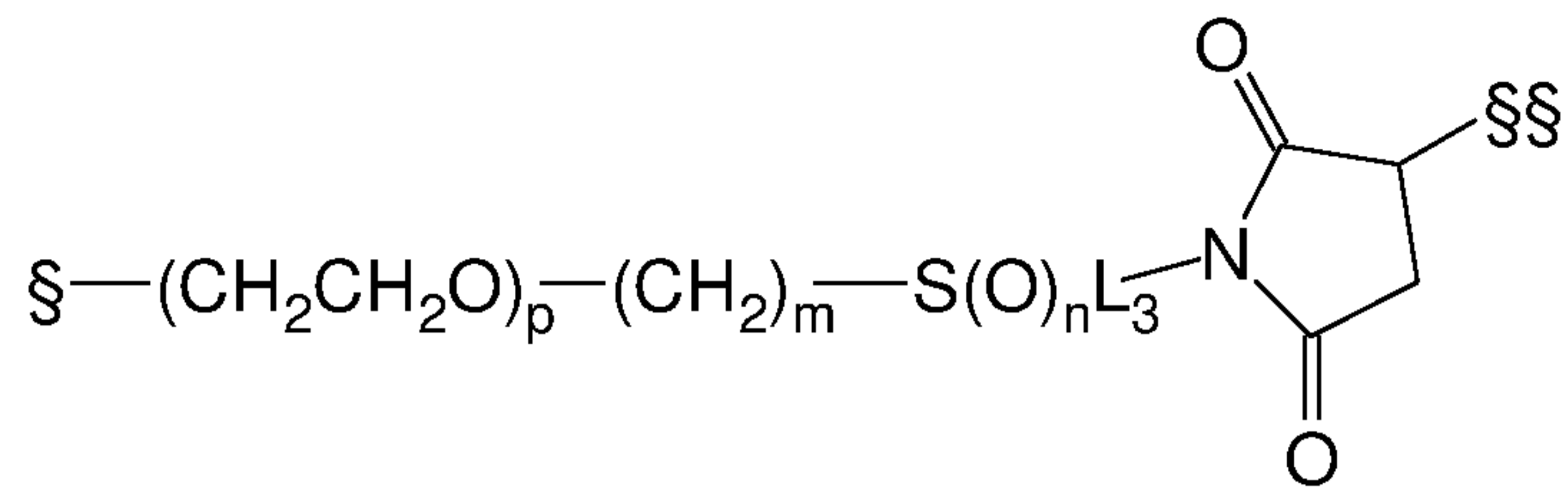
R<sup>22</sup> represents COOH, COOR, COR, CONHR (where R in each case represents C1-3-alkyl), CONH<sub>2</sub>, preferably COOH.

15 Here, the structures of the formula A5 or A6 are generally present together, preferably in a ratio of from 60:40 to 40:60, based on the number of bonds to the binder. The remaining bonds are then present as the structure



20

Other linkers -L- attached to a cysteine side chain or cysteine residue have the formula below:



where

5  $\S$  represents the bond to the active compound molecule and

$\text{SS}$  represents the bond to the binder peptide or protein,

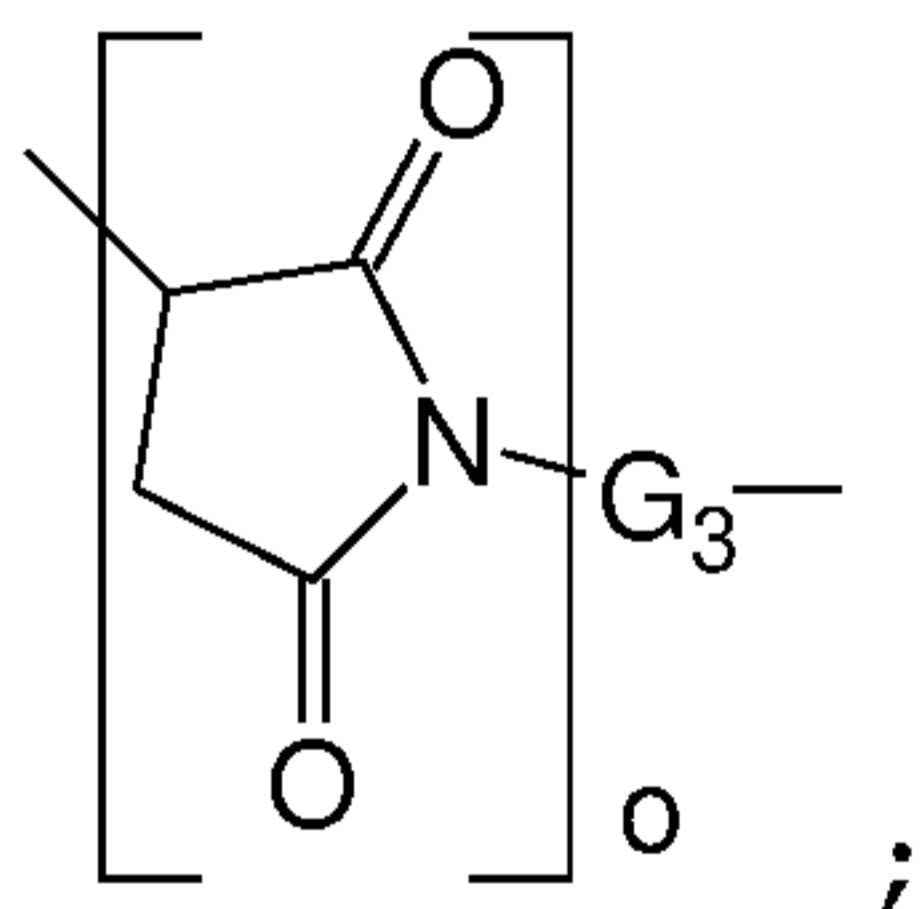
$m$  is 0, 1, 2 or 3;

10

$n$  is 0, 1 or 2;

$p$  is 0 to 20; and

15  $\text{L}_3$  represents



where

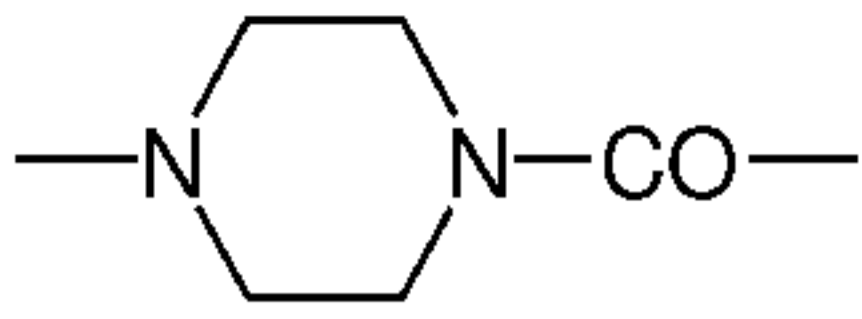
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$o$  is 0 or 1;

and

25  $\text{G}_3$  represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups  $-\text{O}-$ ,  $-\text{S}-$ ,  $-\text{SO}-$ ,  $\text{SO}_2$ ,  $-\text{NH}-$ ,  $-\text{CO}-$ ,  $-\text{NHCO}-$ ,  $-\text{CONH}-$ ,  $-\text{NMe}-$ ,  $-\text{NHNH}-$ ,  $-\text{SO}_2\text{NHNH}-$ ,  
 30  $-\text{CONHNH}-$  and a 3- to 10-membered (preferably 5- to 10-membered) aromatic or non-aromatic heterocycle having up to 4 heteroatoms

selected from the group consisting of N, O and S, -SO- or SO<sub>2</sub>

(preferably ) , where the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulfoxide or sulphonic acid.

5

In the formula above, preferably

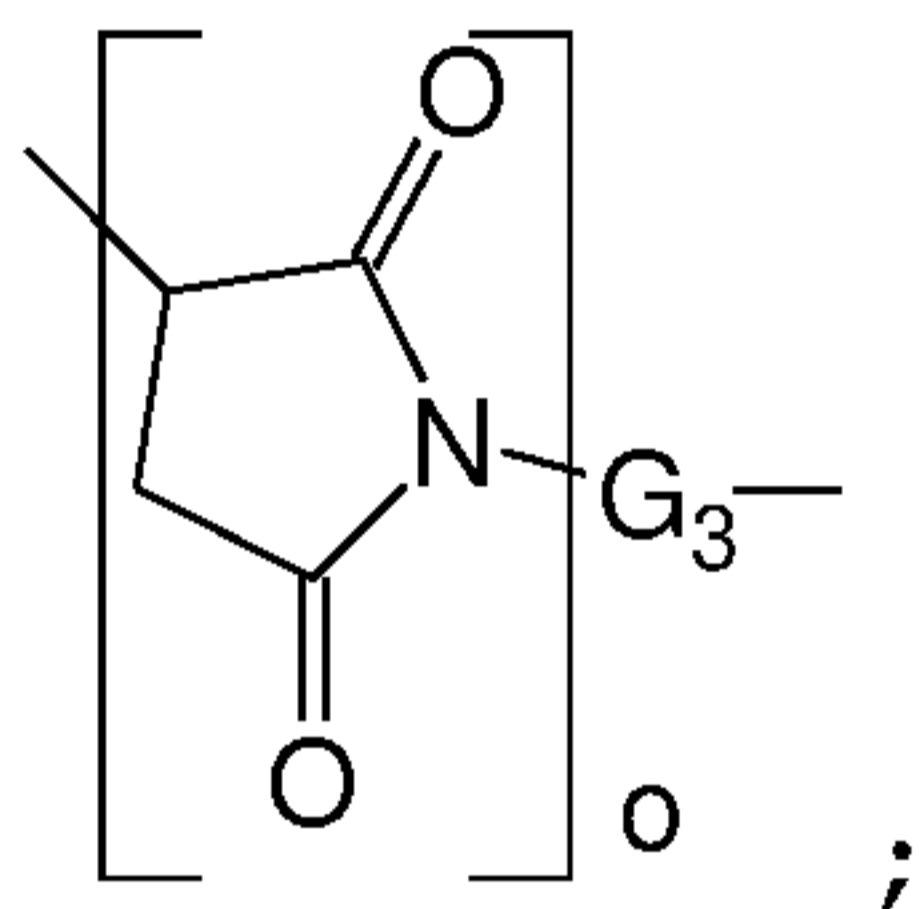
m is 1;

10 p is 0;

n is 0;

and L<sub>3</sub> represents

15



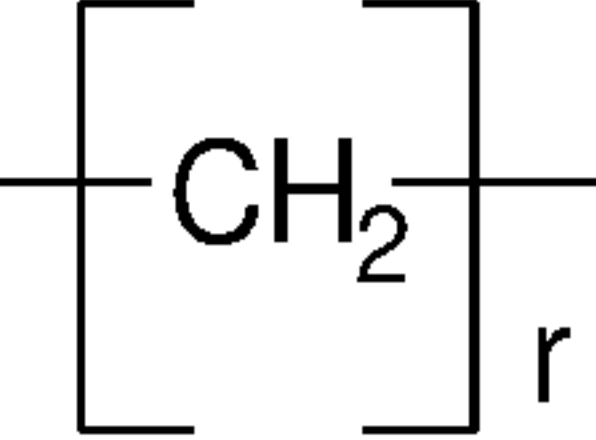
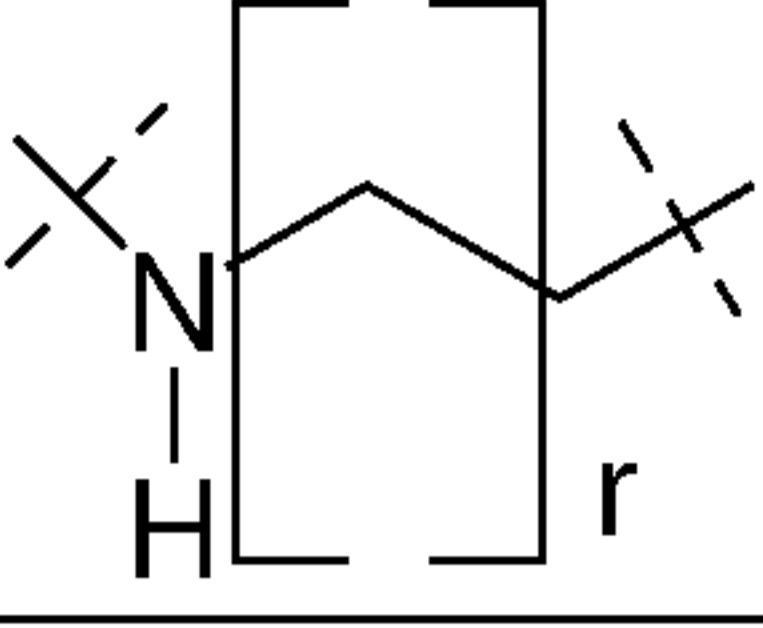
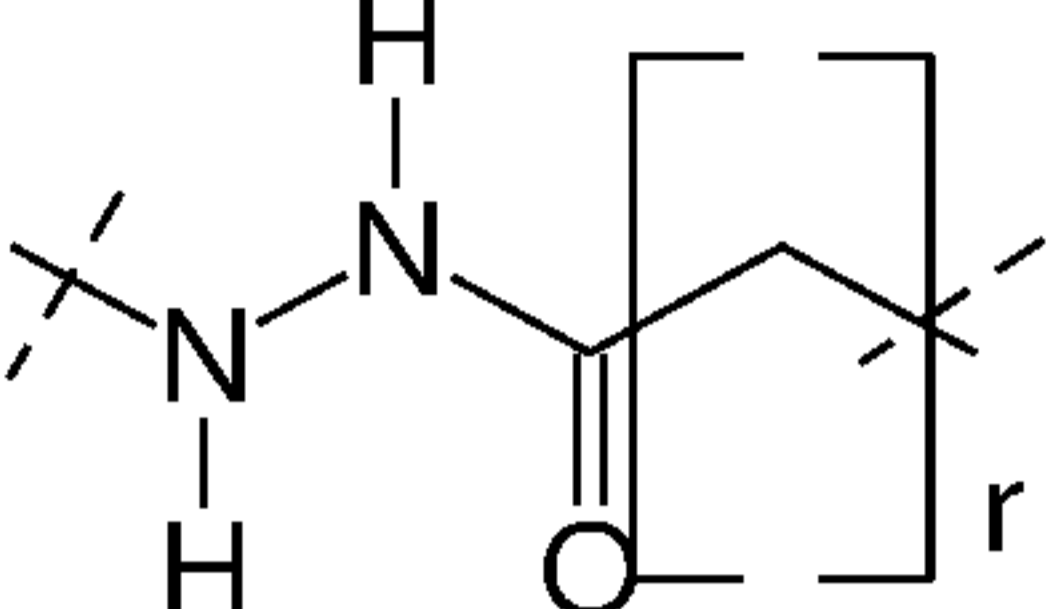
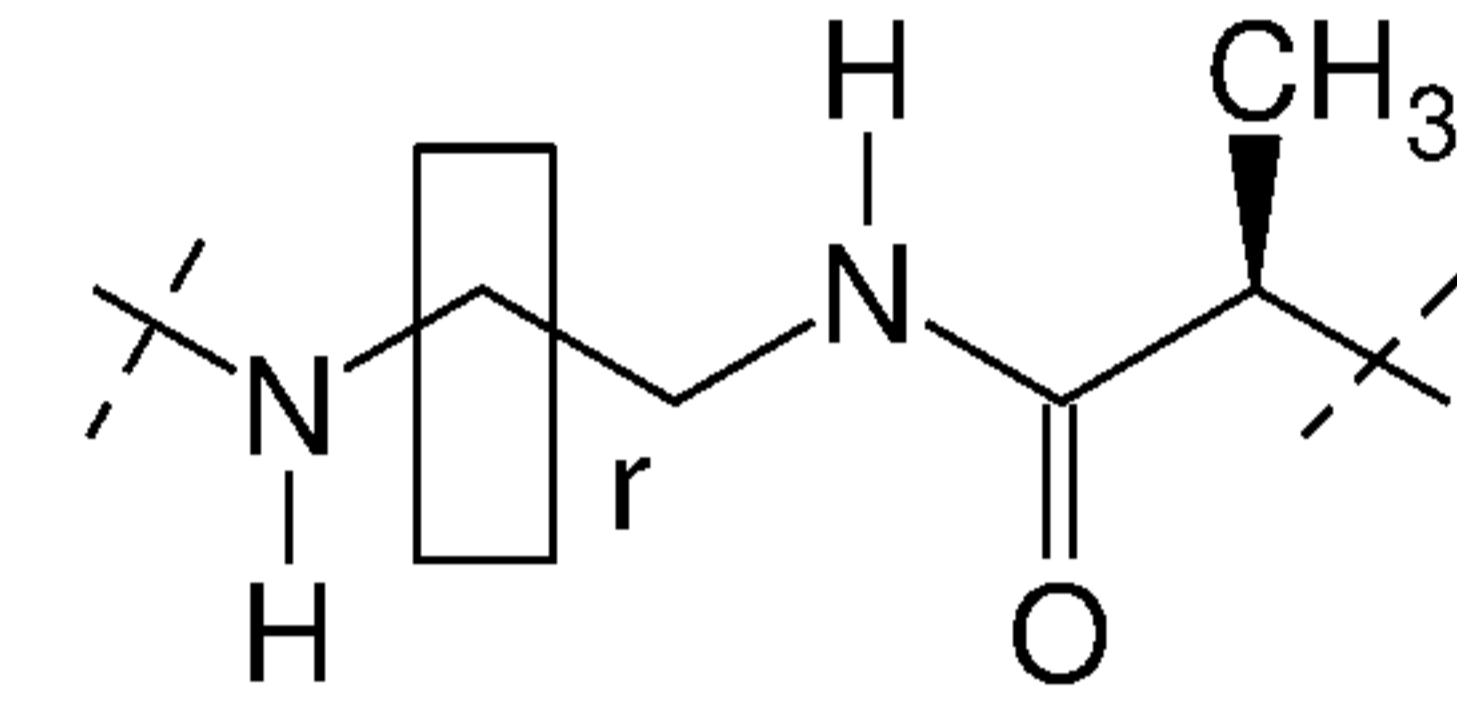
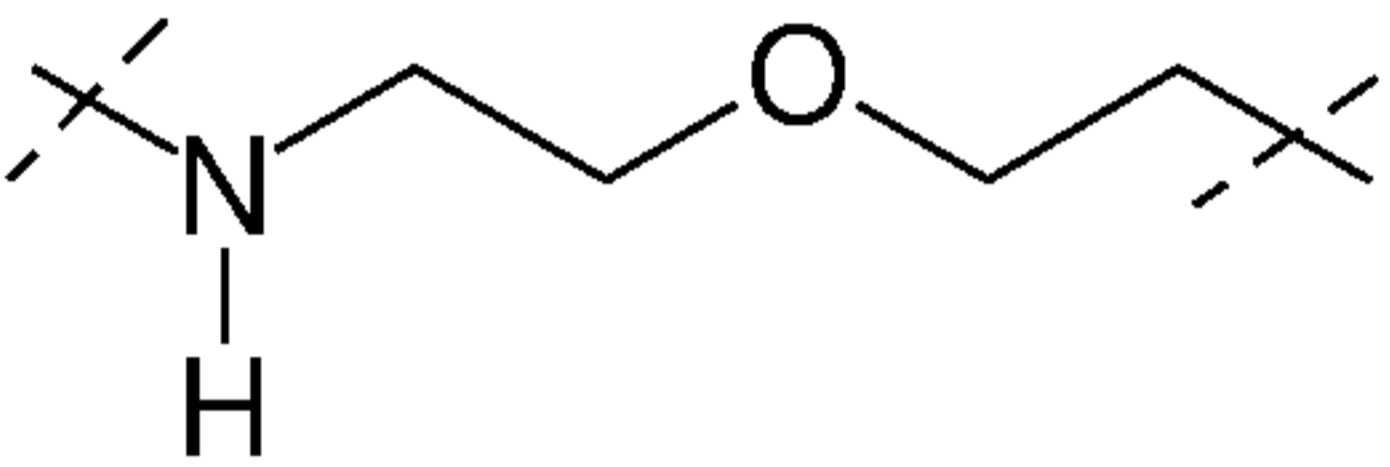
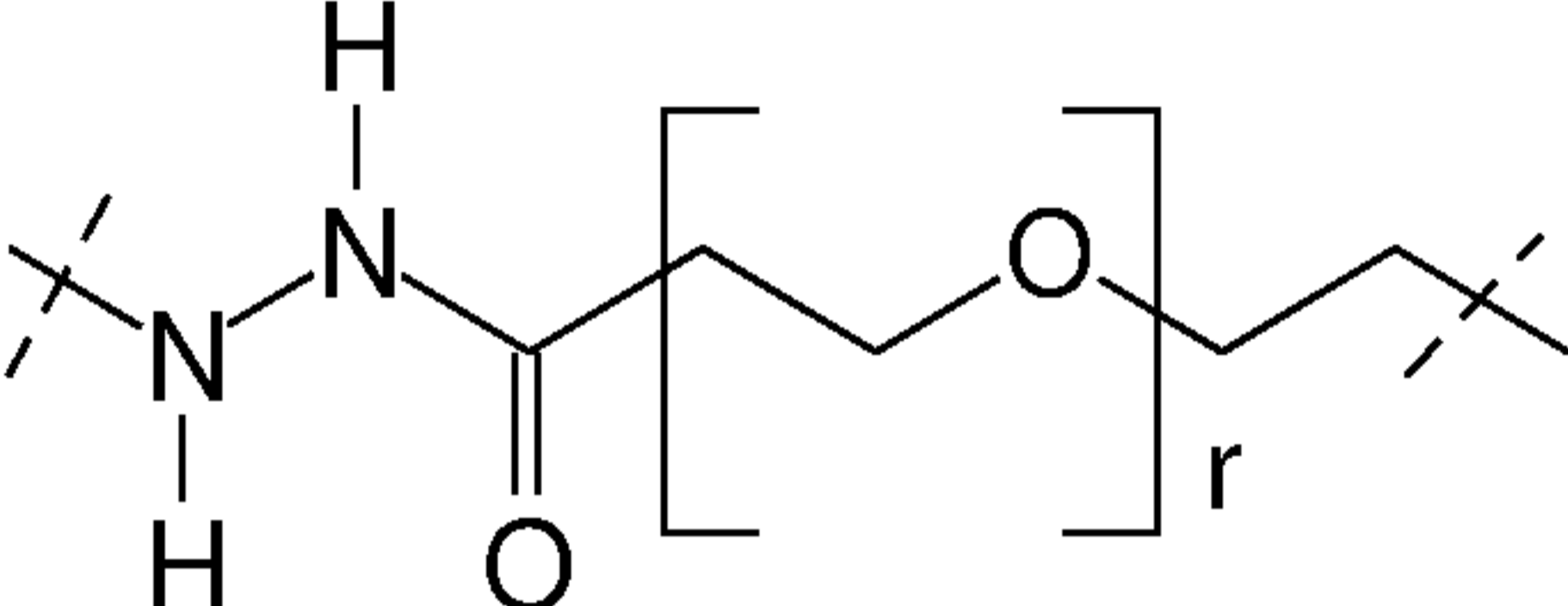
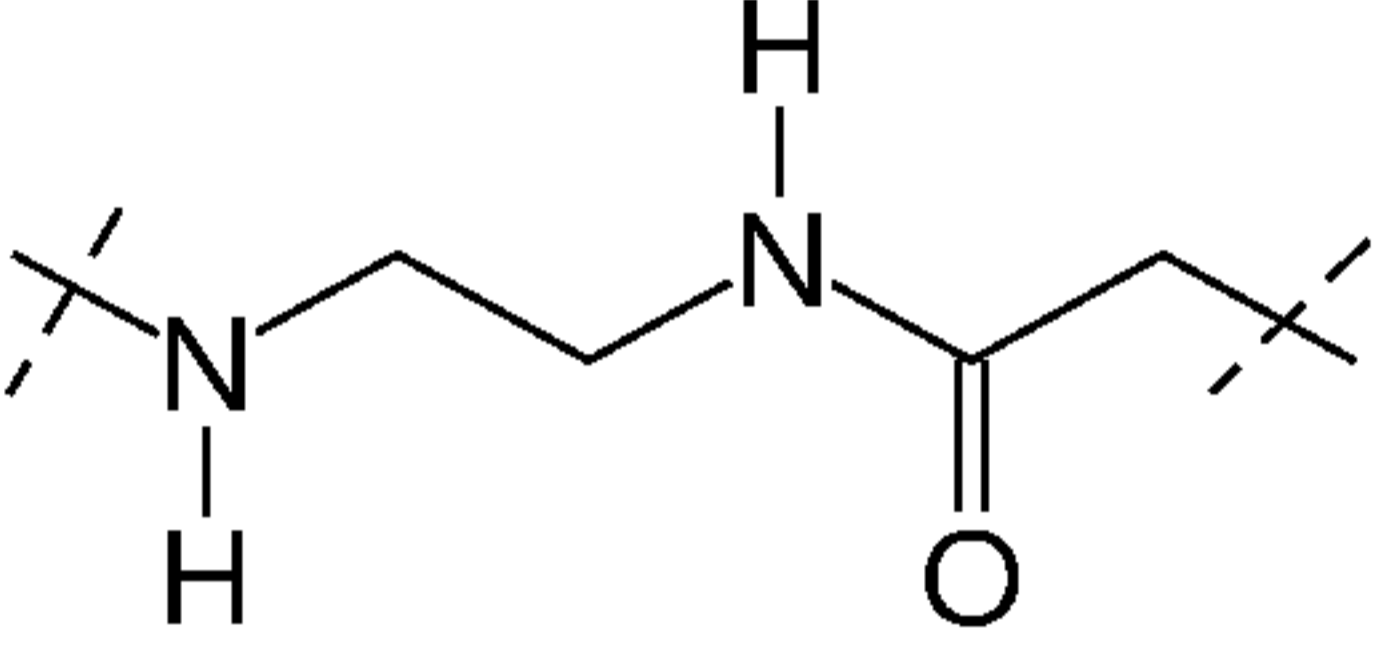
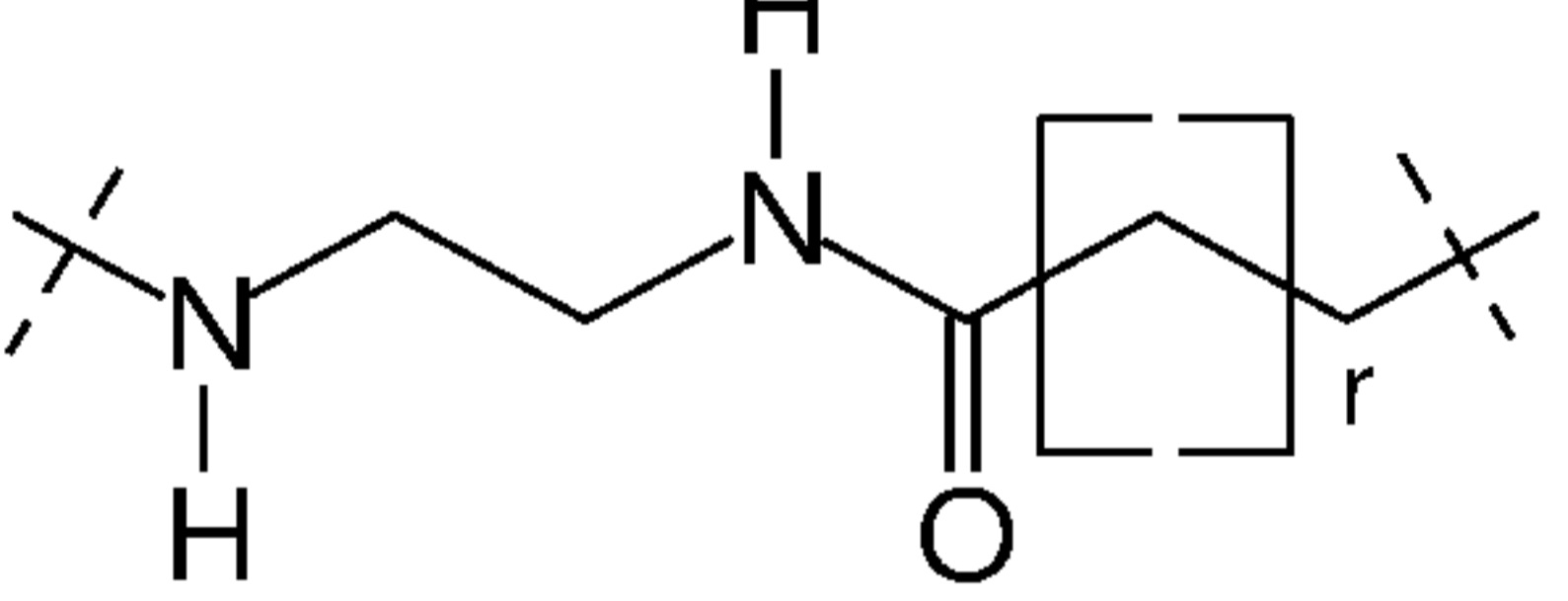
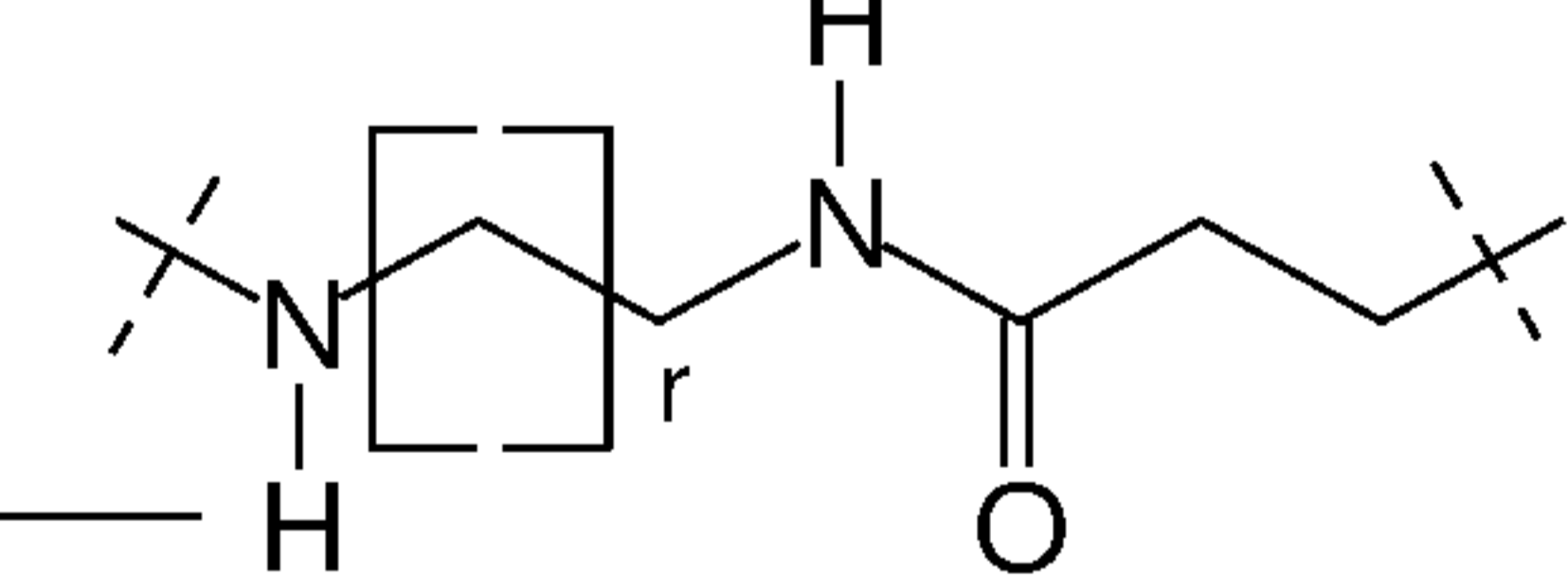
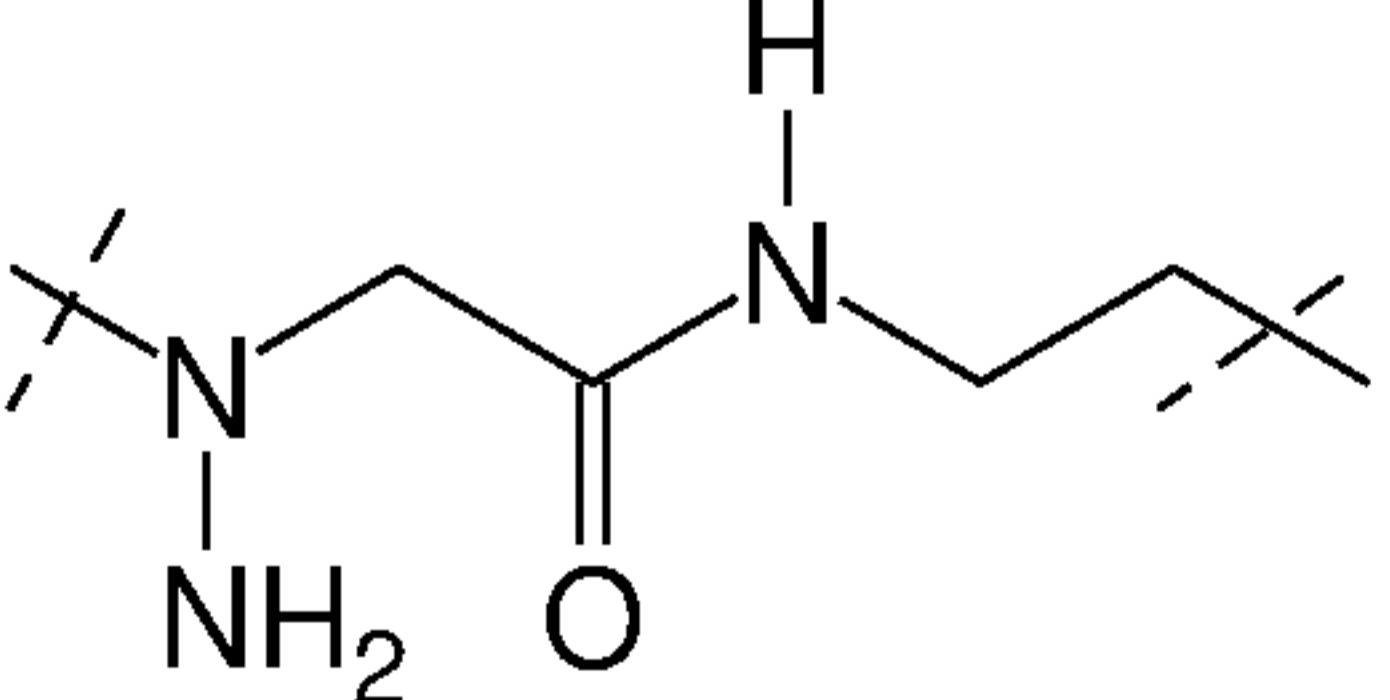
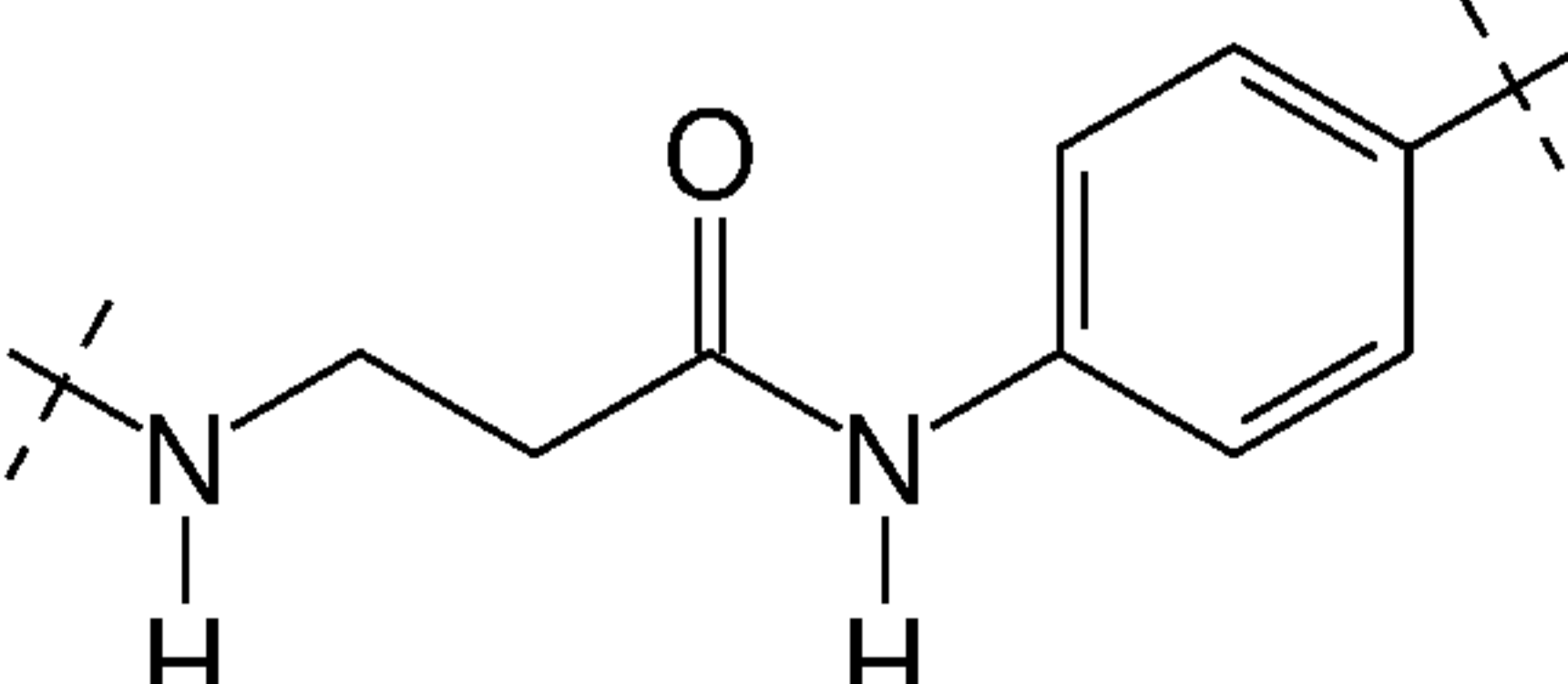
where

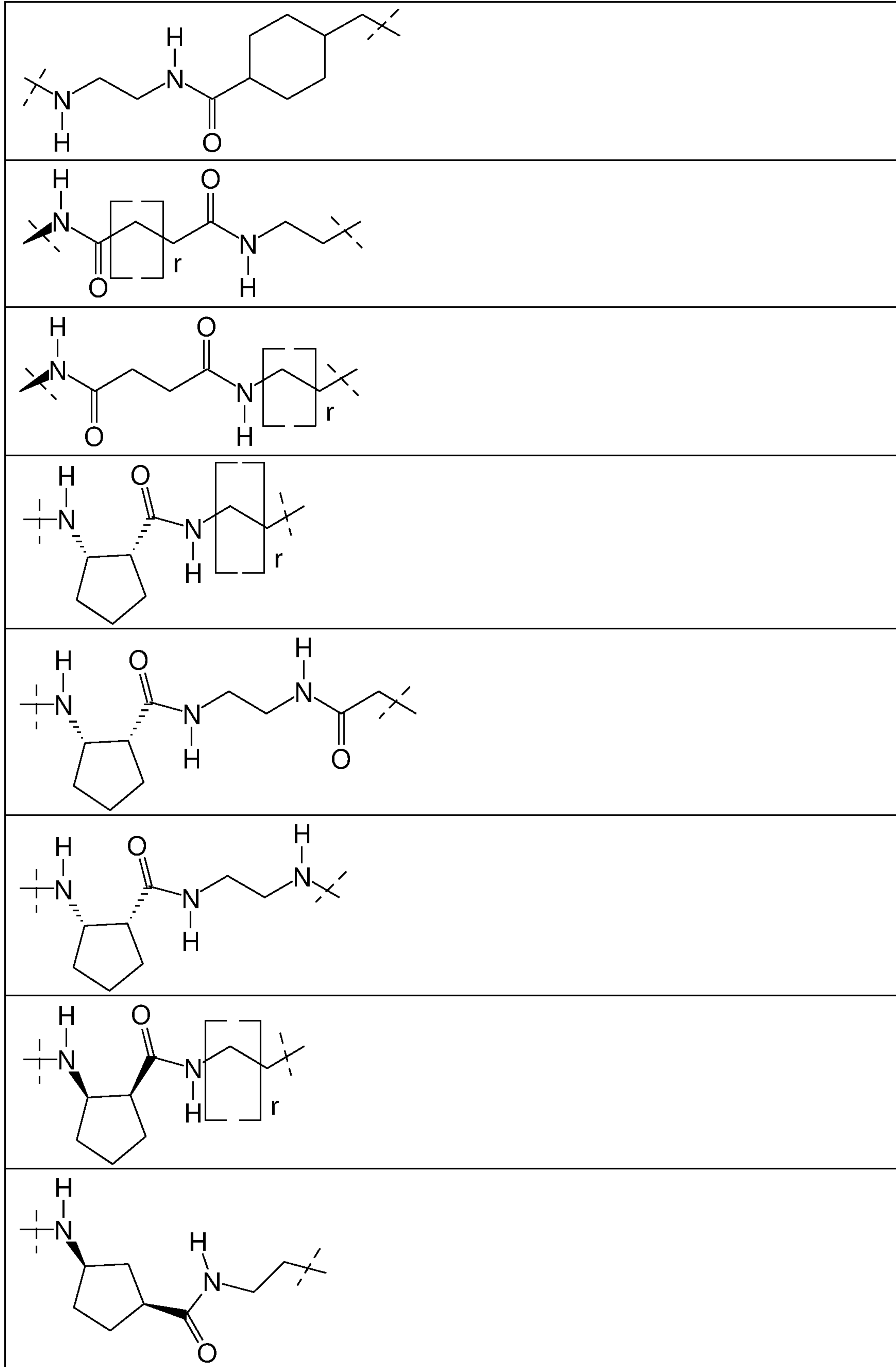
20 o is 0 or 1; and

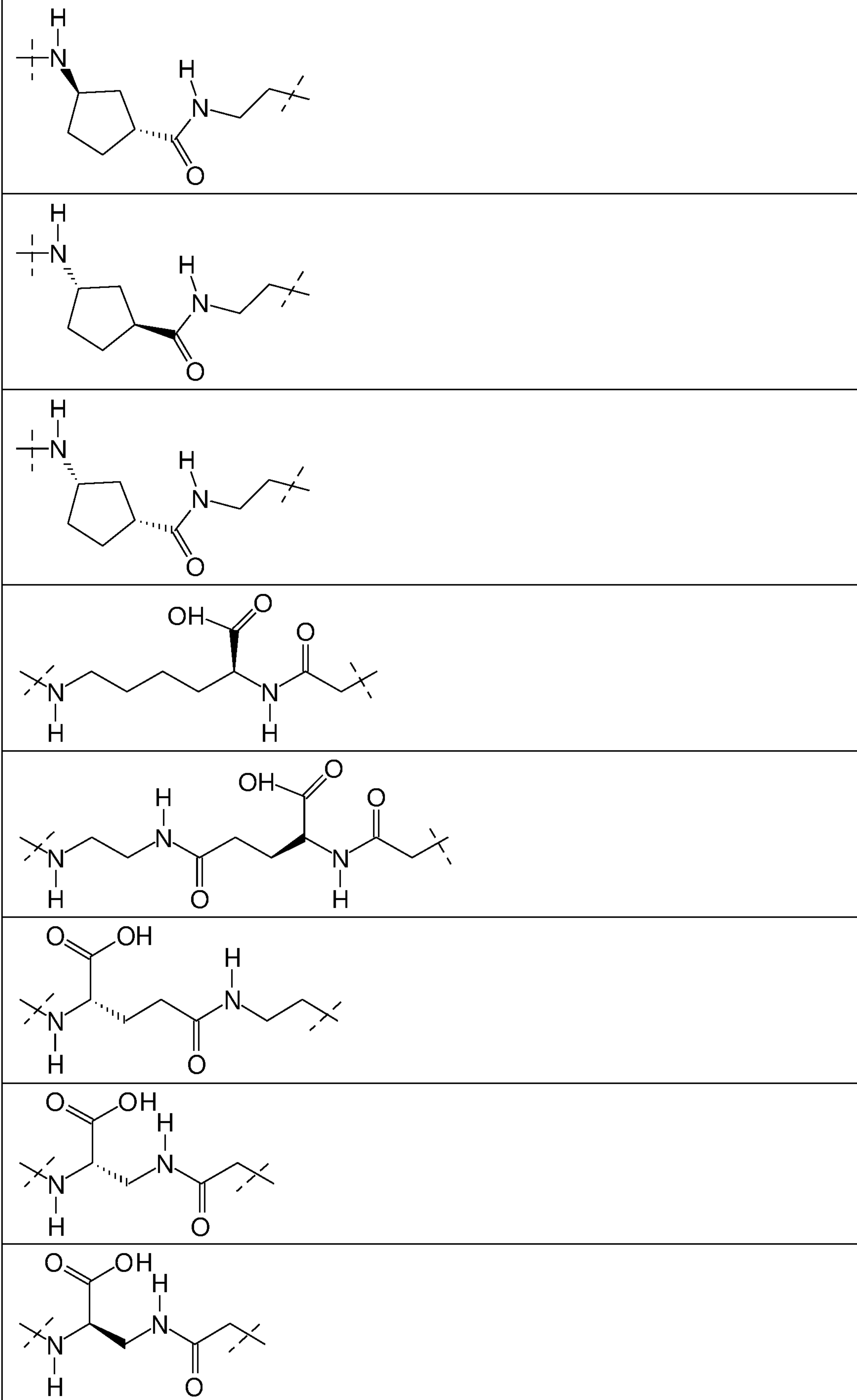
G<sub>3</sub> represents  $-(\text{CH}_2\text{CH}_2\text{O})_s(\text{CH}_2)_t(\text{CONH})_u\text{CH}_2\text{CH}_2\text{O})_v(\text{CH}_2)_w-$ , where

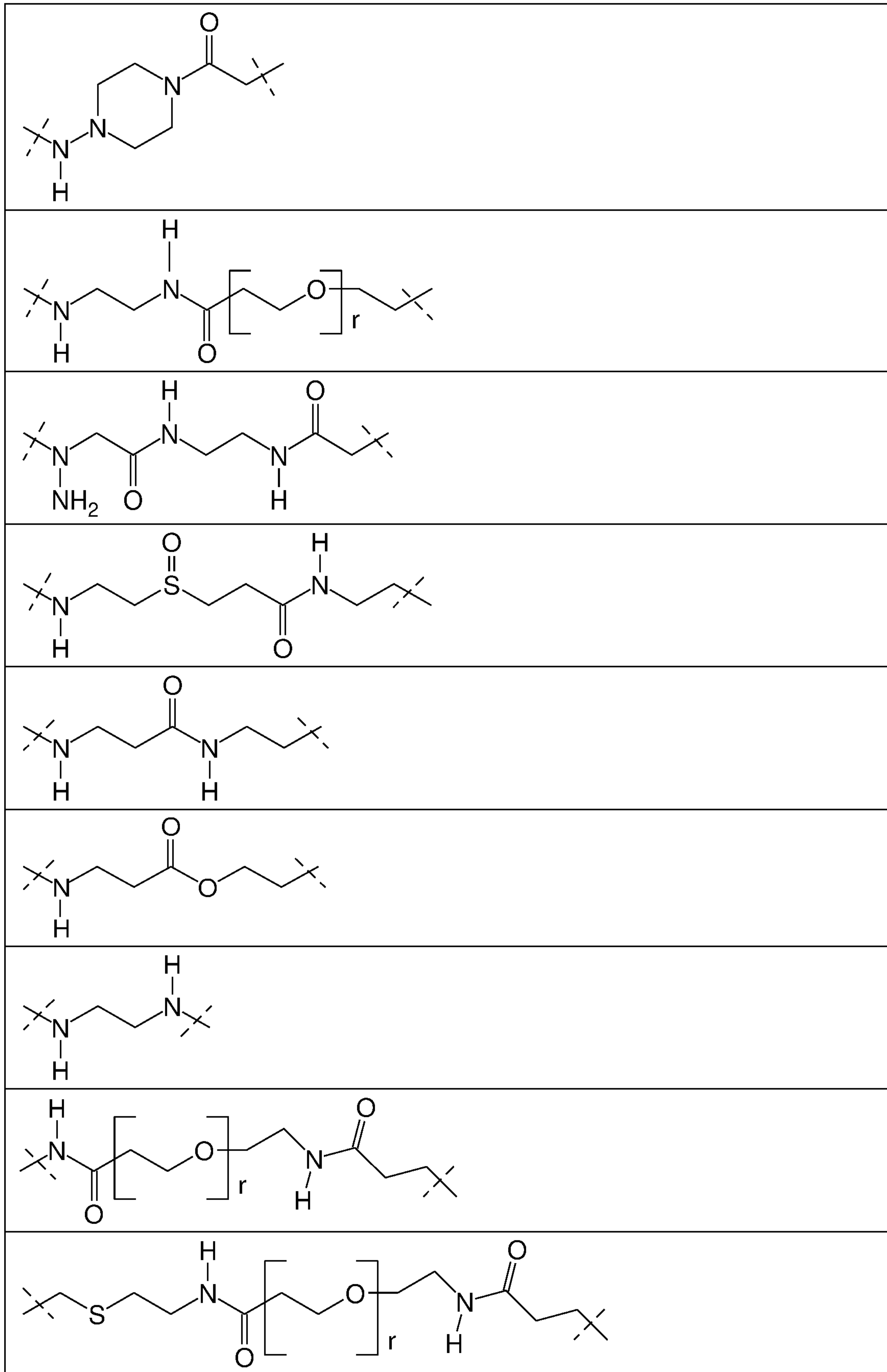
s, t, v and w each independently of one another are from 0 to 25 20 and u is 0 or 1.

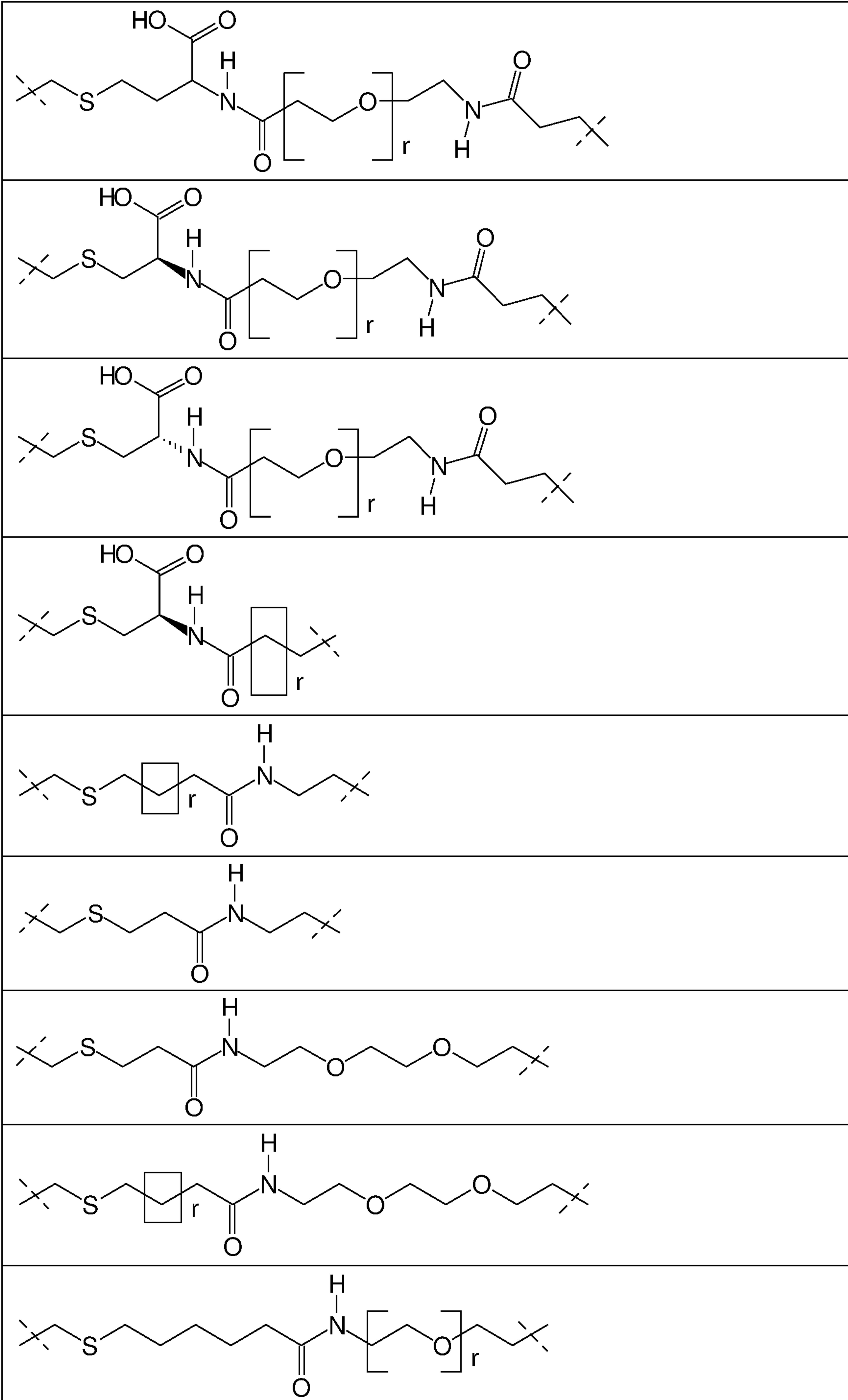
Preferred groups L<sub>1</sub> in the formula  $\text{\$}-(\text{CO})_m\text{-L}_1\text{-L}_2\text{-}\text{\$}$  above are those below, where r in each case independently of one another represents a number from 0 to 20, preferably from 0 to 15, 30 particularly preferably from 1 to 20, especially preferably from 2 to 10:

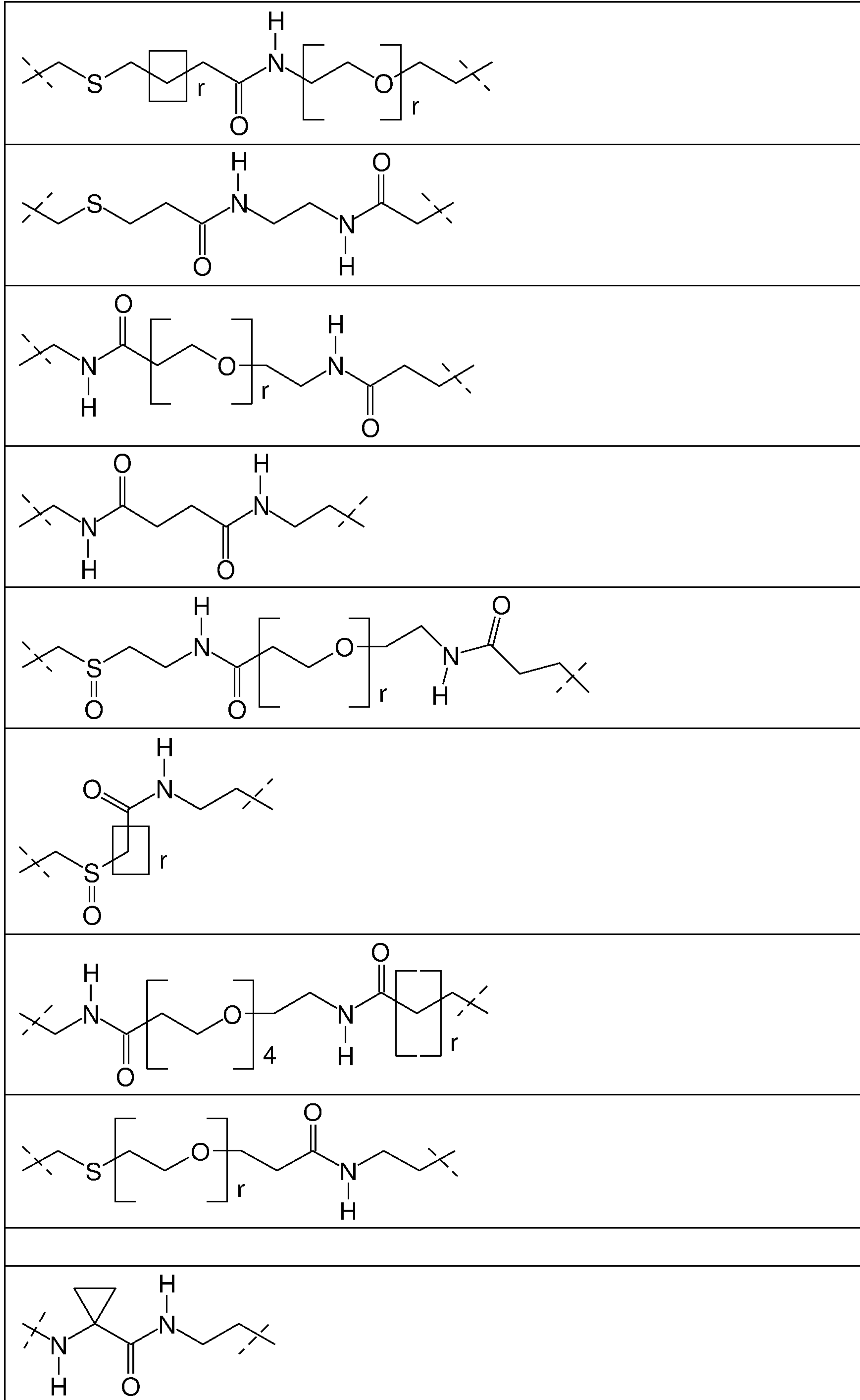


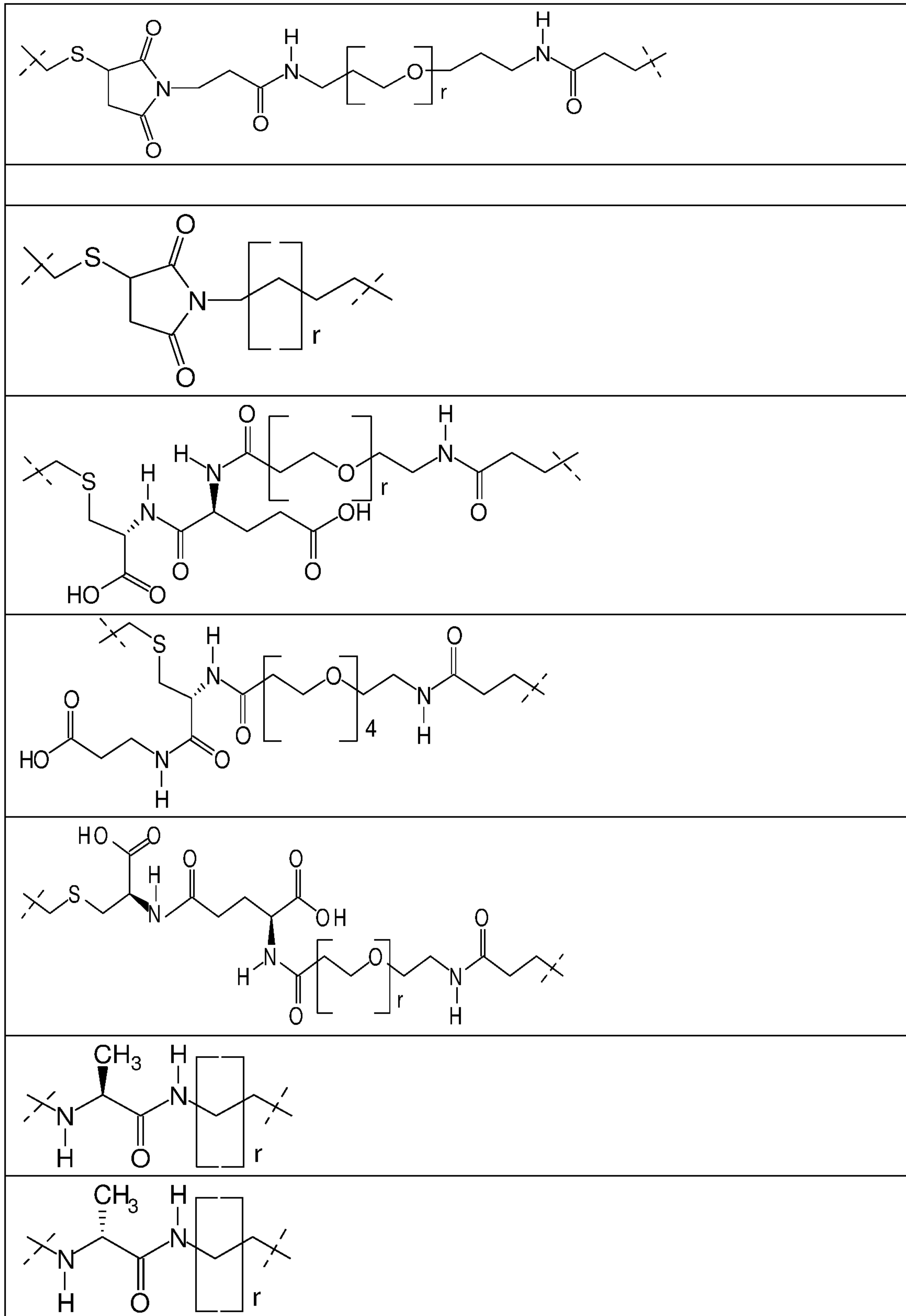












Further examples of L1 are given in Table C, in which this group is highlighted in a box.

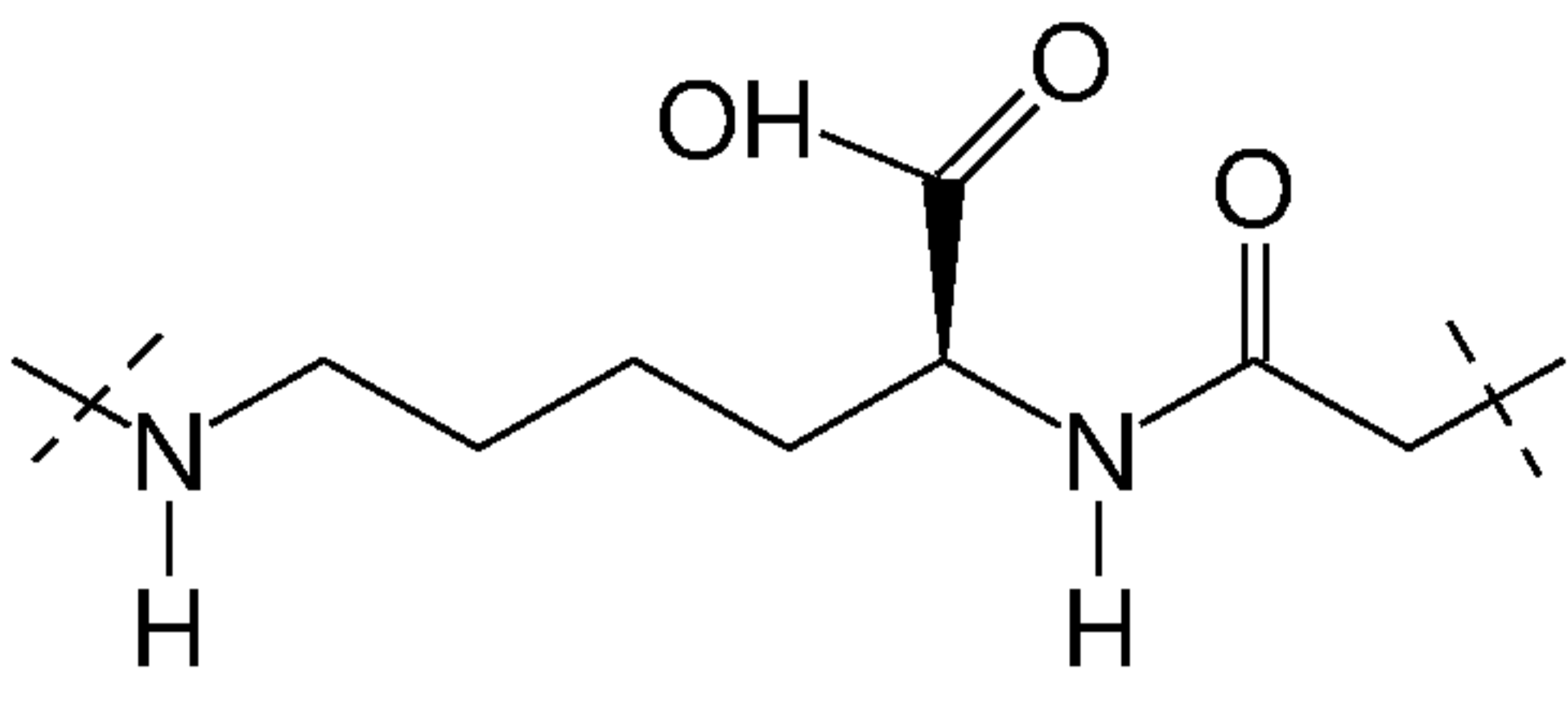
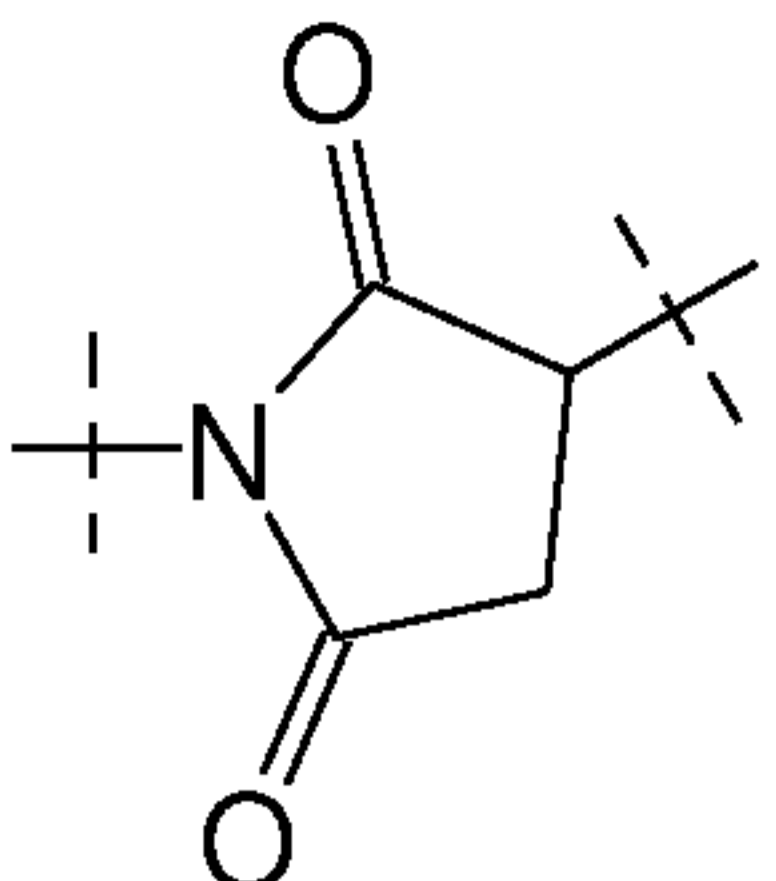
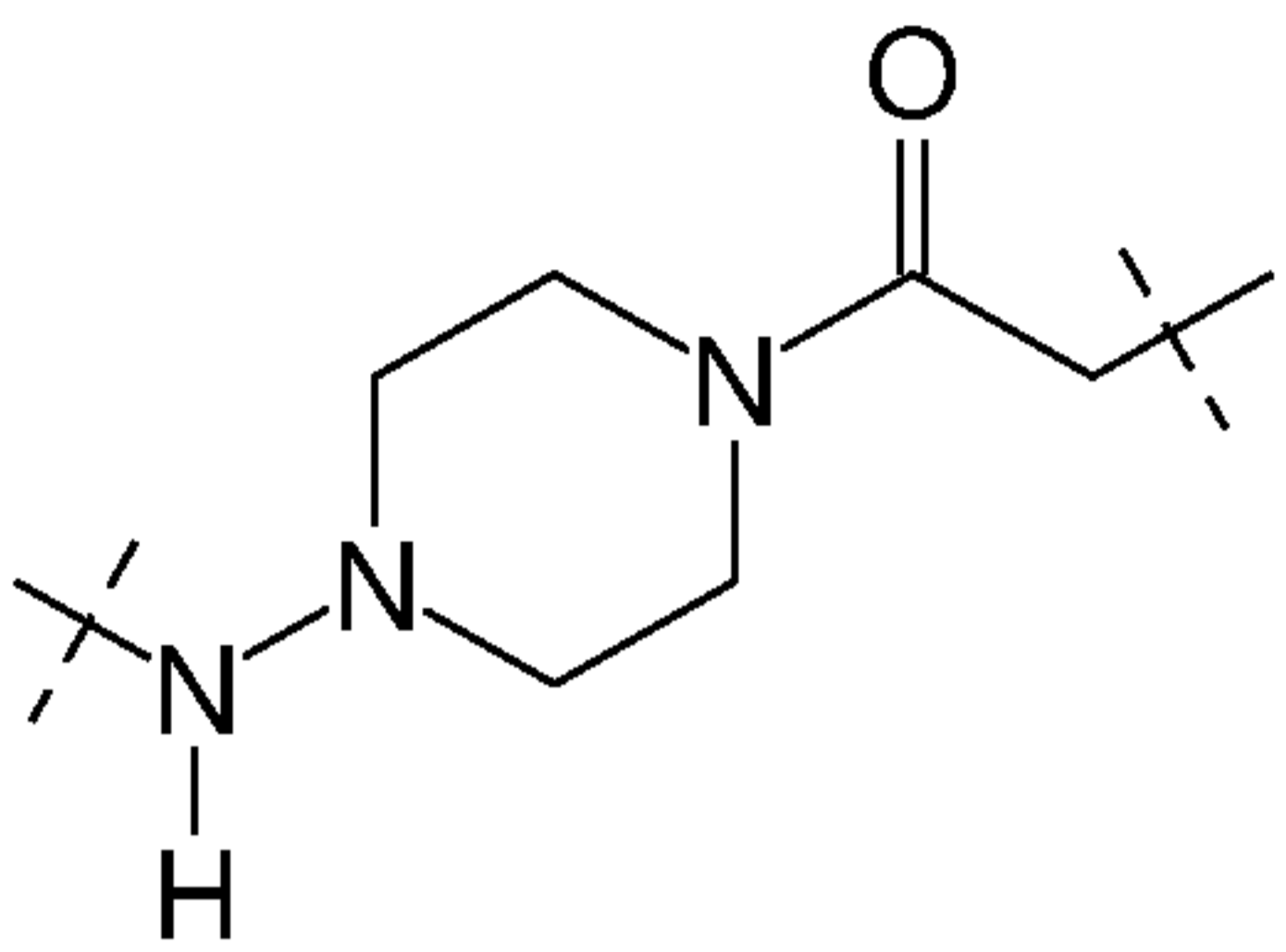
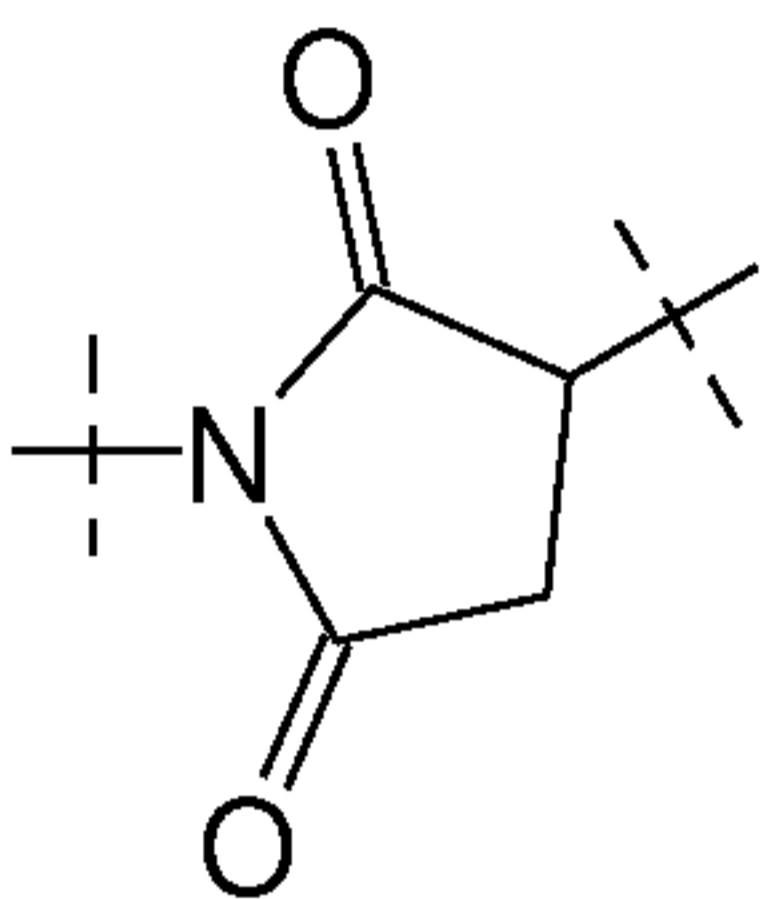
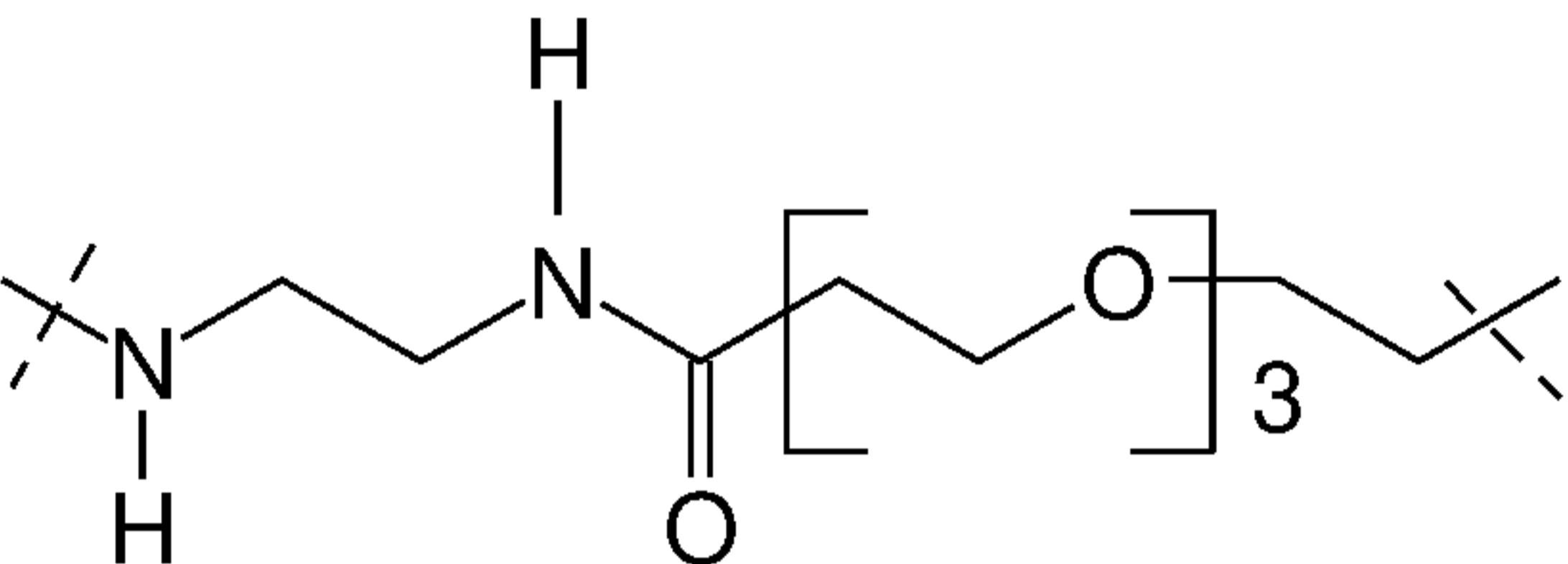
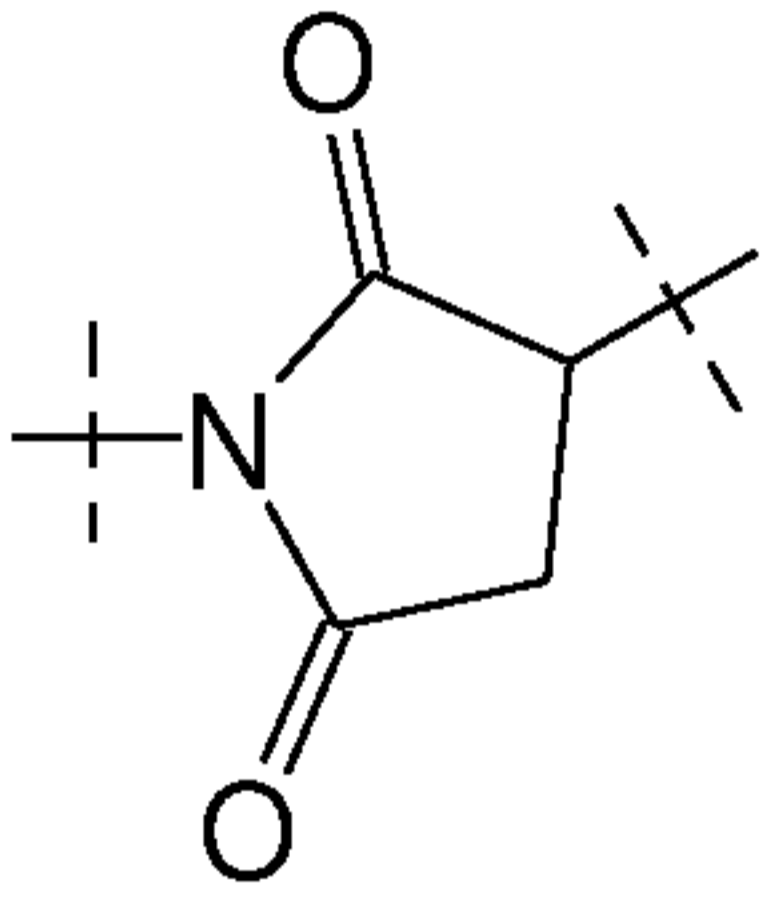
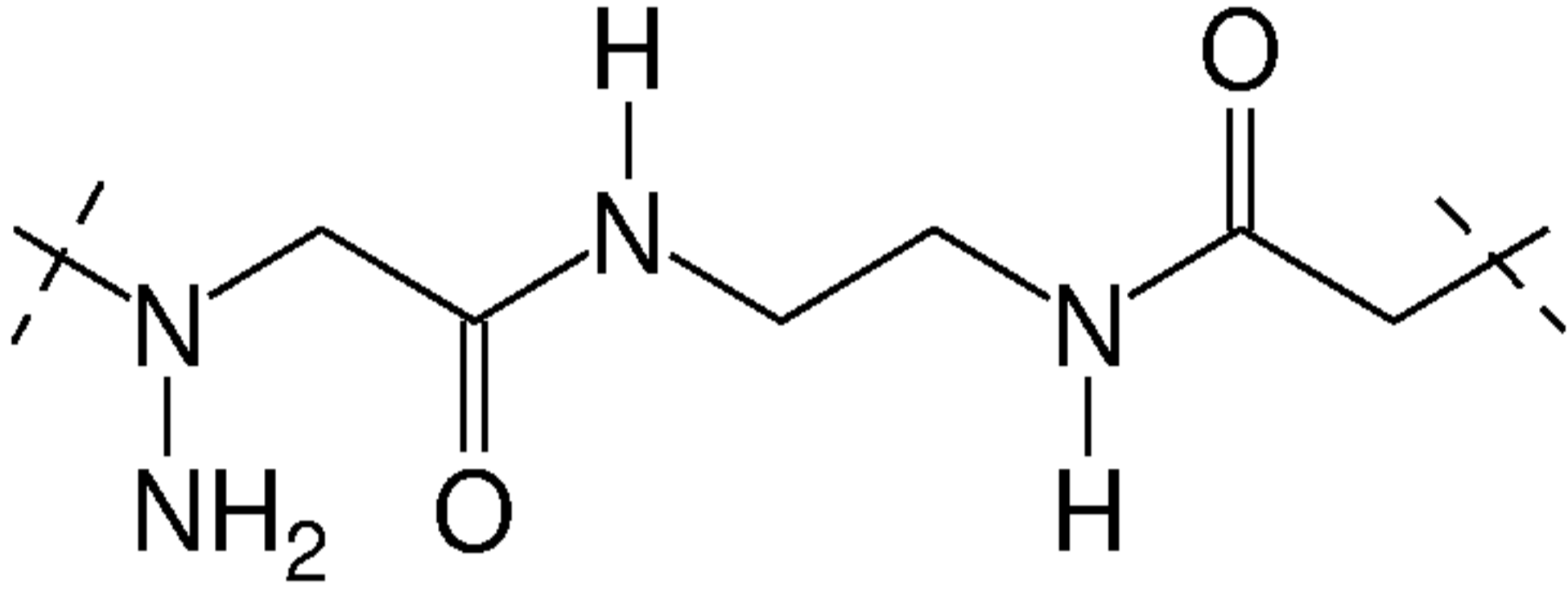
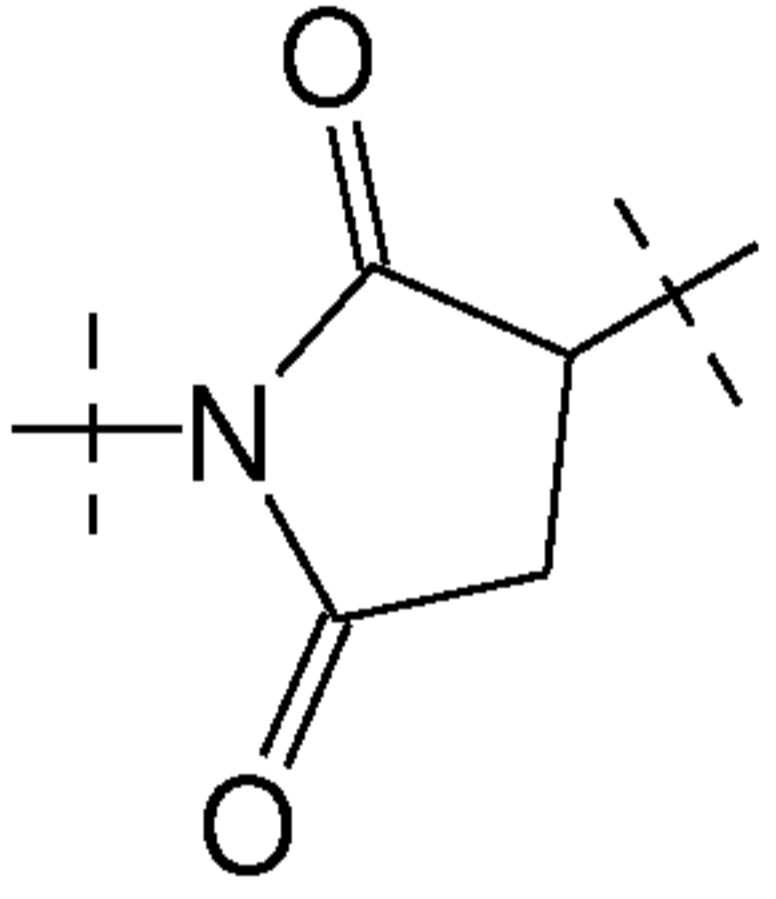
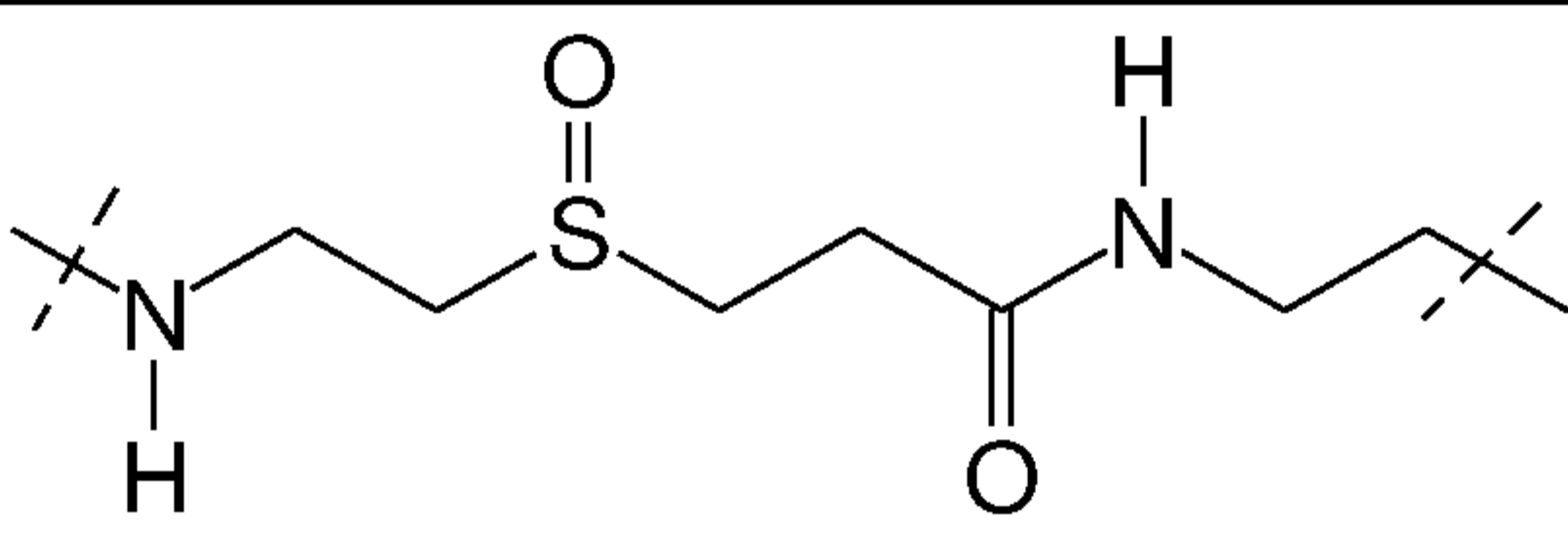
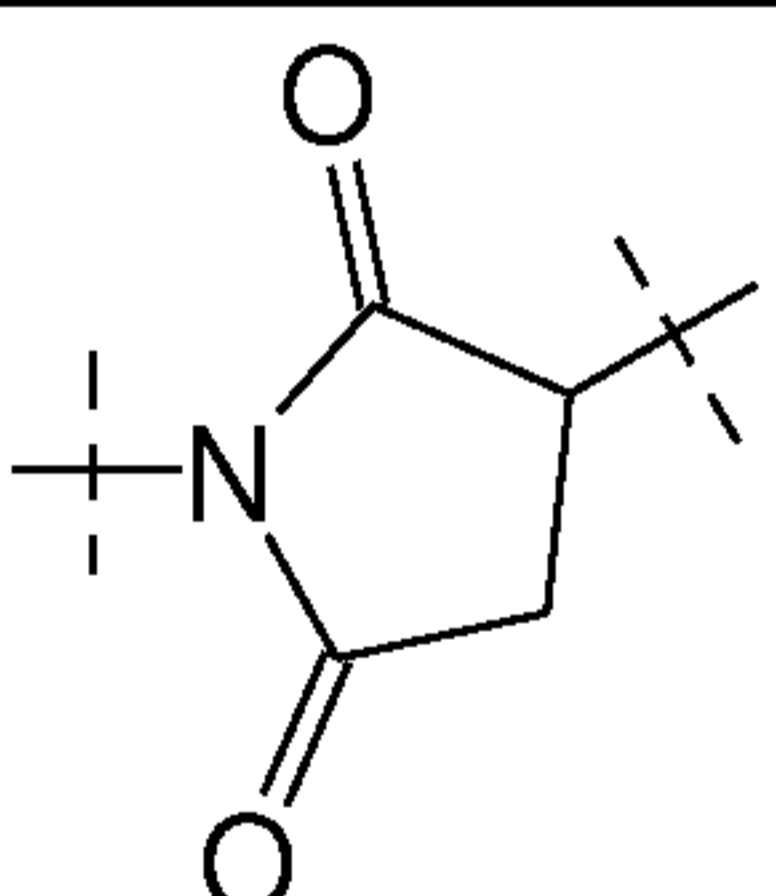
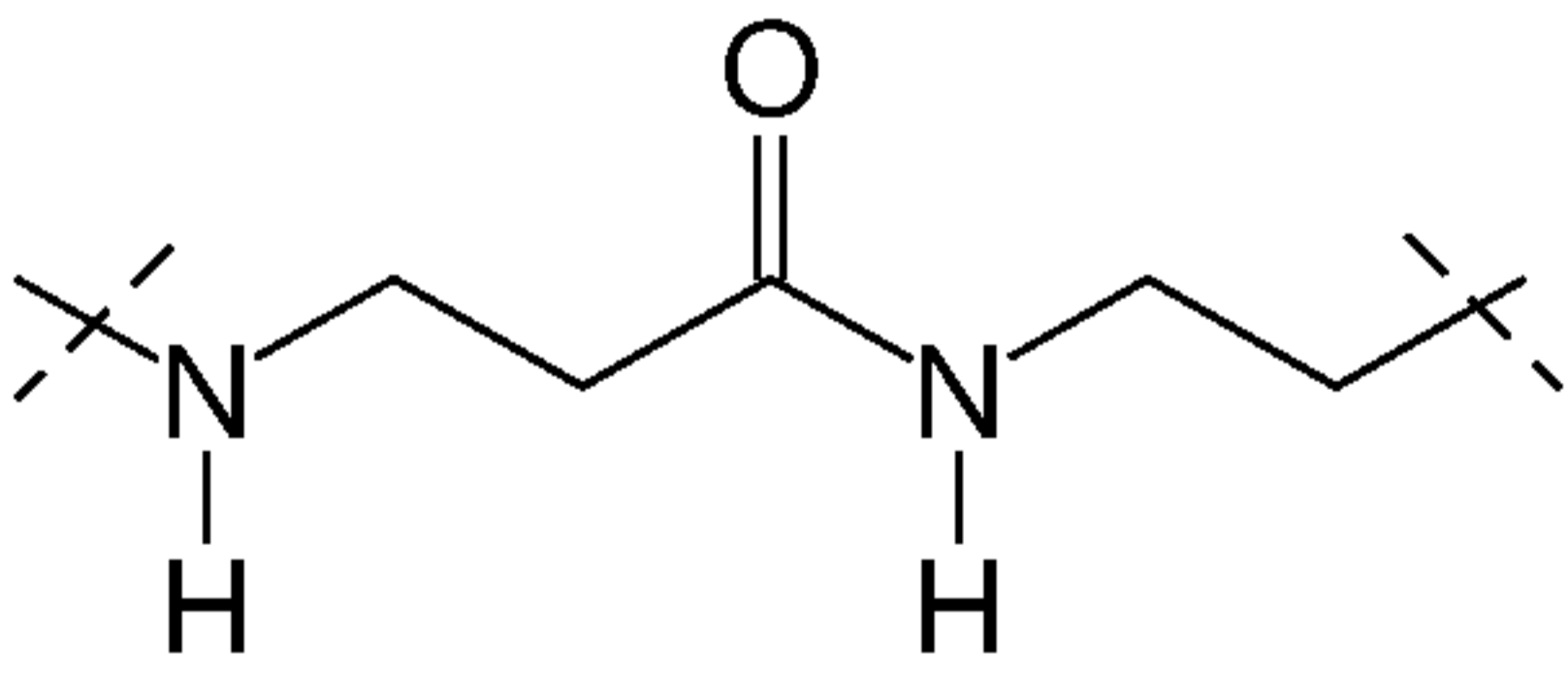
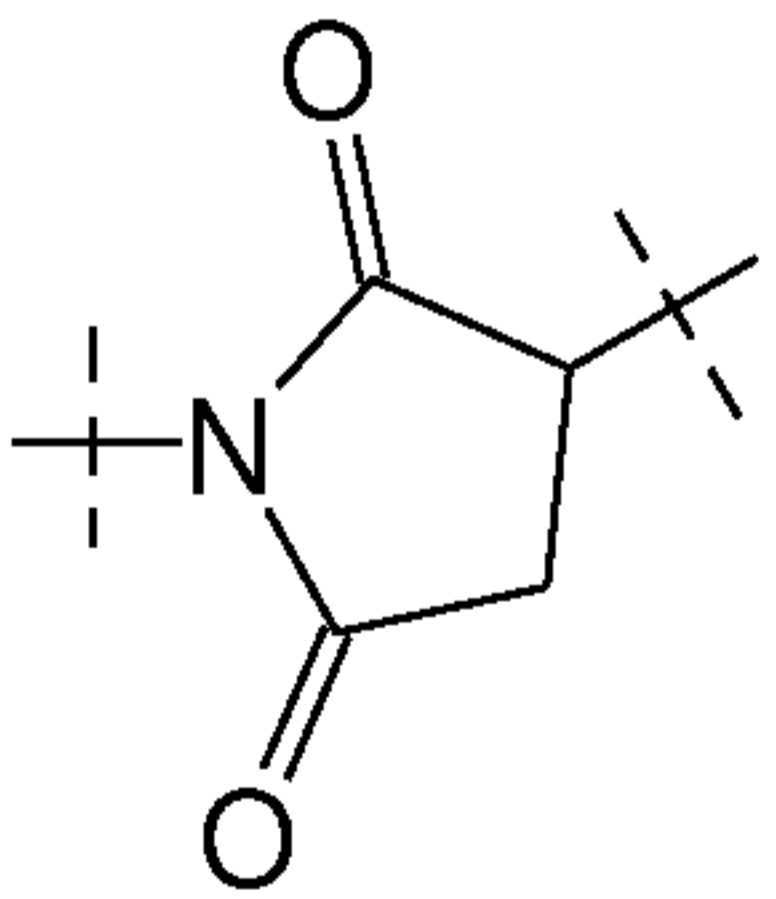
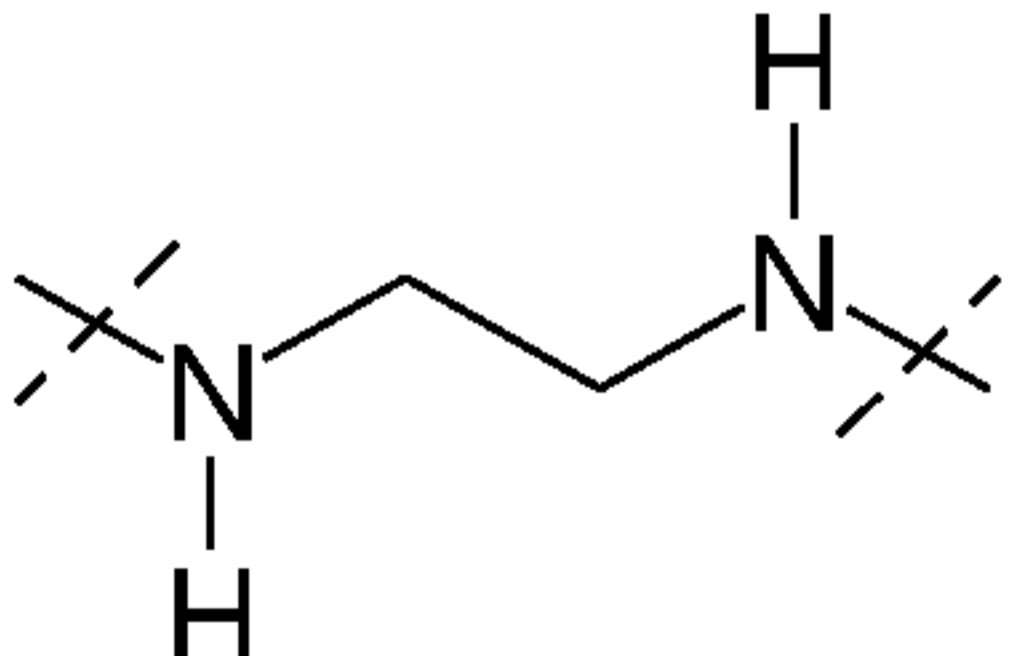
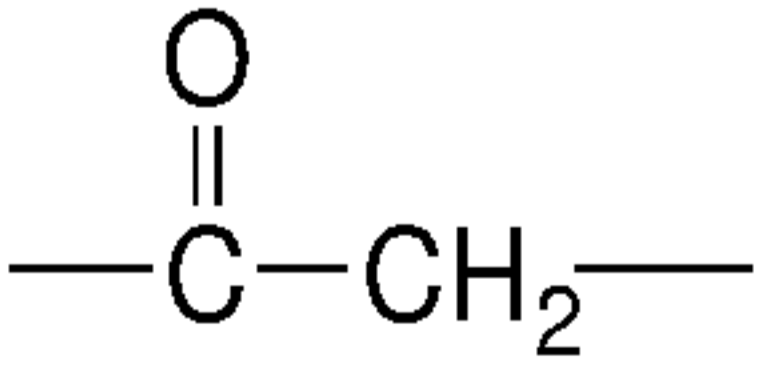
5 Examples of a linker moiety L1 are given in Tables A and A'

below. The table furthermore states with which group L2 these examples of L1 are preferably combined, and also the preferred coupling point ( $R^1$ - $R^5$ ) and the preferred value for m, this is whether there is a carbonyl group in front of L1 or not (cf. §-5 (CO) $_m$ -L1-L2-§§). These linkers are preferably coupled to a cysteine residue. The first column furthermore states the example numbers for the cetuximab ADCs in which the linkers in question are used, but which likewise apply in each row for ADCs with other antibodies. If L2 is a succinimide or derived therefrom, this imide may also be fully or partially in the form of the hydrolysed open-chain succinamide, as described above. Depending on L1, this hydrolysis to open-chain succinamides may be more or less pronounced or not present at all.

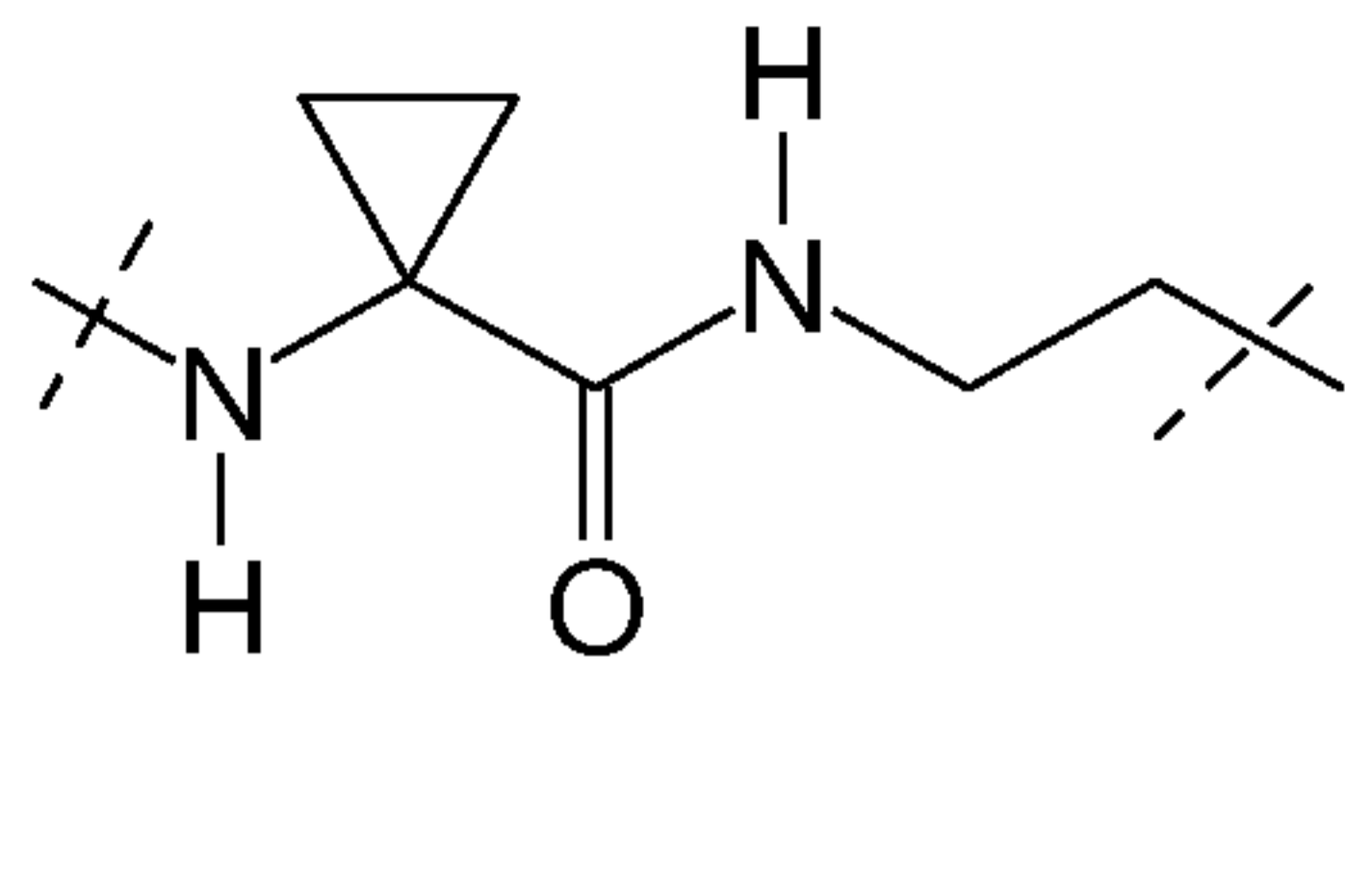
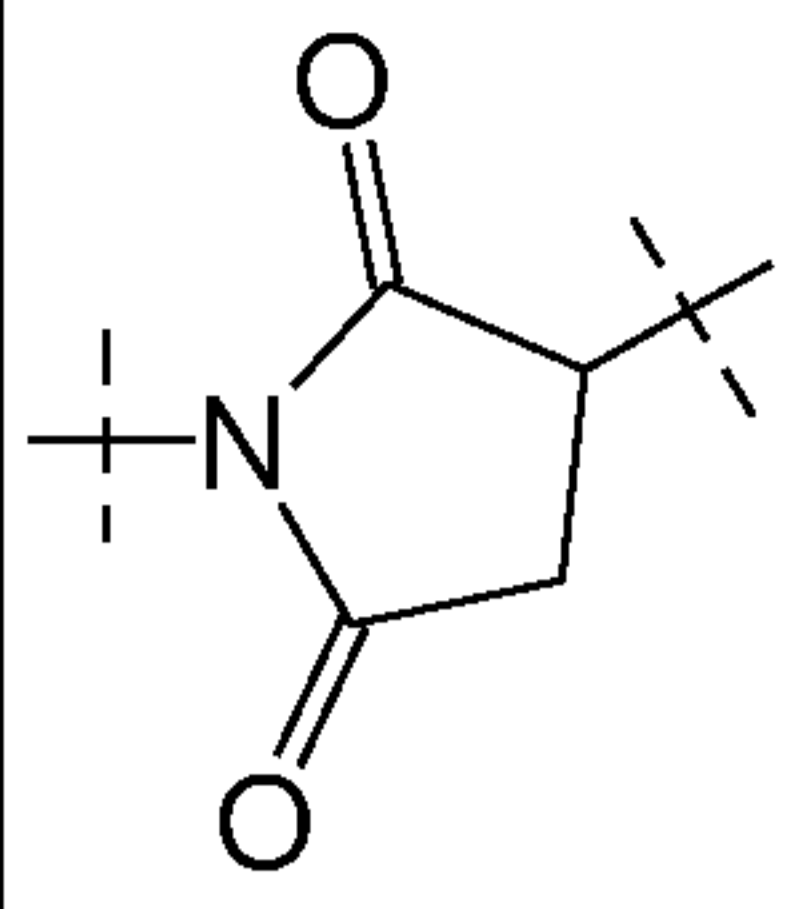
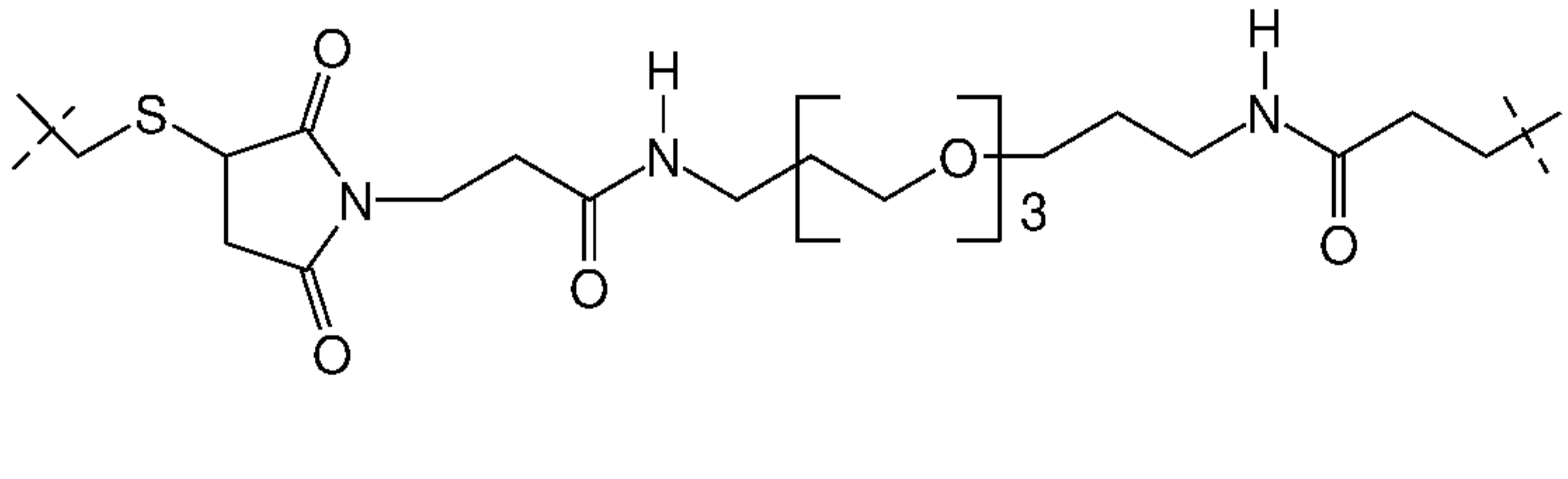
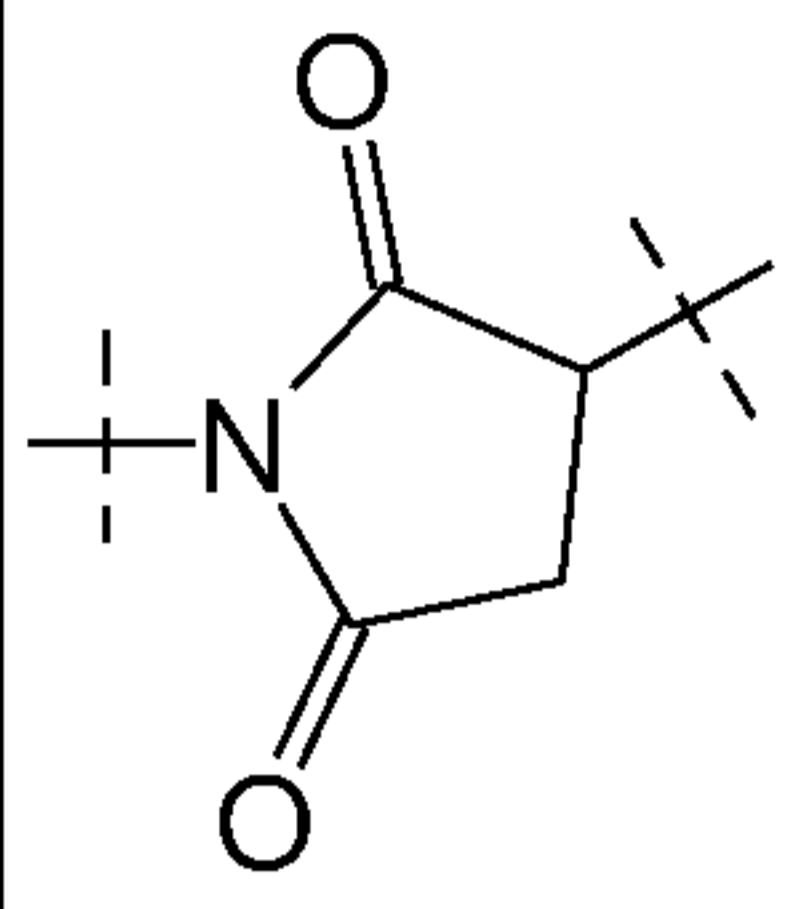
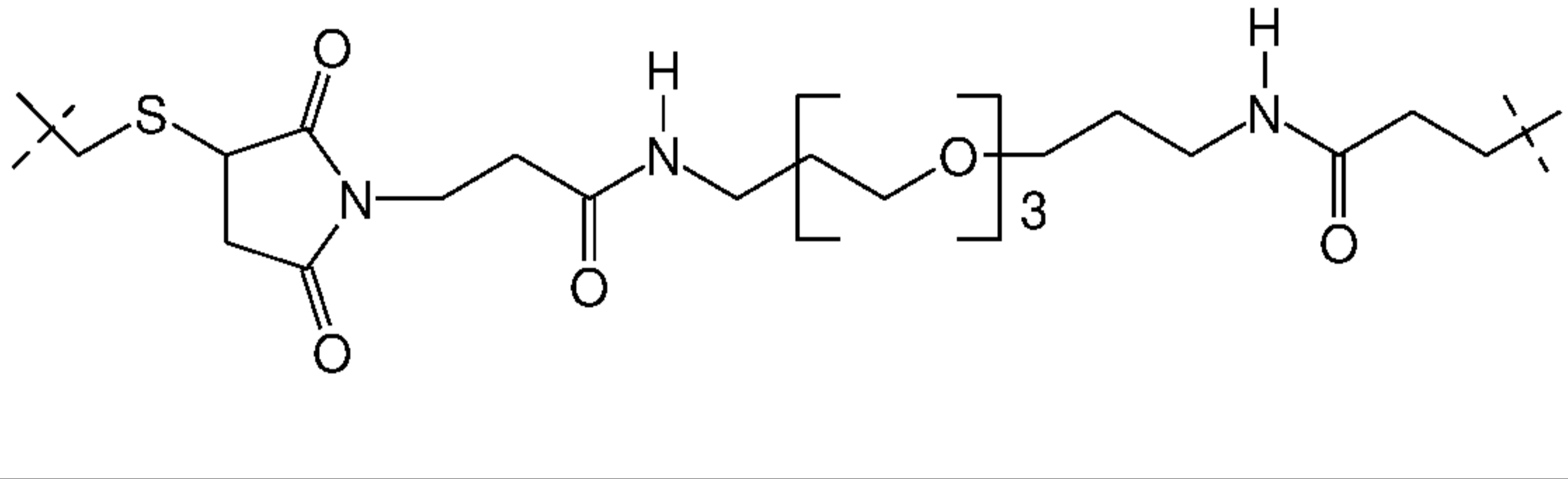
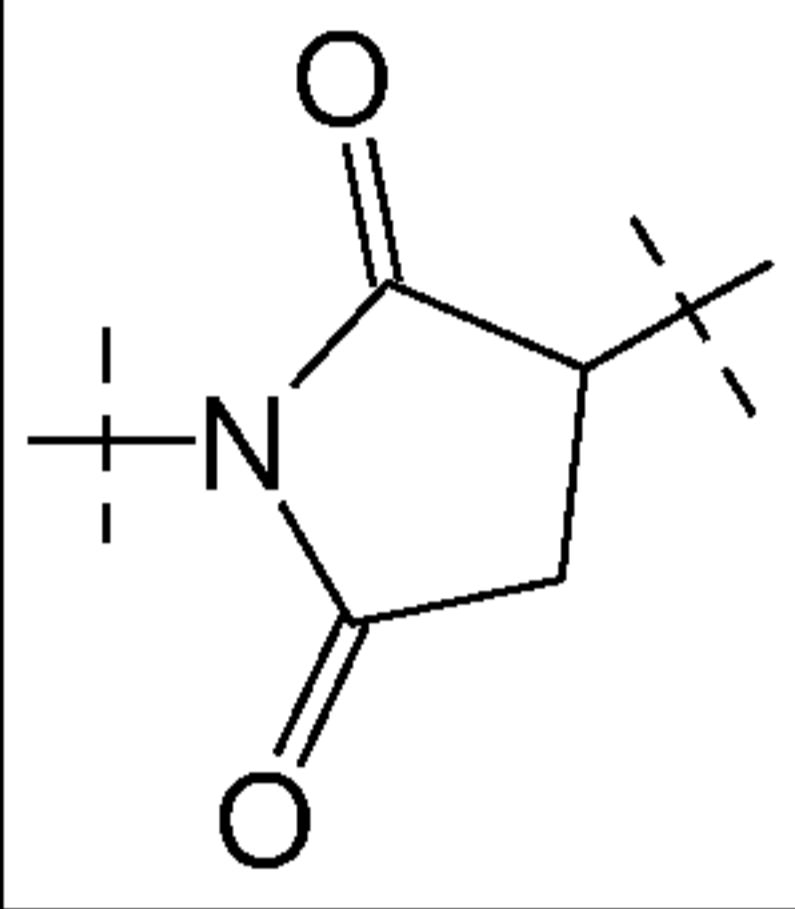
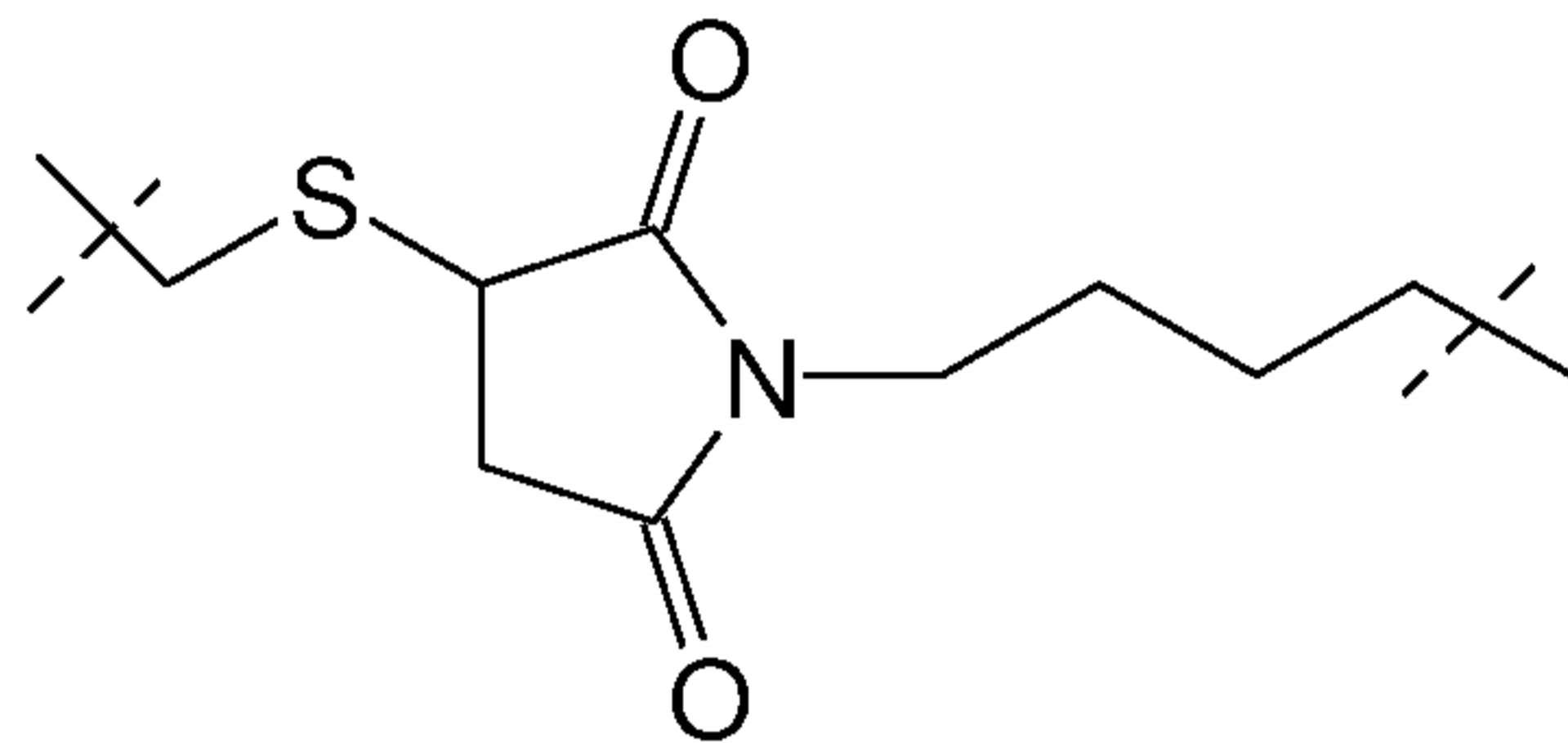
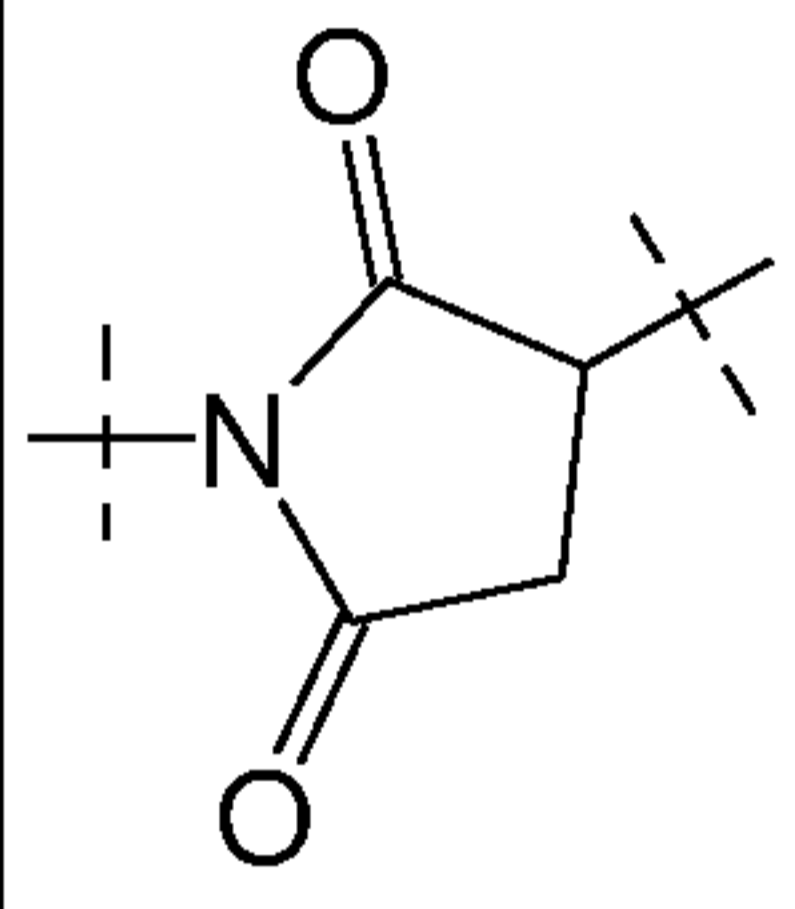
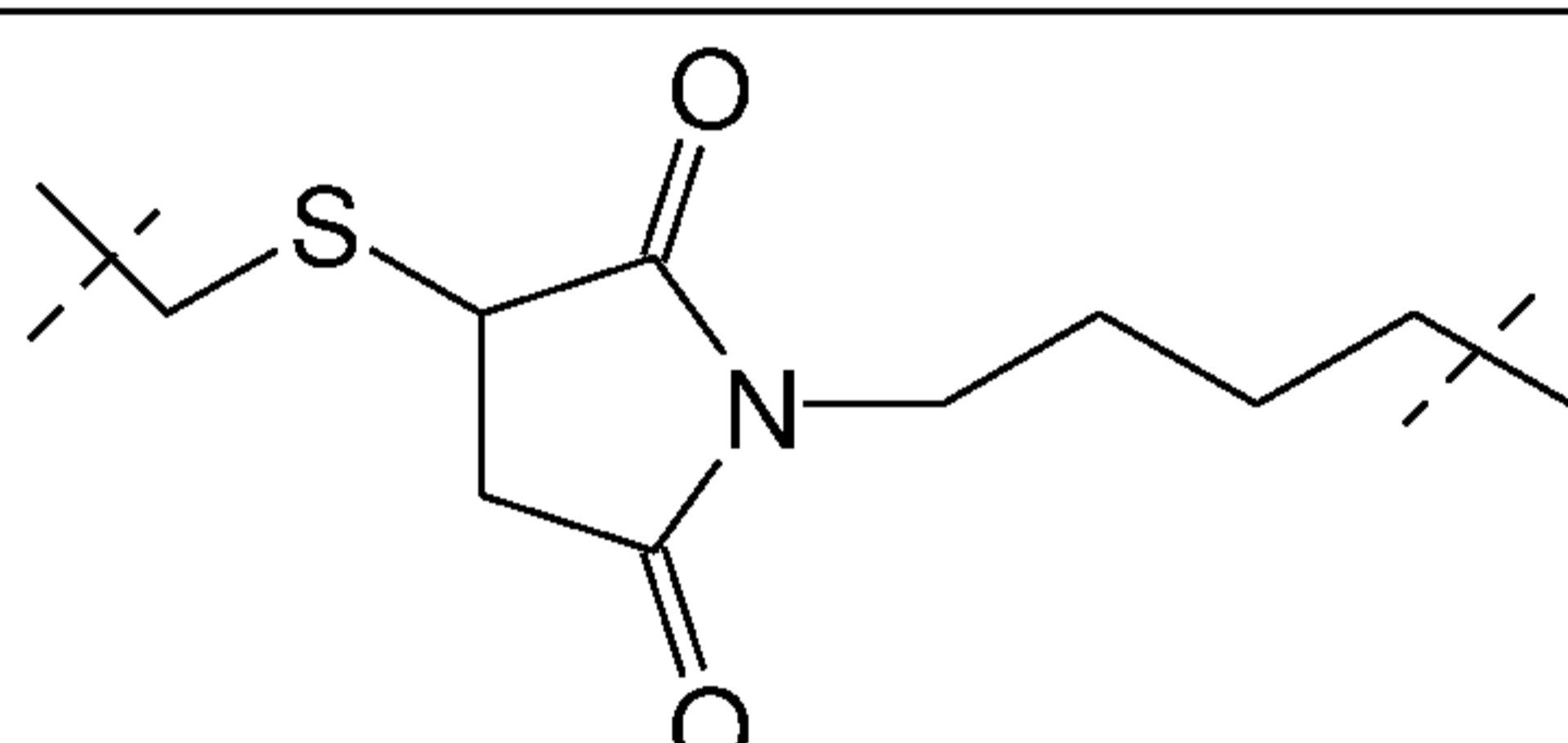
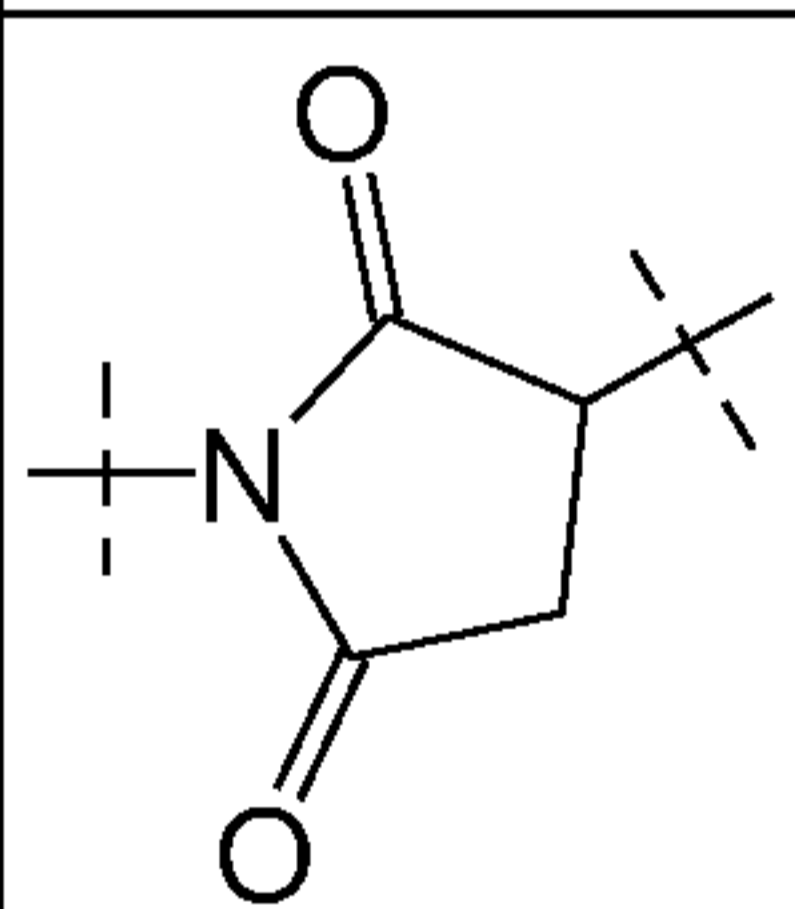
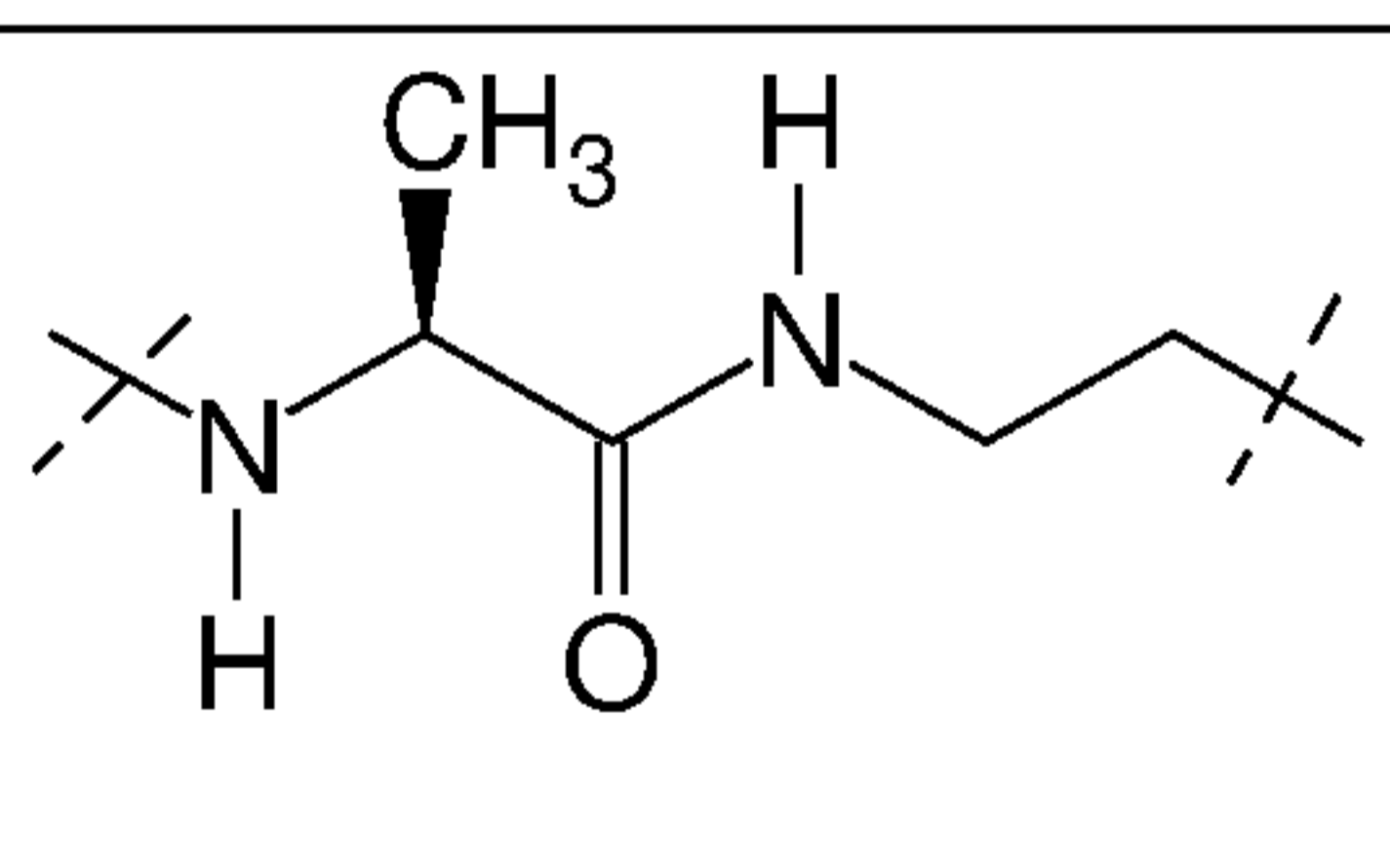
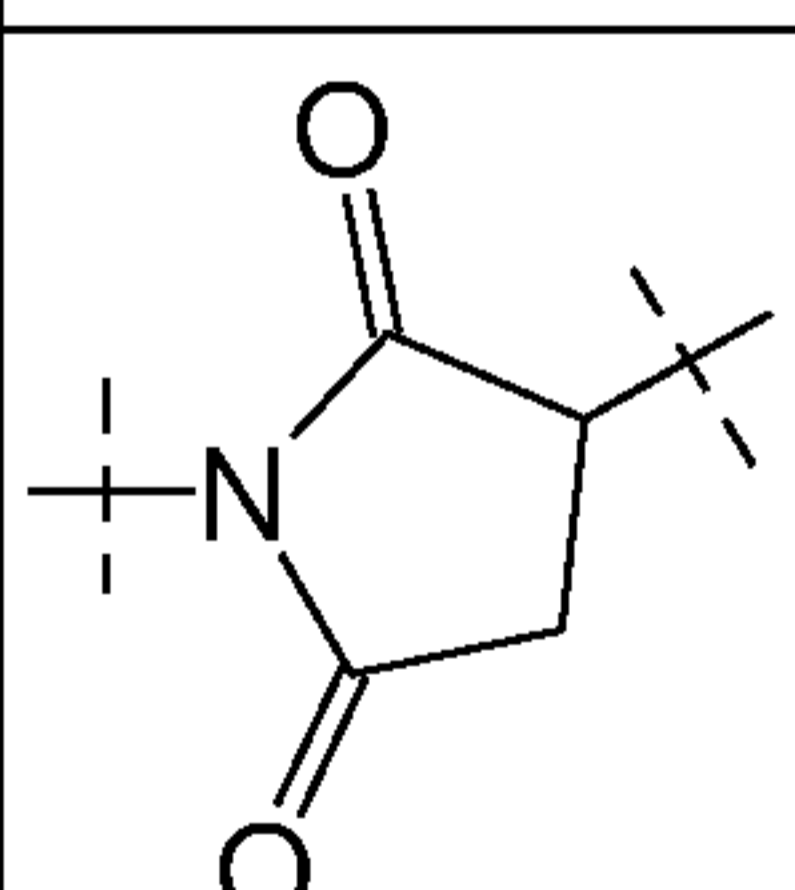
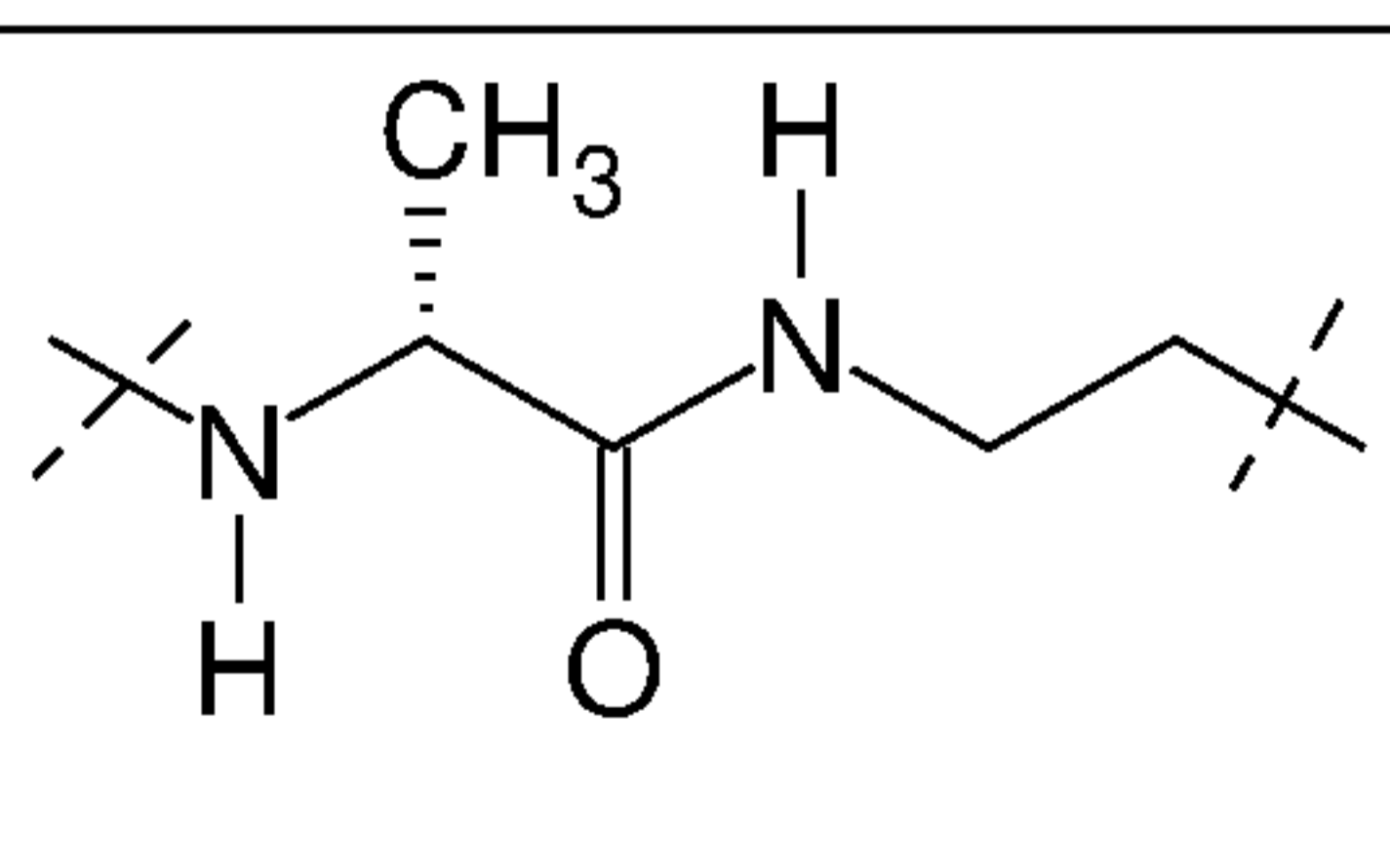
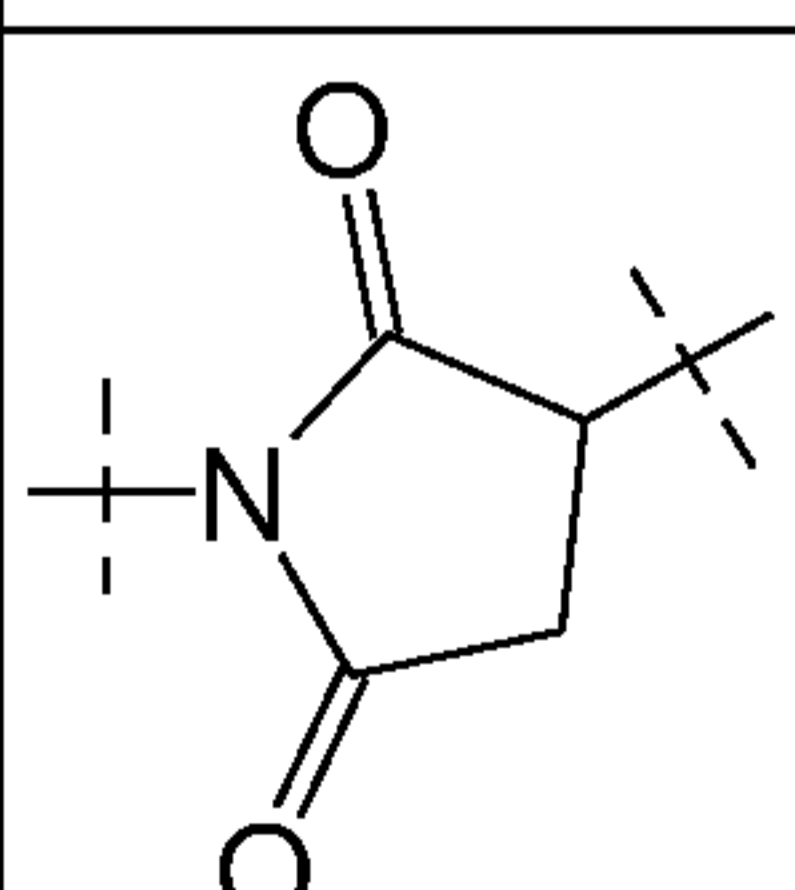
15 Table A

Ex.	Subst.	m	L1	L2
2	$R^1$	1		
3	$R^1$	1		
4	$R^1$	1		
5	$R^1$	1		

6	R <sup>1</sup>	1		
7	R <sup>1</sup>	1		
				See note **
9	R <sup>1</sup>	1		
10	R <sup>1</sup>	1		
11	R <sup>1</sup>	1		
12	R <sup>1</sup>	1		
13	R <sup>1</sup>	1		

19	R <sup>1</sup>	1		 See note **
20	R <sup>1</sup>	1		 See note **
21/ 176	R <sup>1</sup>	1		
22	R <sup>1</sup>	1		 See note **
30	R <sup>1</sup>	1		
32	R <sup>1</sup>	1		
33	R <sup>1</sup>	1		

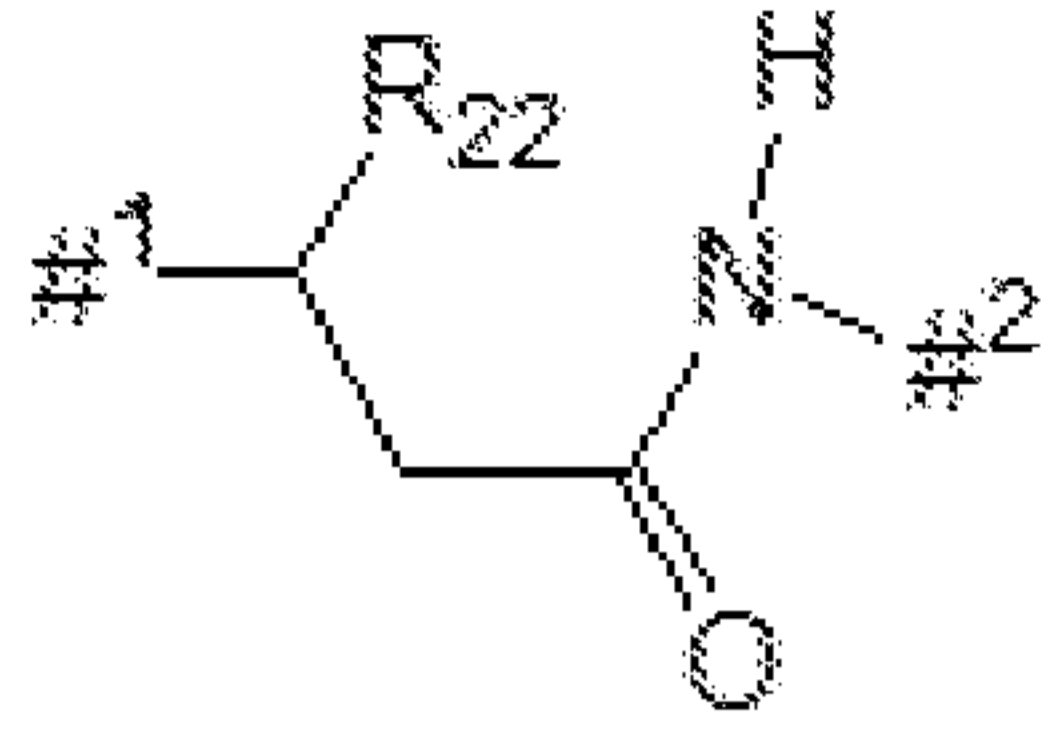
37	$R^2-R^4$ *	0		
38	$R^3$	0		
40	$R^2-R^4$ *	0		
41	$R^3$	0		
42	$R^2-R^4$ *	1		
44	$R^2-R^4$ *	0		
48	$R^3$	0		
49	$R^3$	0		

50	R <sup>1</sup>	1		
51	R <sup>2</sup>	0		
52	R <sup>2</sup>	0		
53	R <sup>2</sup>	0		
54	R <sup>2</sup>	0		
56	R <sup>1</sup>	1		
57	R <sup>1</sup>	1		

\*R<sup>2</sup> and R<sup>4</sup> form a pyrrolidone ring which is substituted by the linker.

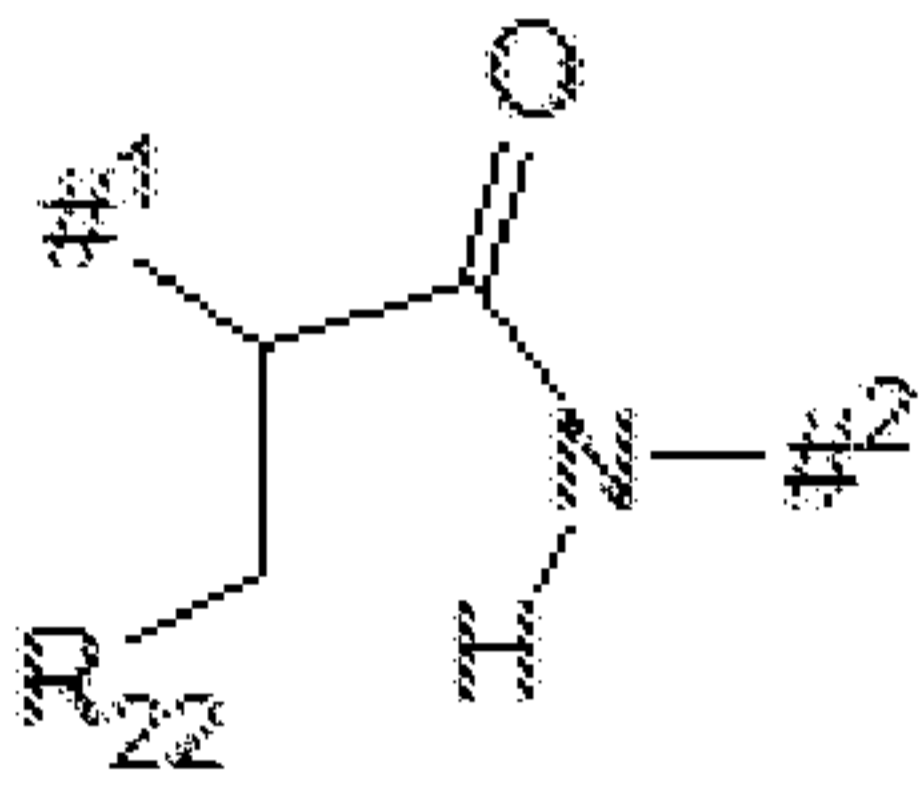
5 \*\*With particular preference, the linkers L1 given in these rows

are attached to a linker L2 selected from:



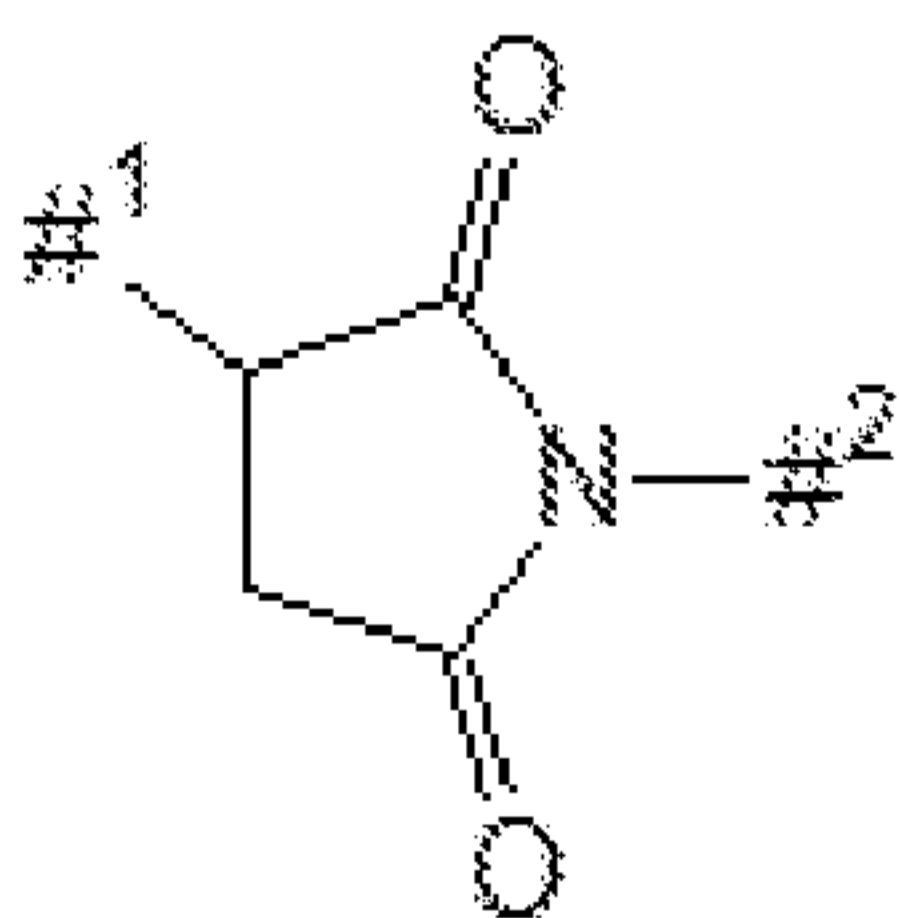
Formula A7

5 and/or



Formula A8

where #<sup>1</sup> denotes the point of attachment to the sulphur atom of  
 10 the binder, #<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>, R<sup>22</sup>  
 preferably represents COOH. In a conjugate according to the  
 invention or in a mixture of the conjugates according to the  
 invention, the bonds to a cysteine residue of the binder are  
 present, to an extent of preferably more than 80%, particularly  
 15 preferably more than 90% (in each case based on the total number  
 of bonds of the linker to the binder), particularly preferably  
 as one of the two structures of the formula A7 or A8. Here, the  
 structures of the formula A7 or A8 are generally present  
 together, preferably in a ratio of from 60:40 to 40:60, based  
 20 on the number of bonds to the binder. The remaining bonds are  
 then present as the structure



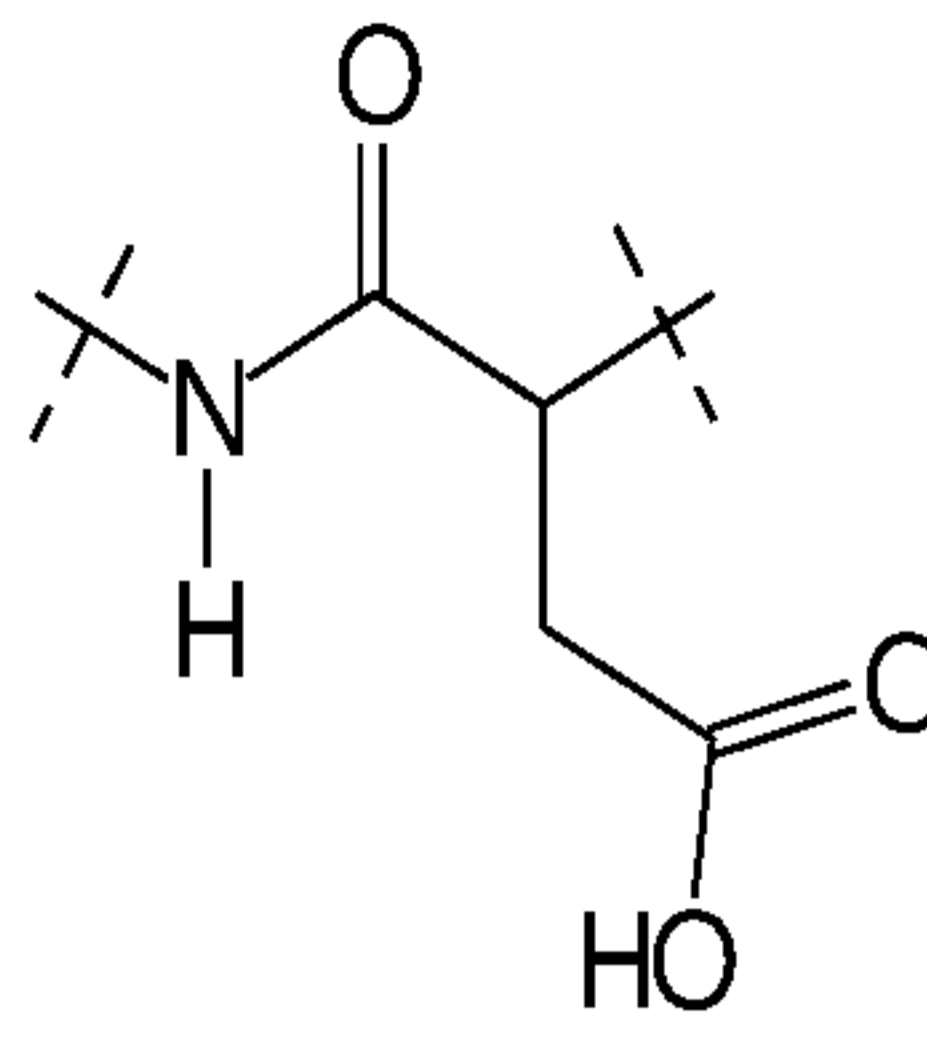
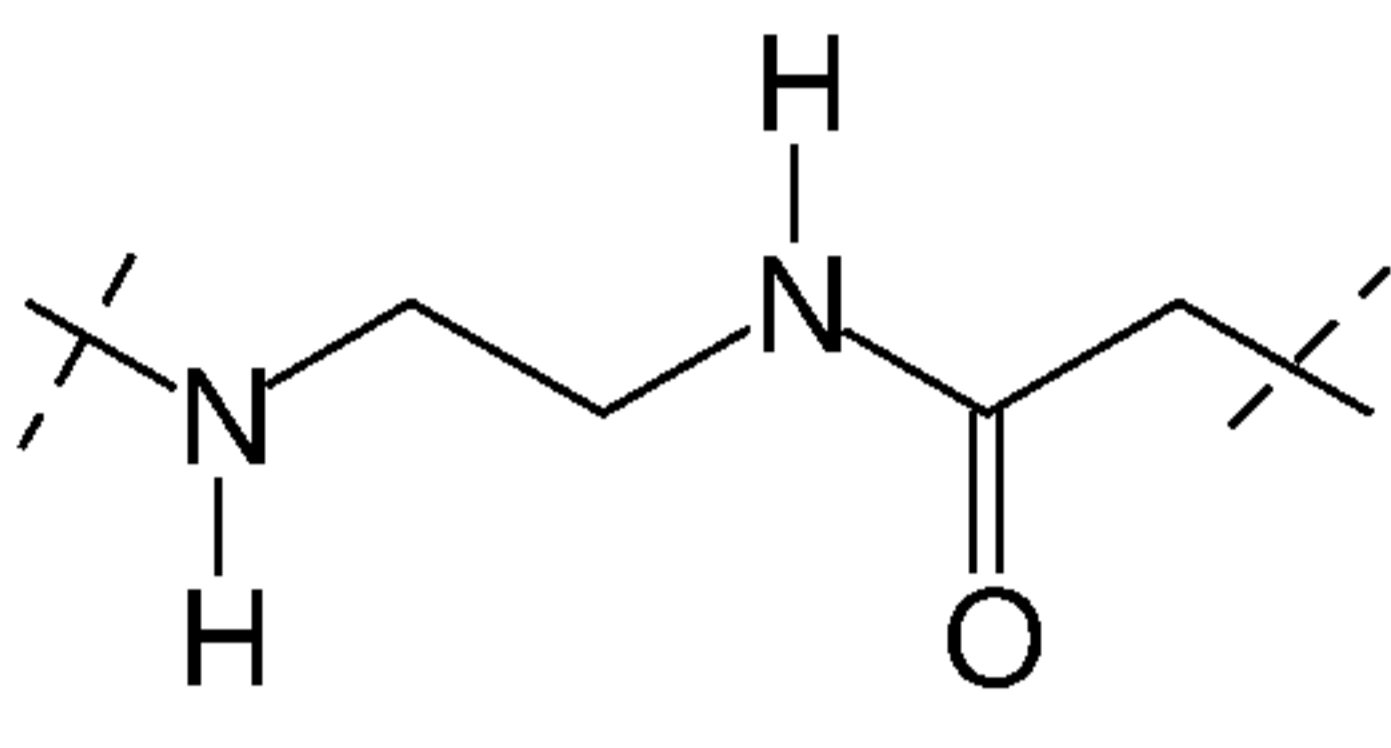
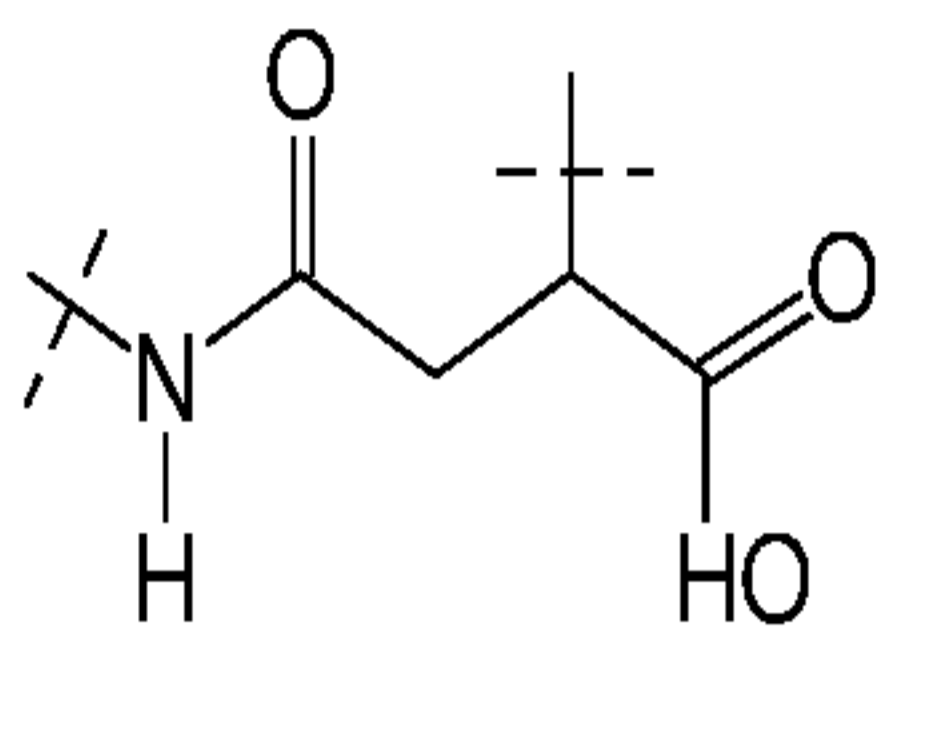
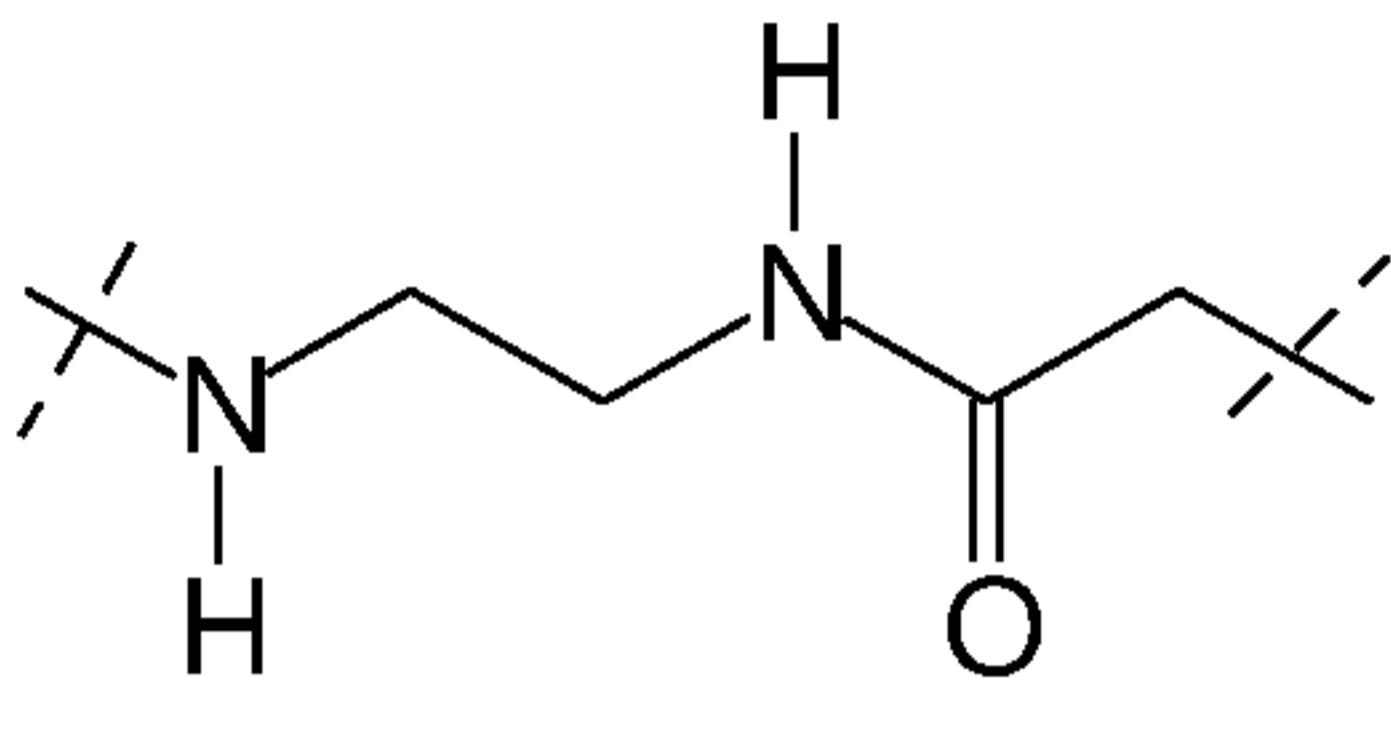
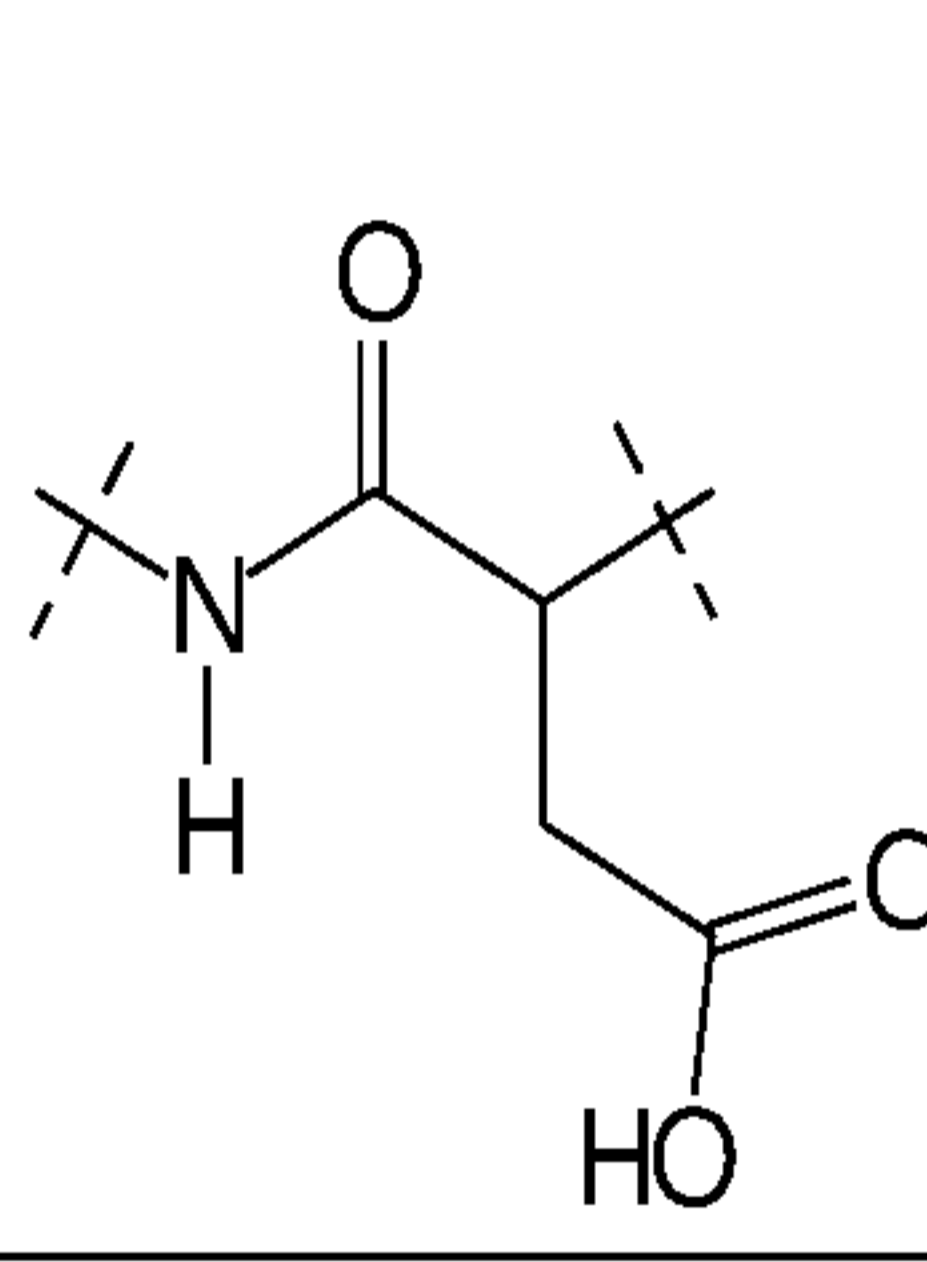
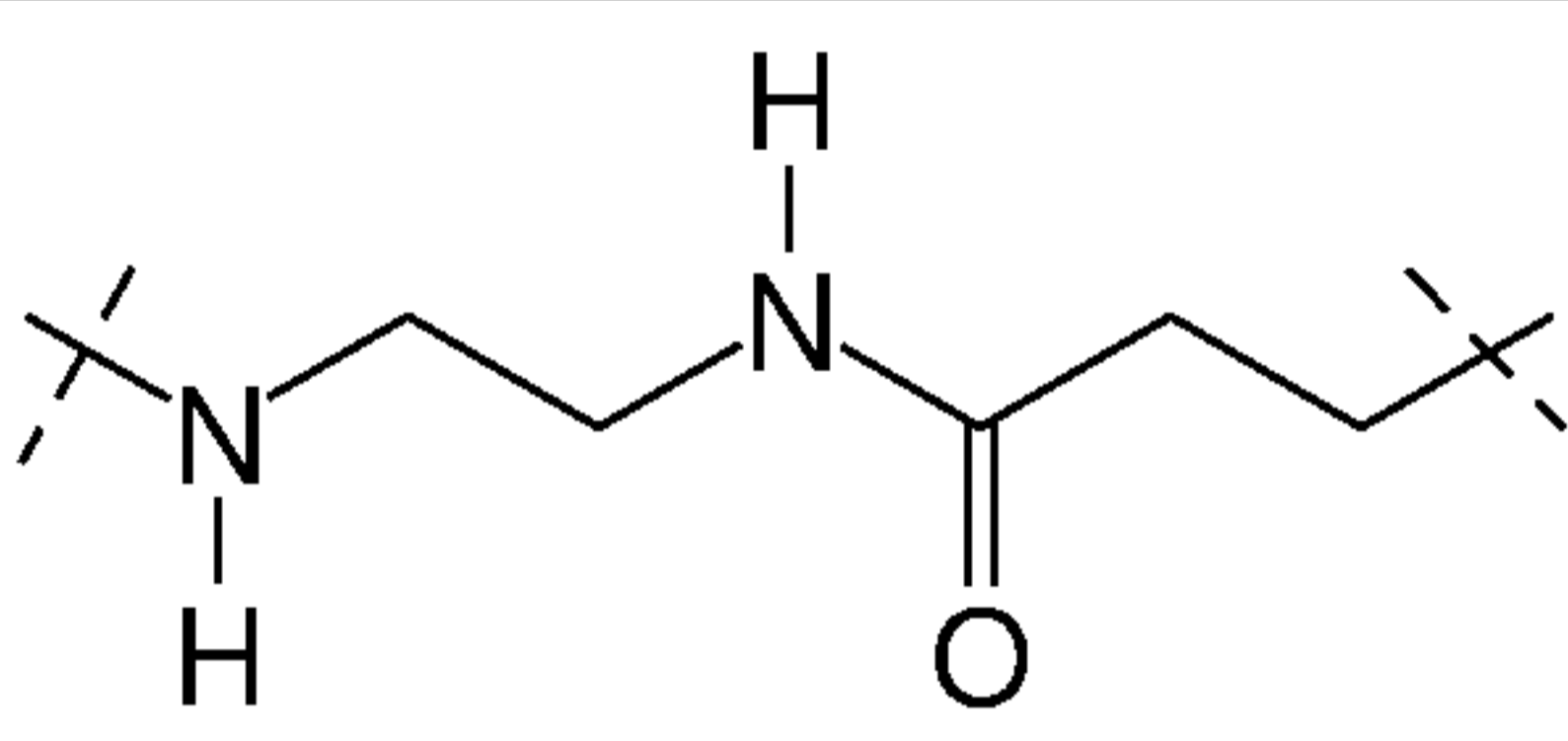
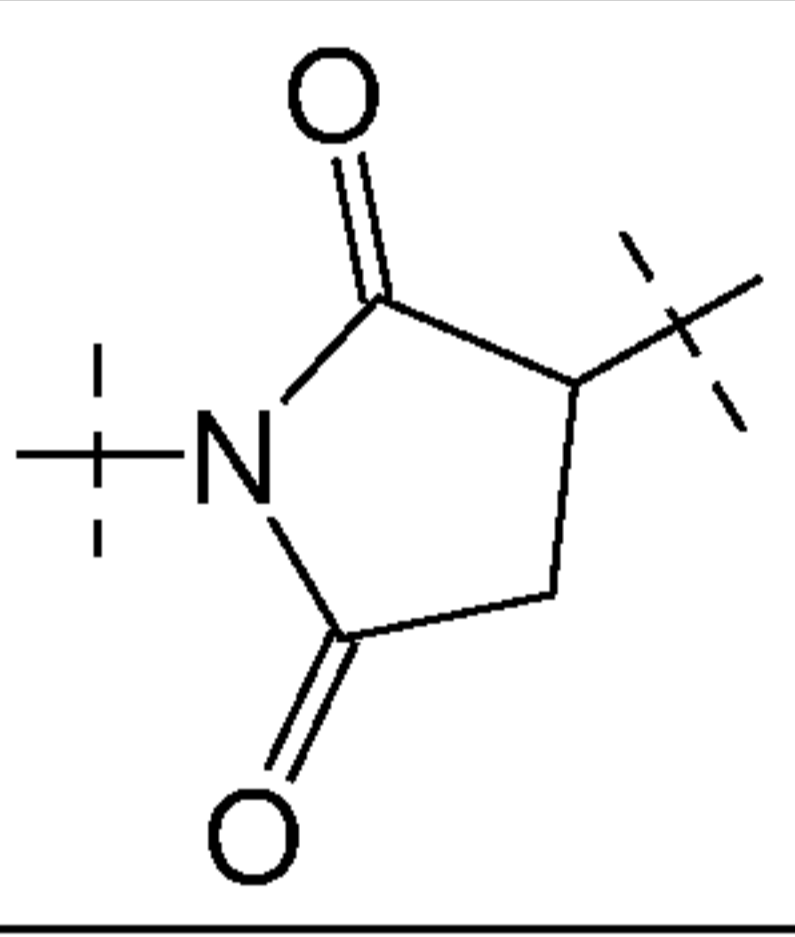
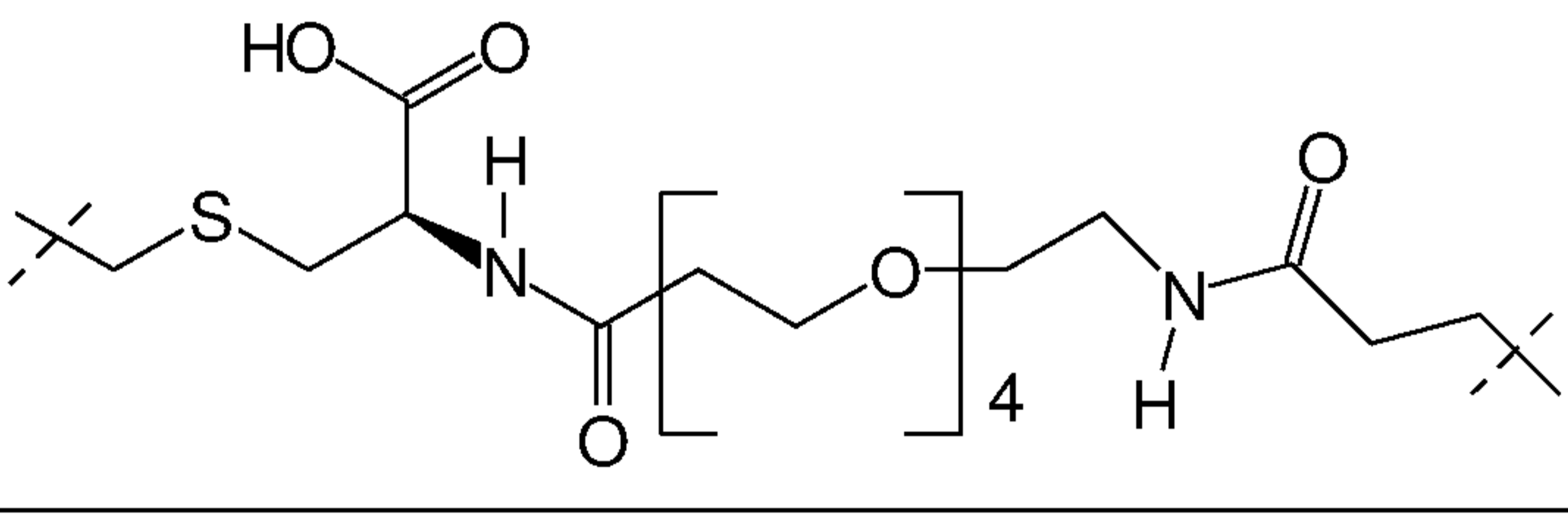
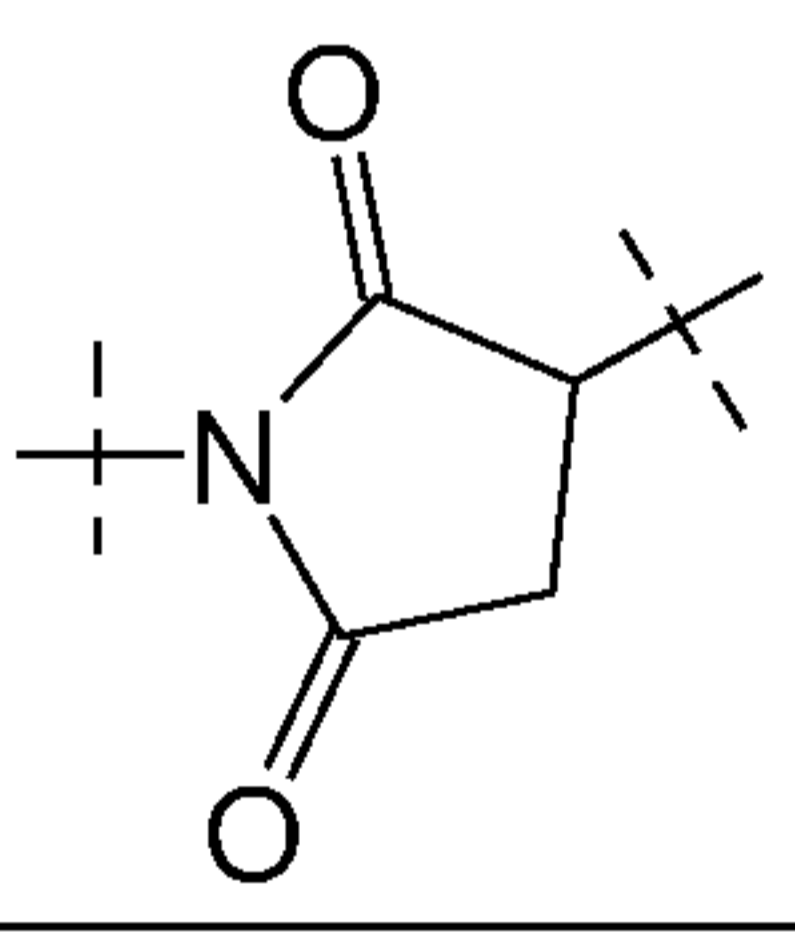
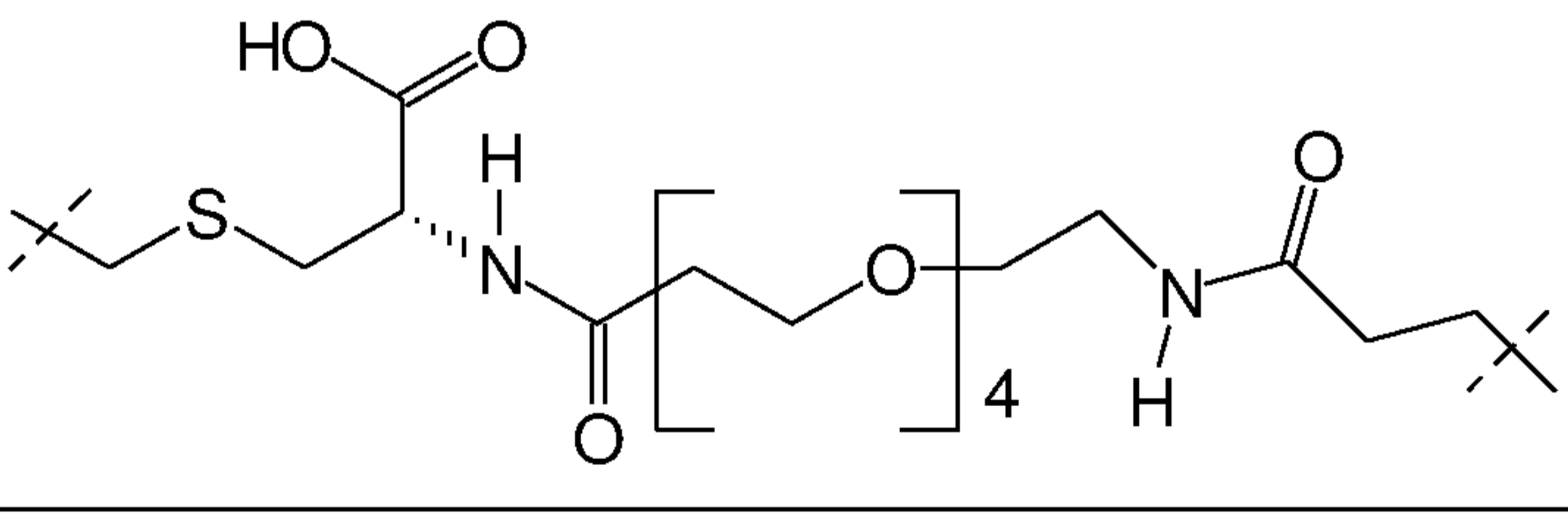
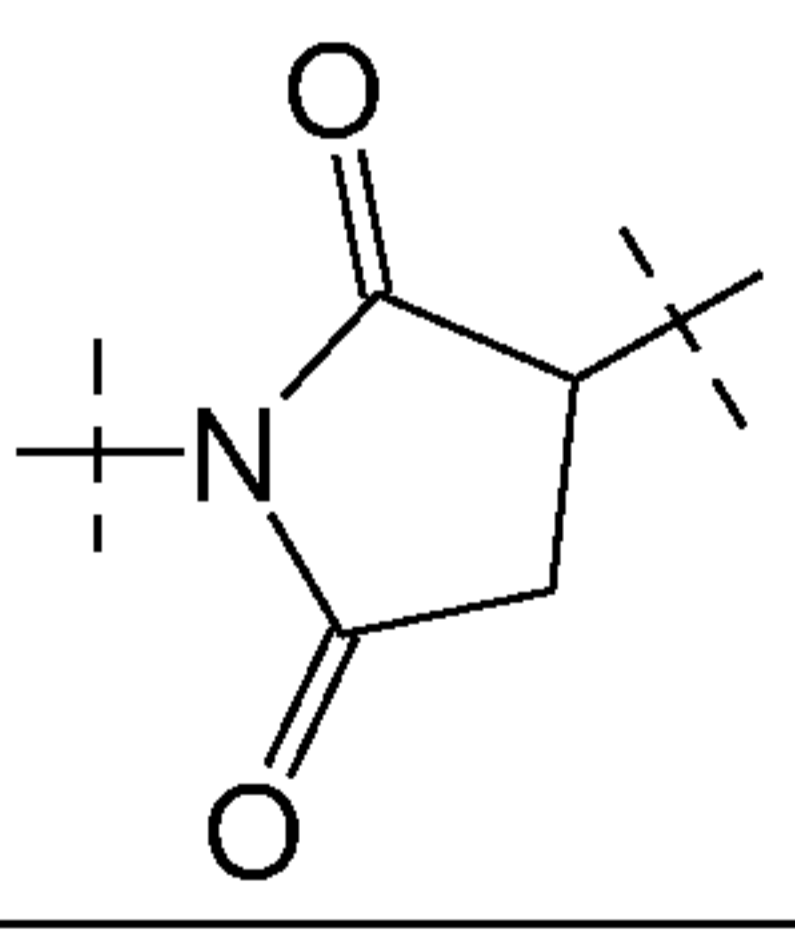
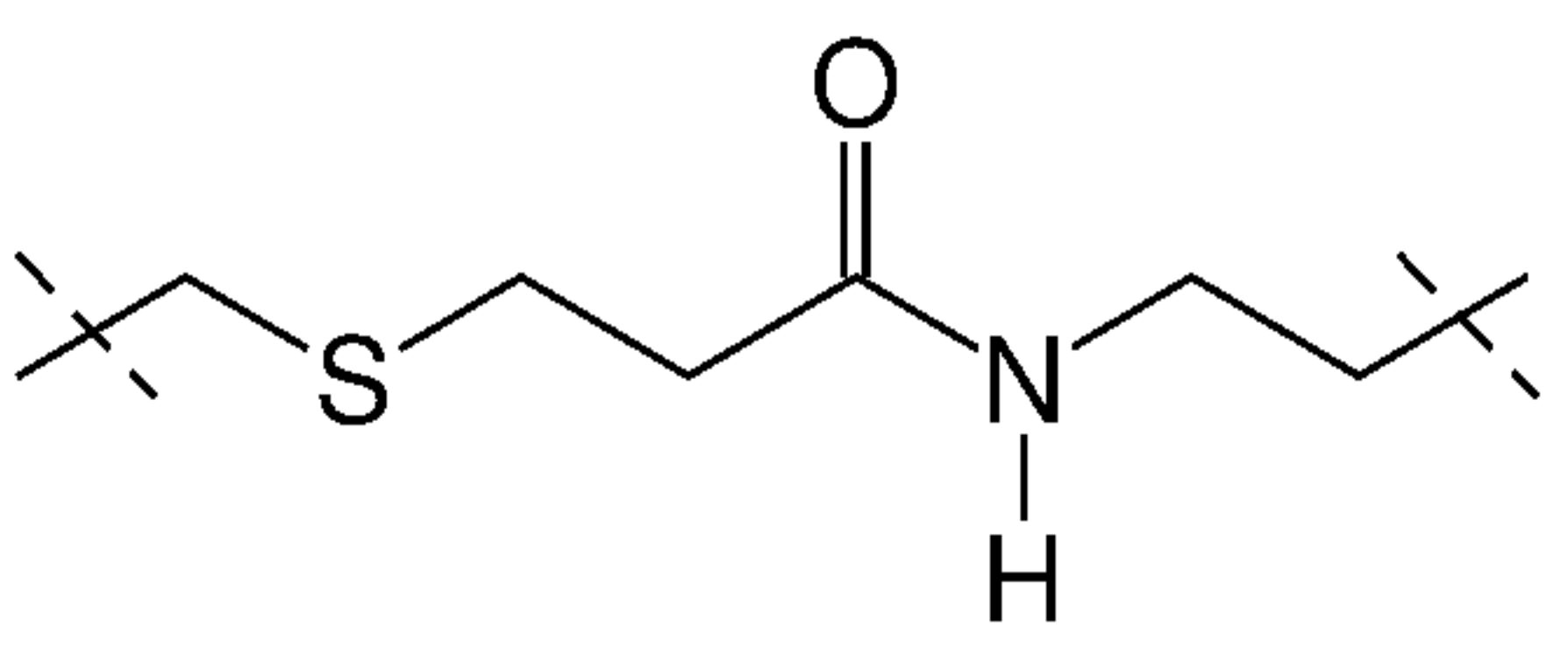
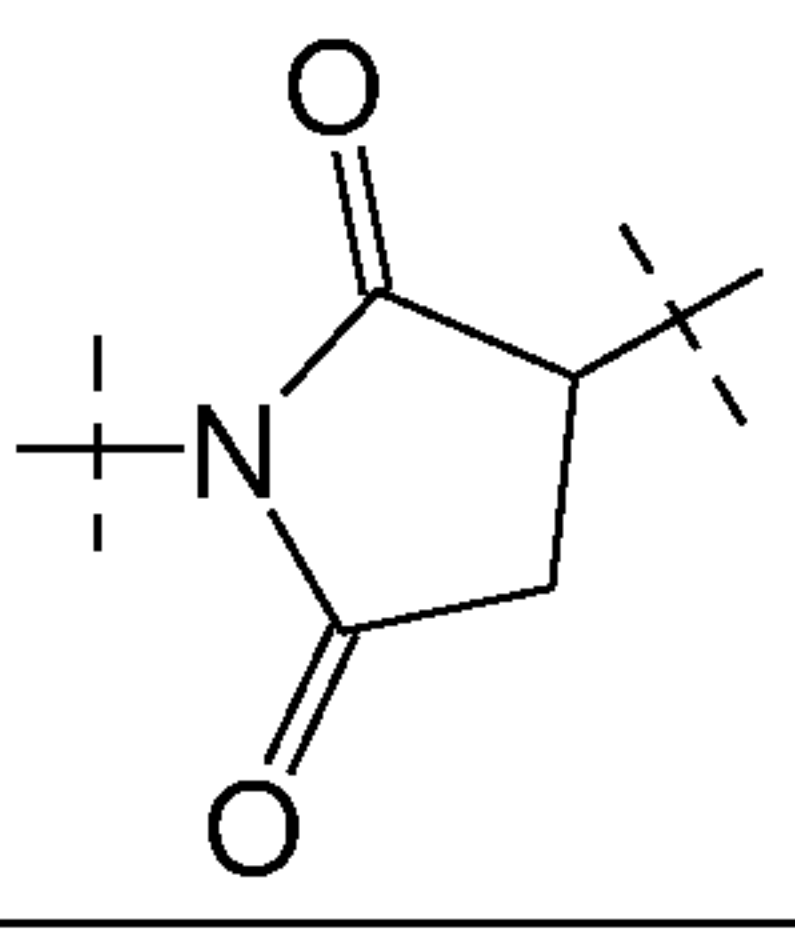
25 Table A`

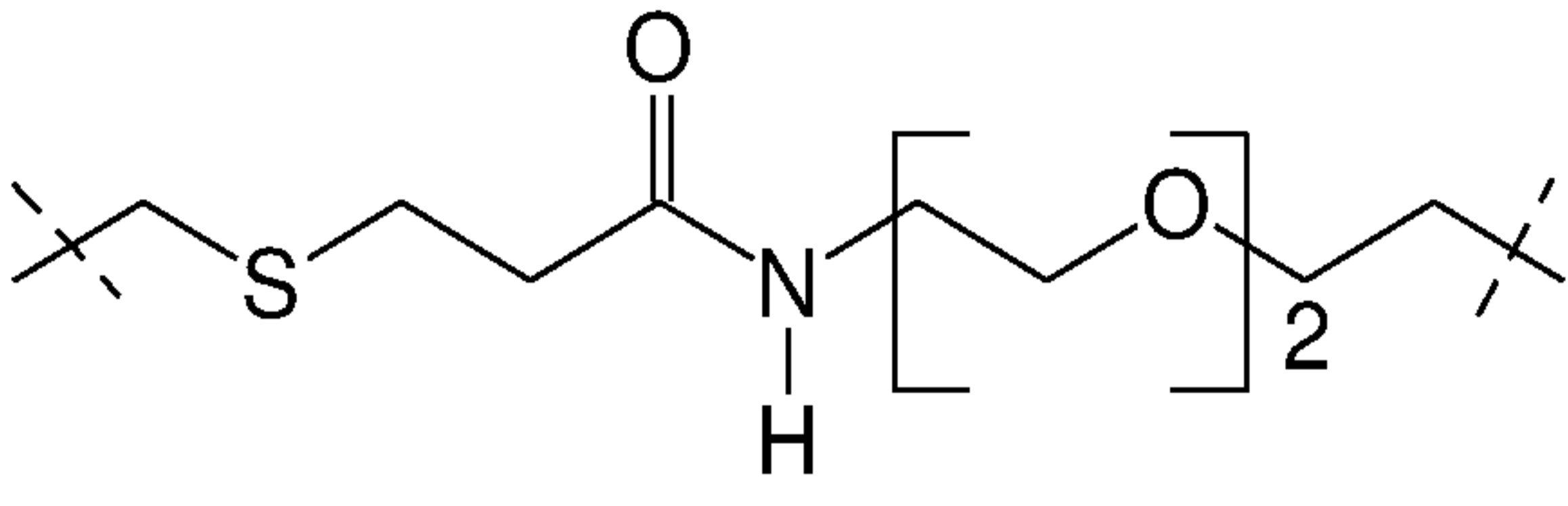
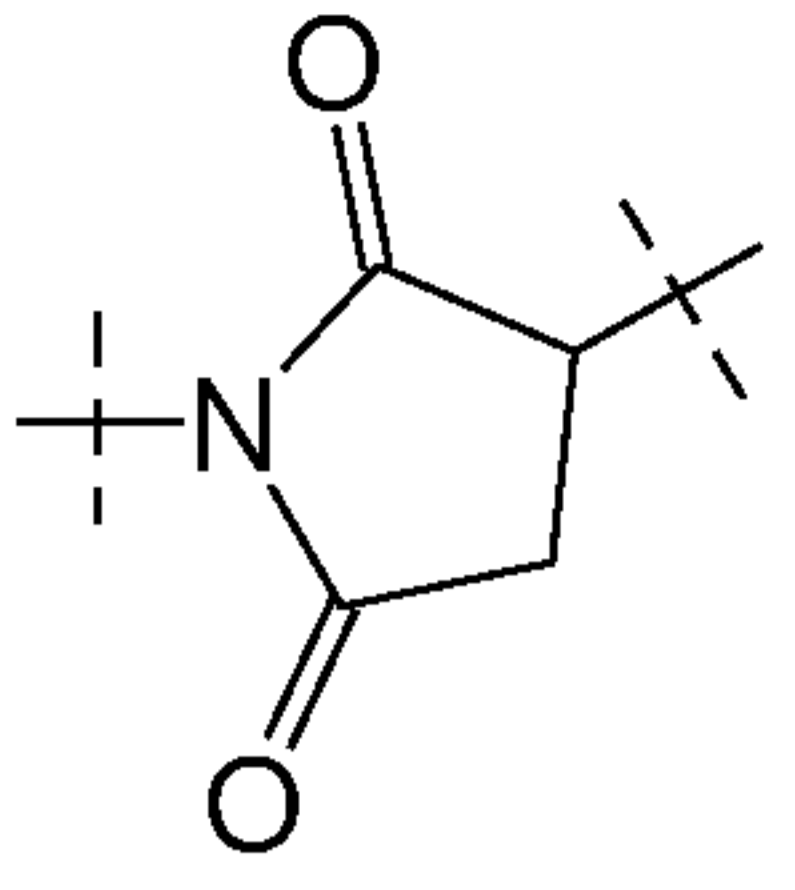
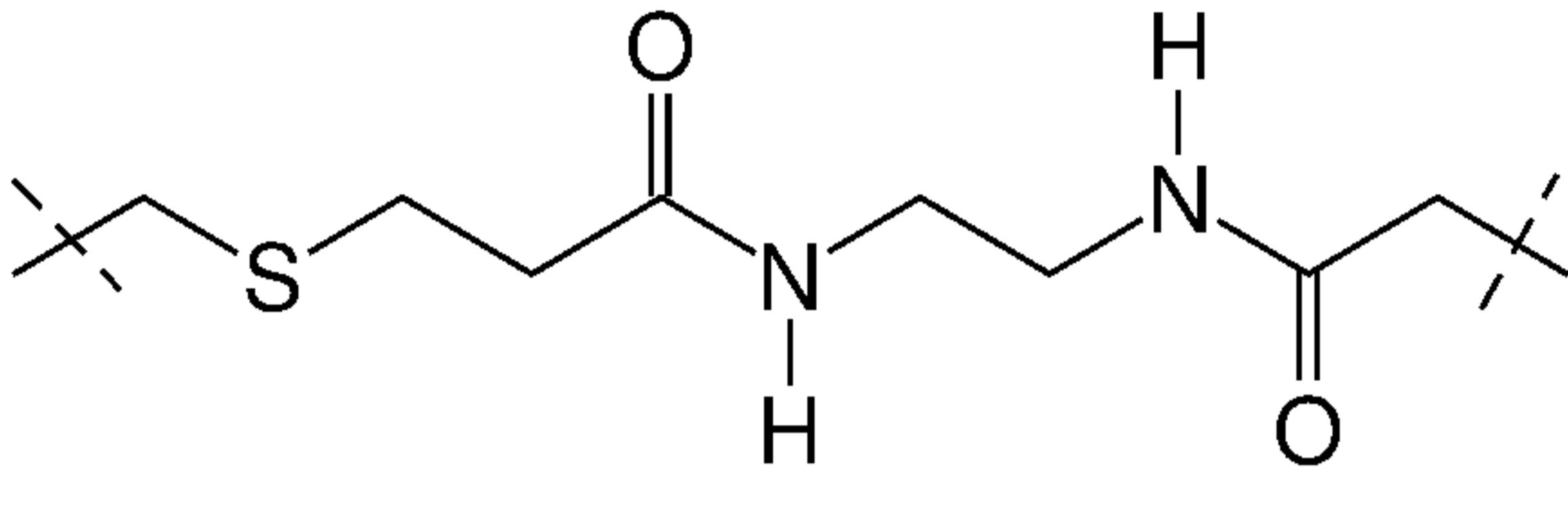
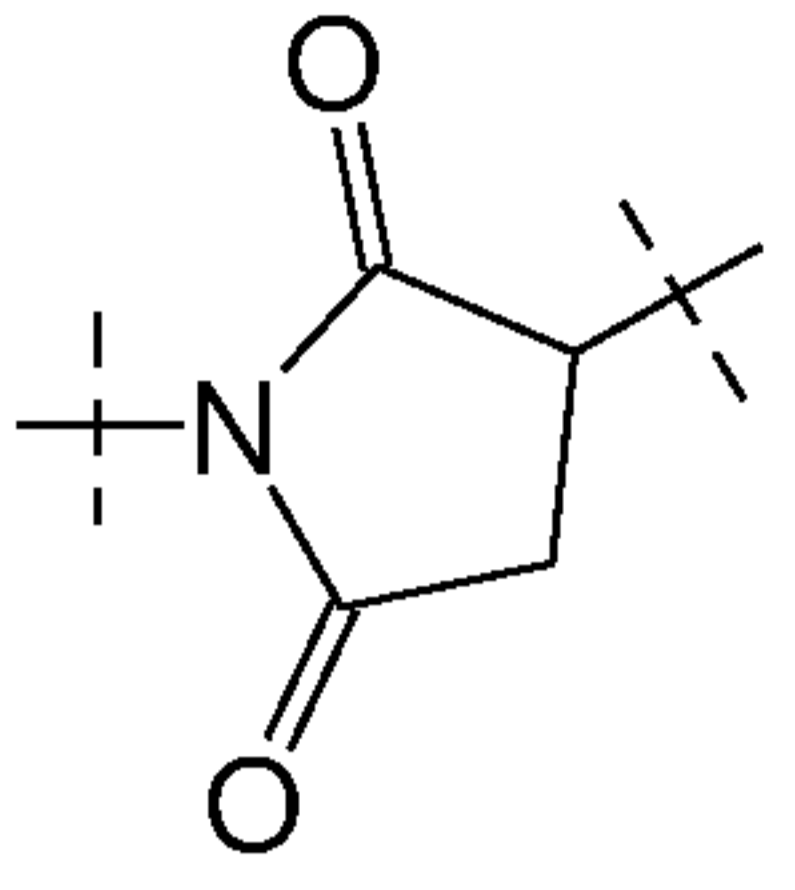
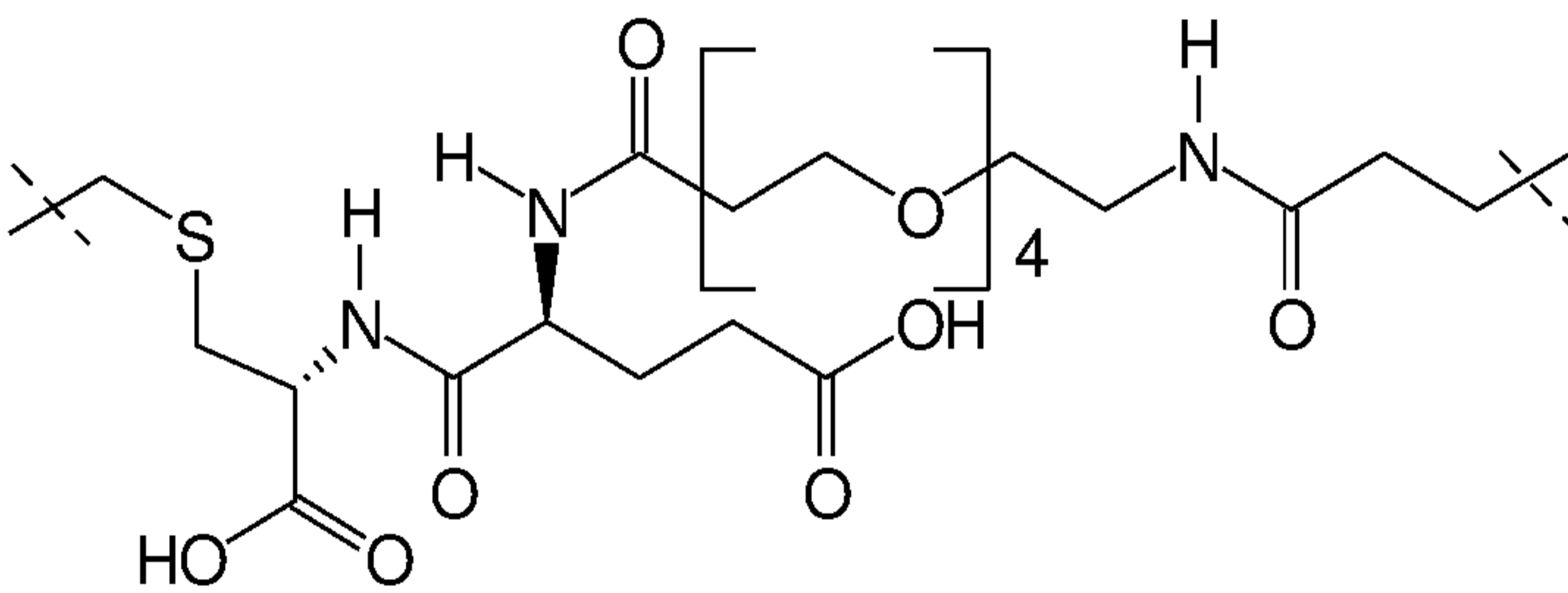
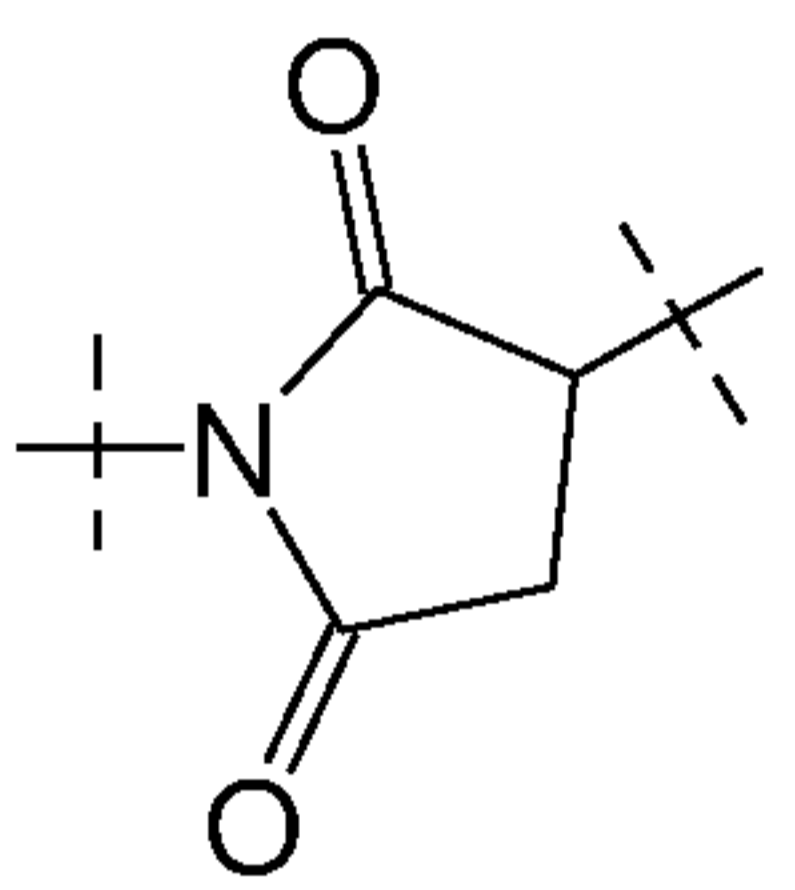
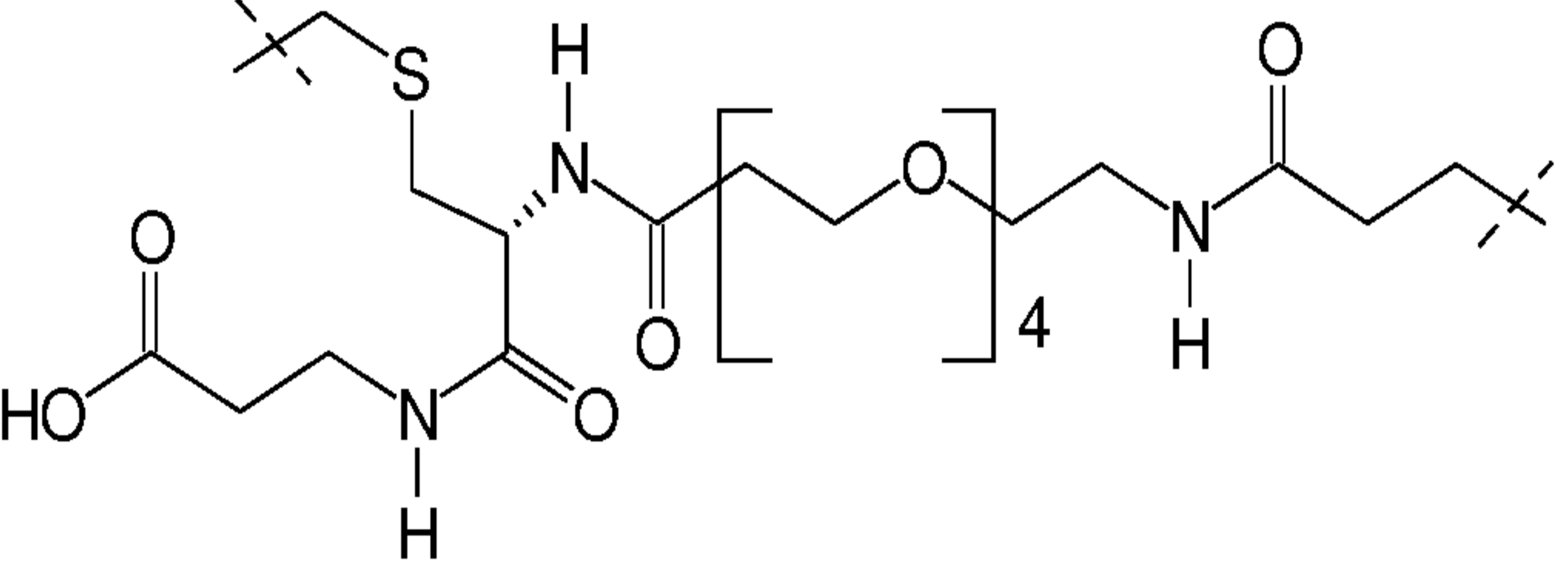
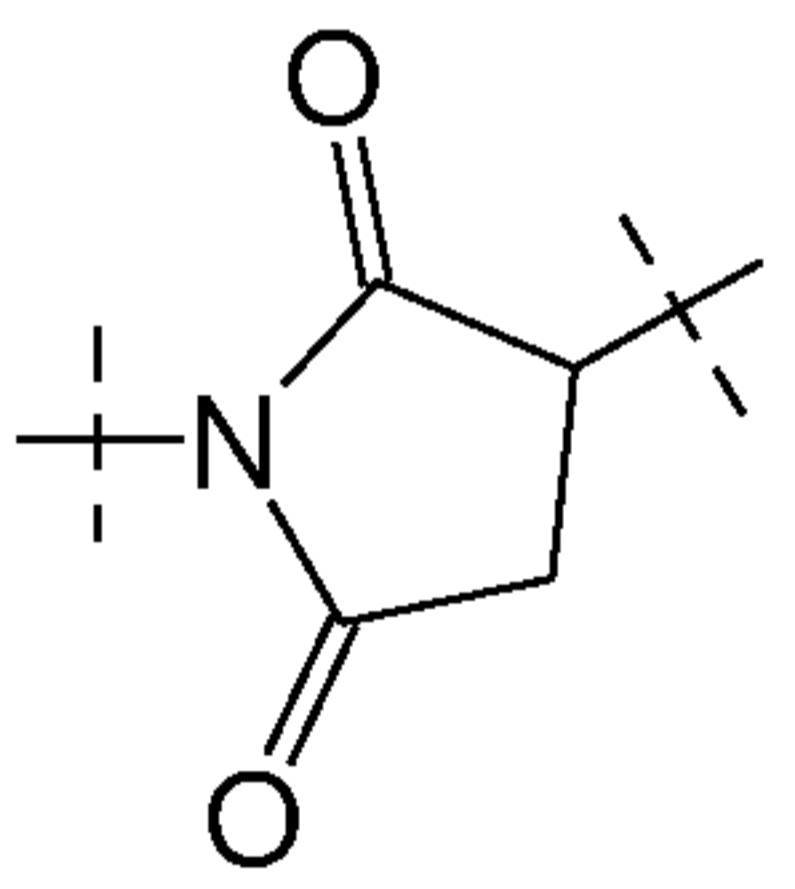
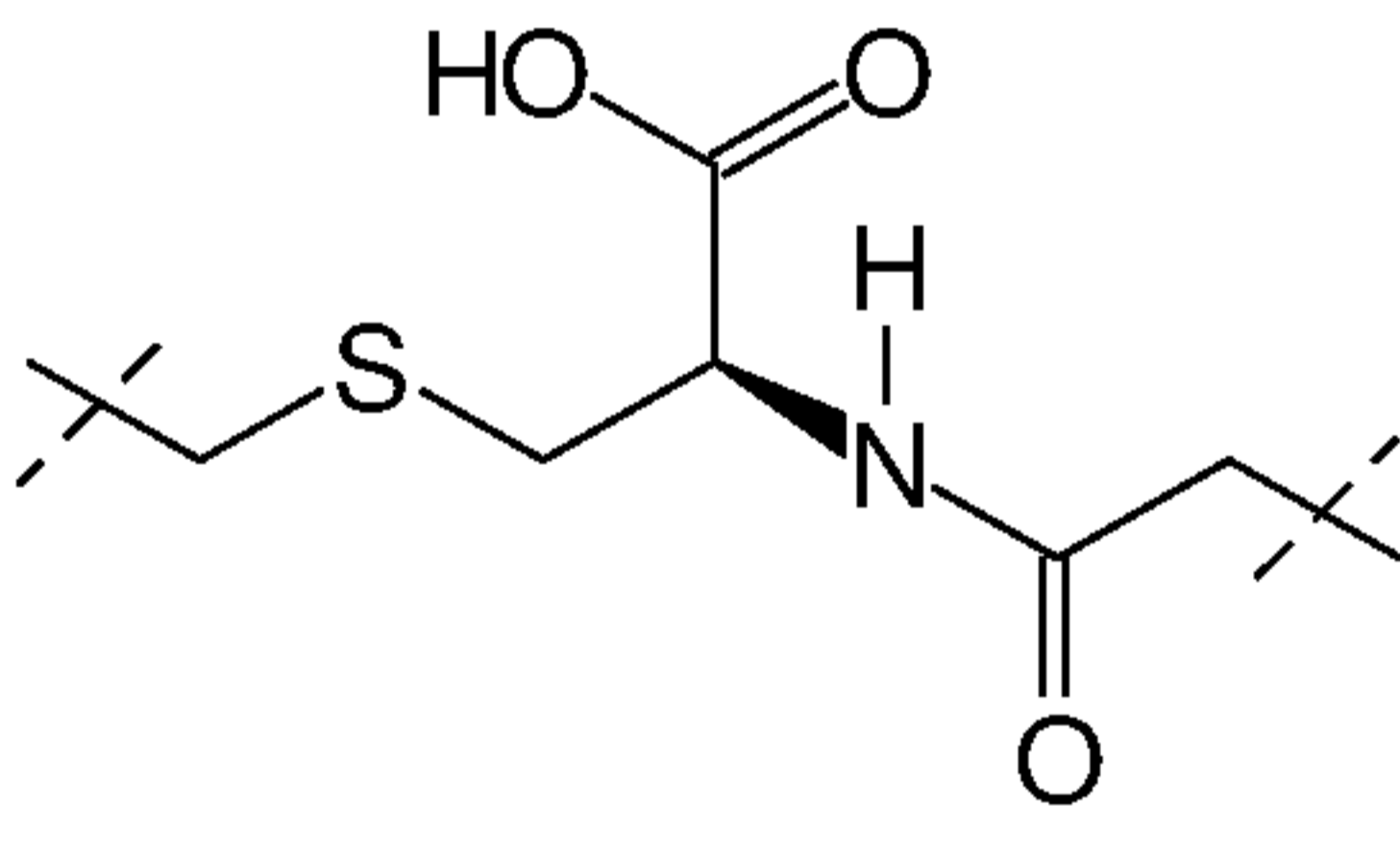
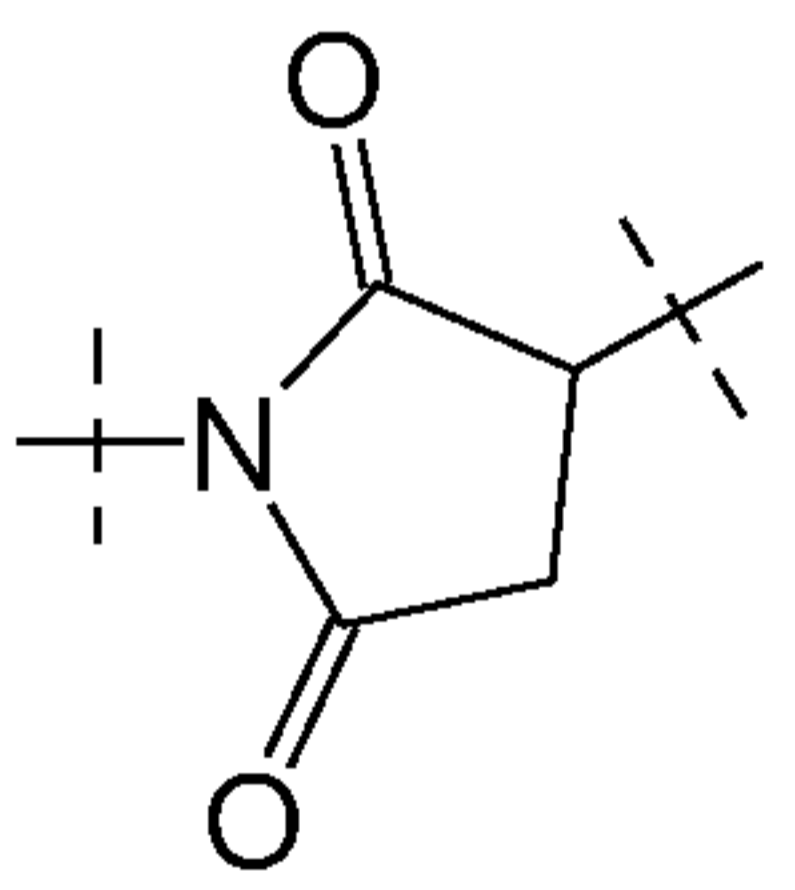
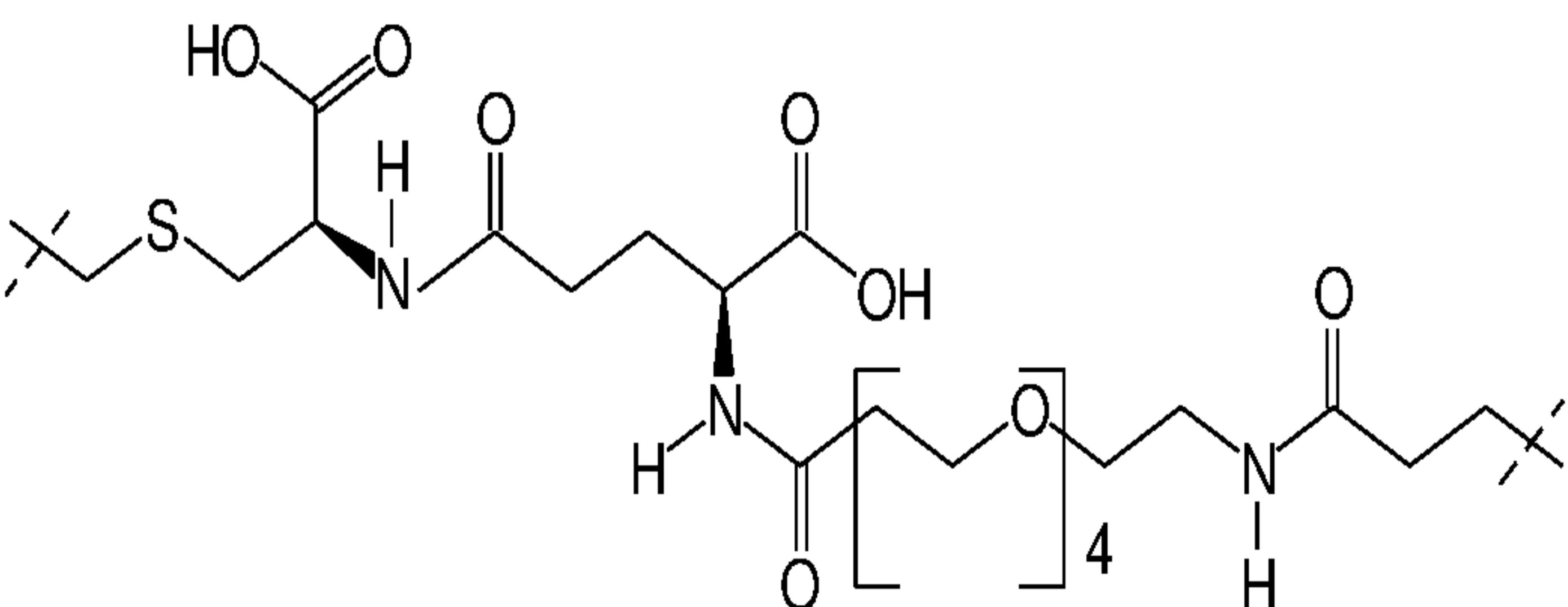
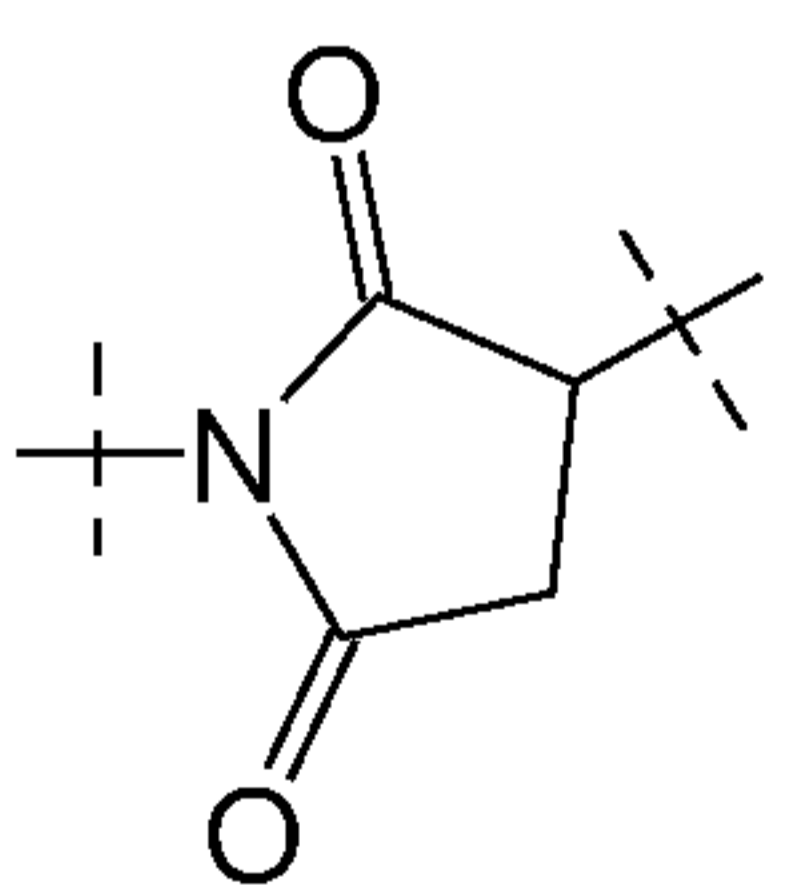
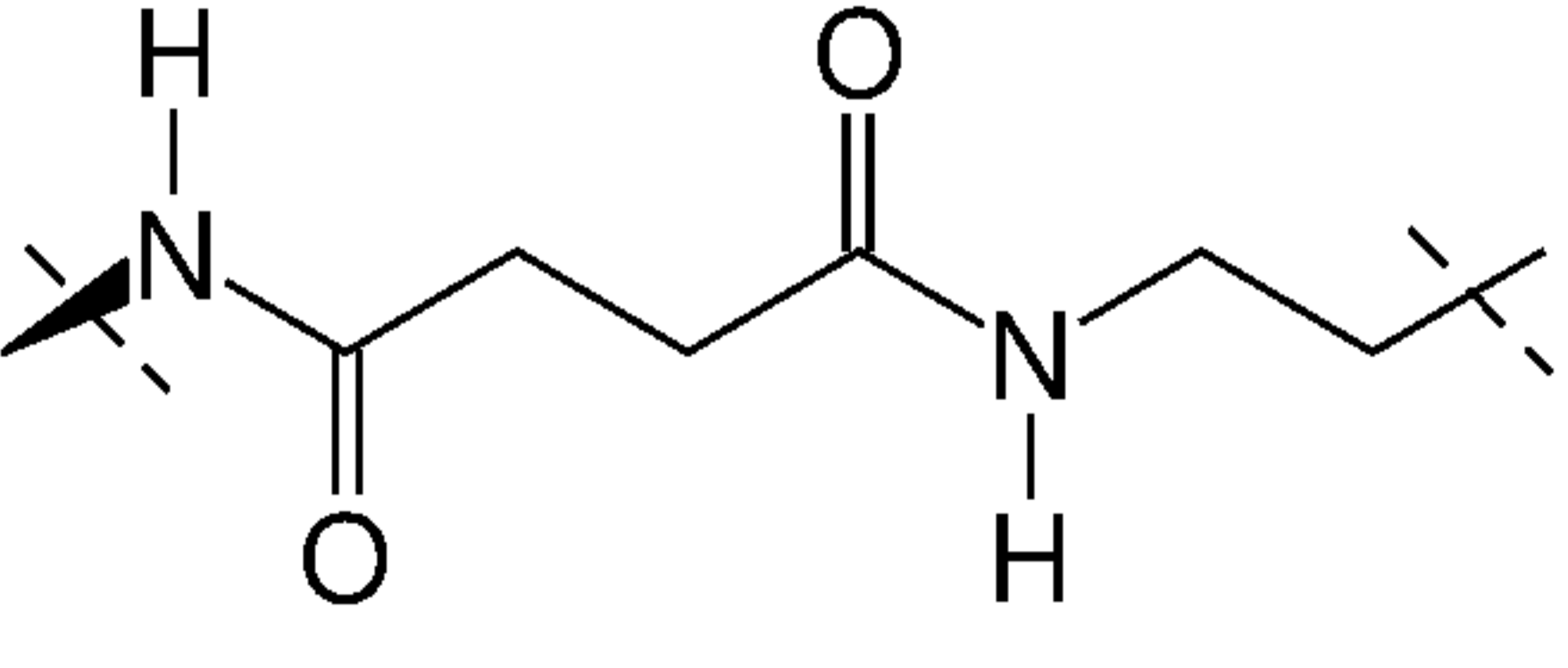
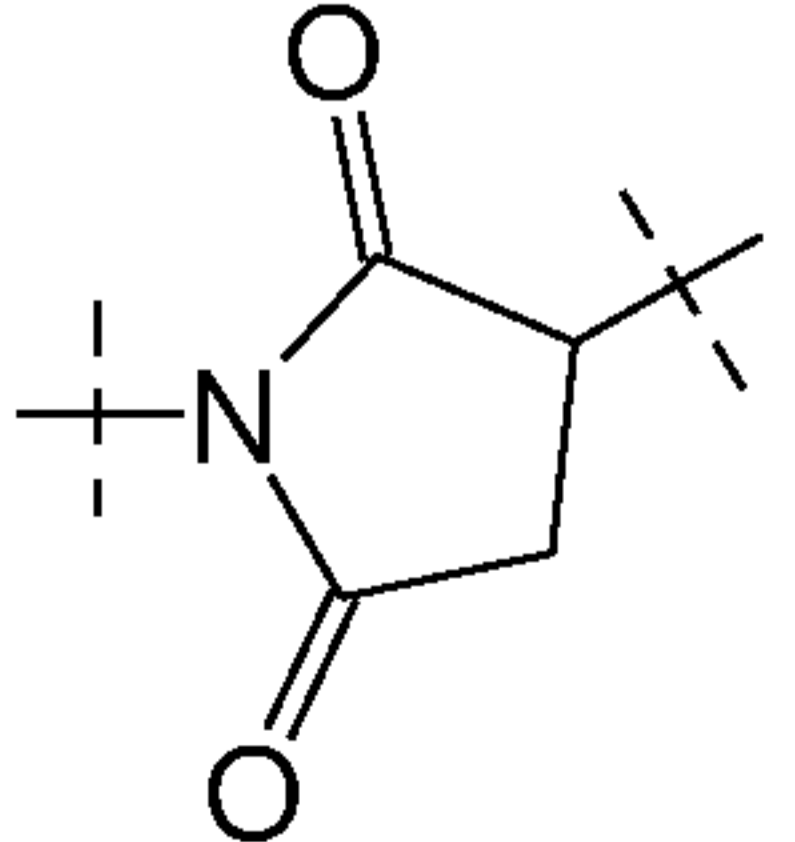


Ex.	Subst	mL1	L1	L2
105	R <sup>1</sup>	1		
111	R <sup>1</sup>	1		
114	R <sup>1</sup>	1		
119 / 158	R <sup>1</sup>	1		
120	R <sup>1</sup>	1		
123 / 157 / 125	R <sup>1</sup>	1		
138 / 142	R <sup>3</sup>	0		

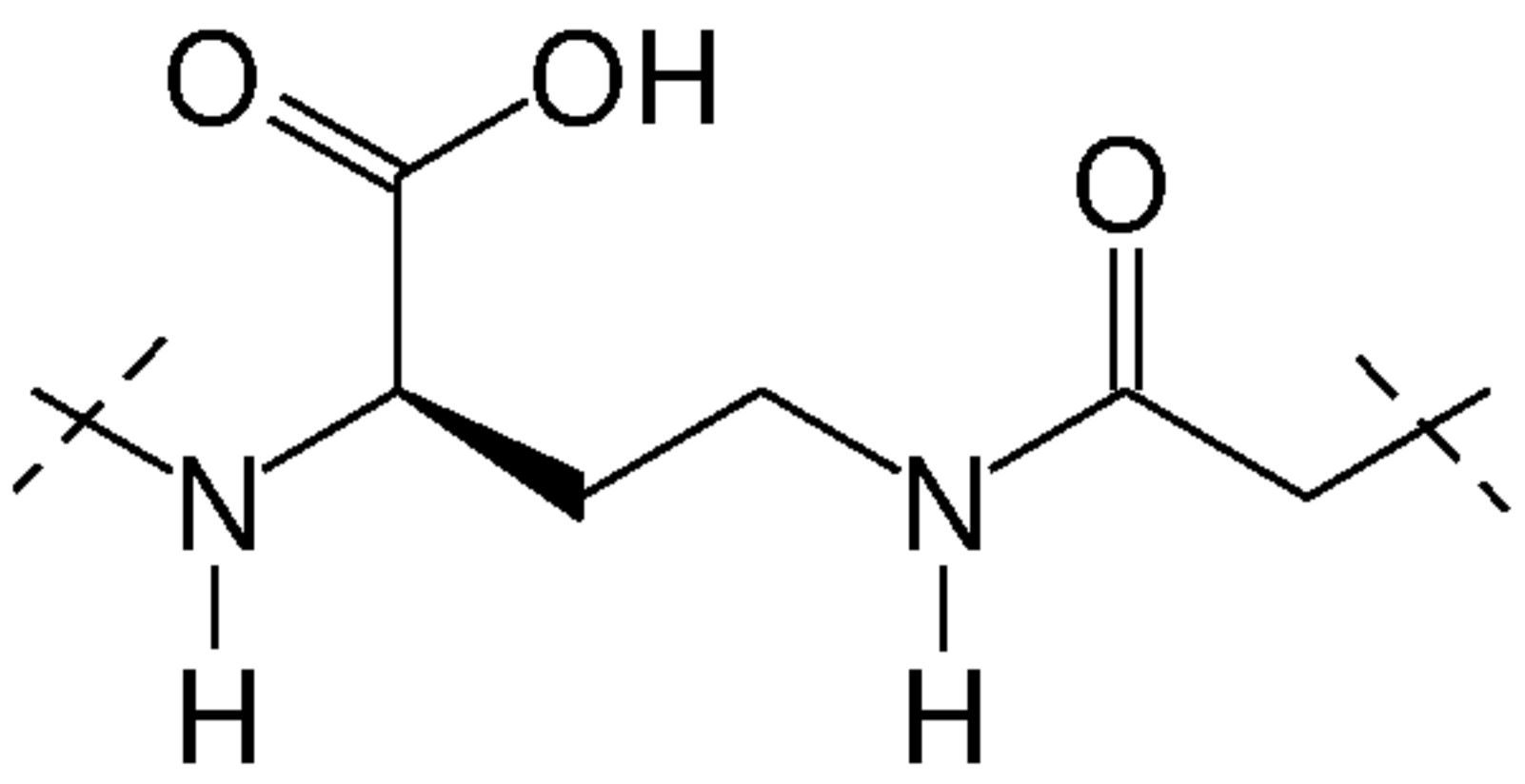
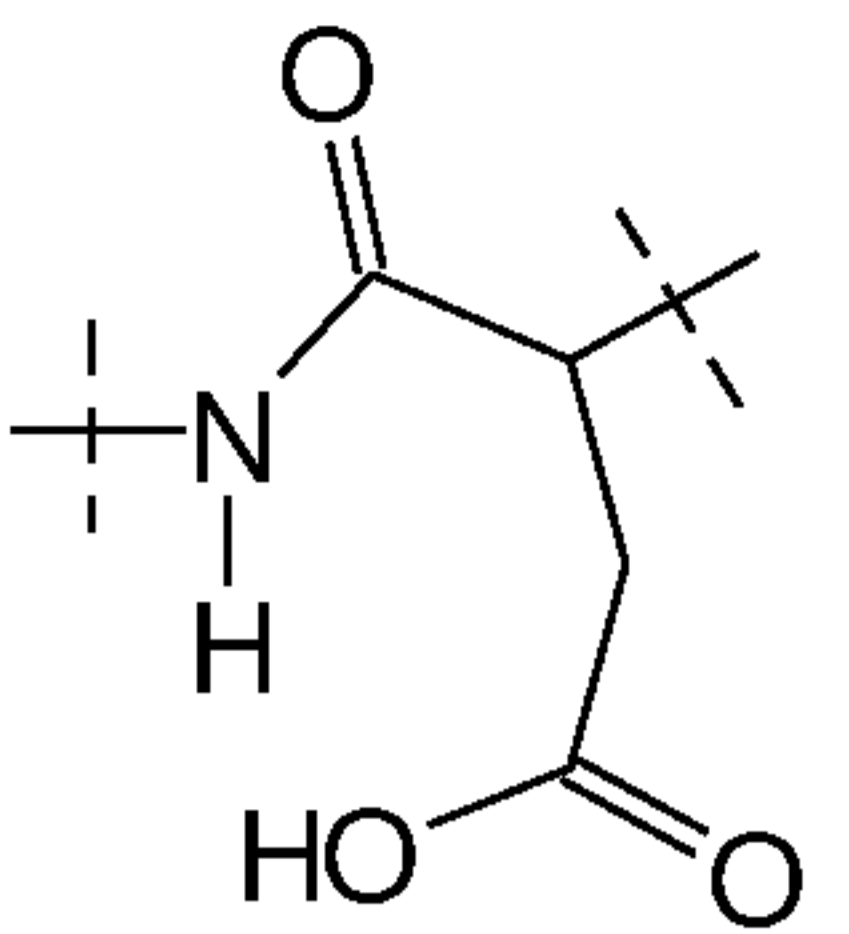
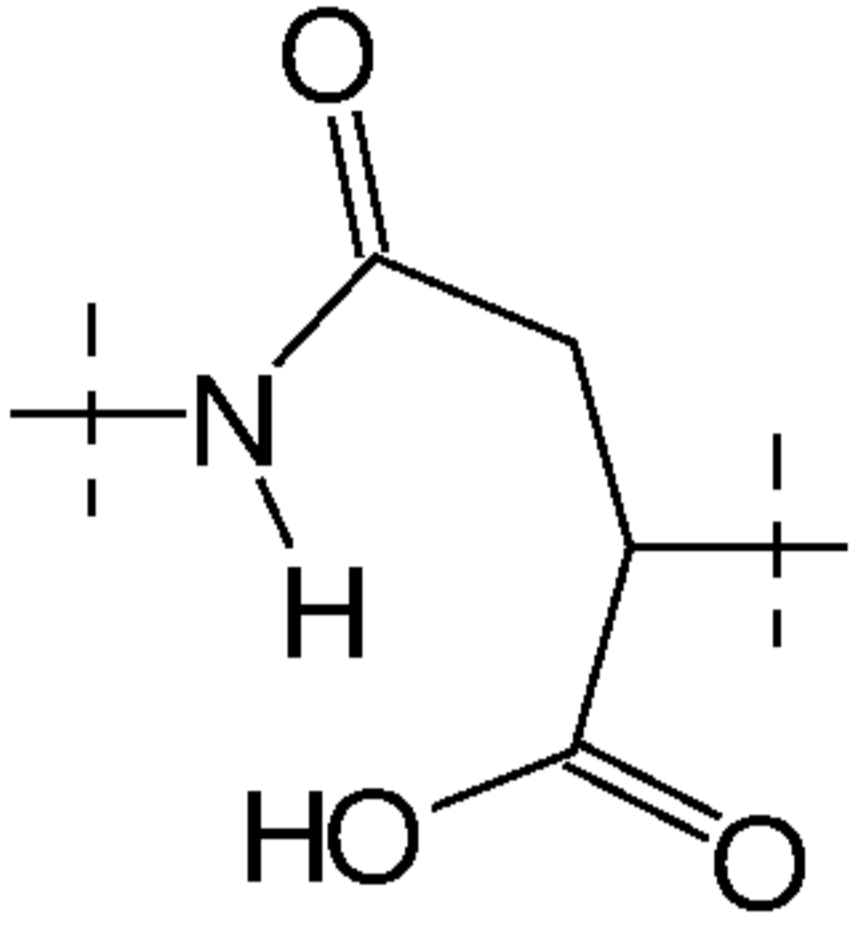
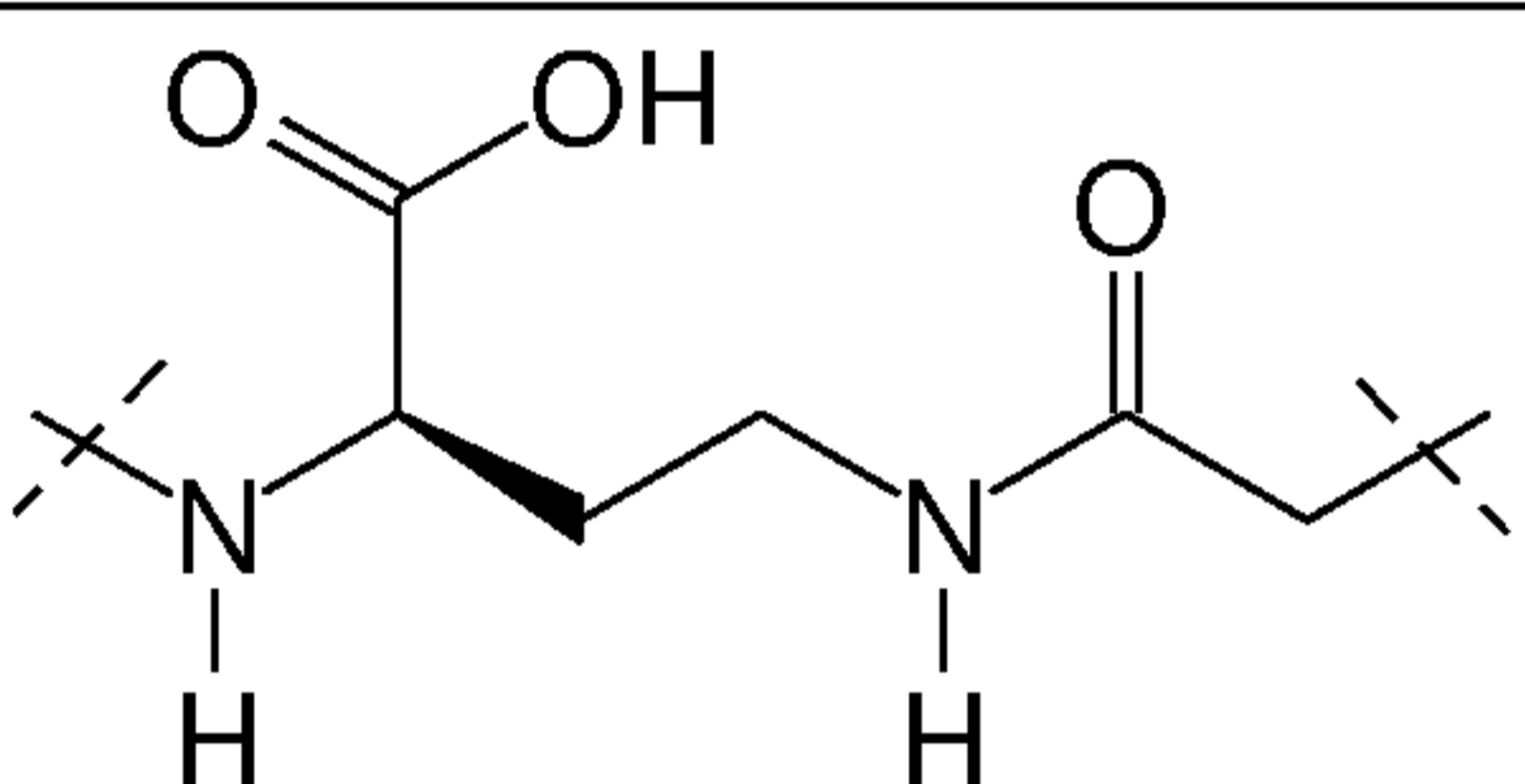
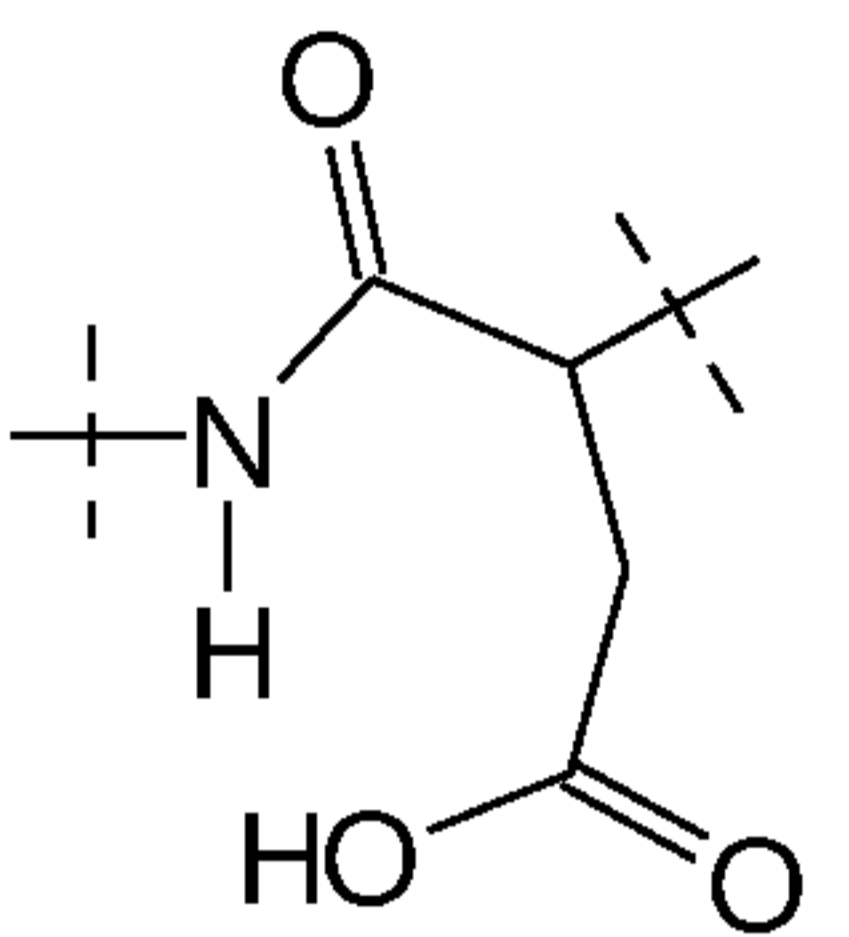
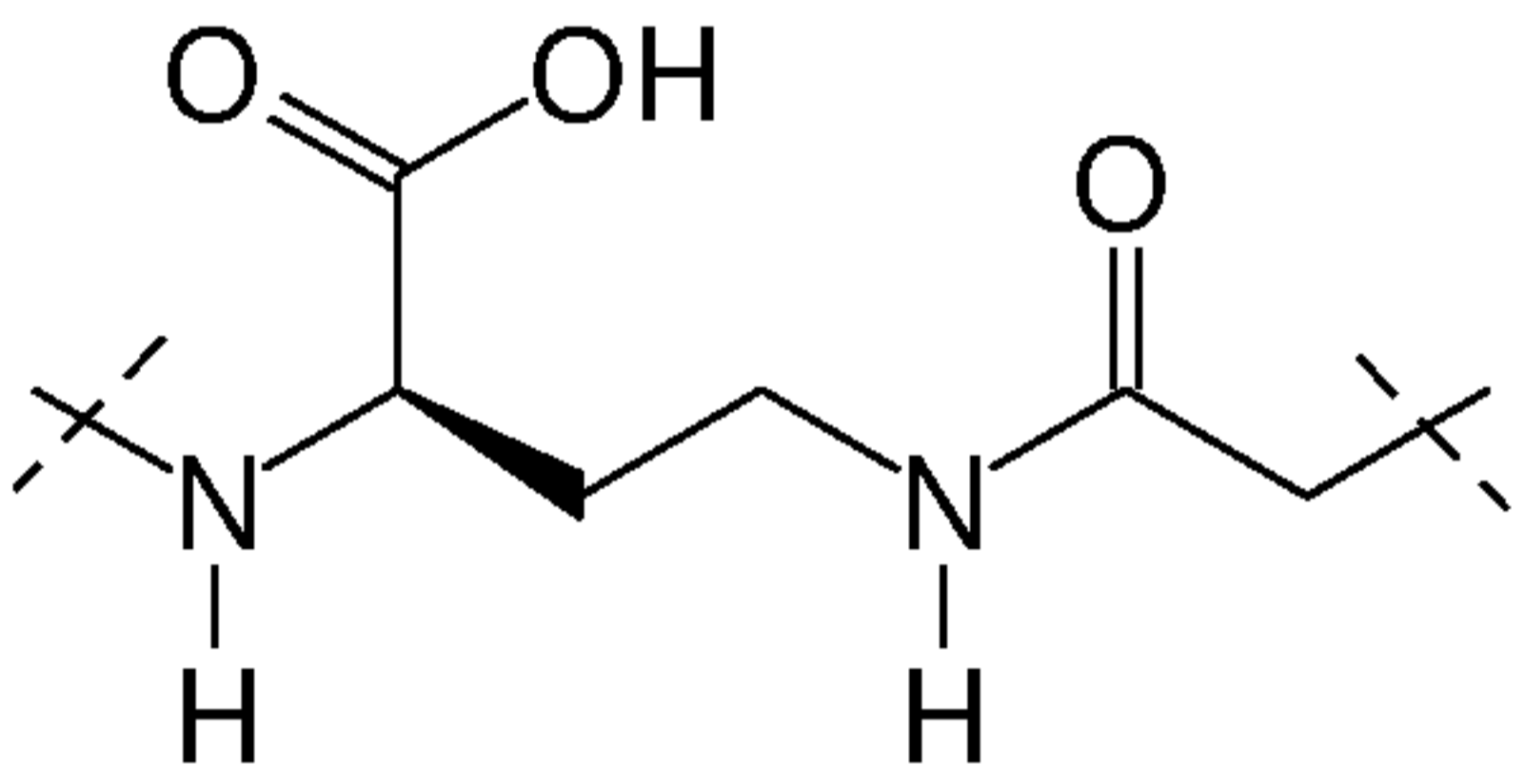
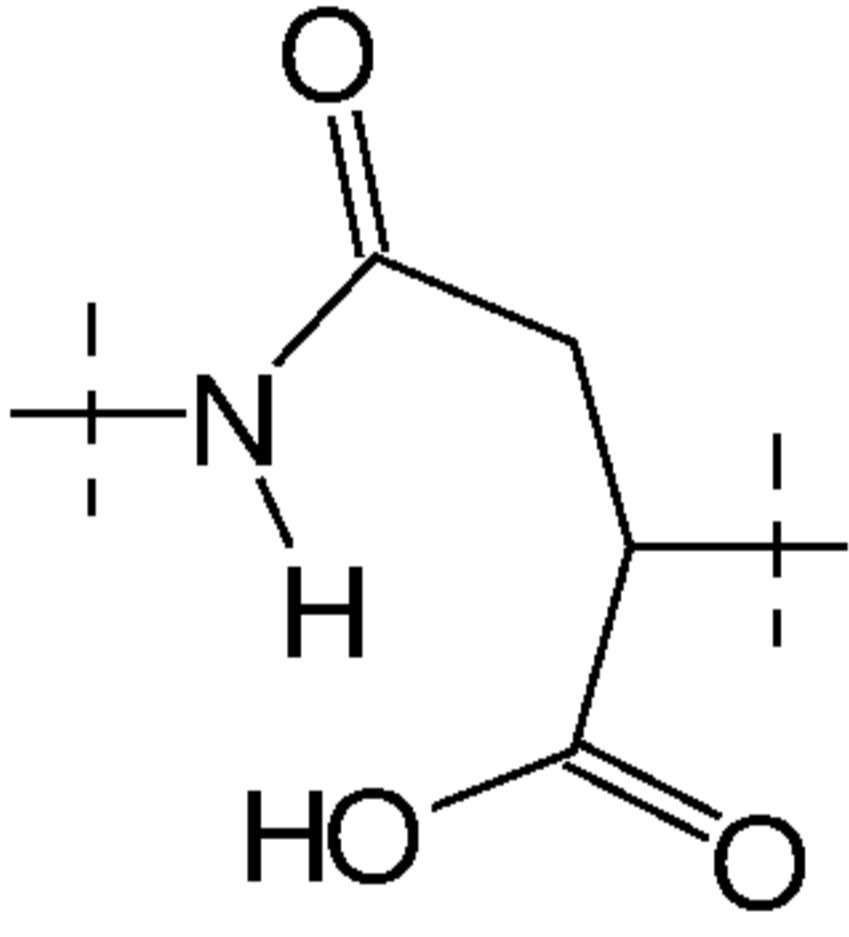
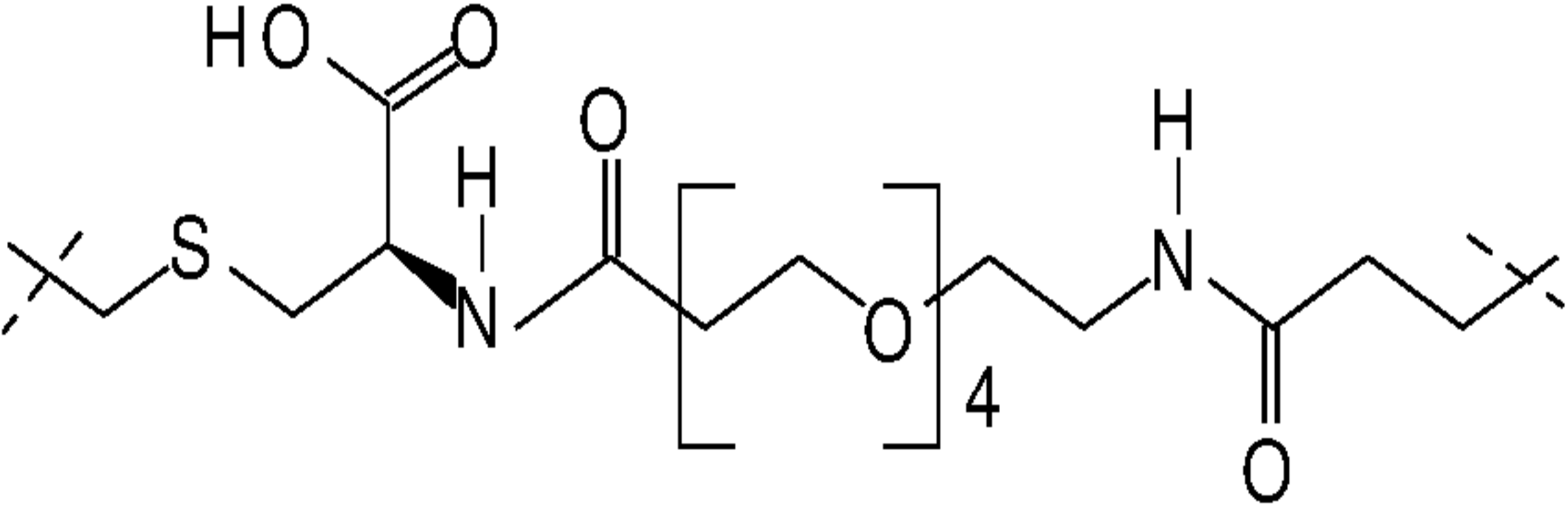
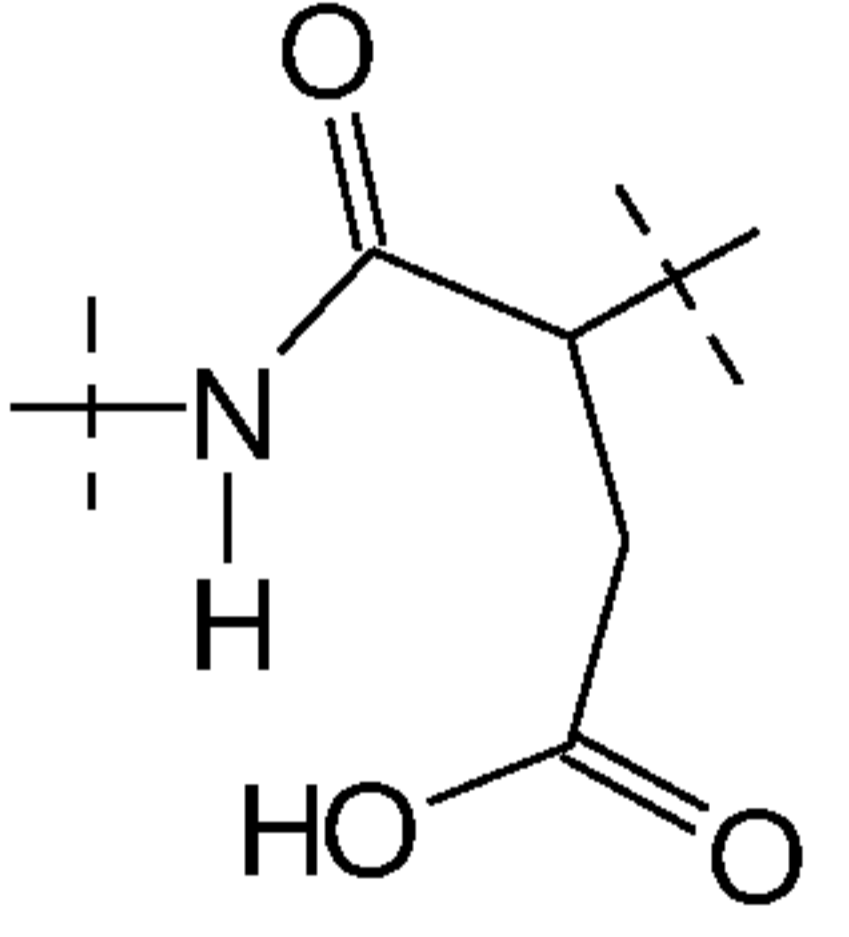
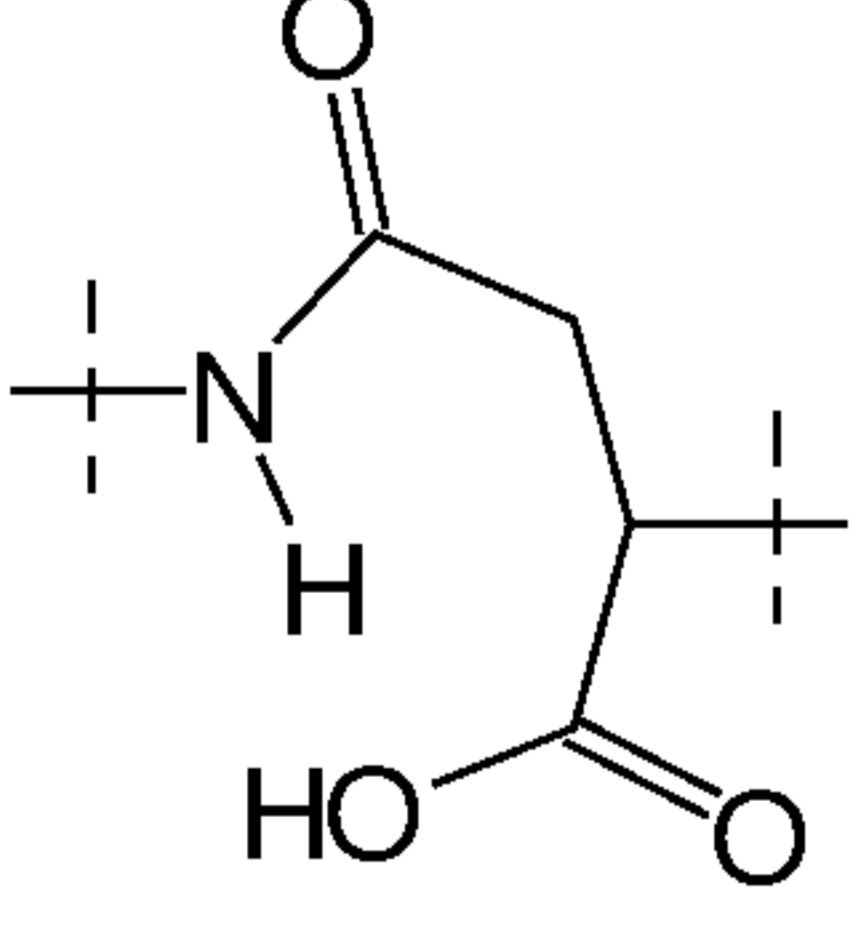
141 / 143	R <sup>3</sup>	0		
144	R <sup>3</sup>	0		
147	R <sup>3</sup>	0		
148	R <sup>3</sup>	0		
175	R <sup>1</sup>	1		
177	R <sup>1</sup>	1		 See note **
178	R <sup>1</sup>	1		

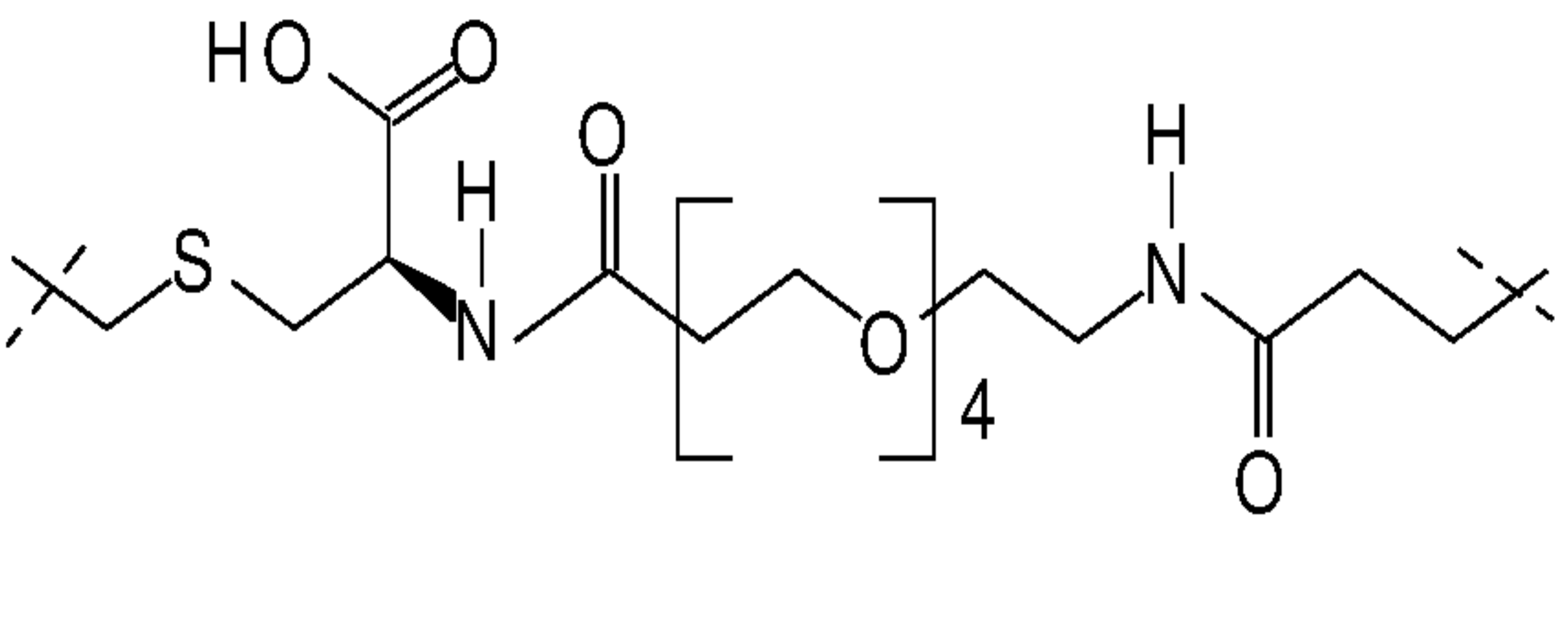
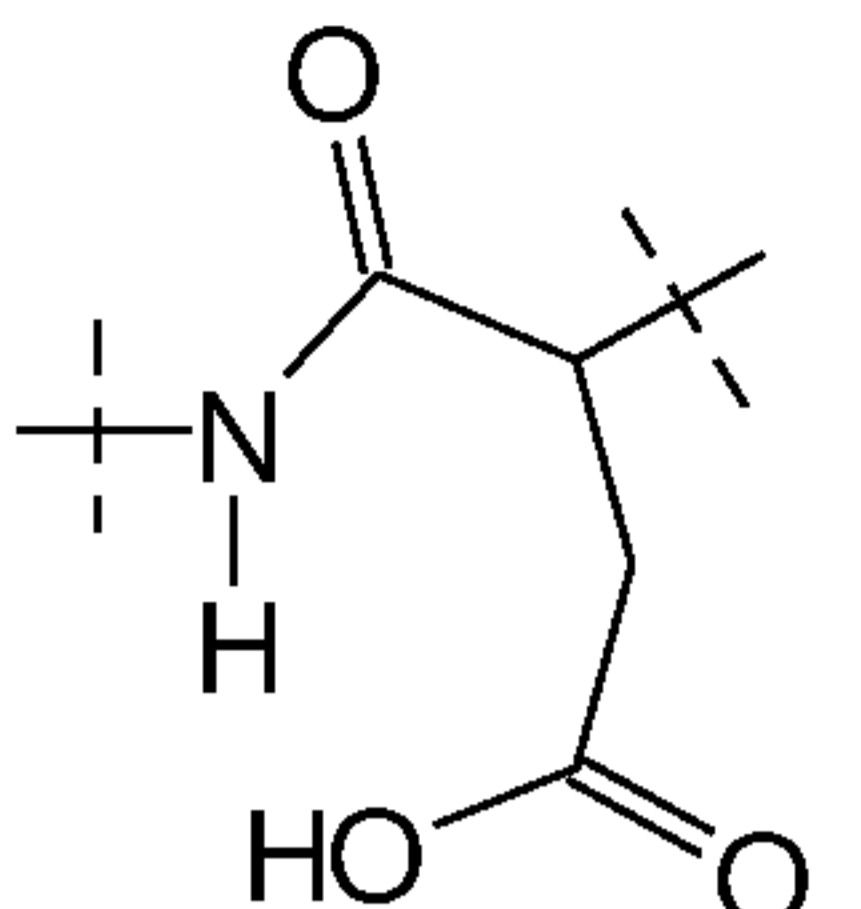
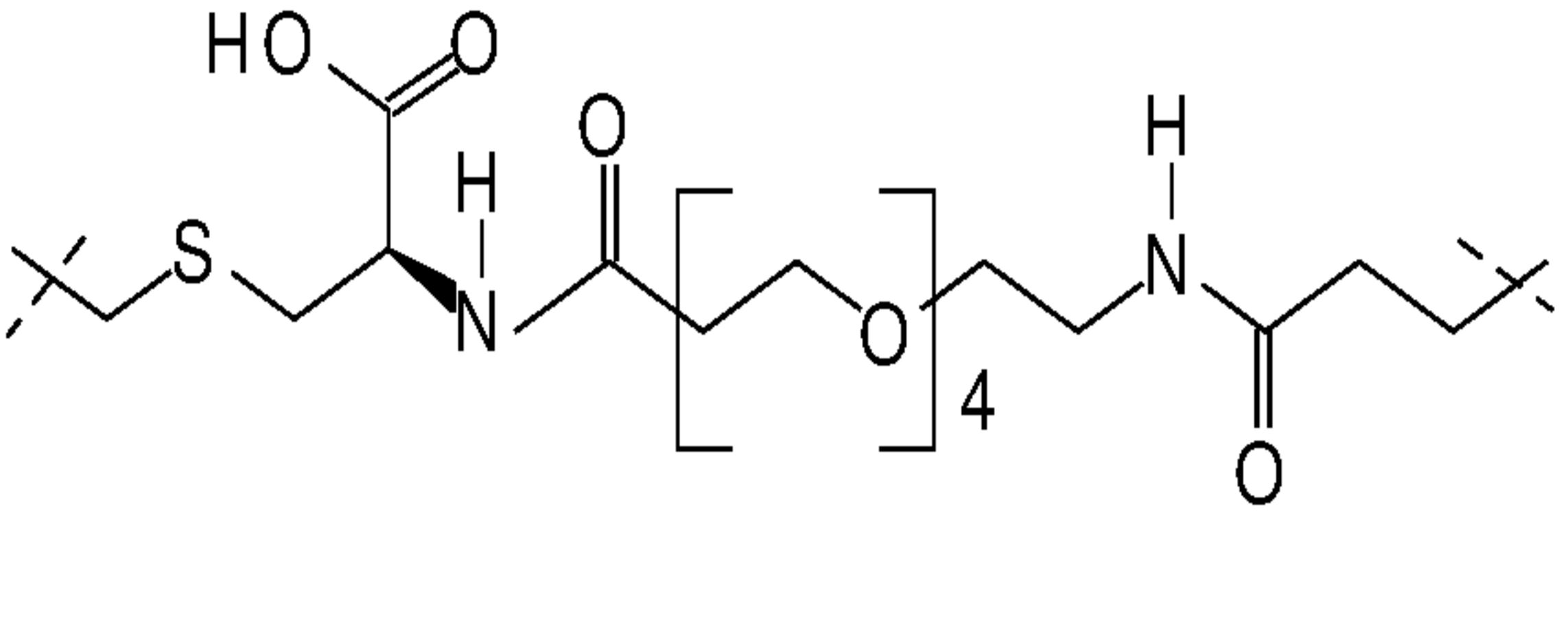
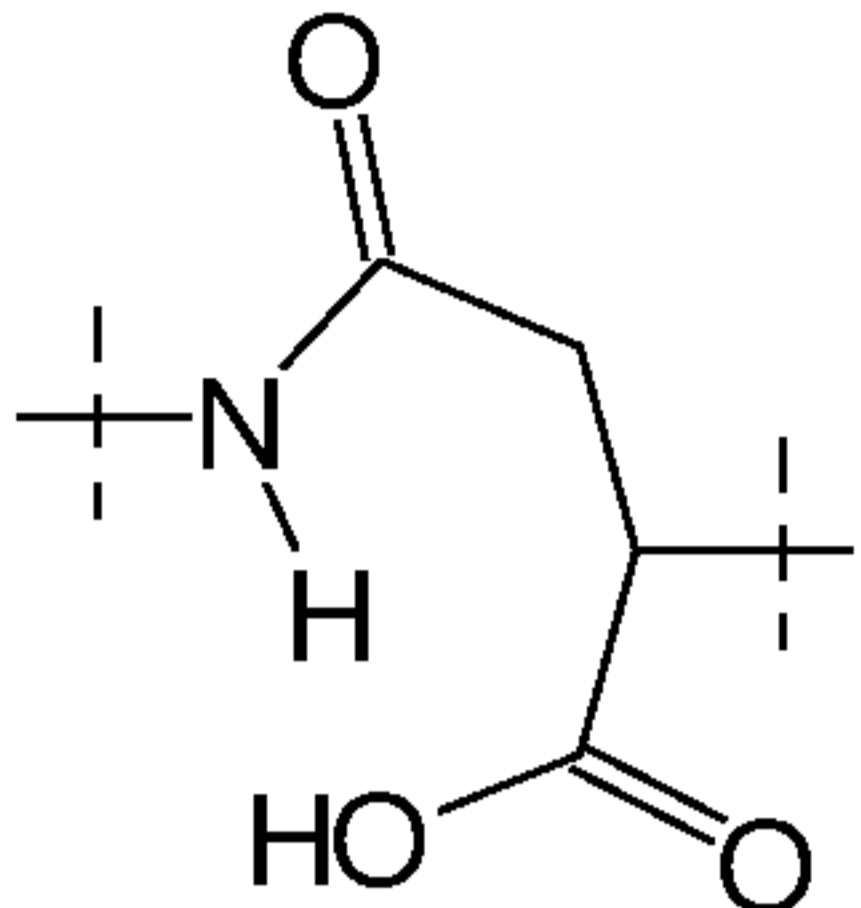
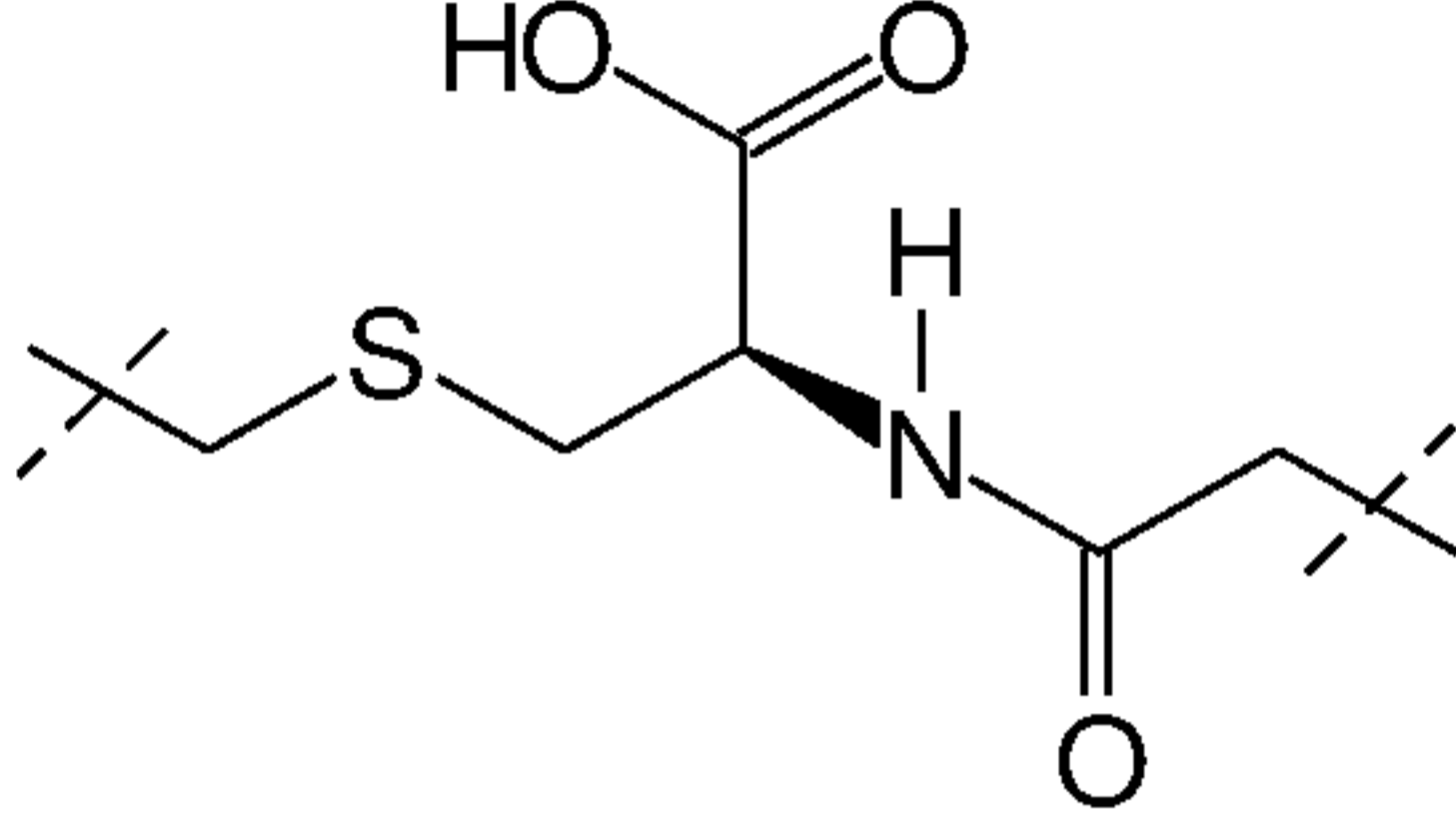
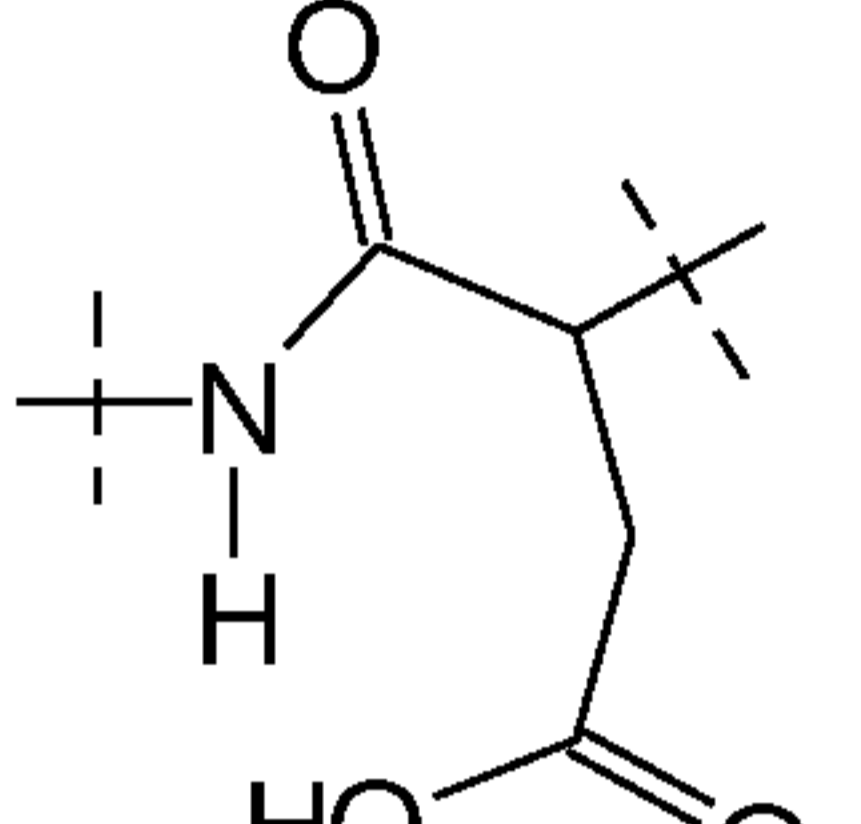
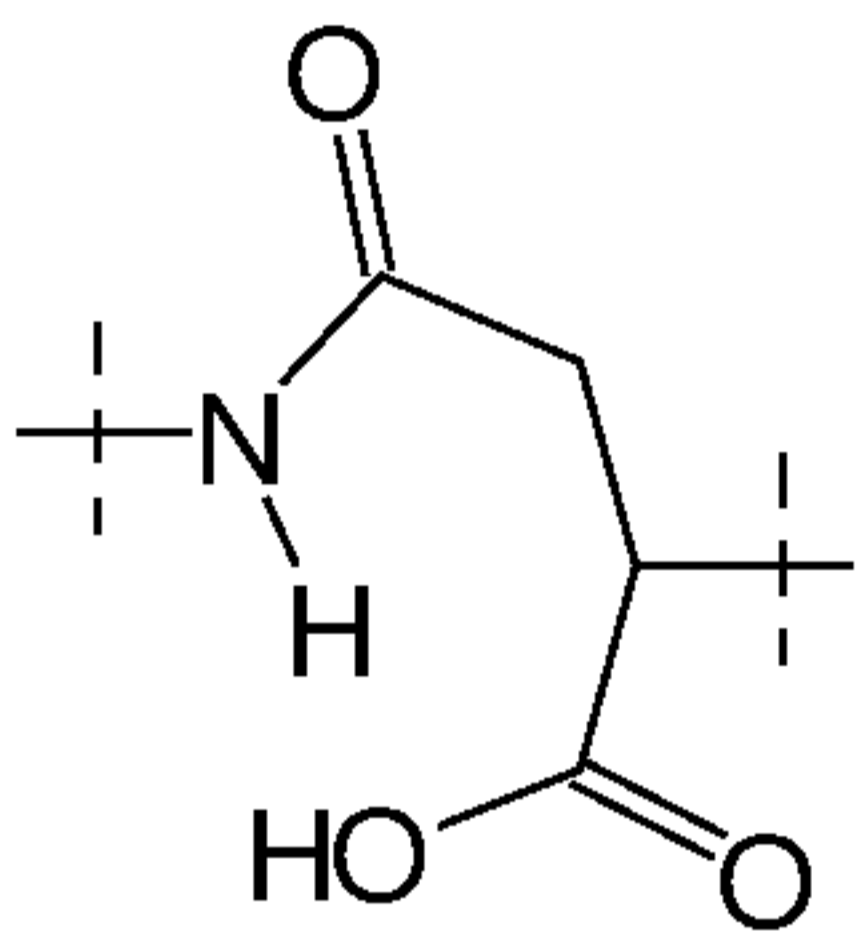
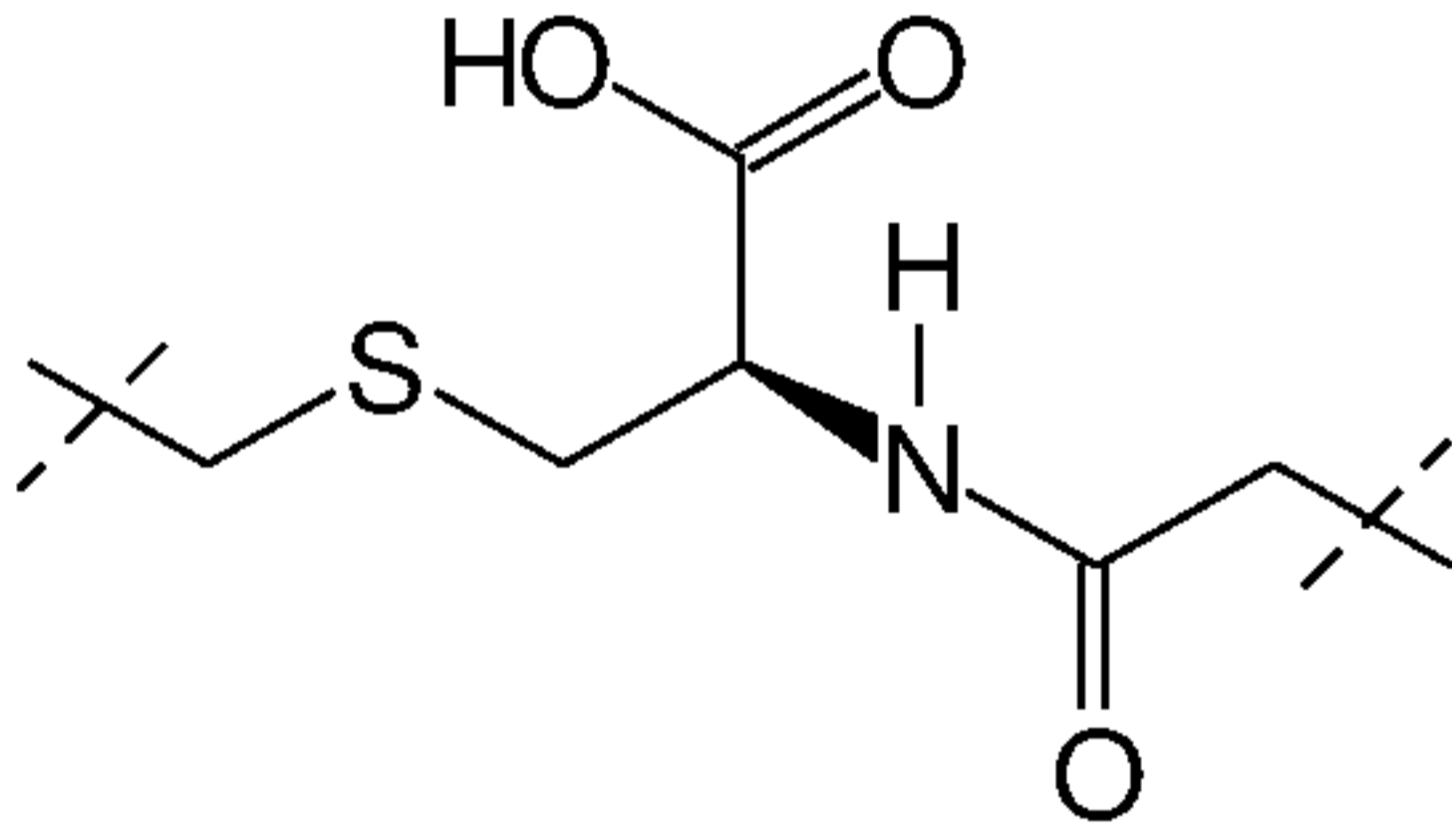
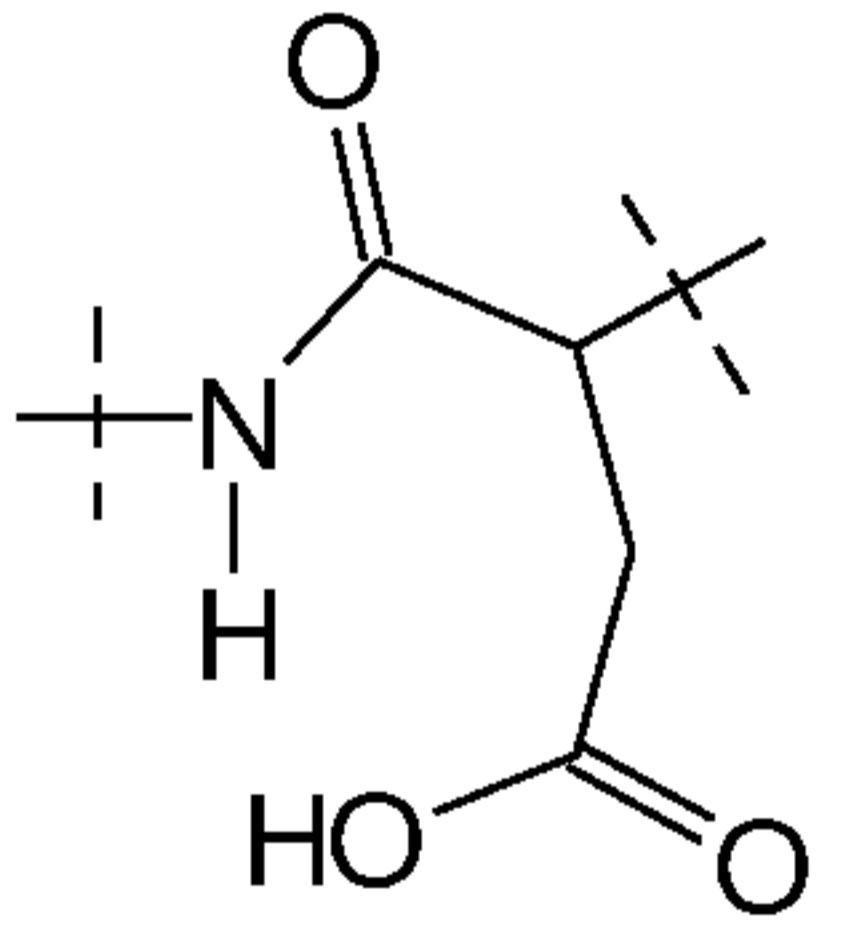
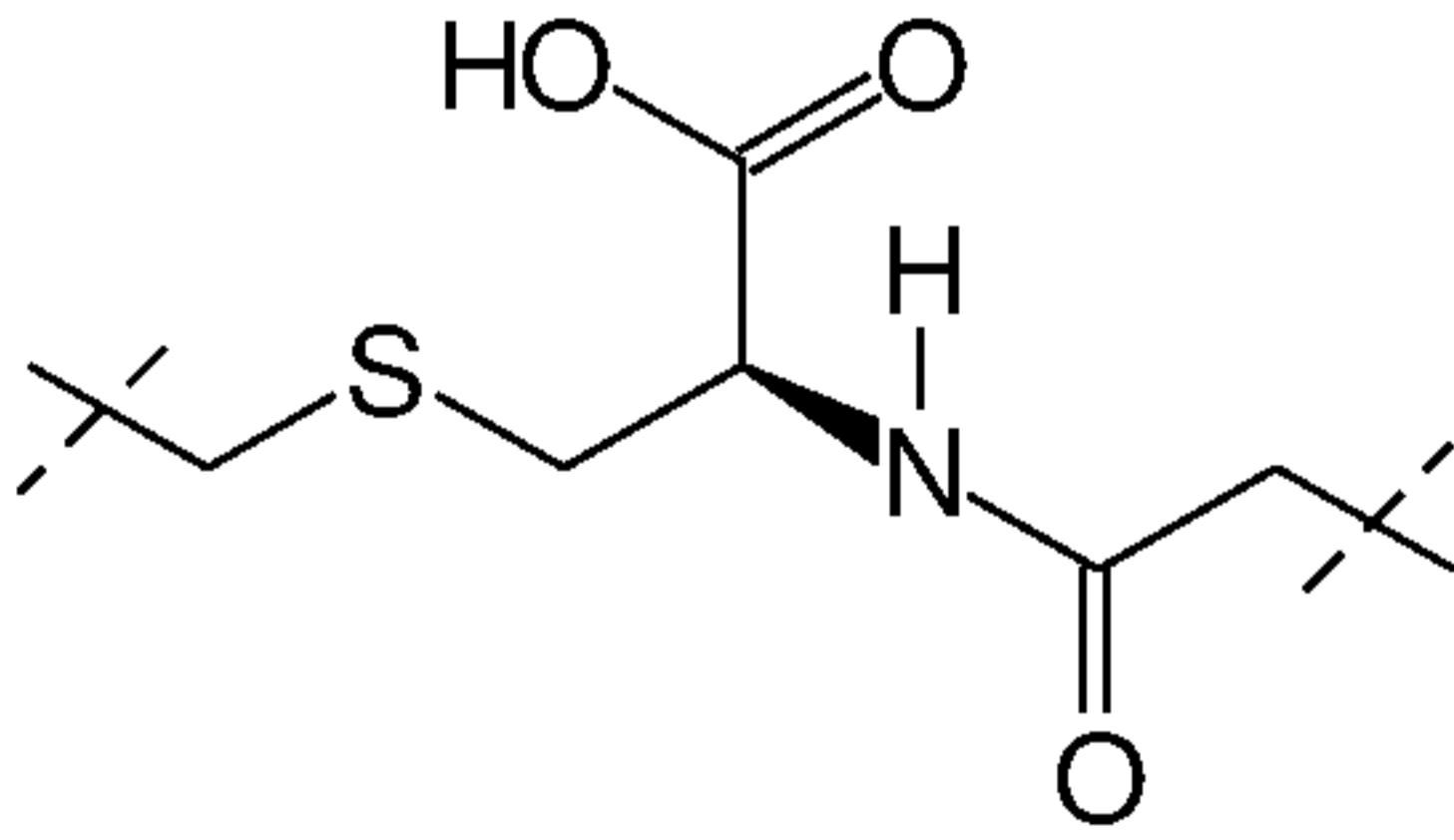
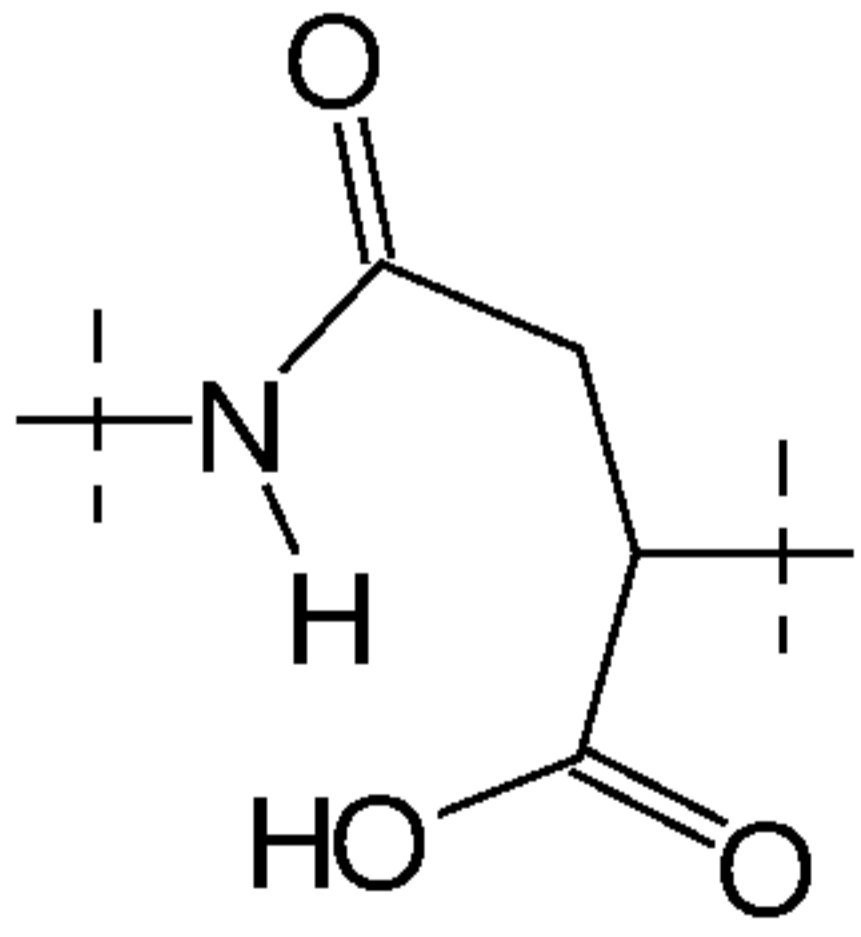
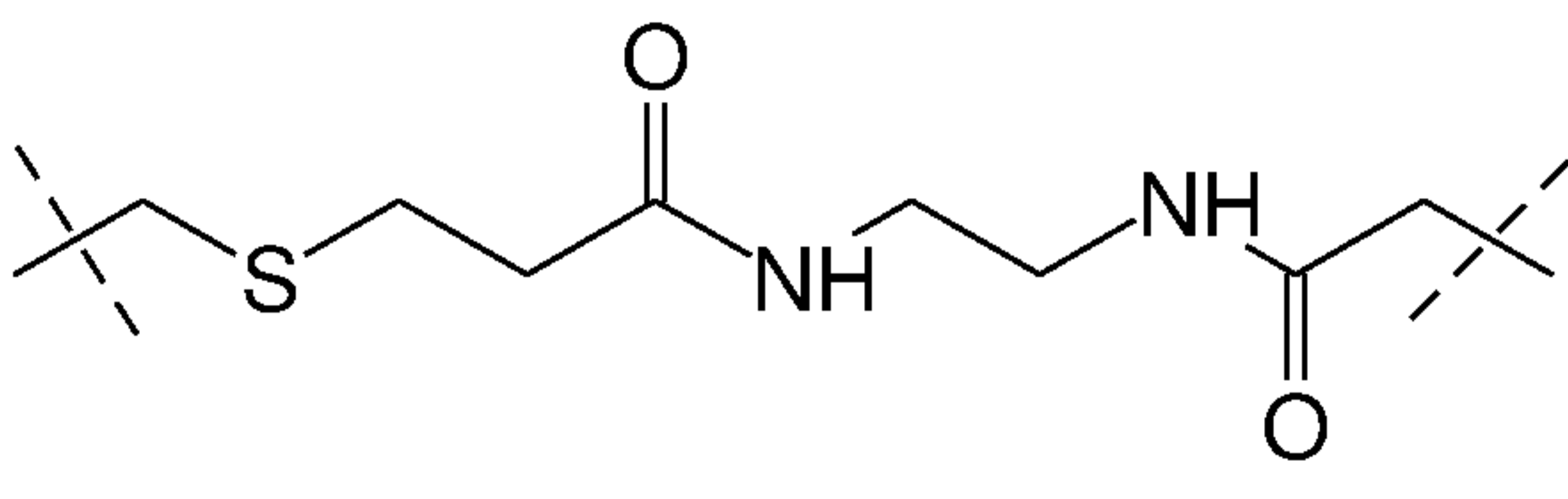
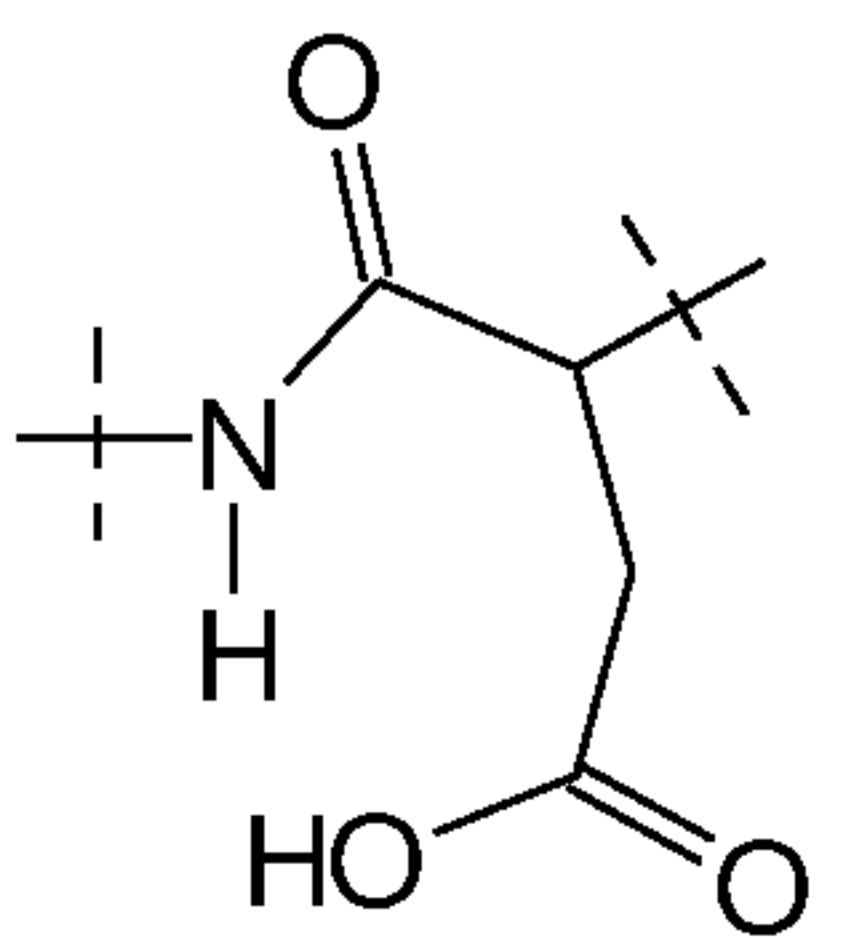
179	R <sup>1</sup>	1		
180	R <sup>1</sup>	1		 See note **
192	R <sup>1</sup>	1		 See note **
193	R <sup>1</sup>	1		 See note **
195	R <sup>1</sup>	0		
196	R <sup>1</sup>	1		
208 / 199	R <sup>1</sup>	1		 and 

				 <p>See note ***</p>
208 / 199	R <sup>1</sup>	1		
208 / 199	R <sup>1</sup>	1		
204	R <sup>1</sup>	1		
209 / 226	R <sup>3</sup>	0		
210	R <sup>3</sup>	0		
211	R <sup>3</sup>	0		

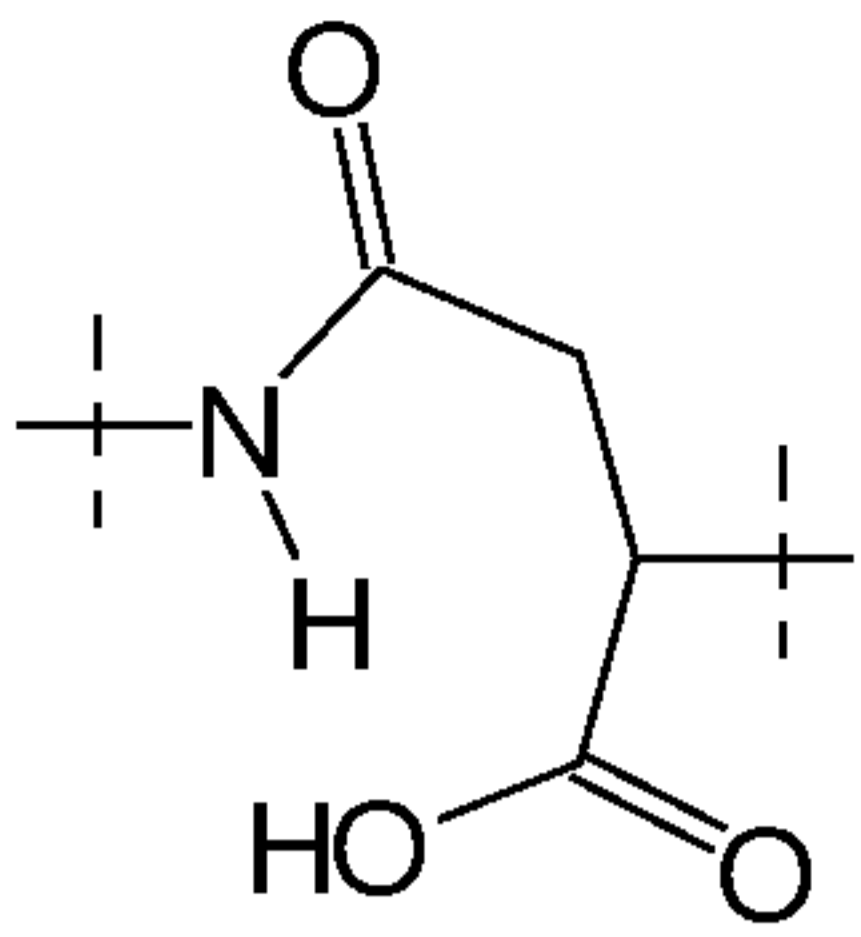
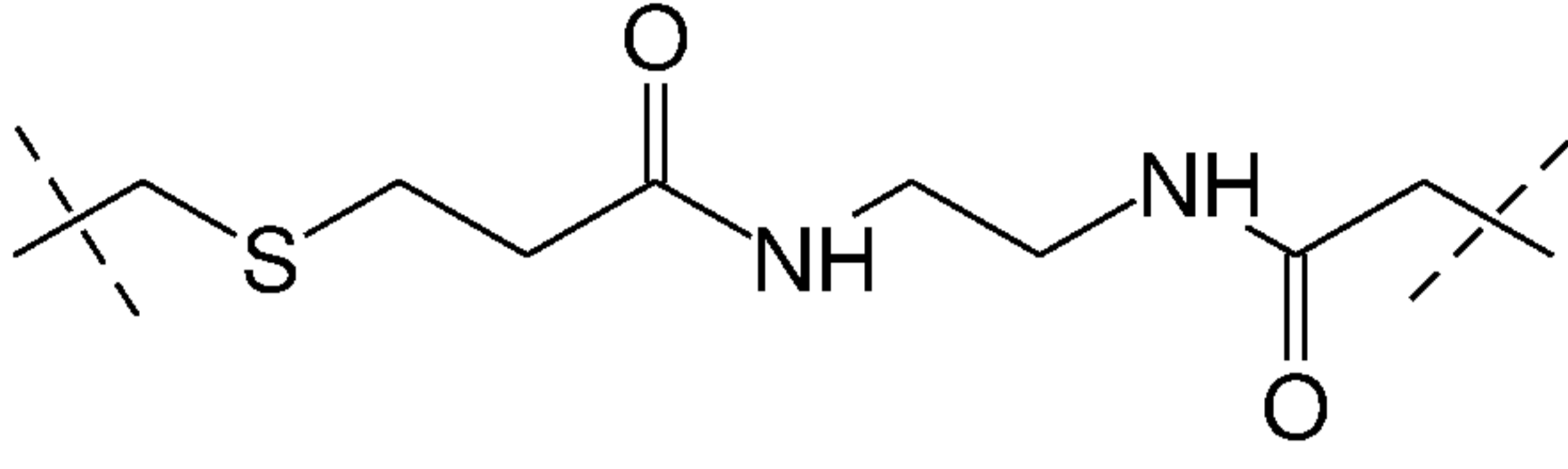
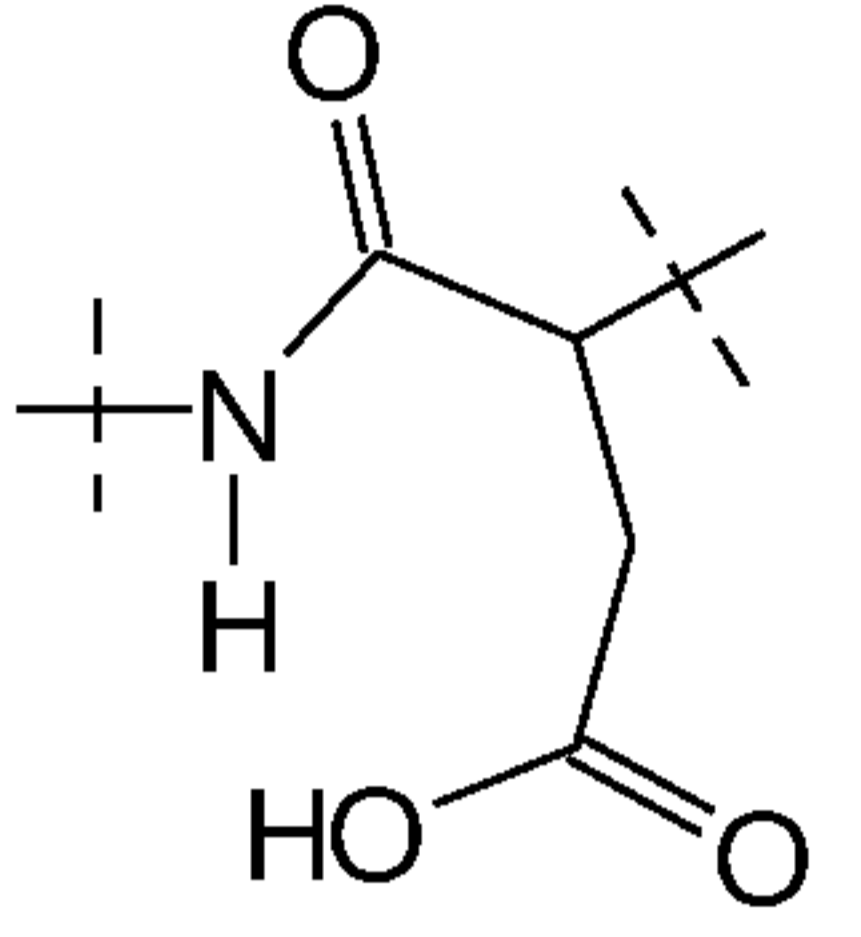
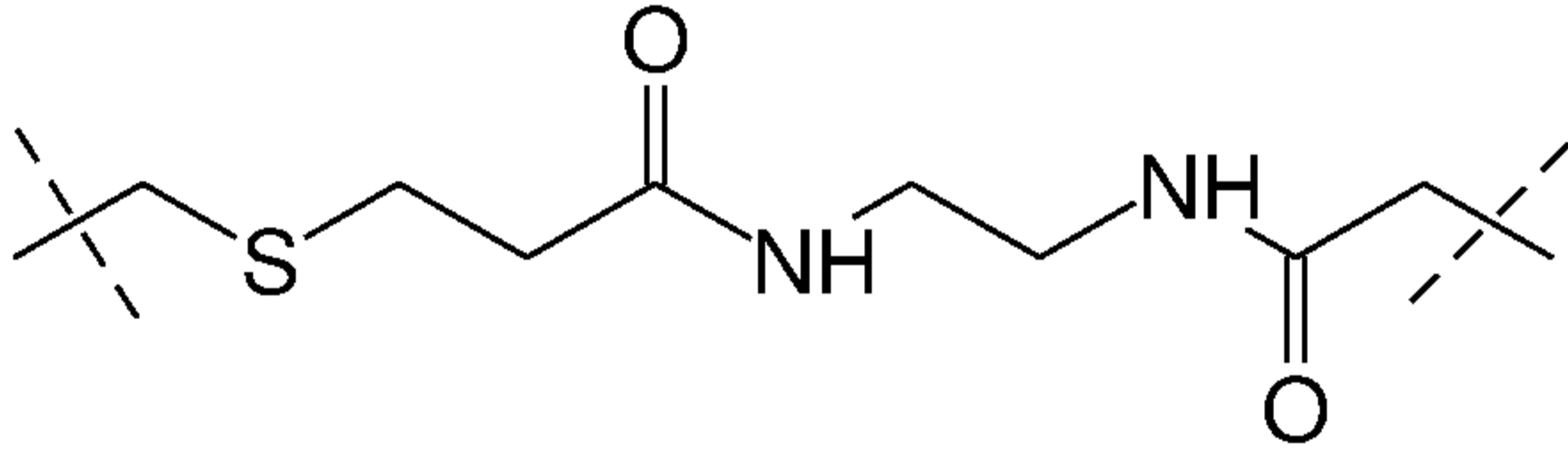
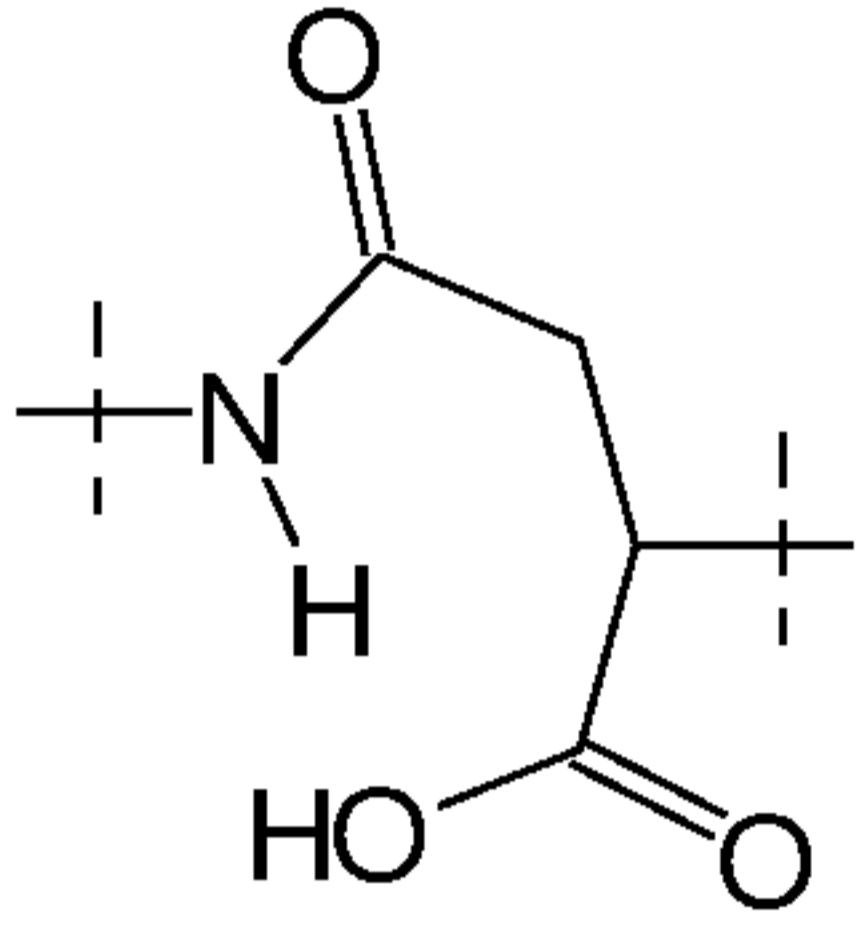
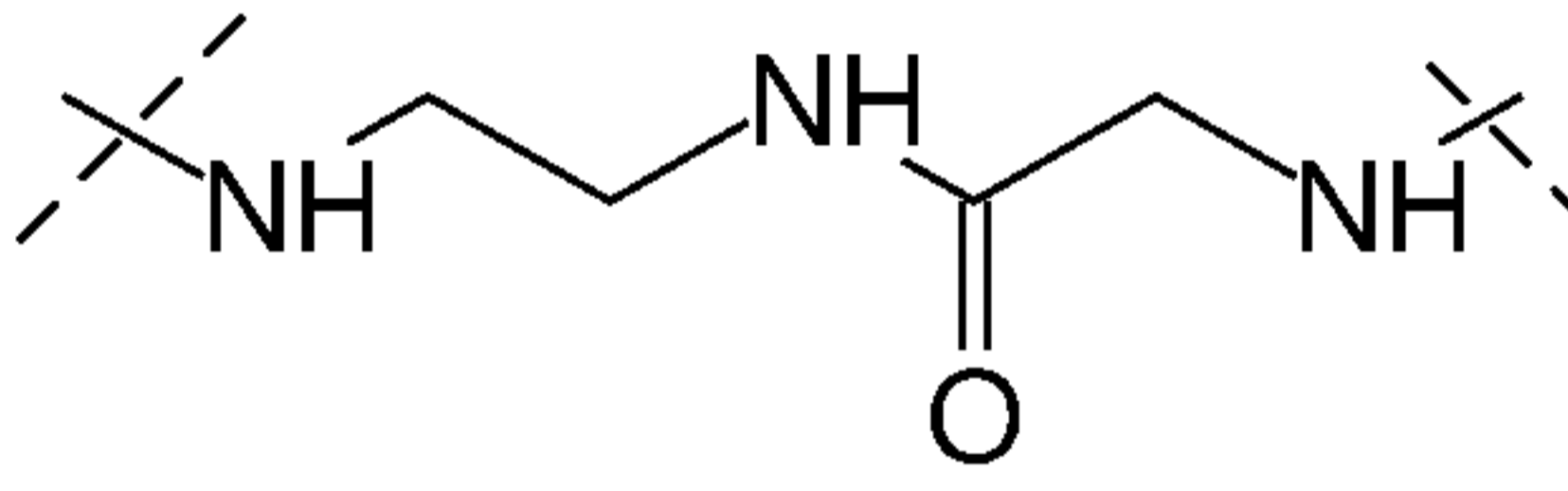
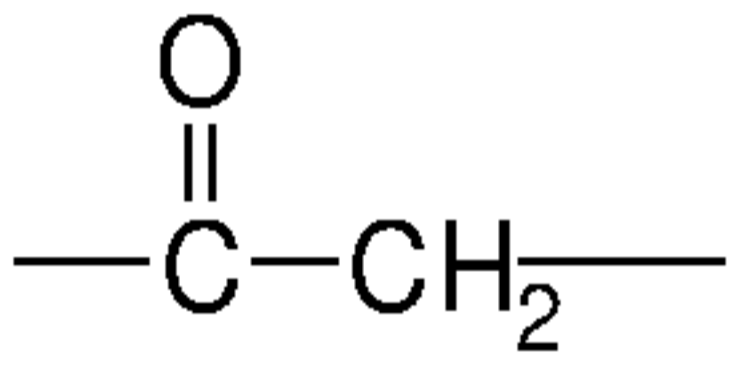
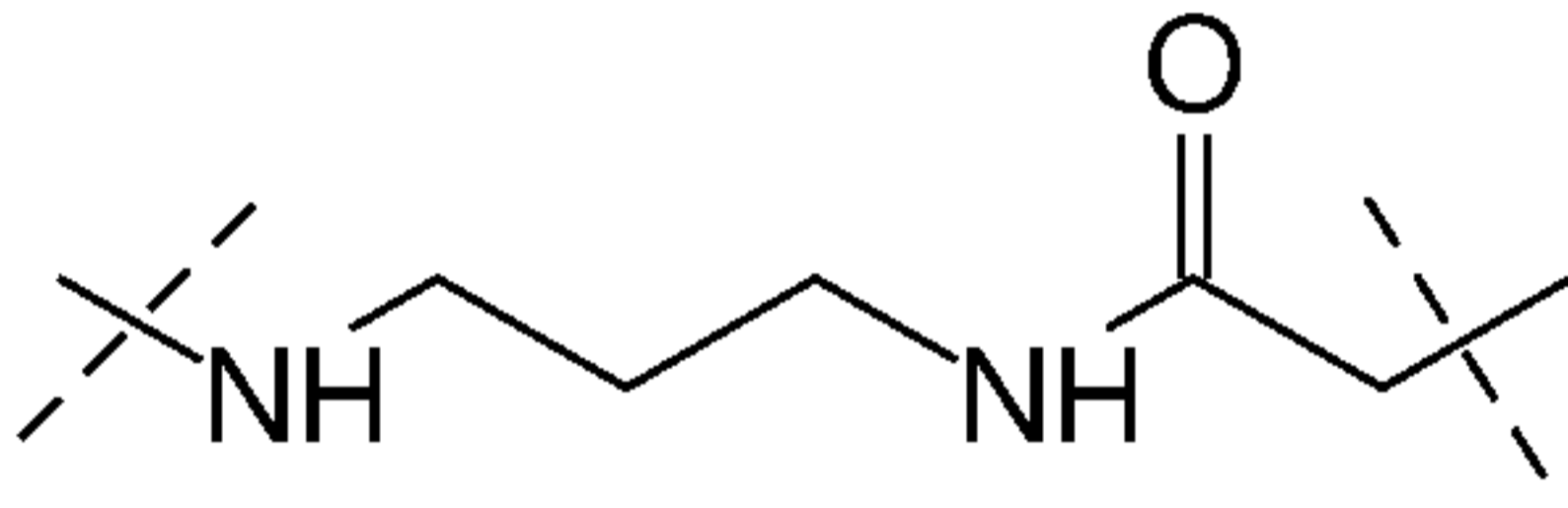
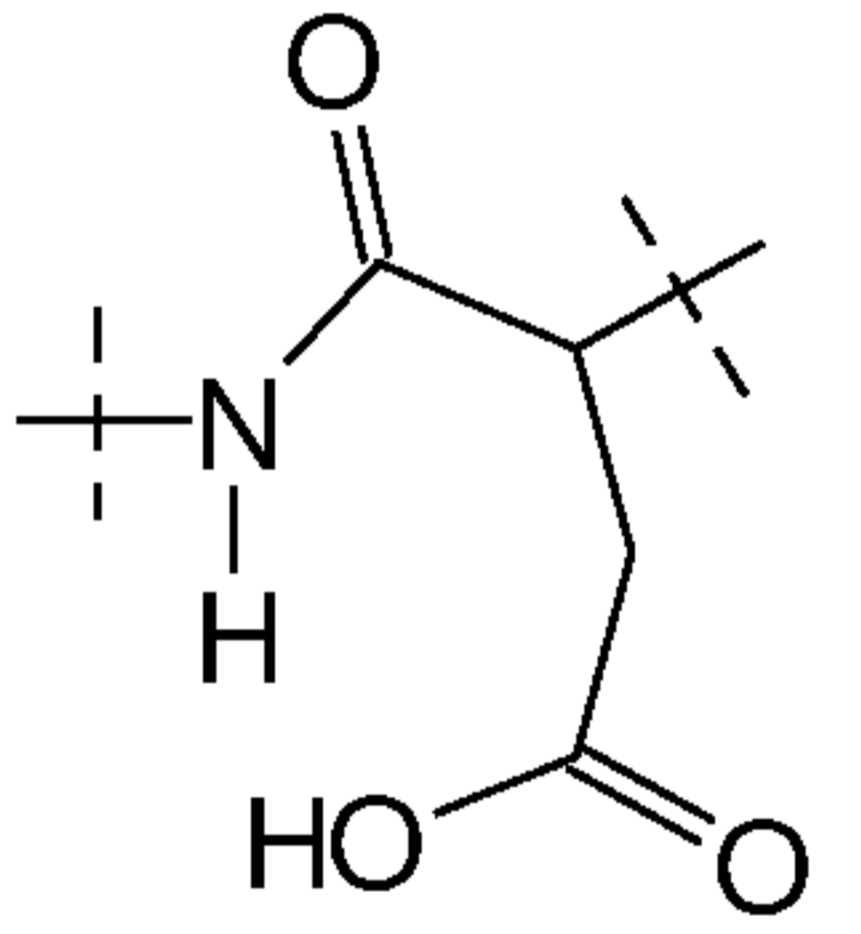
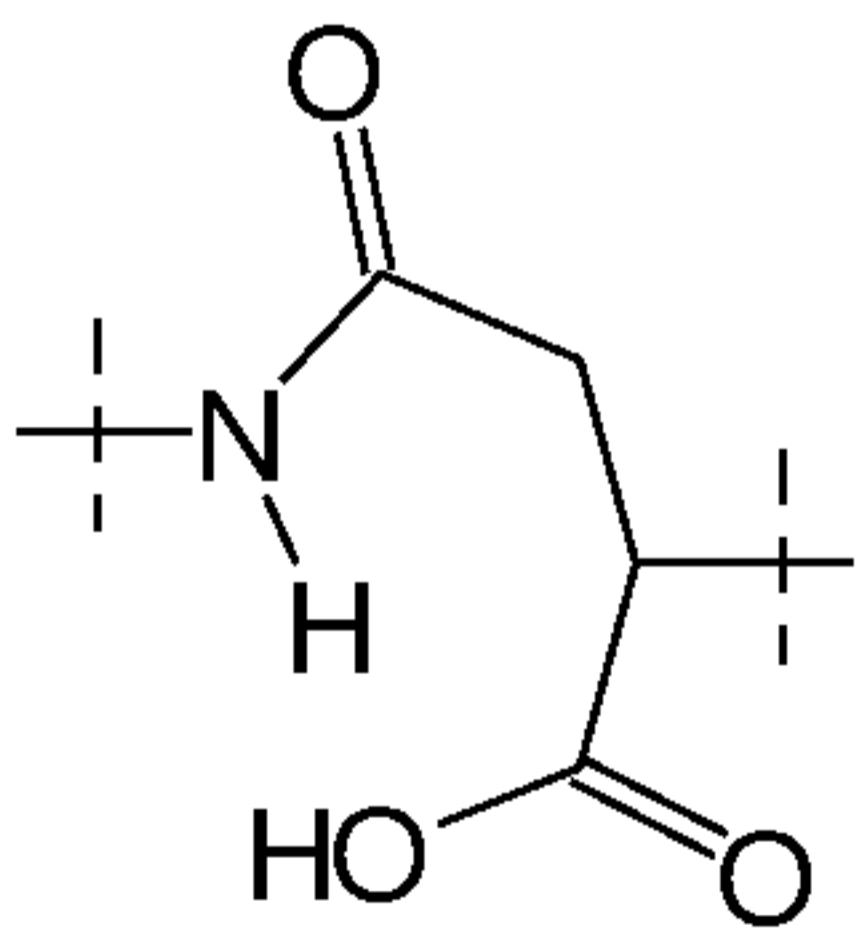
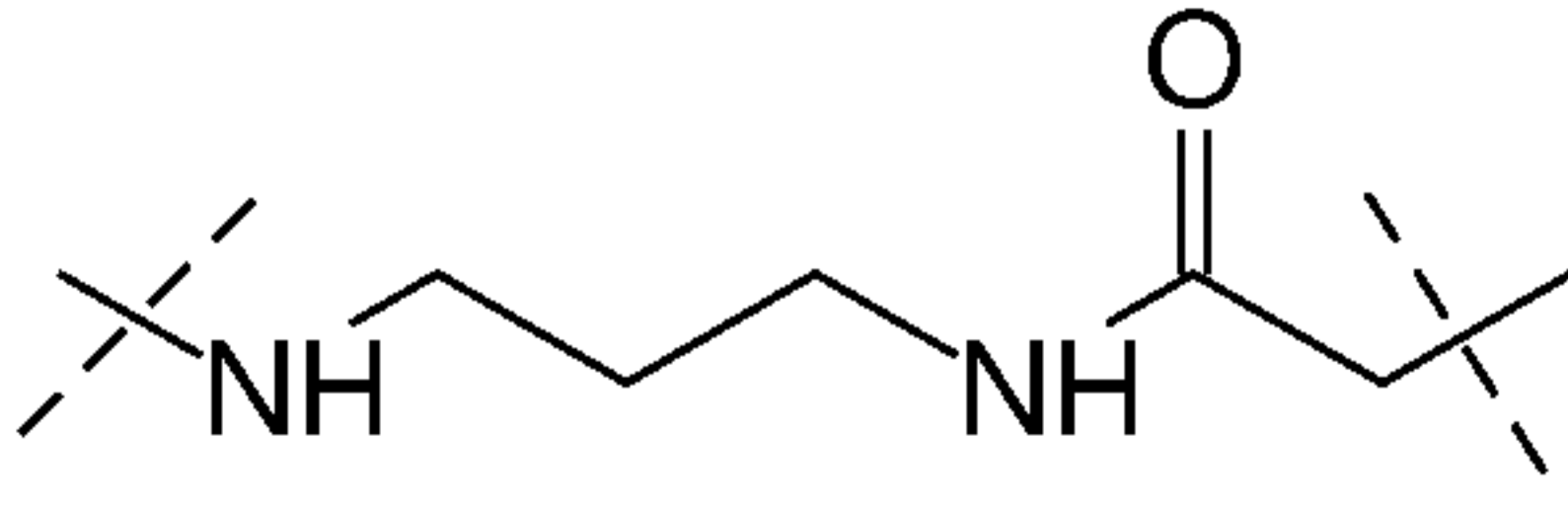
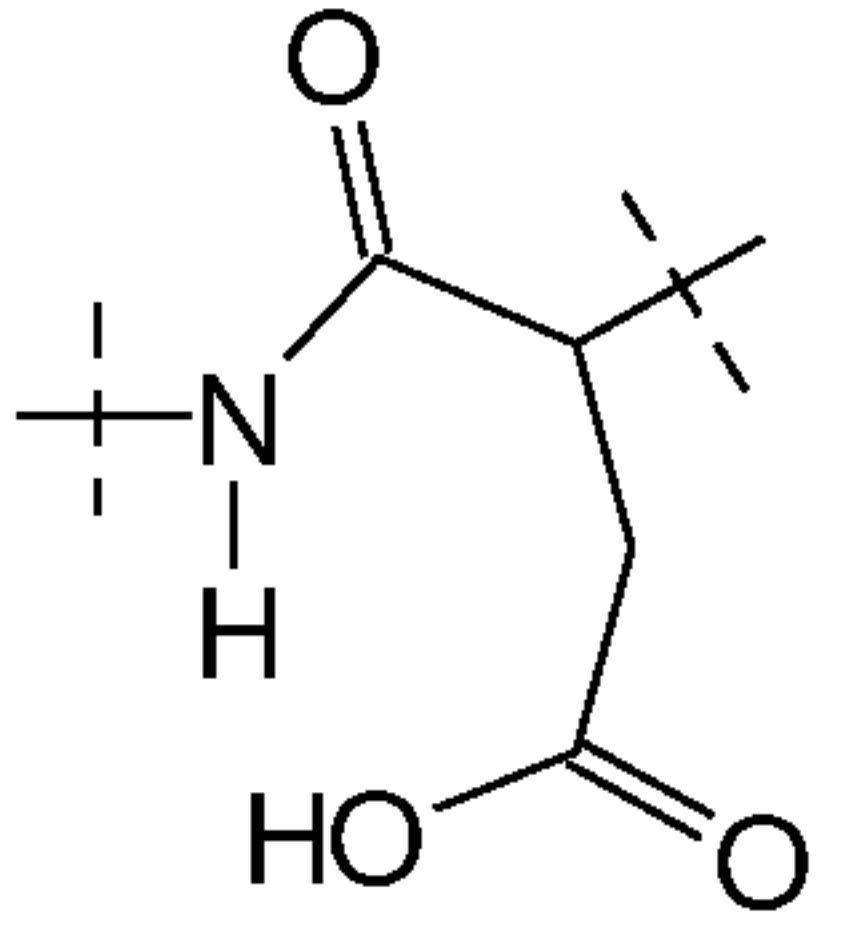
212	R <sup>3</sup>	0		
213	R <sup>3</sup>	0		 See note **
214	R <sup>3</sup>	0		
216	R <sup>3</sup>	0		
217	R <sup>3</sup>	0		 See note **
218	R <sup>3</sup>	0		
238	R <sup>2</sup>	0		

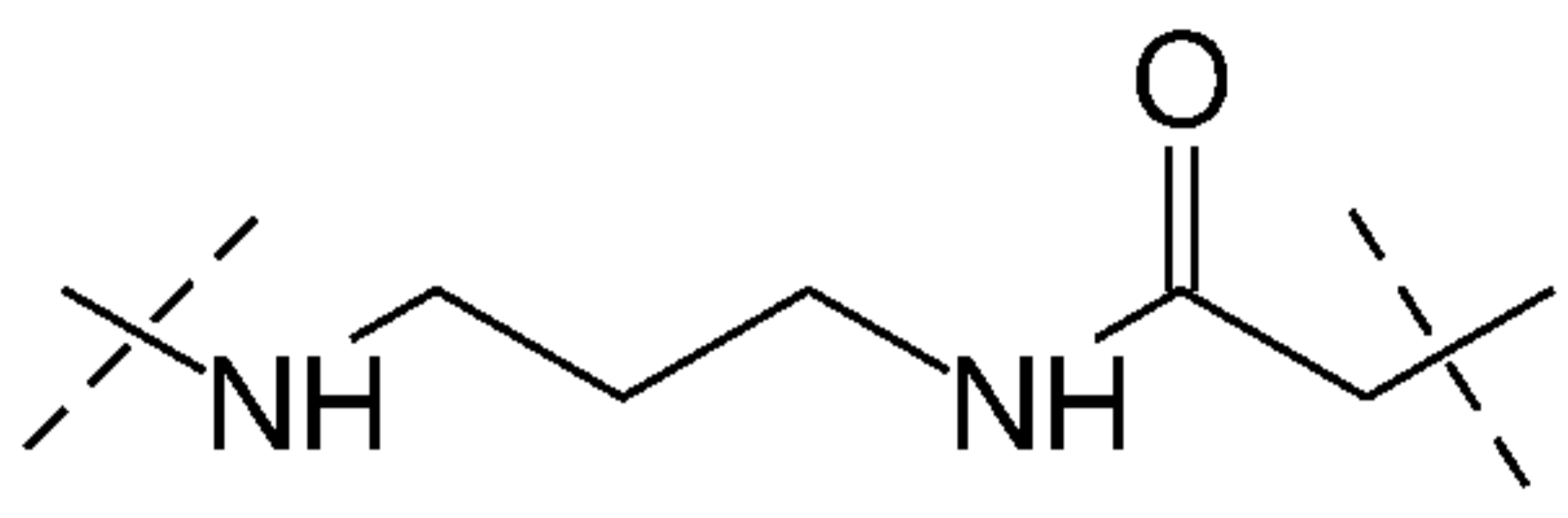
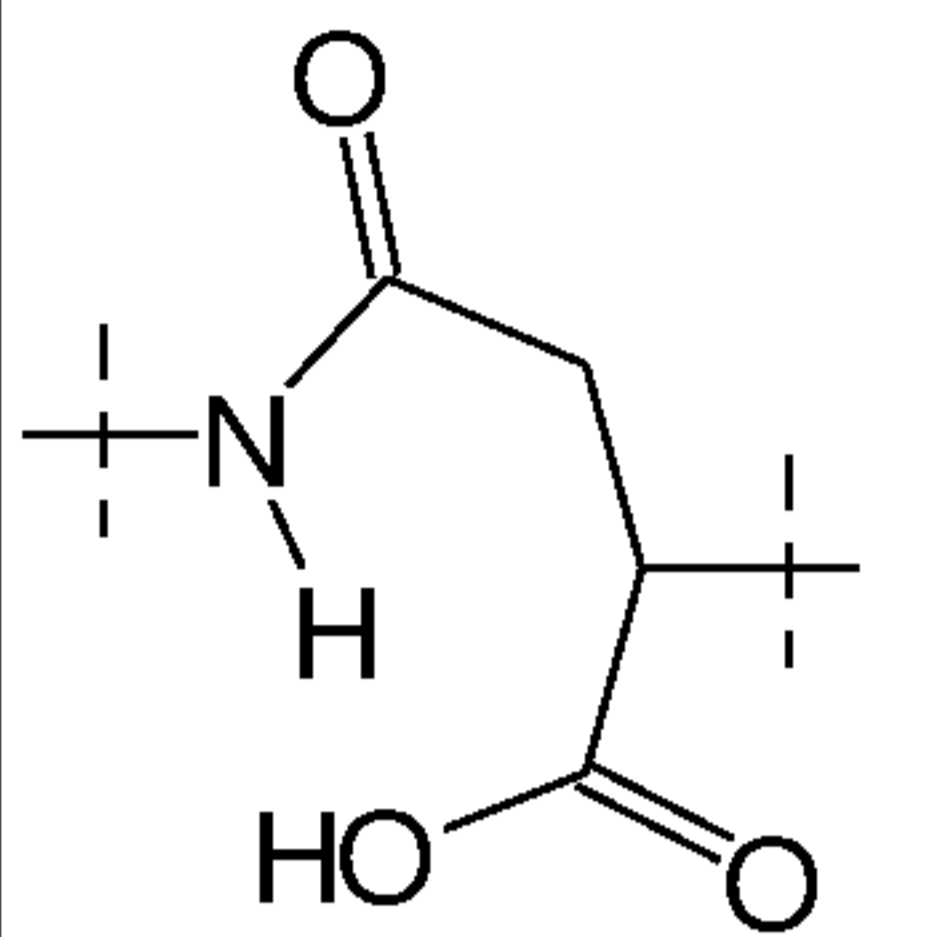
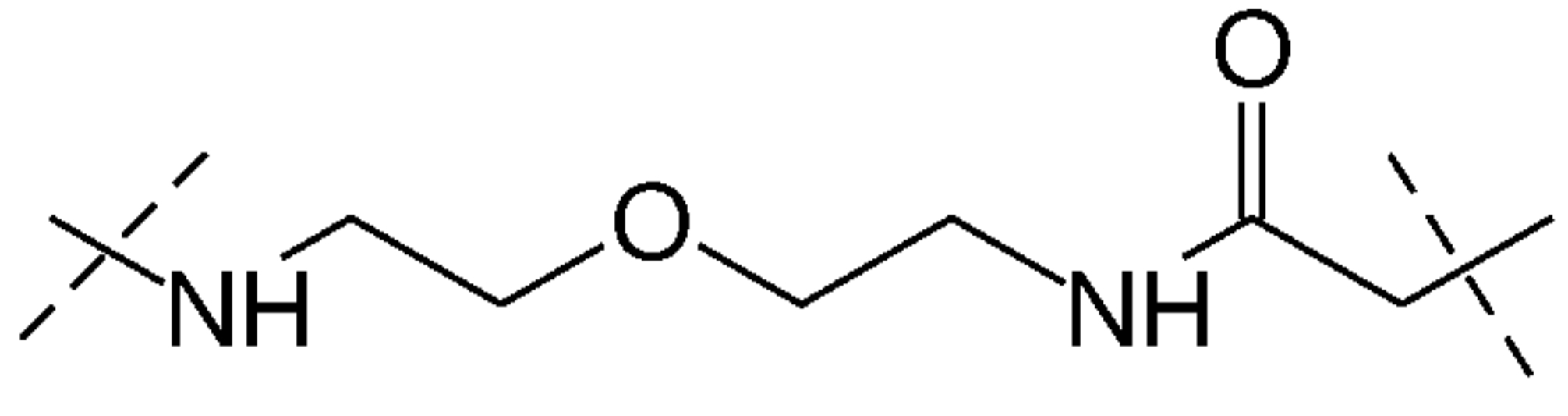
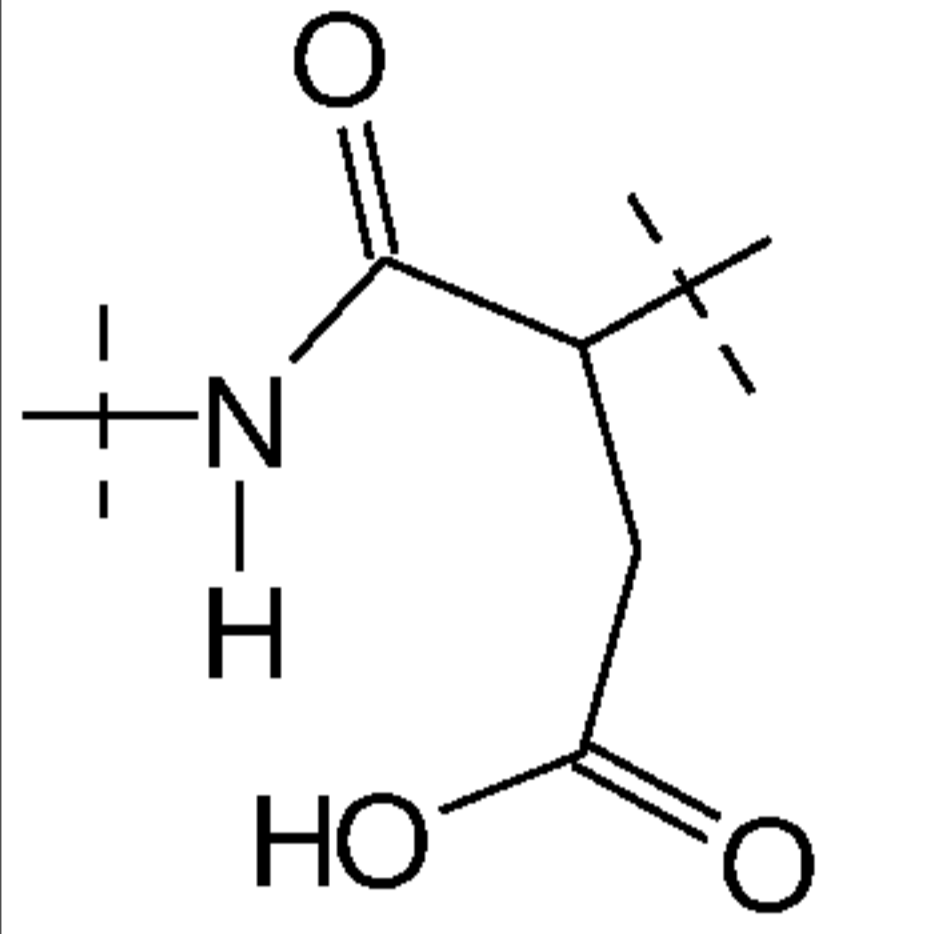
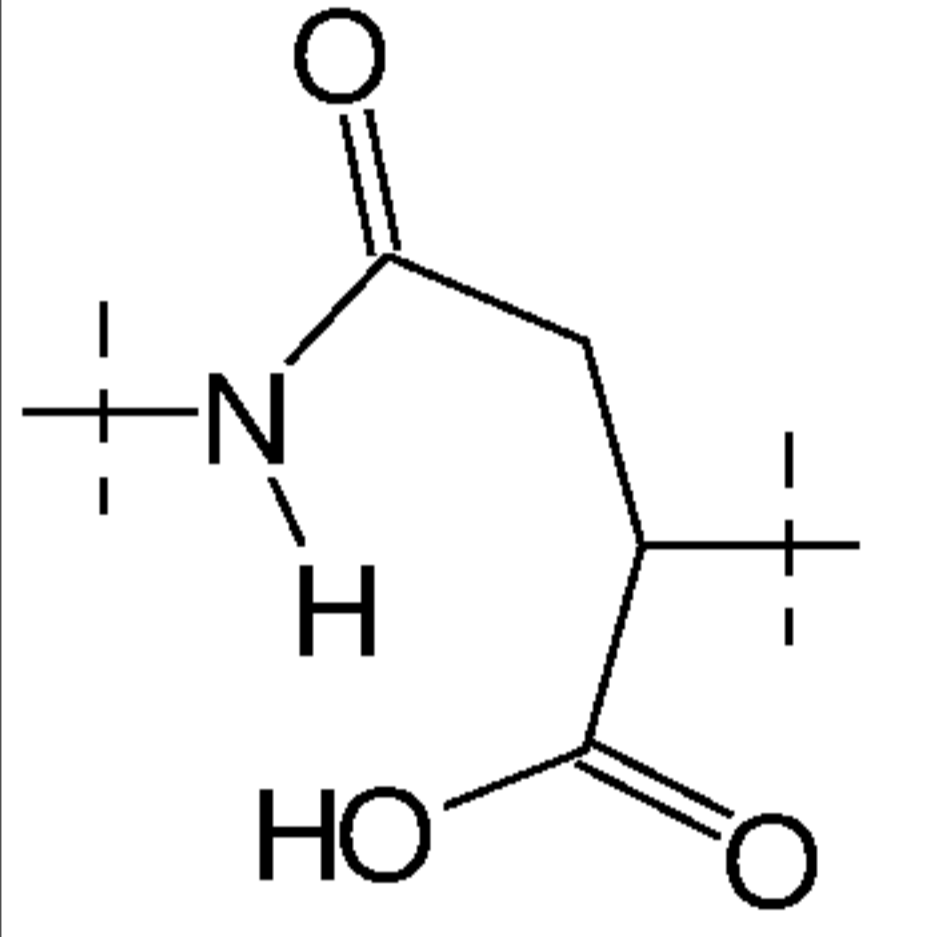
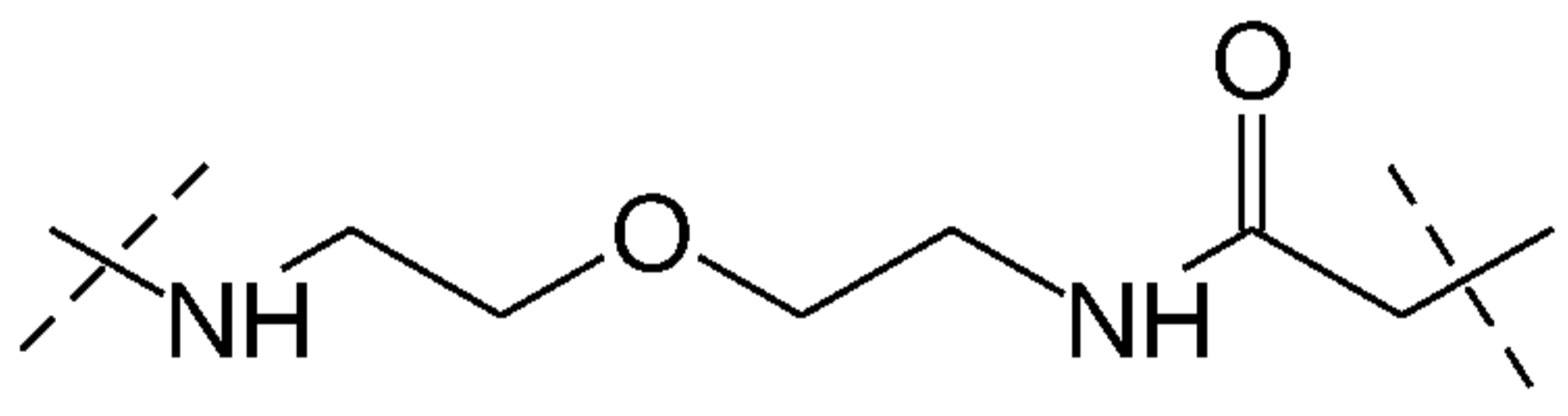
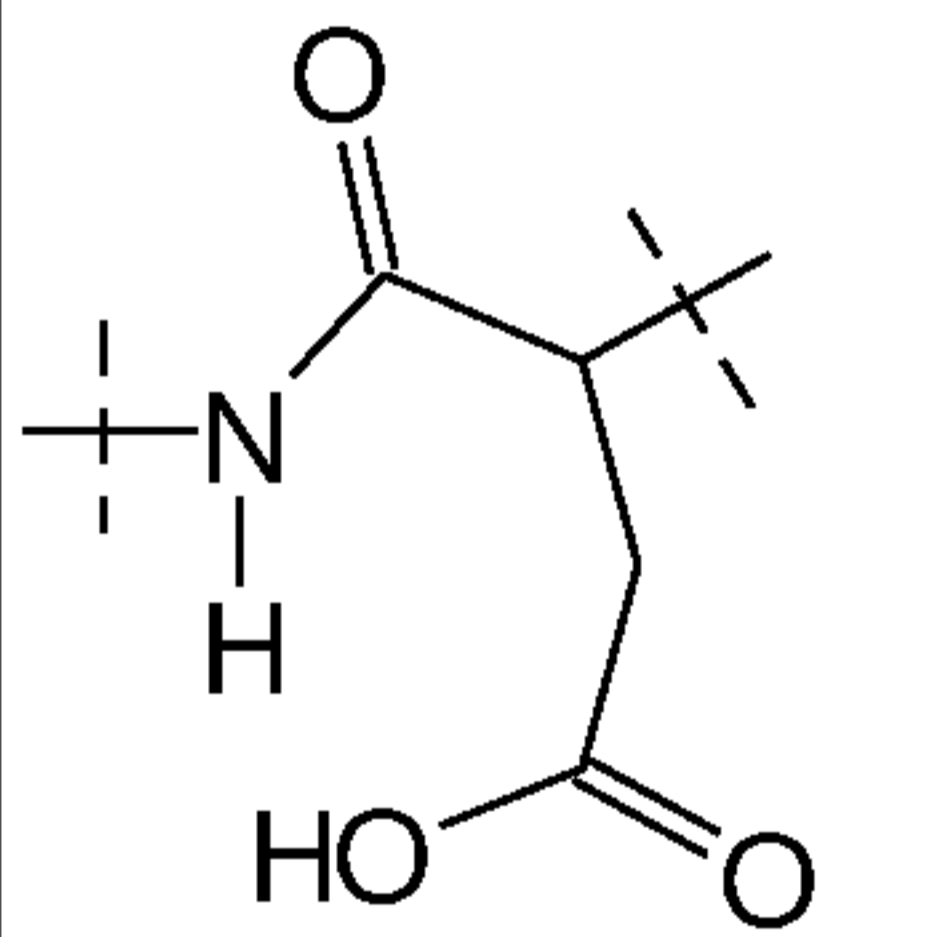
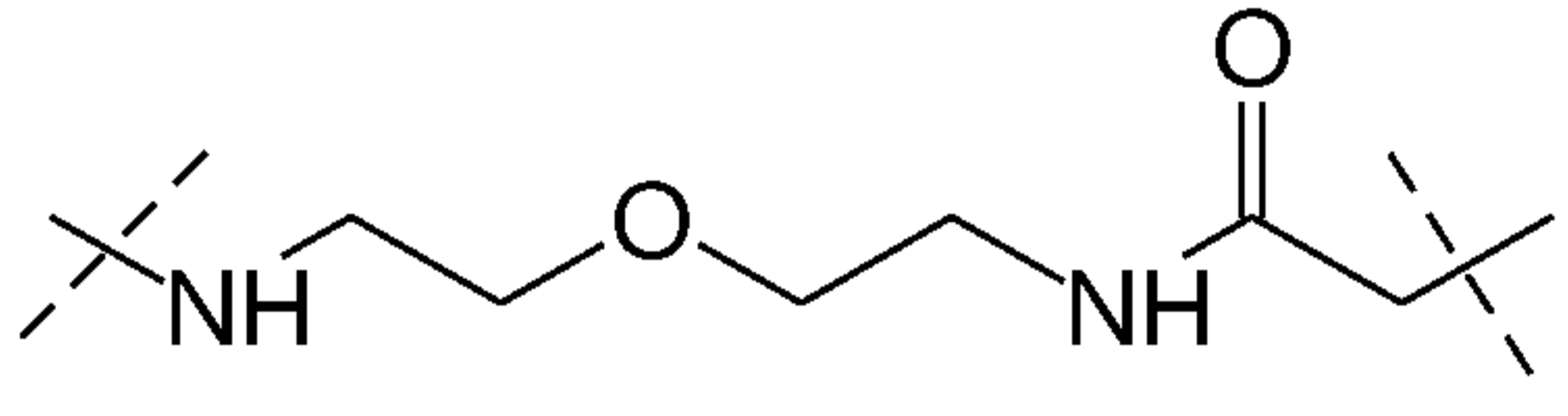
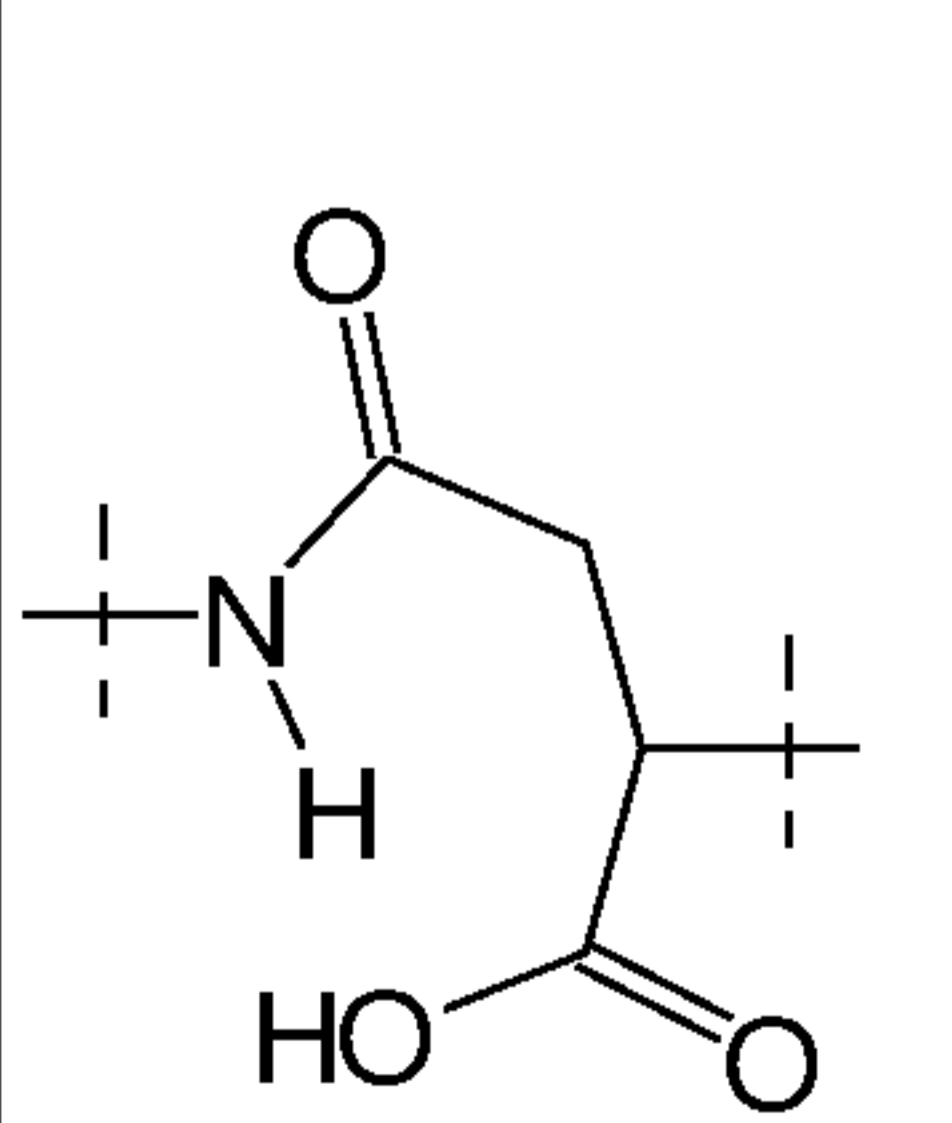
	R <sup>1</sup>	1		<p>where R<sub>22</sub> = - OH or -NH<sub>2</sub></p>
	R <sup>1</sup>	1		<p>where R<sub>22</sub> = - OH or -NH<sub>2</sub></p>
234	R <sup>1</sup>	1		<p>and</p> <p>See note ***</p>
234	R <sup>1</sup>	1		
234	R <sup>1</sup>	1		

236	R <sup>1</sup>	1		 and  See note ***
236	R <sup>1</sup>	1		
236	R <sup>1</sup>	1		
237	R <sup>3</sup>	0		 and  See note ***

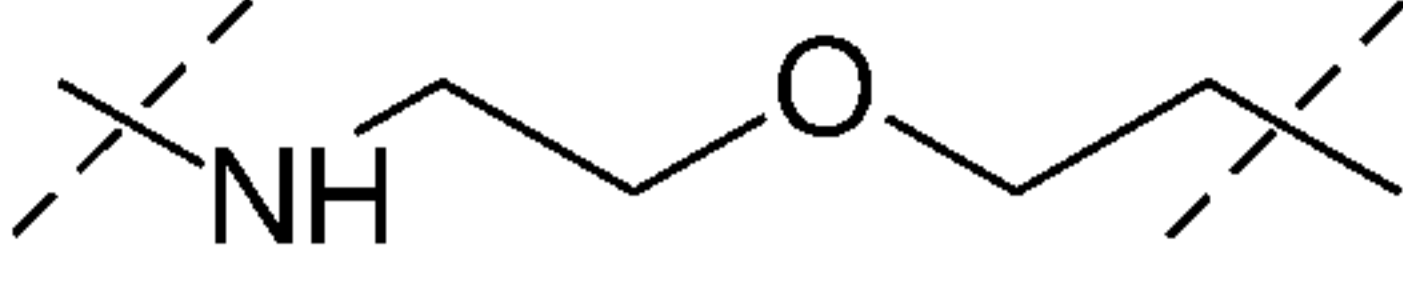
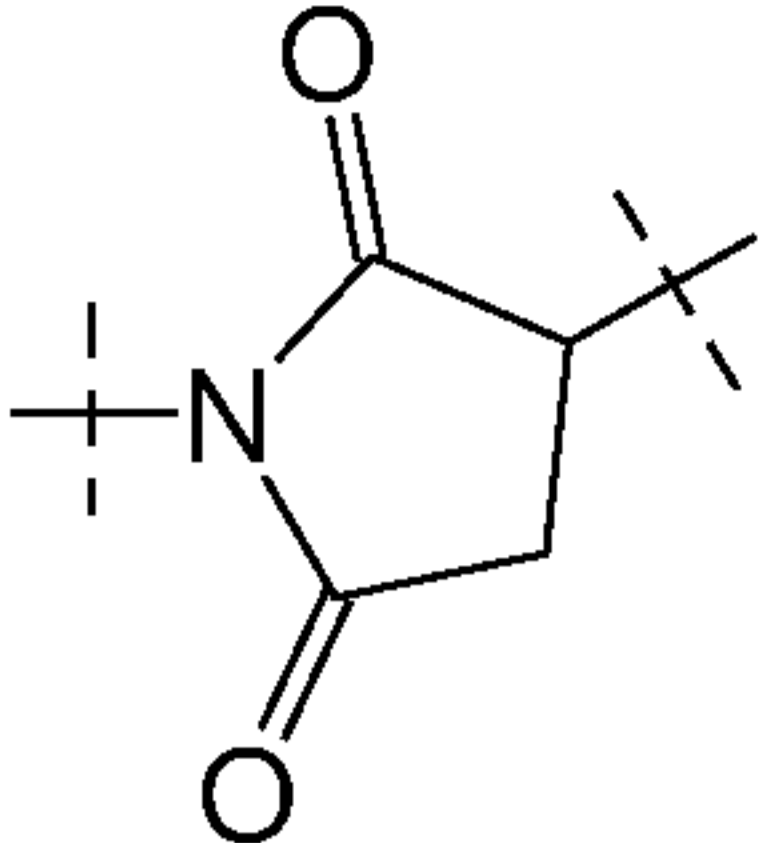
237	R <sup>3</sup>	0		
237	R <sup>3</sup>	0		
239	R <sup>3</sup>	0		 and  See note ***
239	R <sup>3</sup>	0		
239	R <sup>3</sup>	0		
240	R <sup>3</sup>	0		



				and  See note ***
240	R <sup>3</sup>	0		
240	R <sup>3</sup>	0		
241	R <sup>1</sup>	1		
242	R <sup>1</sup>	1		 and  See note ***
242	R <sup>1</sup>	1		

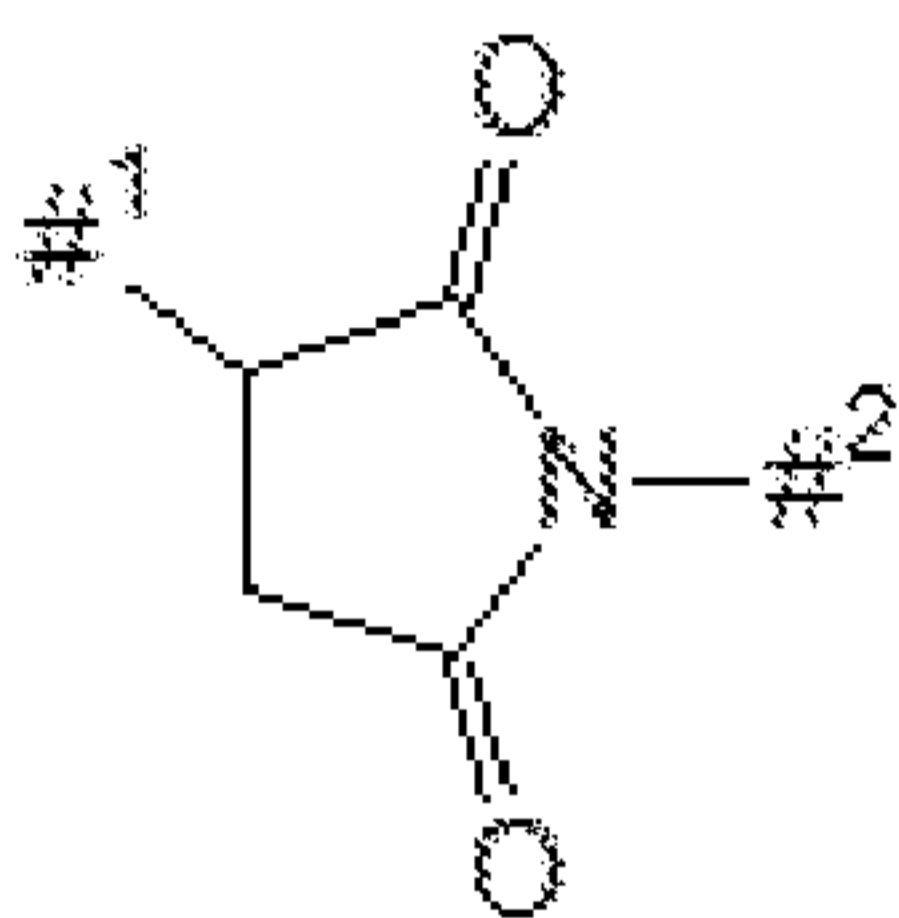
242	R <sup>1</sup>	1		
243	R <sup>1</sup>	1		 <p data-bbox="1541 1151 1619 1190">and</p>  <p data-bbox="1541 1528 1745 1626">See note ***</p>
243	R <sup>1</sup>	1		
243	R <sup>1</sup>	1		

244	R3	0		
245	R1	0		<p>and</p> <p>See note ***</p>
245	R1	0		
245	R1	0		
247	R1	1		

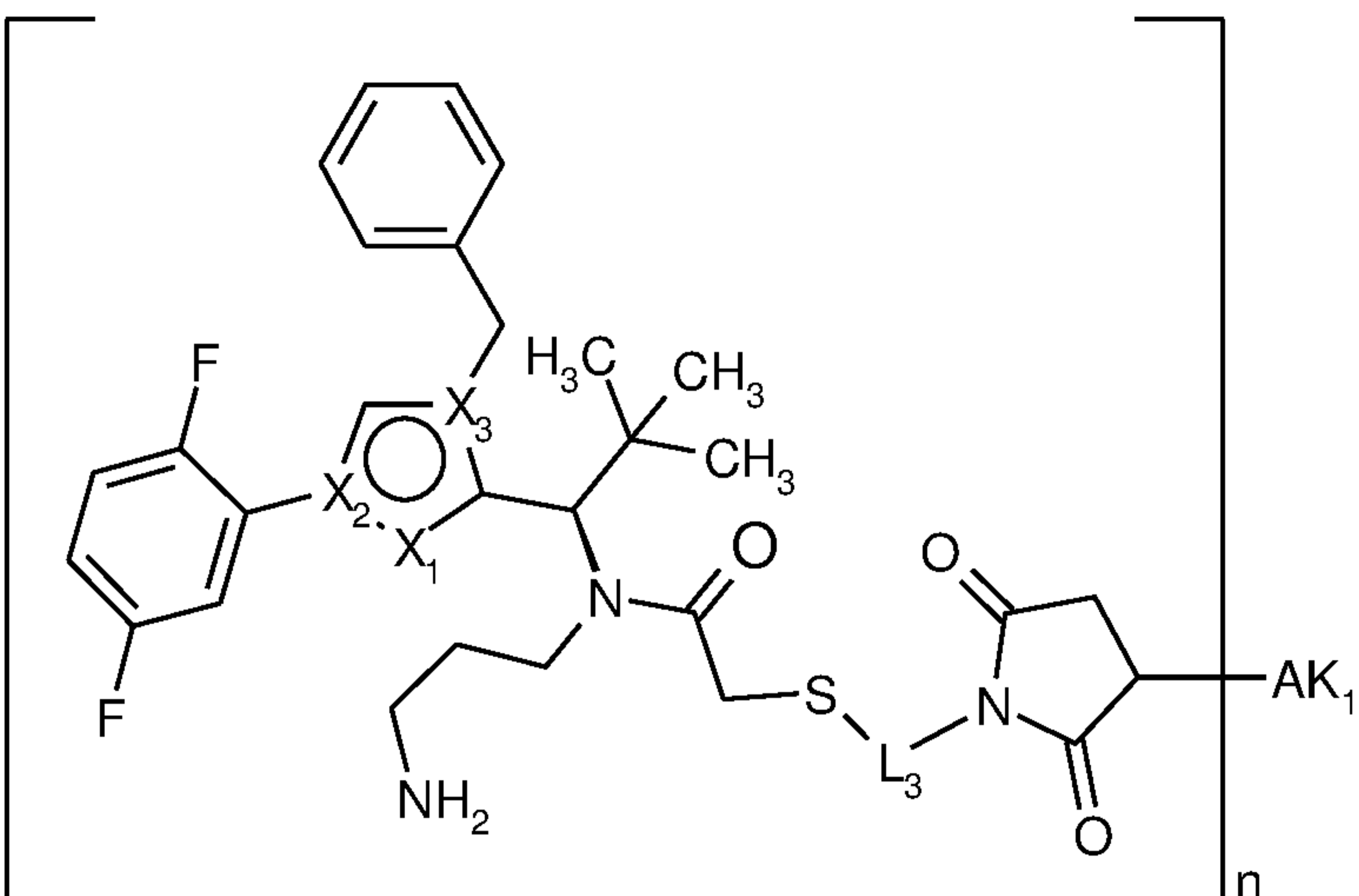
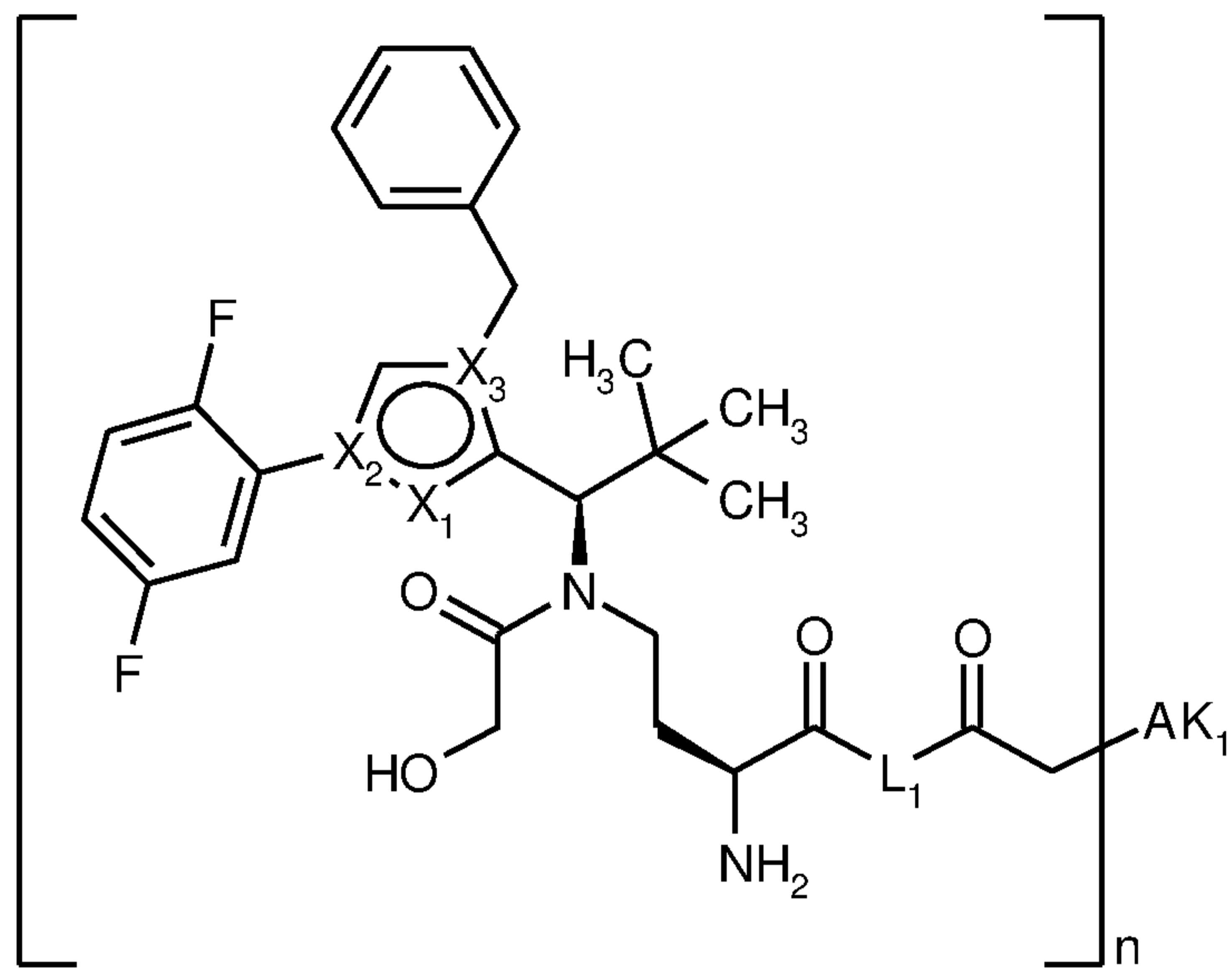
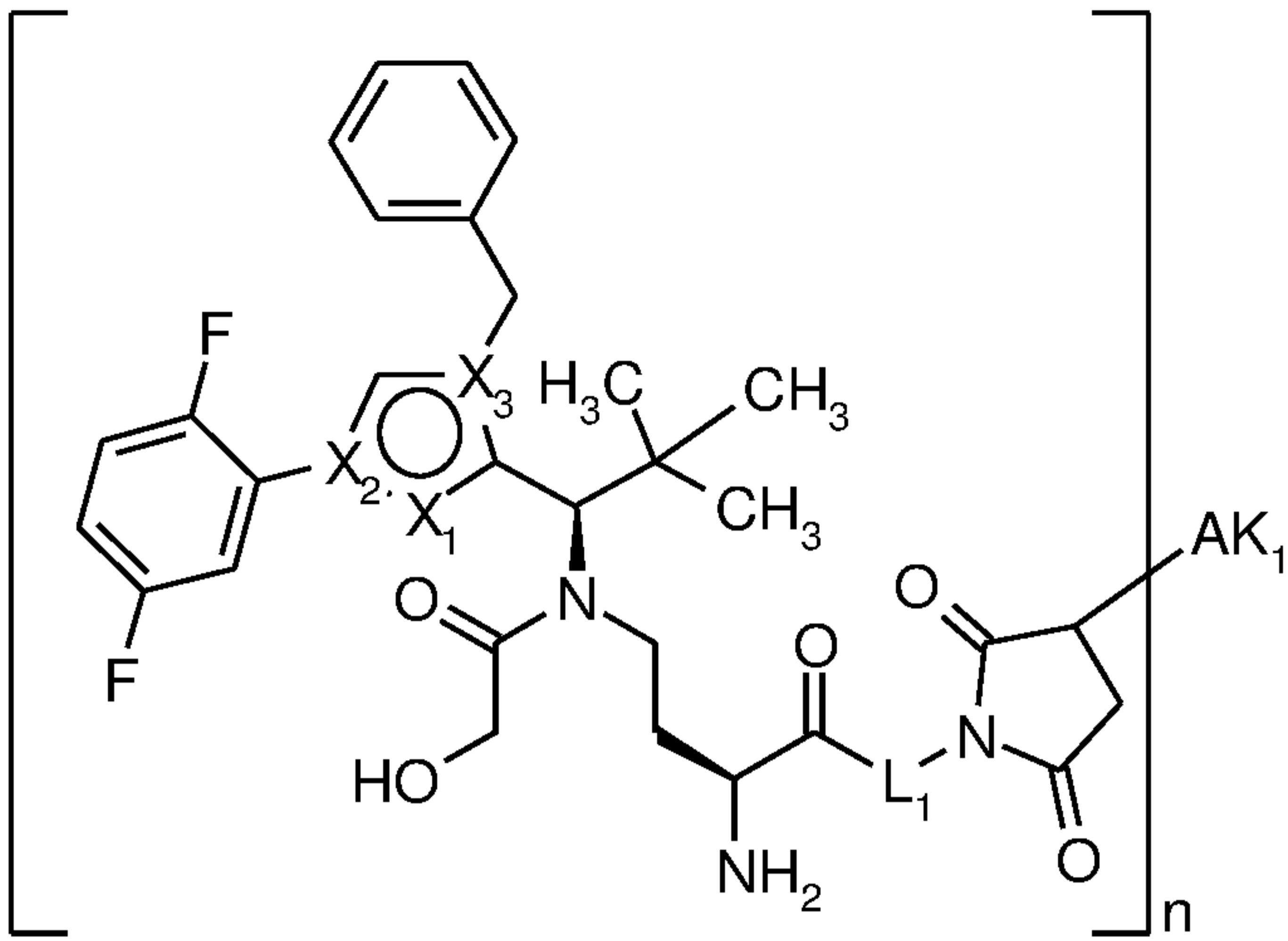
248	R1	1		
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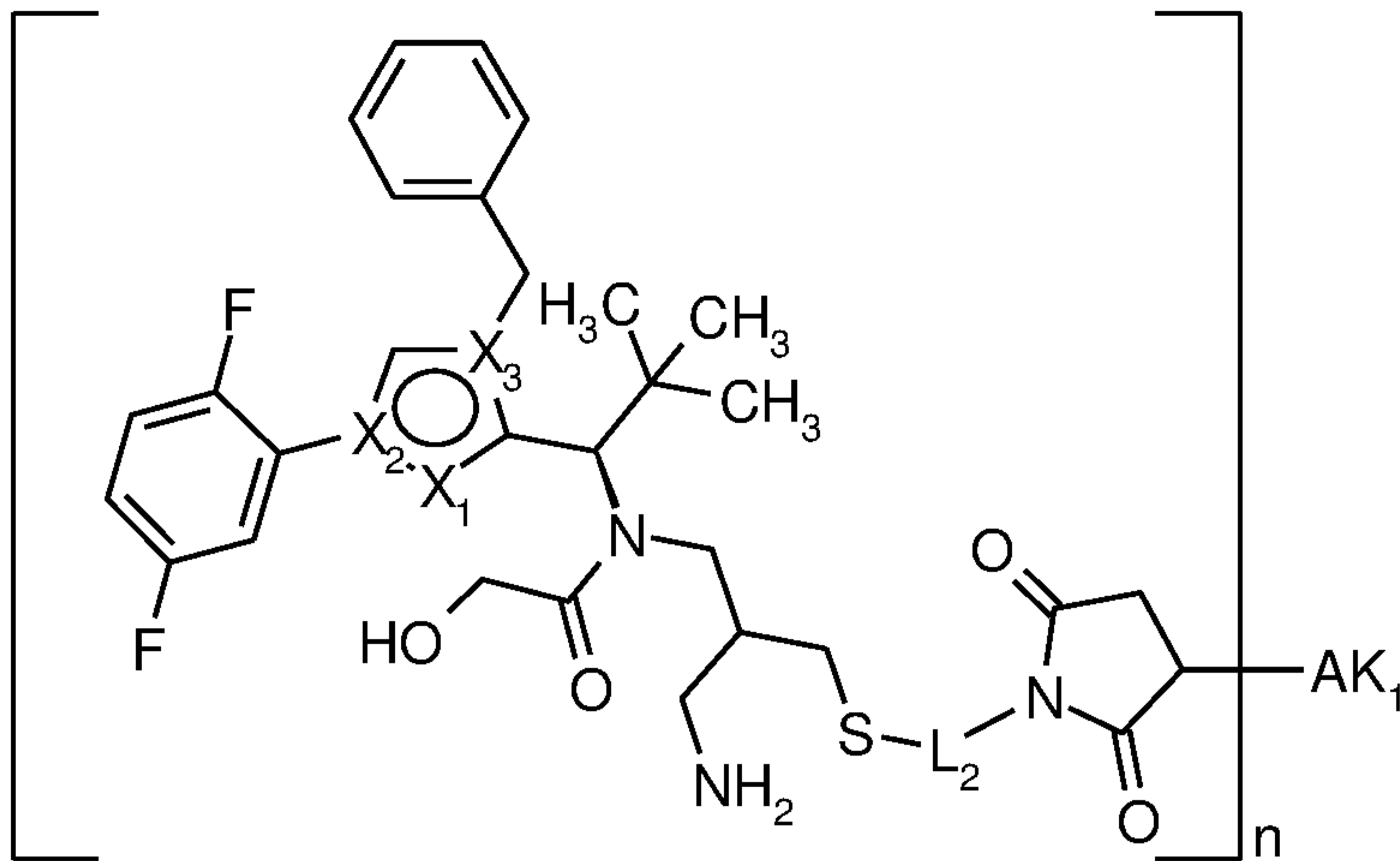
\*\* : See note \*\* for Table A.

\*\*\* : When this structure L2 is present, there may simultaneously  
5 be a structure L2 of the formula below:



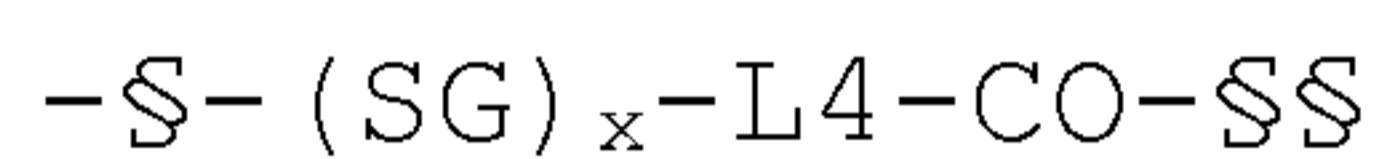
Examples of conjugates having corresponding linkers have the  
10 following structures, where X1, X2, X3 and L1 have the meanings  
given above, L2 and L3 have the same meaning as L1, AK1  
represents an antibody attached via a cysteine residue and n is  
a number from 1 to 10. With particular preference, AK1 is a  
15 human, humanized or chimeric monoclonal antibody or an antigen-  
binding fragment thereof, in particular an anti-TWEAKR antibody  
or an antigen-binding fragment thereof or an anti-EGFR antibody  
or an antigen-binding fragment thereof. Particular preference  
is given to an anti-TWEAKR antibody which binds specifically to  
amino acid D in position 47 (D47) of TWEAKR (SEQ ID NO:169), in  
20 particular the anti-TWEAKR antibody TPP-2090, or the anti-EGFR  
antibodies cetuximab or nimotuzumab.





If the linker is attached to a lysine side chain or a lysine residue, it preferably has the formula below:

5



where

10  $\$$  represents the bond to the active compound molecule and

$\$\$$  represents the bond to the binder peptide or protein,

x represents 0 or 1,

15

SG represents a cleavable group, preferably a 2-8 oligopeptide, particularly preferably a dipeptide,

and

20

L4 represents a single bond or a group  $-(CO)_y-G4-$ , where y represents 0 or 1, and

25 G4 represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups  $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-NHCO-$ ,  $-CONH-$ ,  $-NMe-$ ,  $-NHNH-$ ,  $-SO_2NHNH-$ ,  $-CONHNH-$  and a 5- to 10-membered aromatic or

non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, -SO- or -SO<sub>2</sub>-

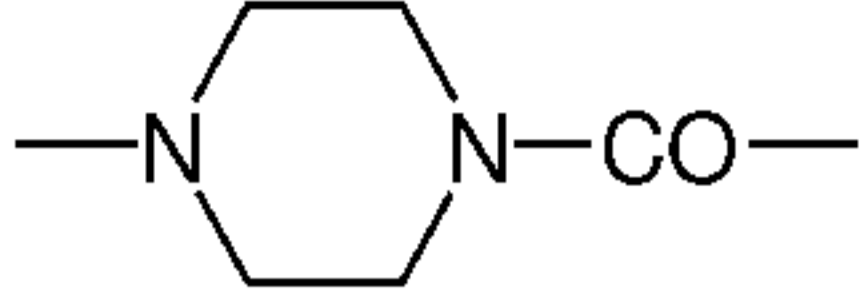
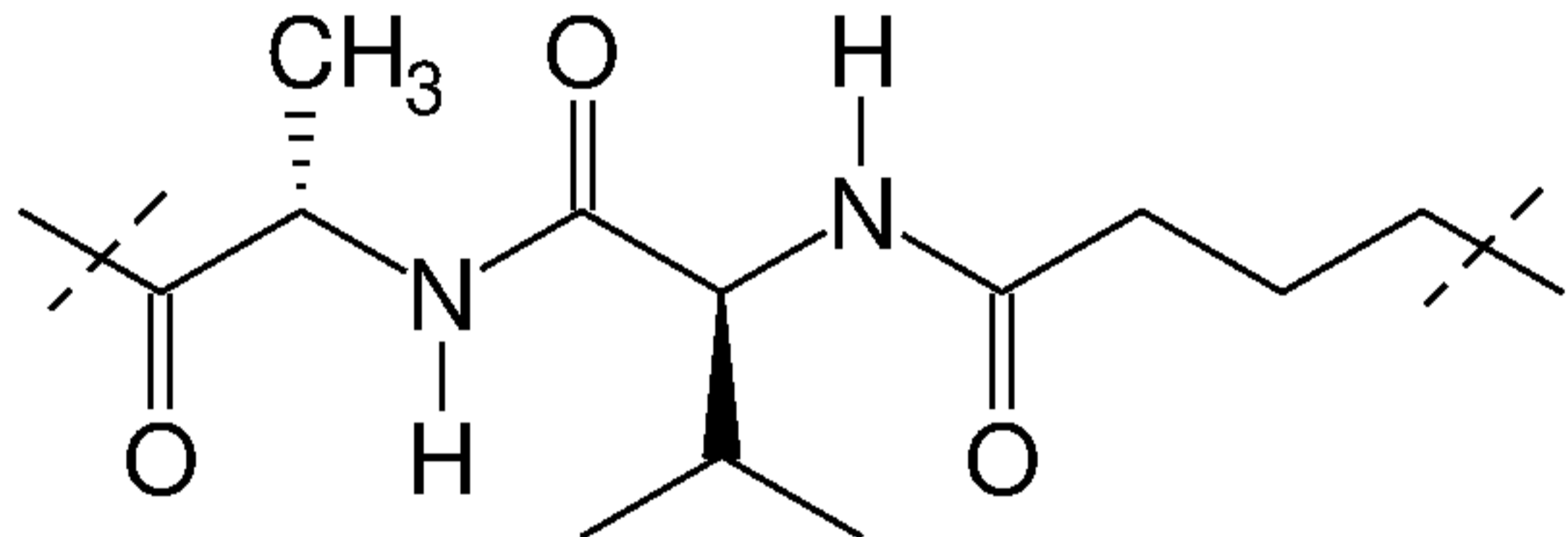
(preferably ) , where the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphamide, sulphone, sulphoxide or sulphonic acid.

Table B below gives examples of linkers to a lysine residue. The table furthermore gives the preferred coupling point (R<sup>1</sup>-R<sup>5</sup>). The first column furthermore states the example numbers in which the corresponding linkers are used.

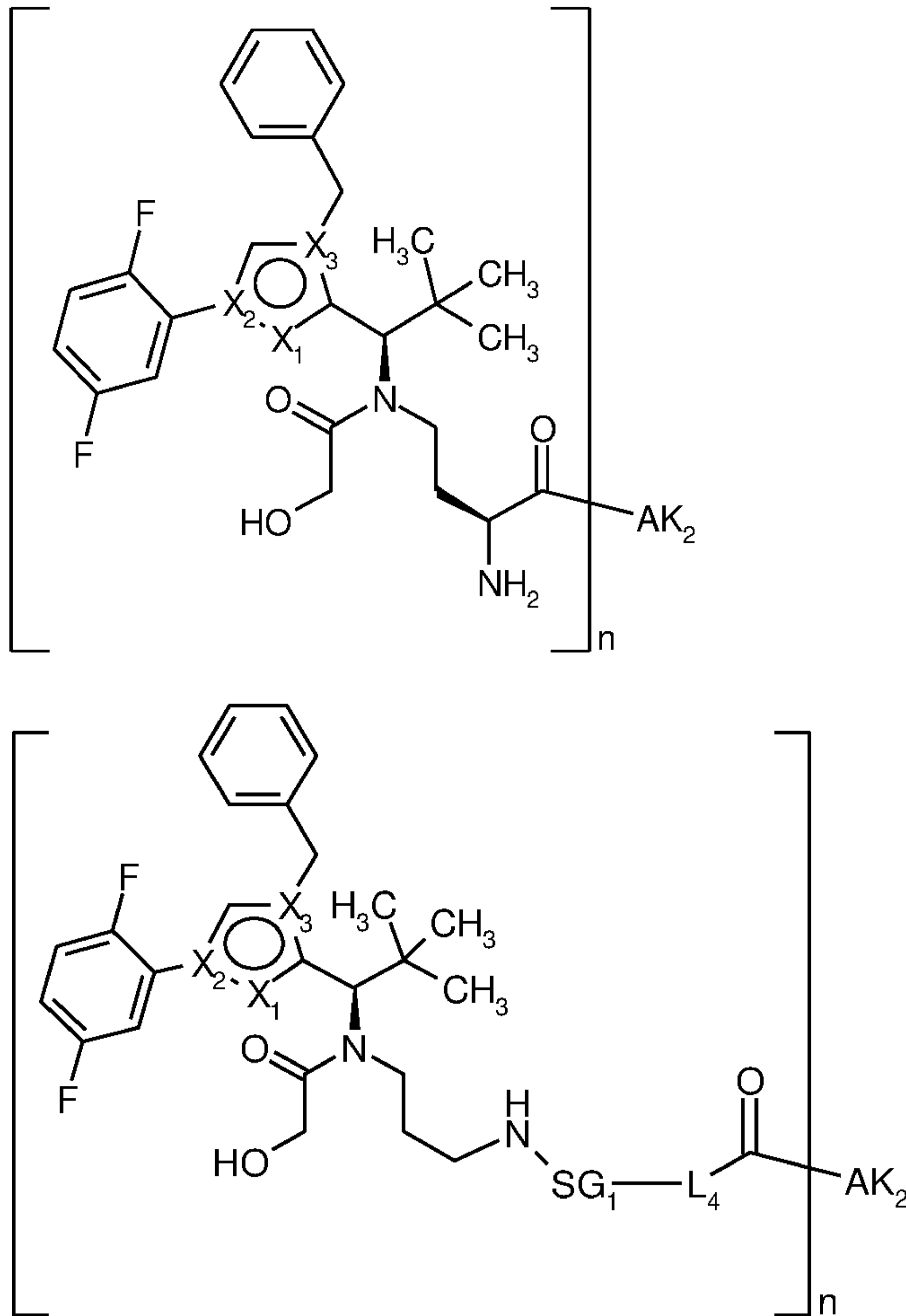
Table B: lysine linker

-§-(SG)<sub>x</sub>-L4-CO-§§

15

Ex.	Subst.	(SG) <sub>x</sub> -L4
1A	R <sup>1</sup>	single bond
58/ 194	R <sup>4</sup>	

Examples of conjugates having corresponding linkers have the following structures, where X1, X2, X3 and L4 have the meaning given above, AK2 represents an antibody attached via a lysine residue and n is a number from 1 to 10. With particular preference, AK2 is a human, humanized or chimeric monoclonal antibody or an antigen-binding fragment thereof, in particular an anti-TWEAKR antibody or an antigen-binding fragment thereof or an anti-EGFR antibody or an antigen-binding fragment thereof. Particular preference is given to an anti-TWEAKR antibody which binds specifically to amino acid D in position 47 (D47) of TWEAKR (SEQ ID NO:169), in particular the anti-TWEAKR antibody TPP-2090, or the anti-EGFR antibodies cetuximab or nimotuzumab.



Preference according to the invention is furthermore given to the basic structure (i), (ii) or (iv), where SG<sub>1</sub> or SG represents a group which can be cleaved by cathepsin and L<sub>1</sub> and L<sub>2</sub> have the meanings given above. Particular preference is given to the following groups:

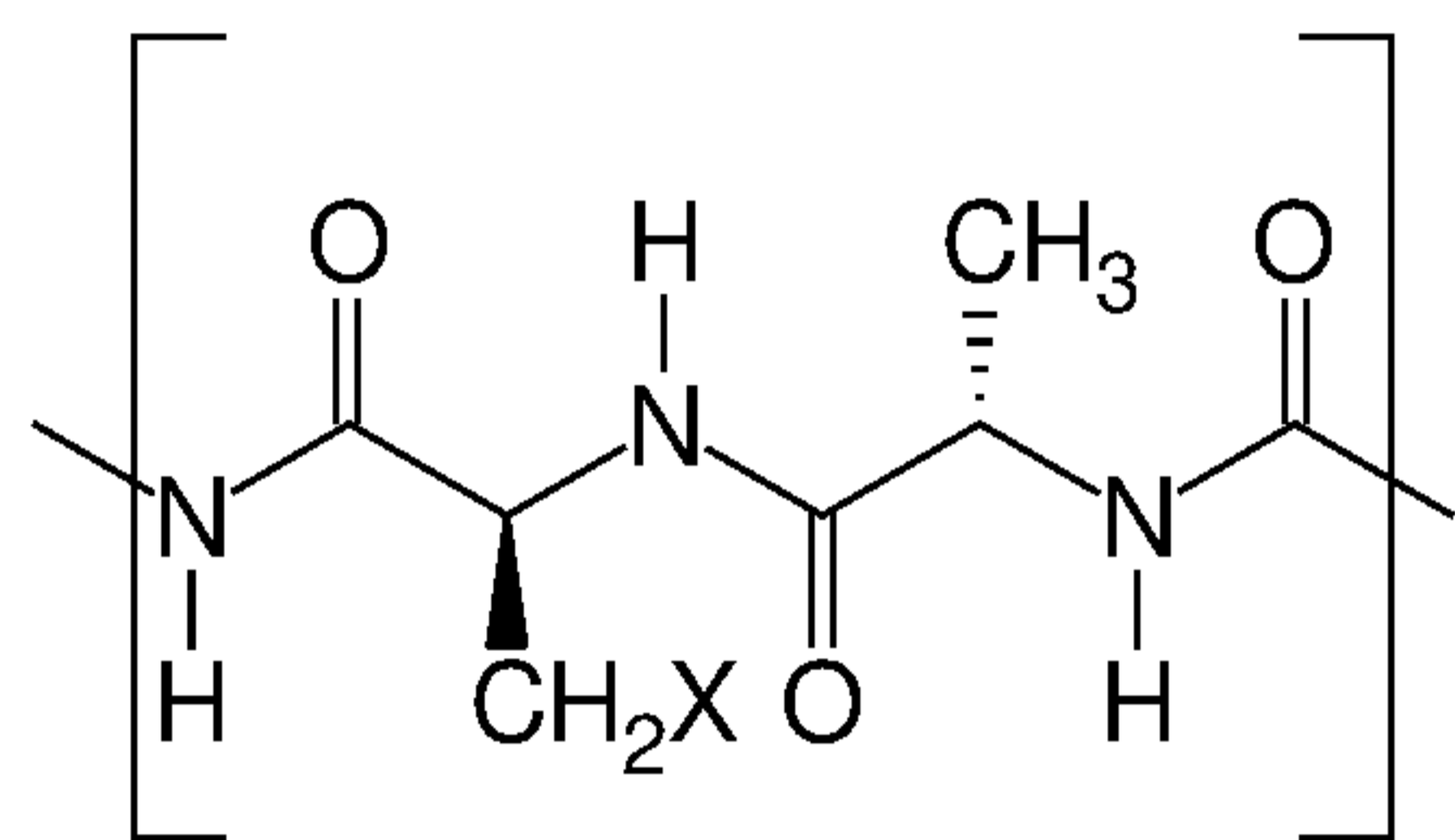
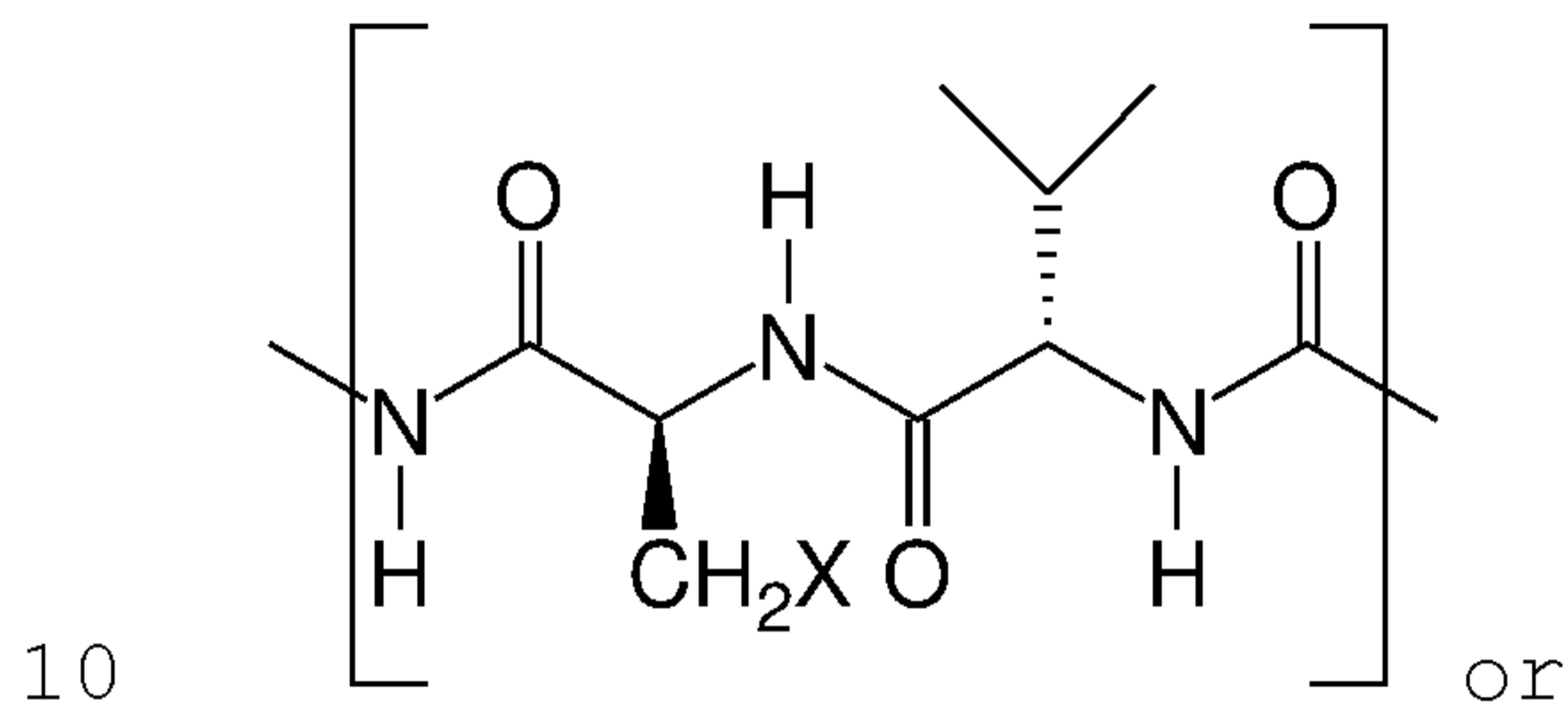
- -Val-Ala-CONH- (hereby cleavage of the amide bond at the C-terminal amide of alanine)
- -NH-Val-Lys-CONH- (cleavage of the amide bond at the C-terminal amide of lysine)
- -NH-Val-Cit-CONH- (cleavage of the amide bond at the C-terminal amide of citrulline)
- -NH-Phe-Lys-CONH- (cleavage of the amide bond at the C-terminal amide of lysine)



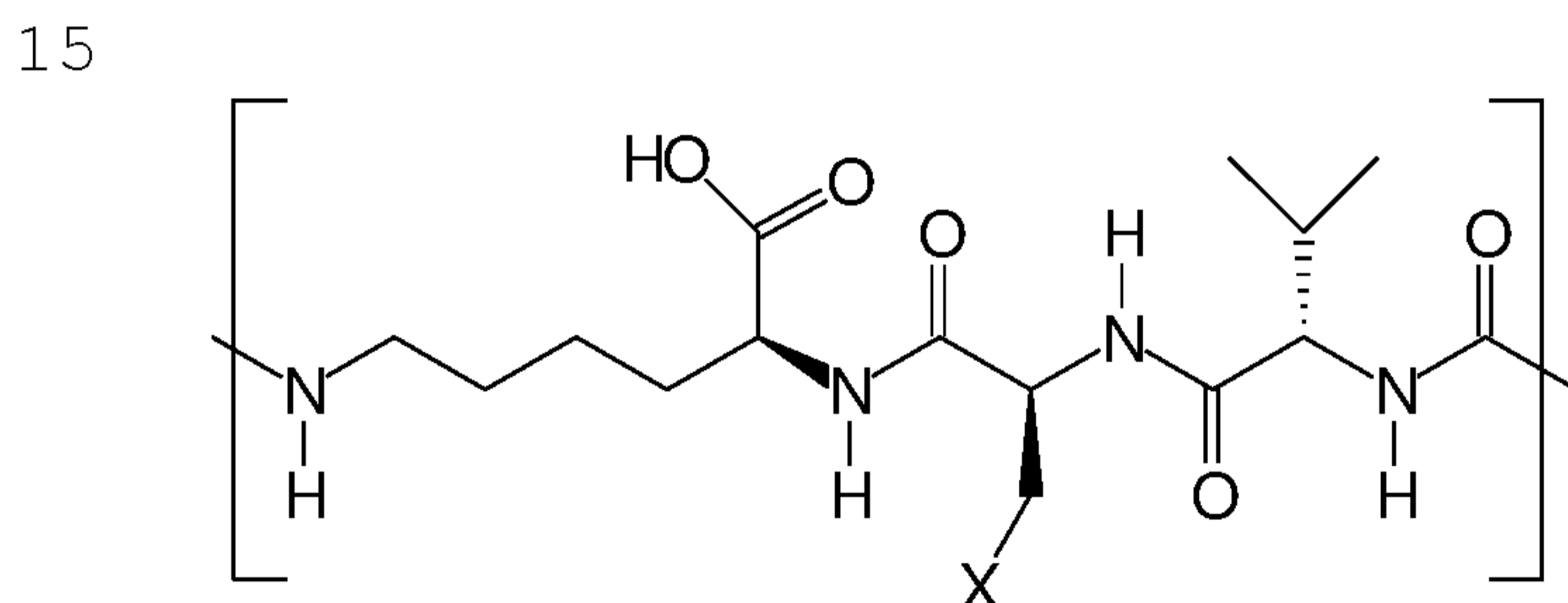
• -NH-Ala-Lys-CONH- (cleavage of the amide bond at the C-terminal amide of lysine)

5 • -NH-Ala-Cit-CONH- (cleavage of the amide bond at the C-terminal amide of citrulline)

SG1 or SG is particularly preferably



or



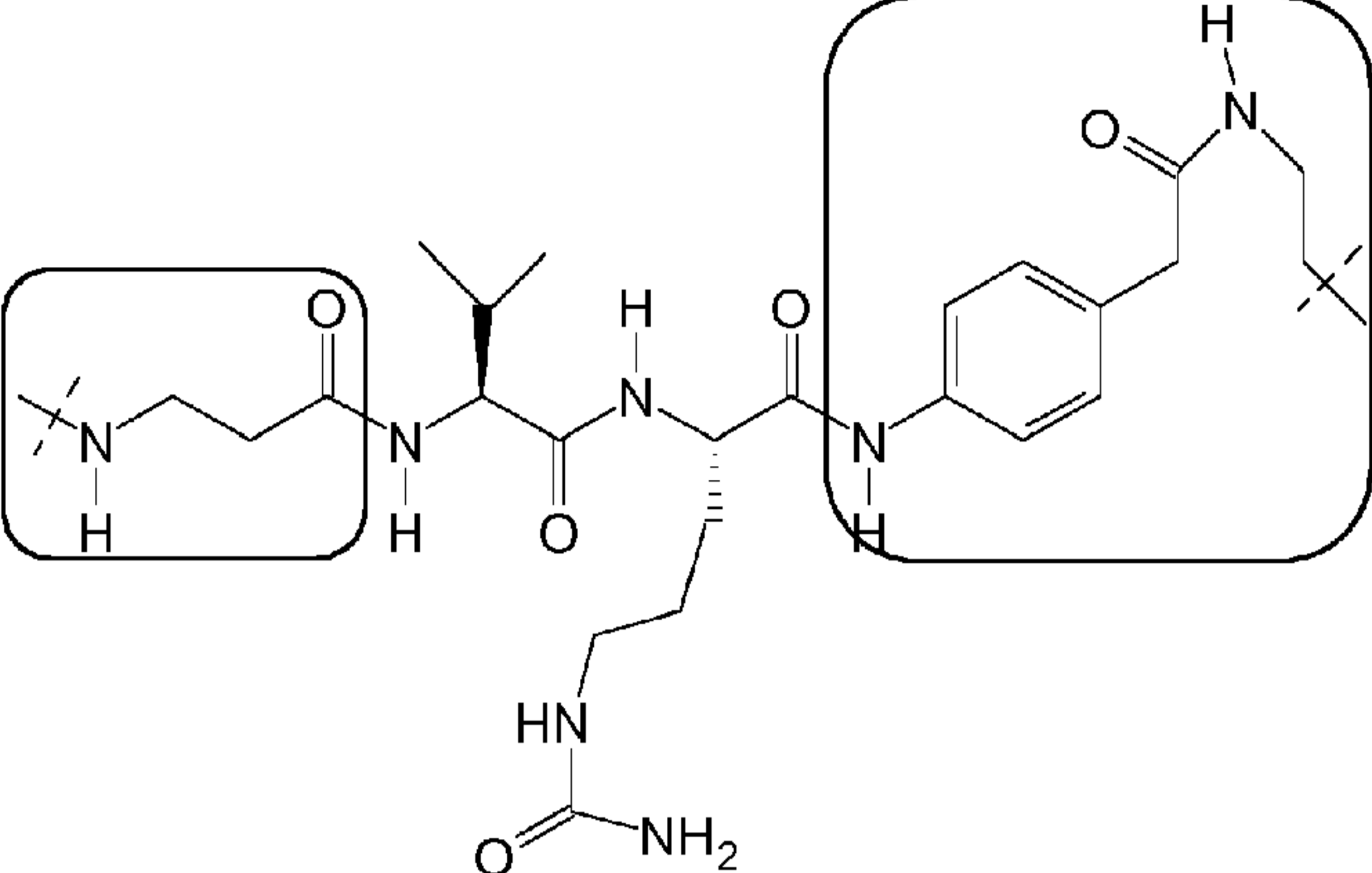
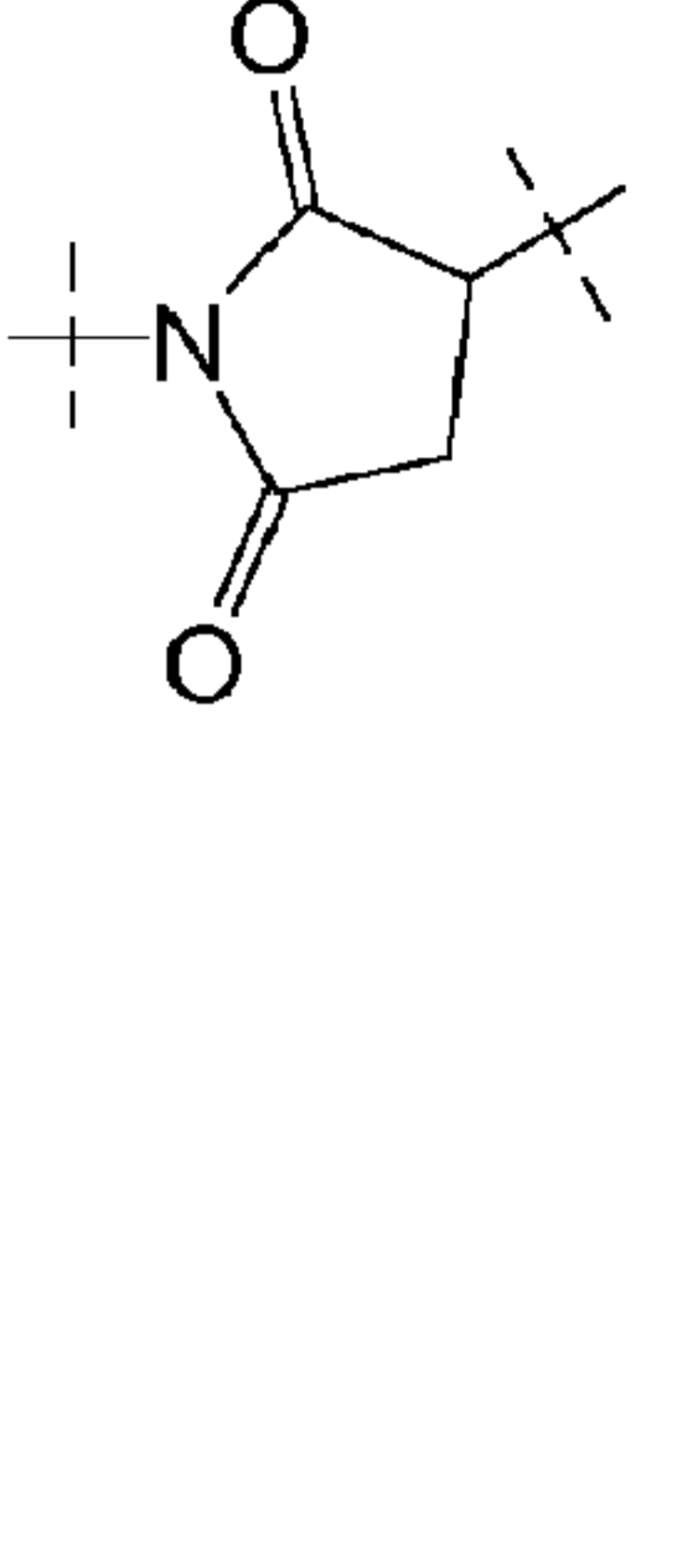
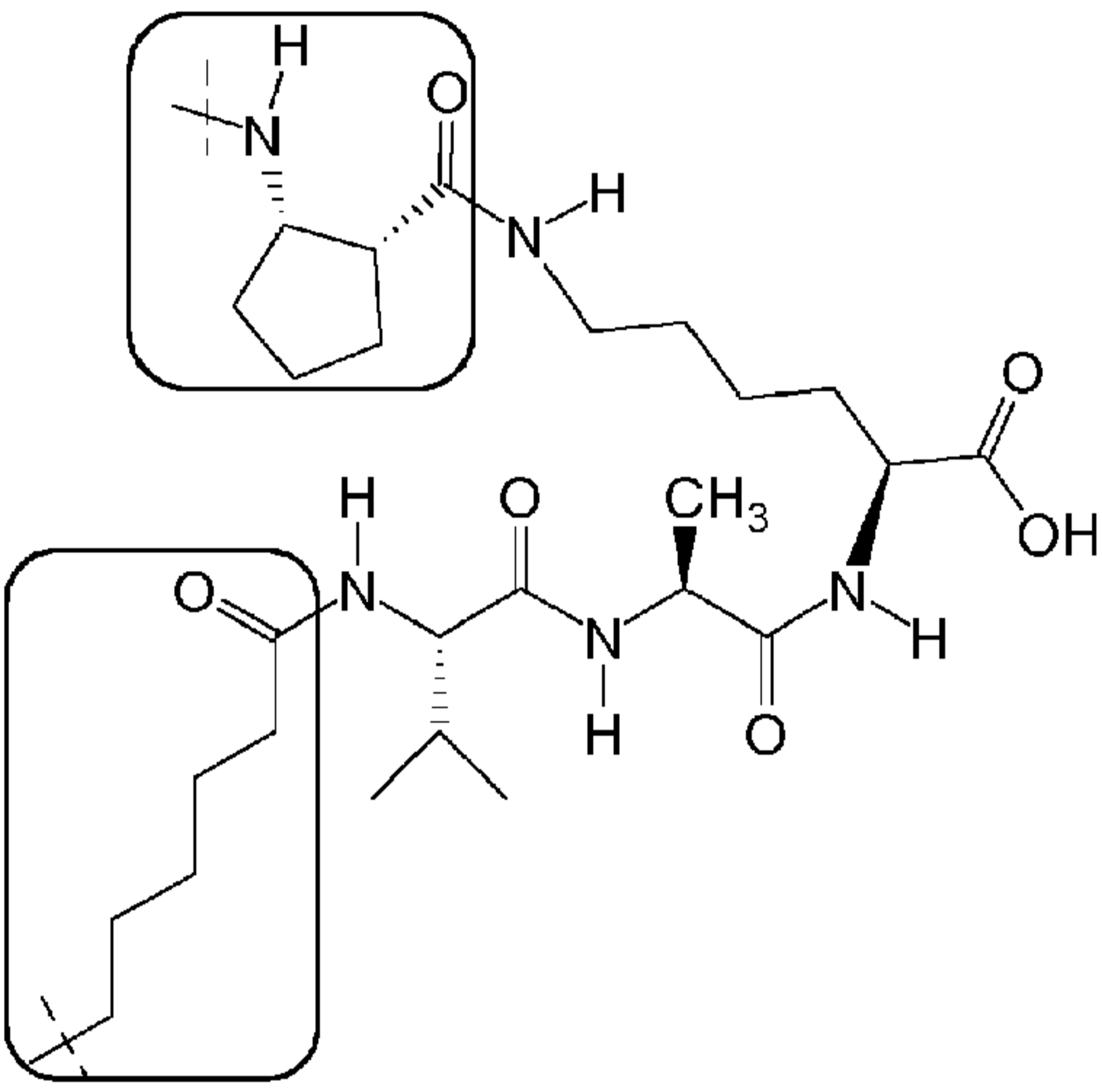
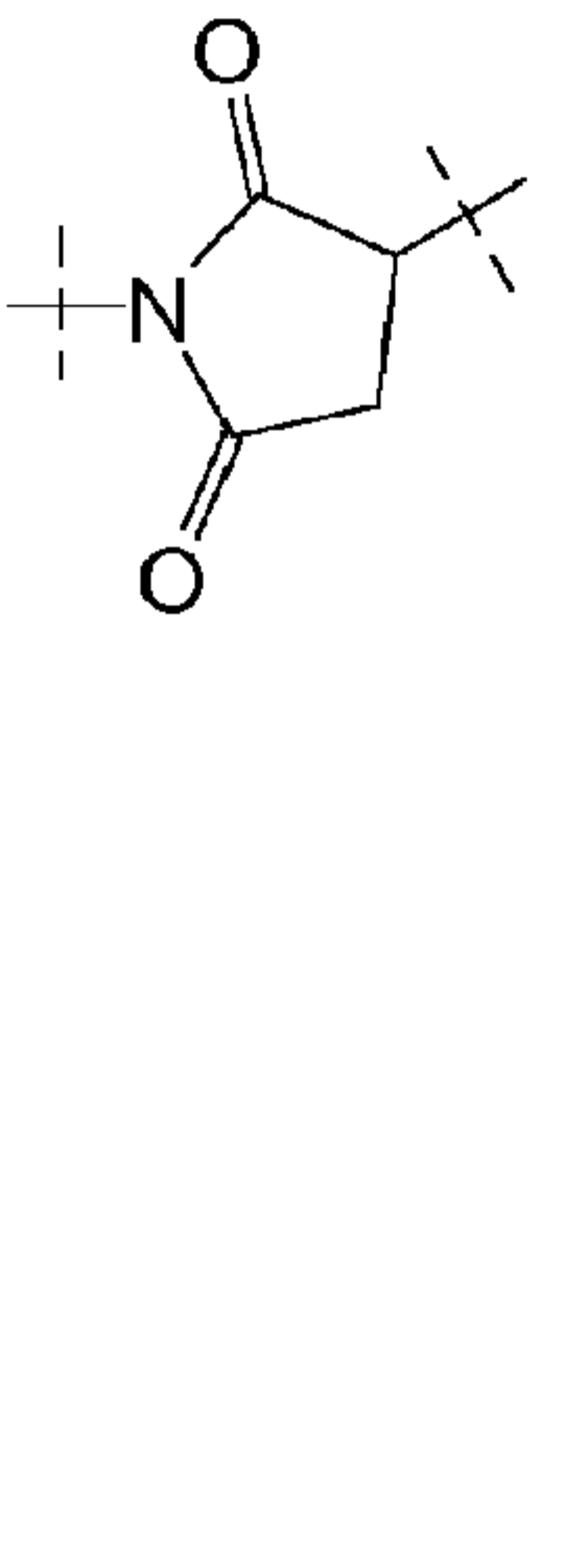
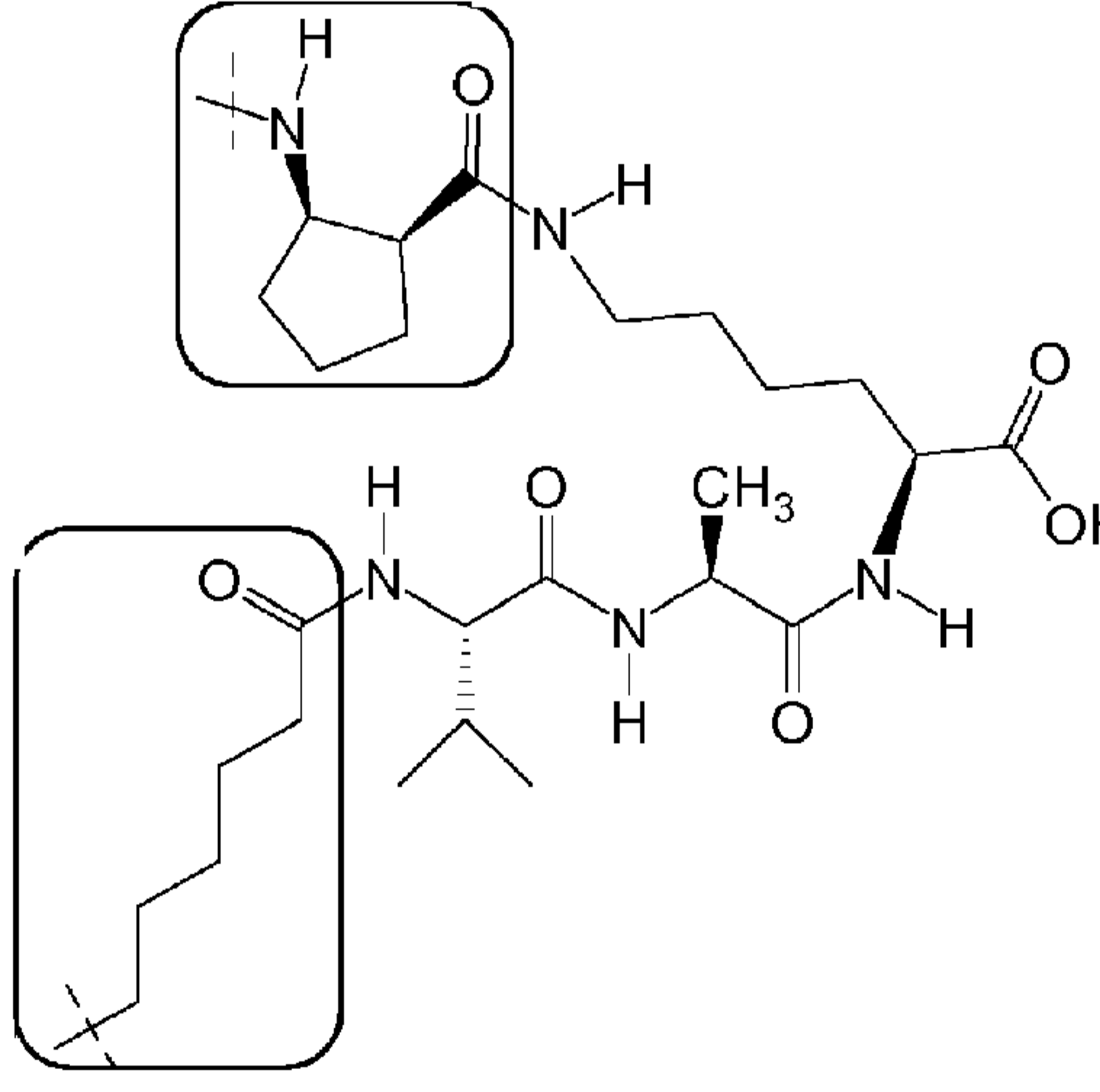
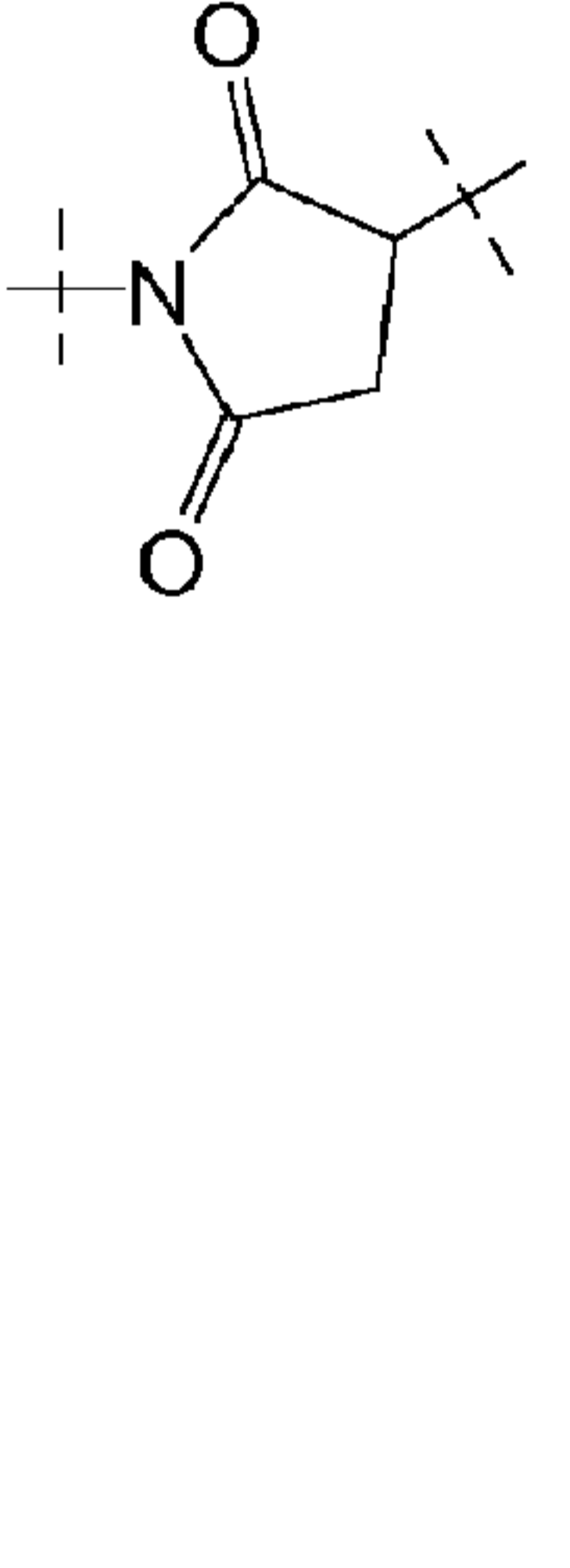
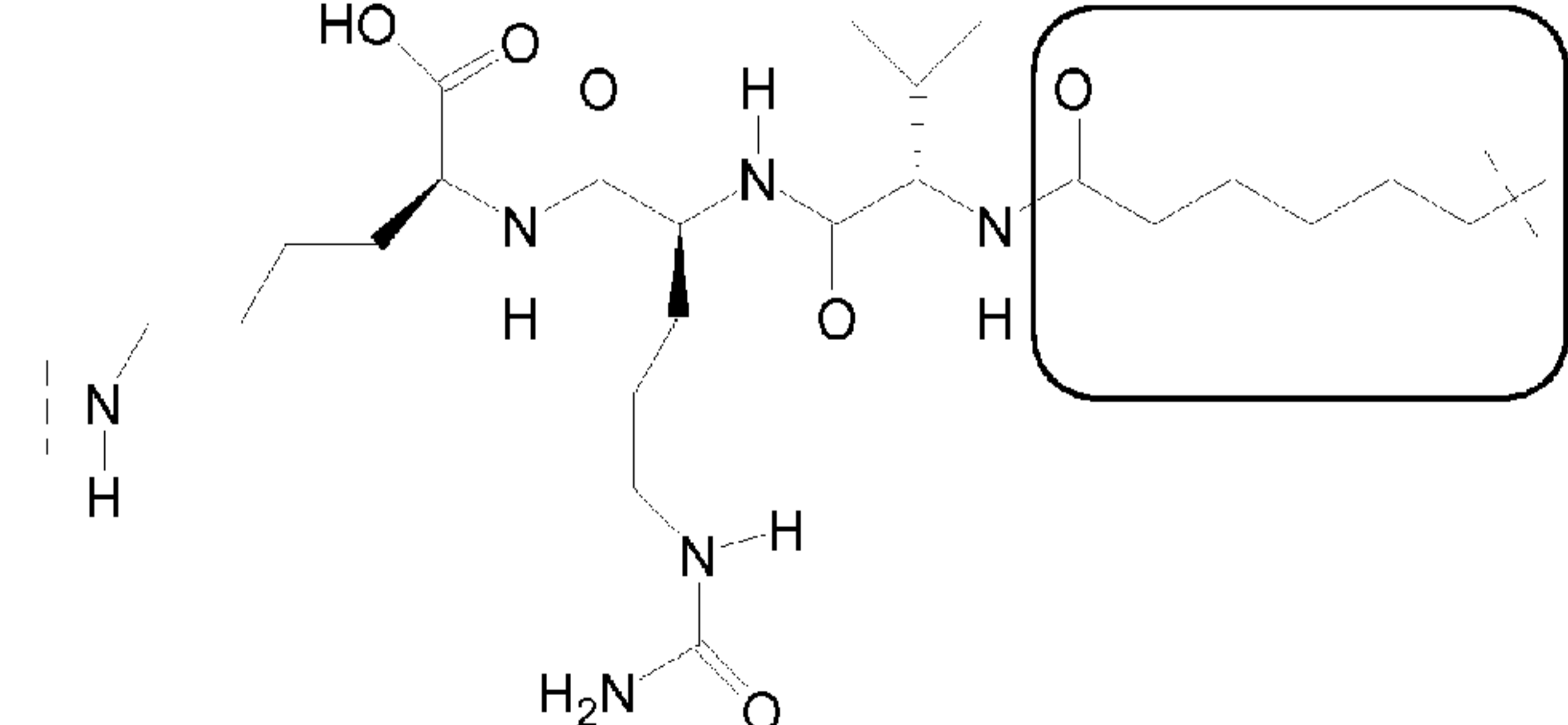
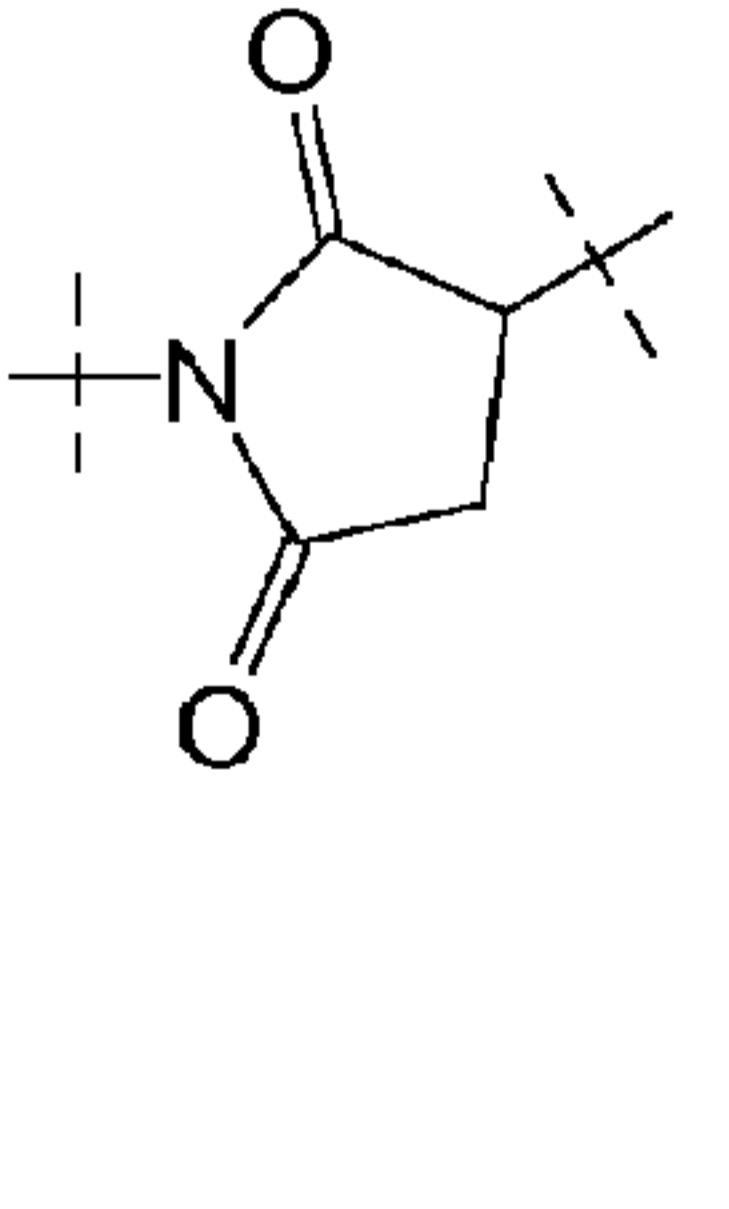
where X represents H or a C<sub>1-10</sub>-alkyl group which may optionally be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, NH<sub>2</sub>, -NH-CNNH<sub>2</sub> or sulphonic acid.

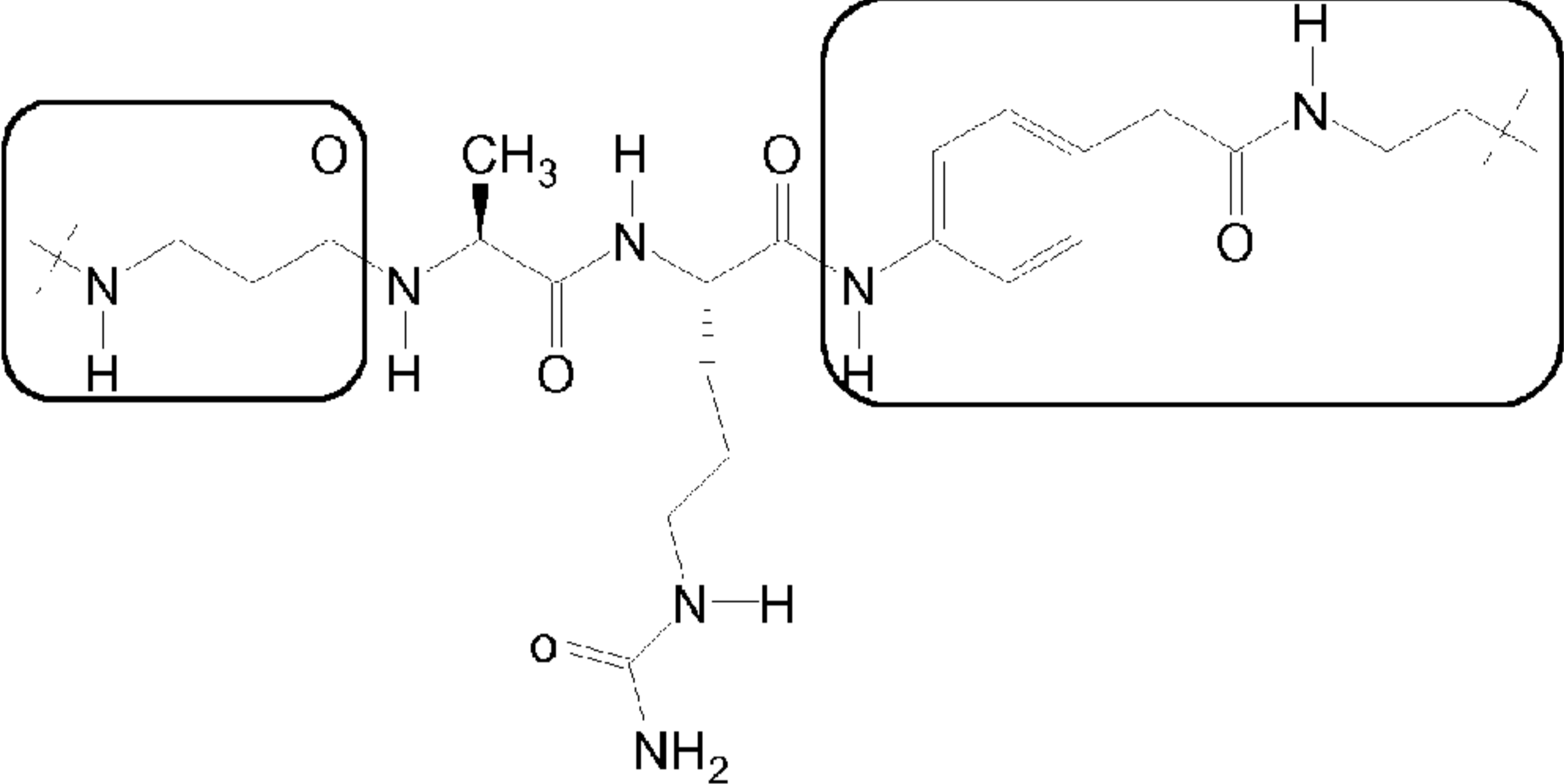
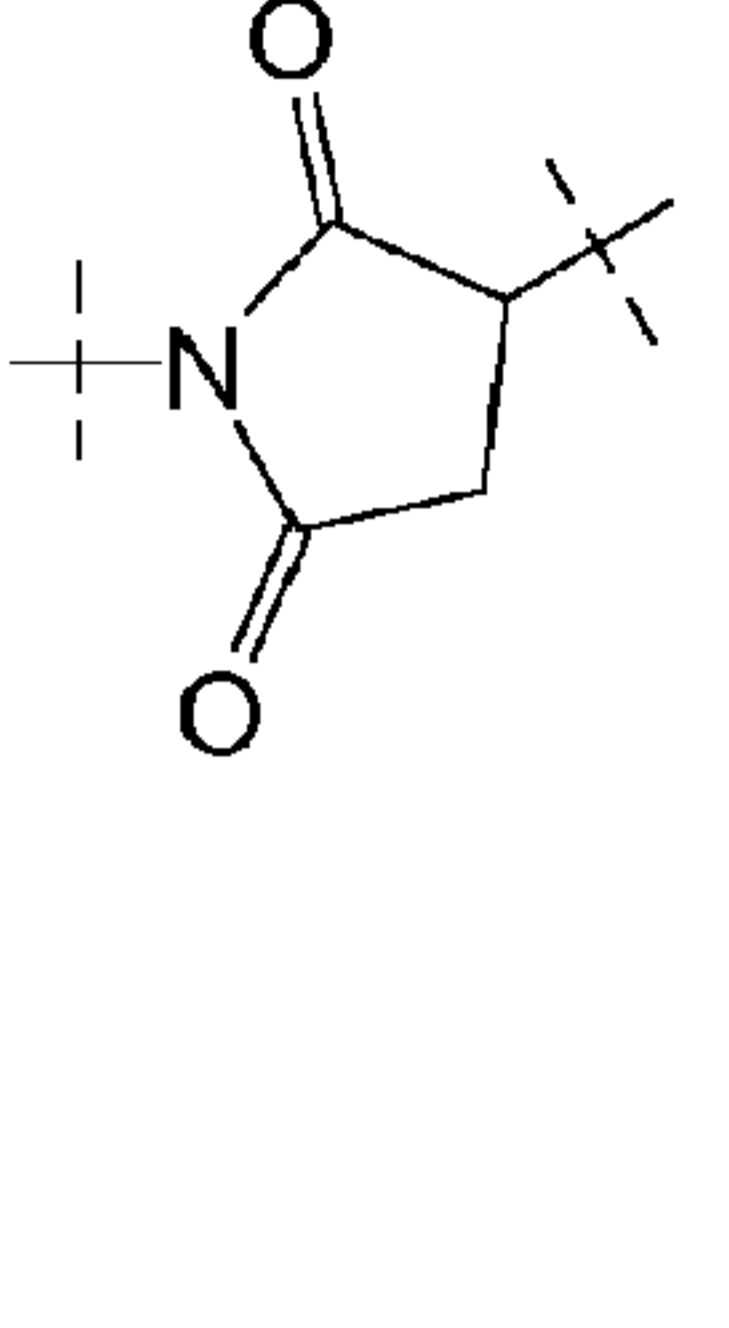
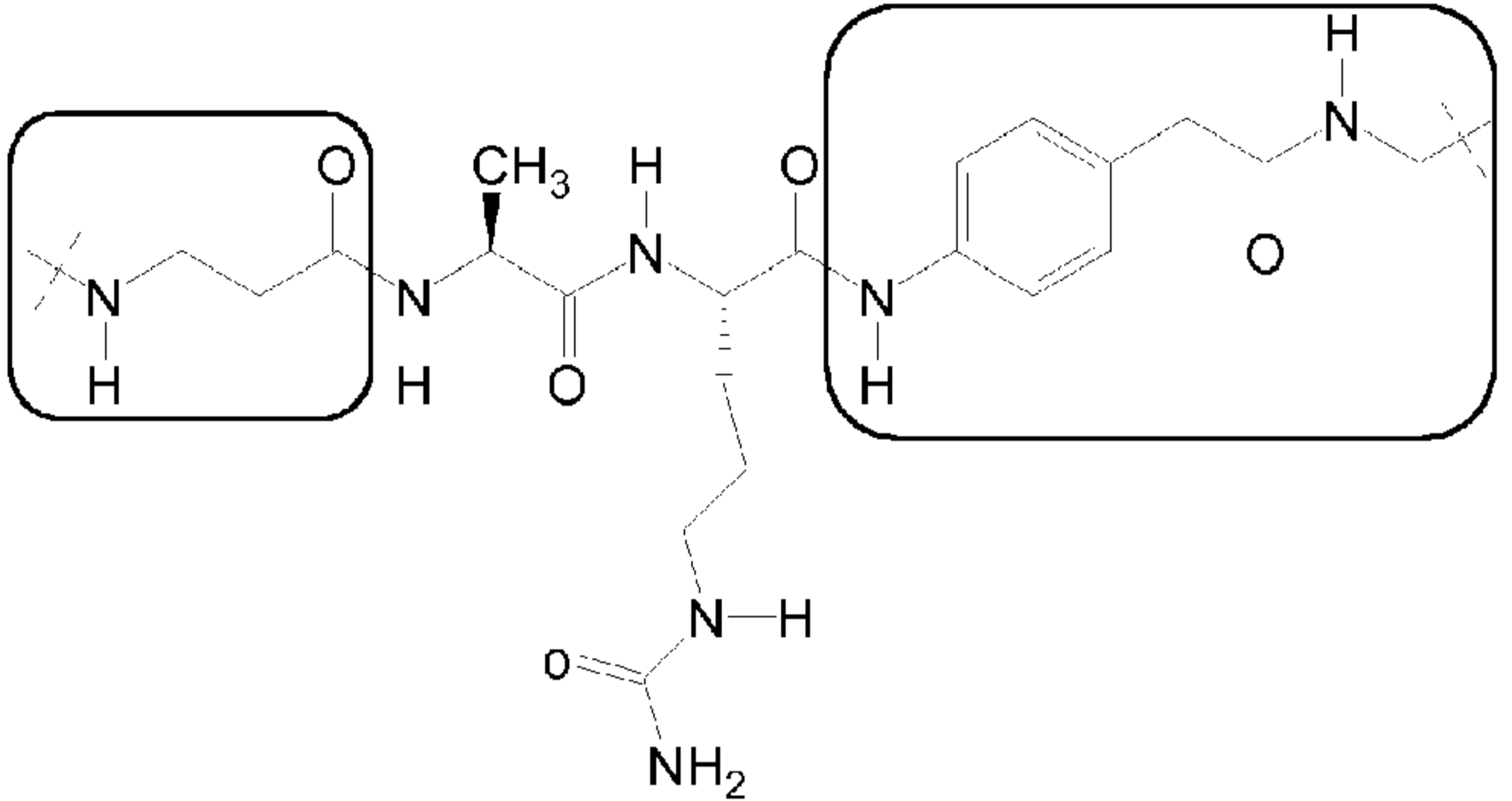
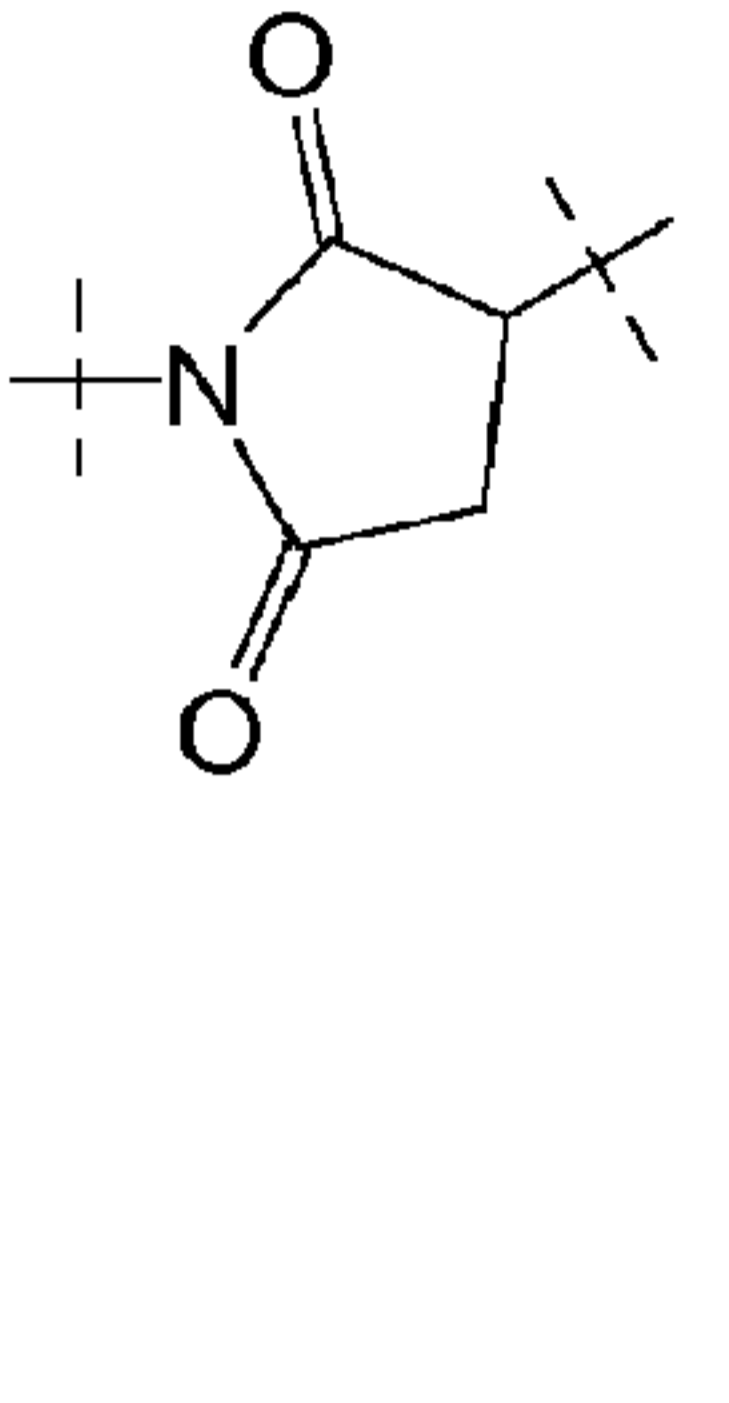
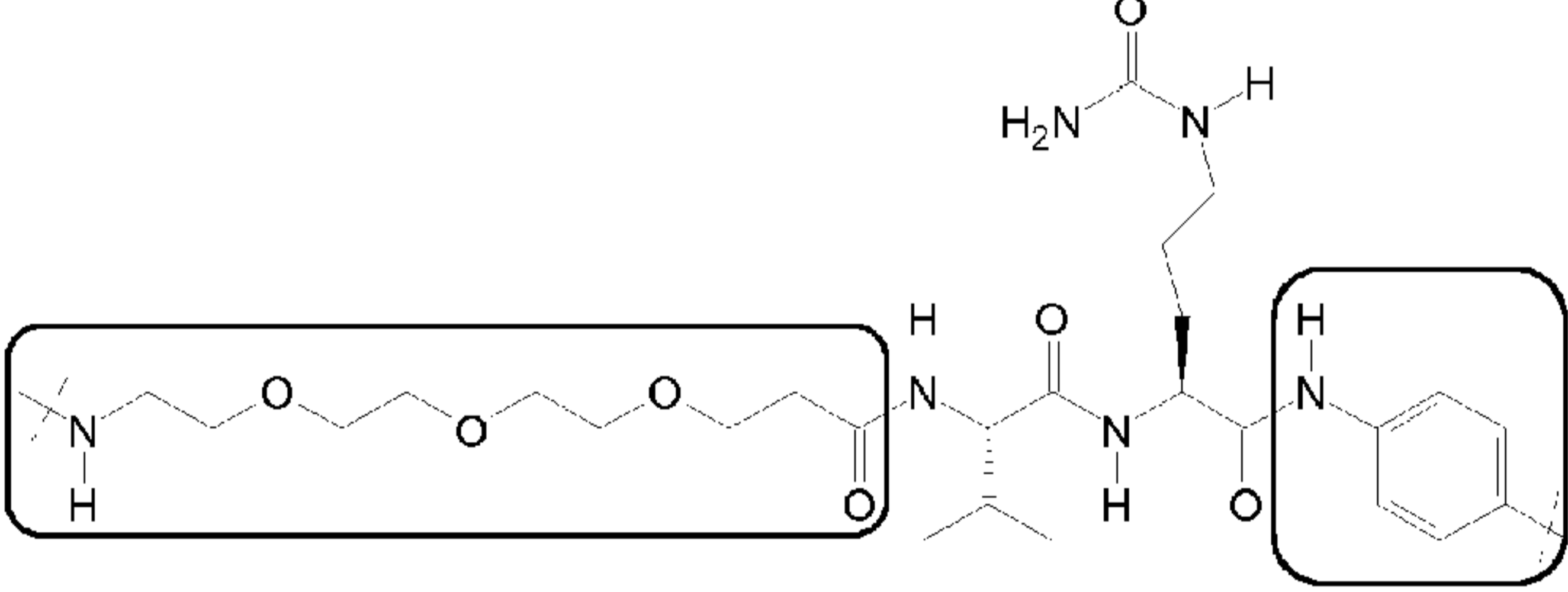
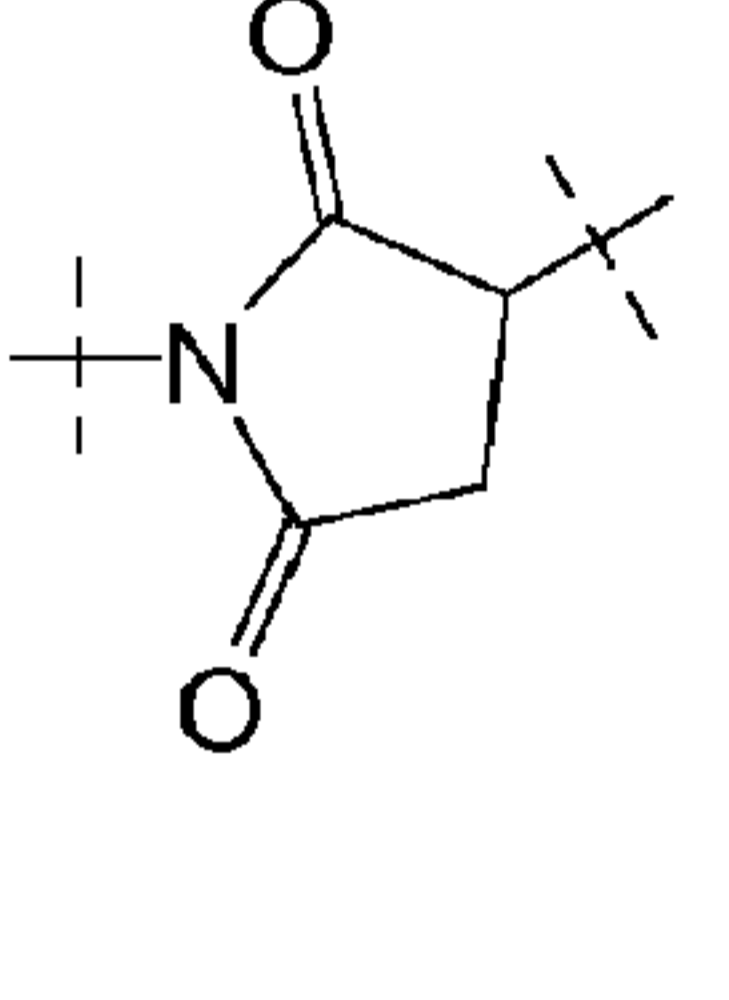
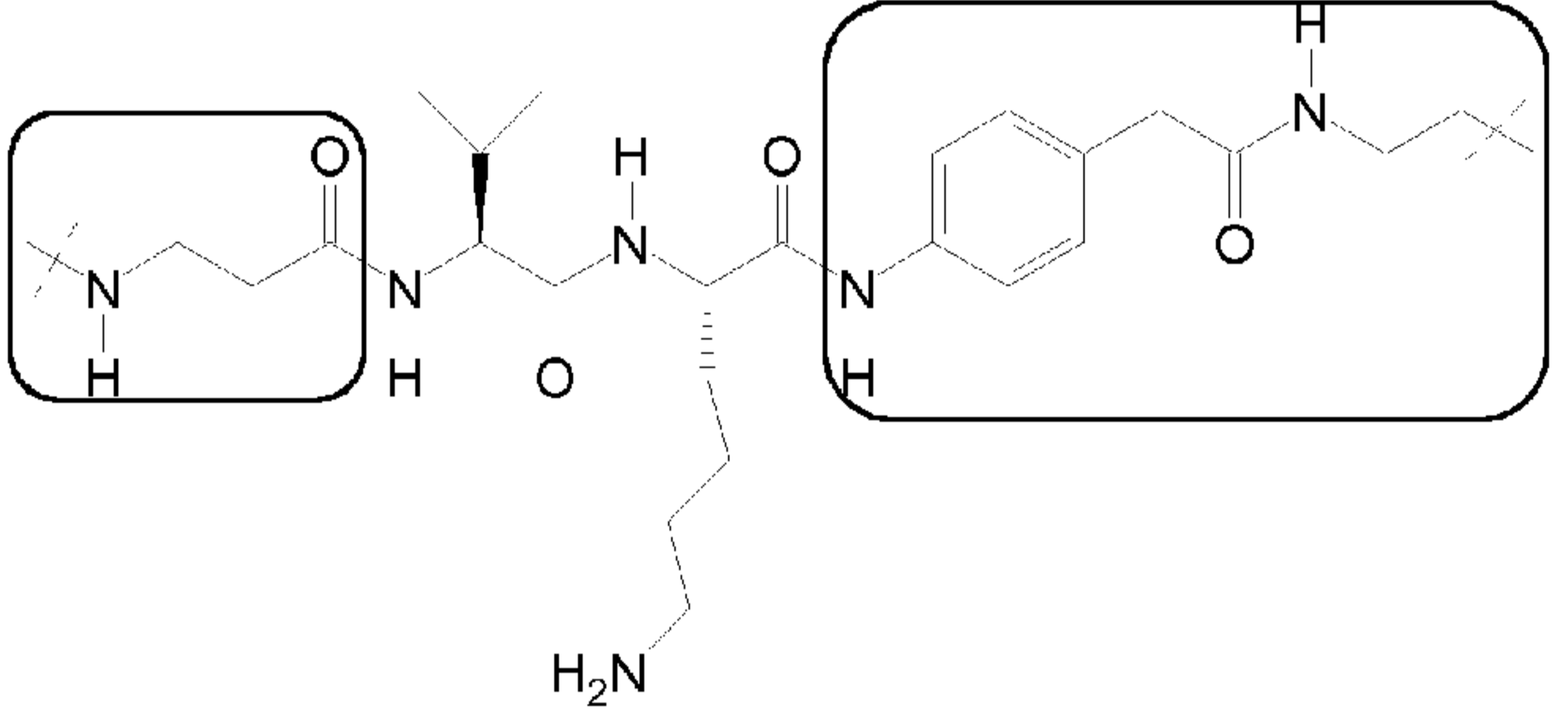
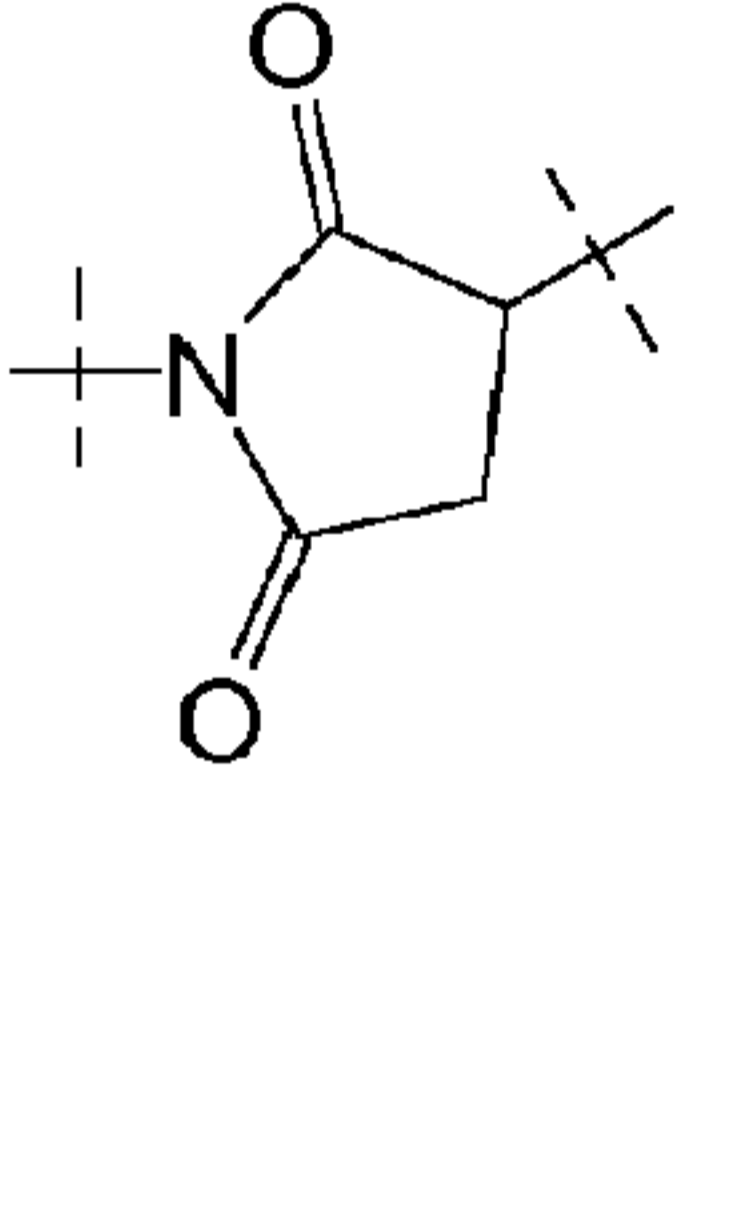
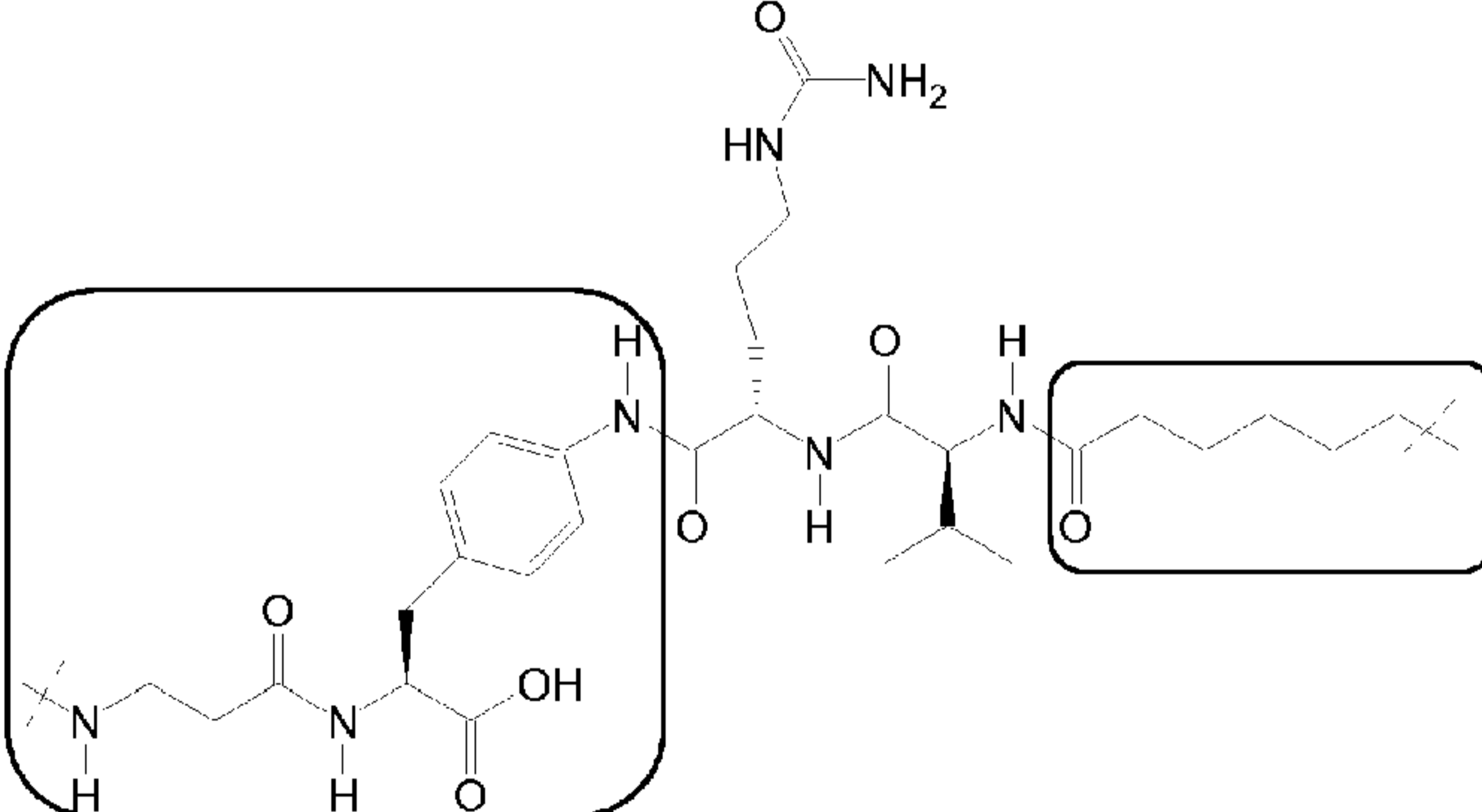
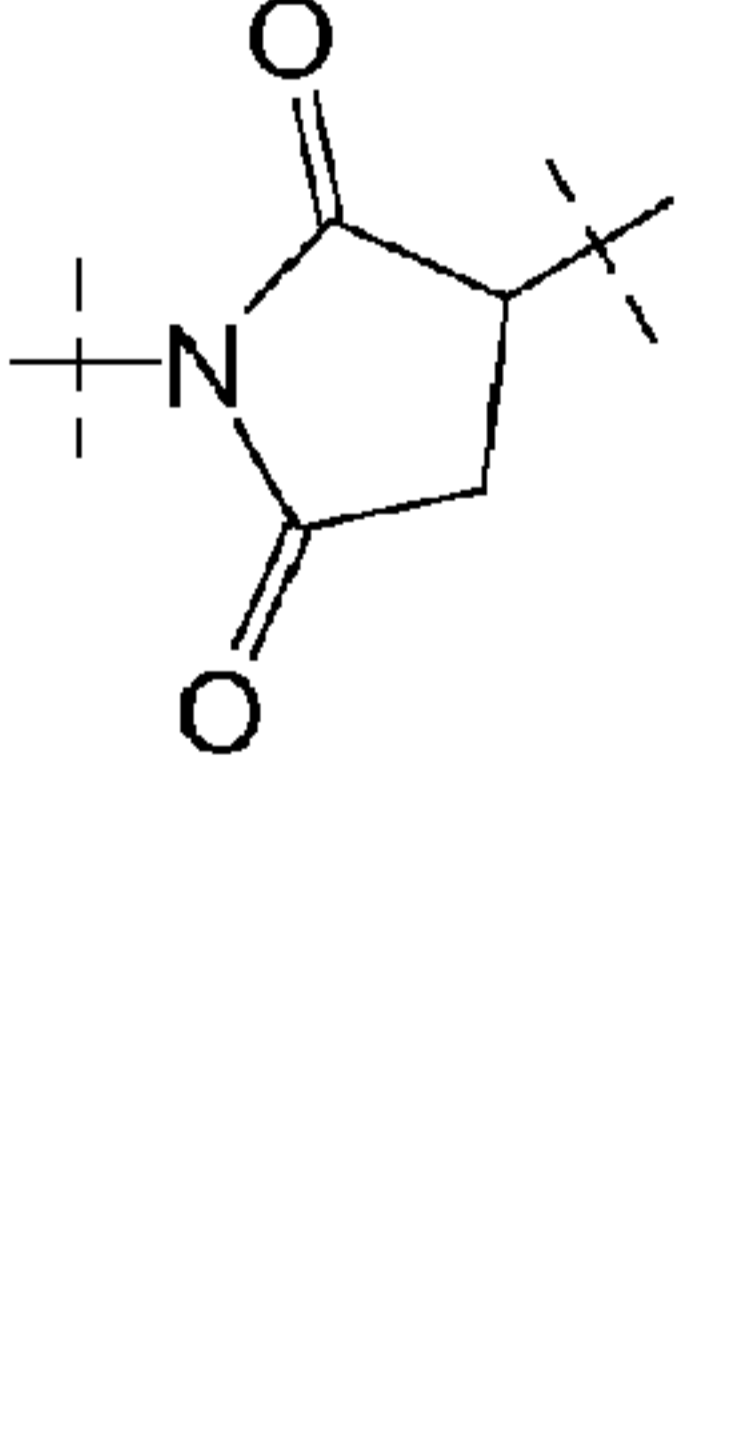
20

Table C below gives examples of a linker moiety -SG1-L1- or -L1-SG-L1-, where SG1 and SG are groups which can be cleaved by cathepsin. Table C furthermore states with which group L2 these examples of -SG1-L1- and -L1-SG-L1- are preferably combined, and also the preferred coupling point (R<sup>1</sup>-R<sup>5</sup>) and the preferred

25



16A	R <sup>1</sup>	1		
17A/ 168A	R <sup>1</sup>	1		
18A	R <sup>1</sup>	1		
23A/ 164A	R <sup>1</sup>	1		

24A	R <sup>1</sup>	1		
25A	R <sup>1</sup>	1		
26A	R <sup>1</sup>	1		
27A	R <sup>1</sup>	1		
28A	R <sup>1</sup>	1		

29A	R <sup>1</sup>	1		
31A	R <sup>5</sup> (m )	0		
34A	R <sup>1</sup>	1		
35A	R <sup>4</sup>	0		
36A	R <sup>4</sup>	0		
39A/ 215A	R <sup>4</sup>	0		
43A	R <sup>4</sup>	0		

45A	R <sup>4</sup>	0		
46A	R <sup>4</sup>	0		
47A	R <sup>4</sup>	0		
55A	R <sup>3</sup>	0		
55B	R <sup>3</sup>	0		
110A /156 A	R <sup>1</sup>	1		
128A	R <sup>1</sup>	1		

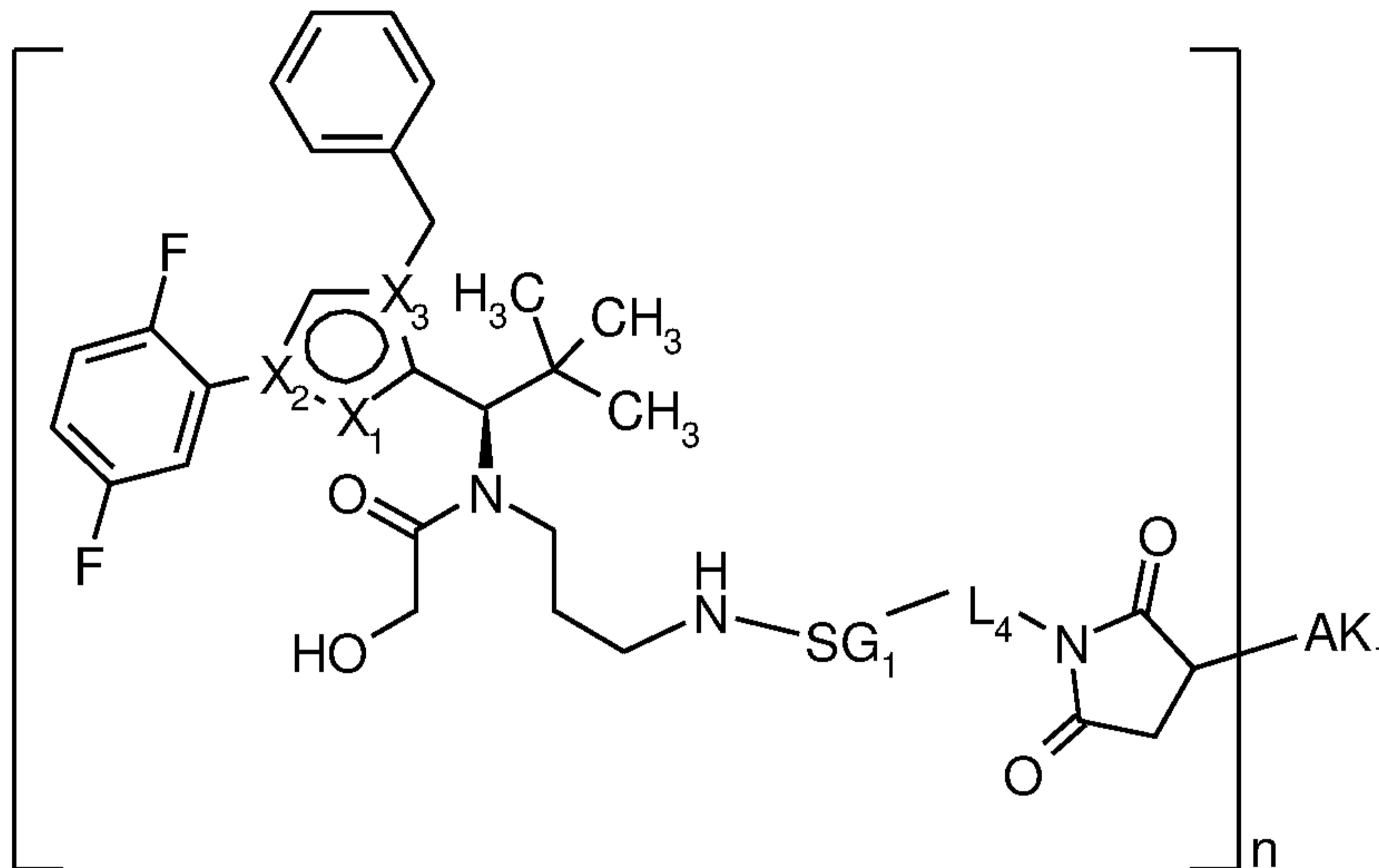
129A /167 A	R <sup>1</sup>	1		
150A	R <sup>3</sup>	0		
154A /155 A/16 0	R <sup>1</sup>	1		
165A	R <sup>1</sup>	1		
166A	R <sup>1</sup>	1		
169A	R <sup>1</sup>	1		
170A	R <sup>1</sup>	1		

171A	R <sup>1</sup>	1		
172A /173 A/17 4A	R <sup>1</sup>	1		
205A	R <sup>1</sup>	1		
206A	R <sup>1</sup>	1		
207A	R <sup>1</sup>	1		
235A	R <sup>3</sup>	0		

Examples of conjugates having basic structure (i) have the



following structure, where X1, X2 and X3 have the meanings given above, L4 has the same meaning as L1, AK1 represents an antibody attached via a cysteine residue and n is a number from 1 to 10. Particularly preferably, AK1 is an anti-TWEAKR antibody, in particular an anti-TWEAKR antibody which binds specifically to amino acid D in position 47 (D47) of TWEAKR (SEQ ID NO:169), in particular the anti-TWEAKR antibody TPP-2090.

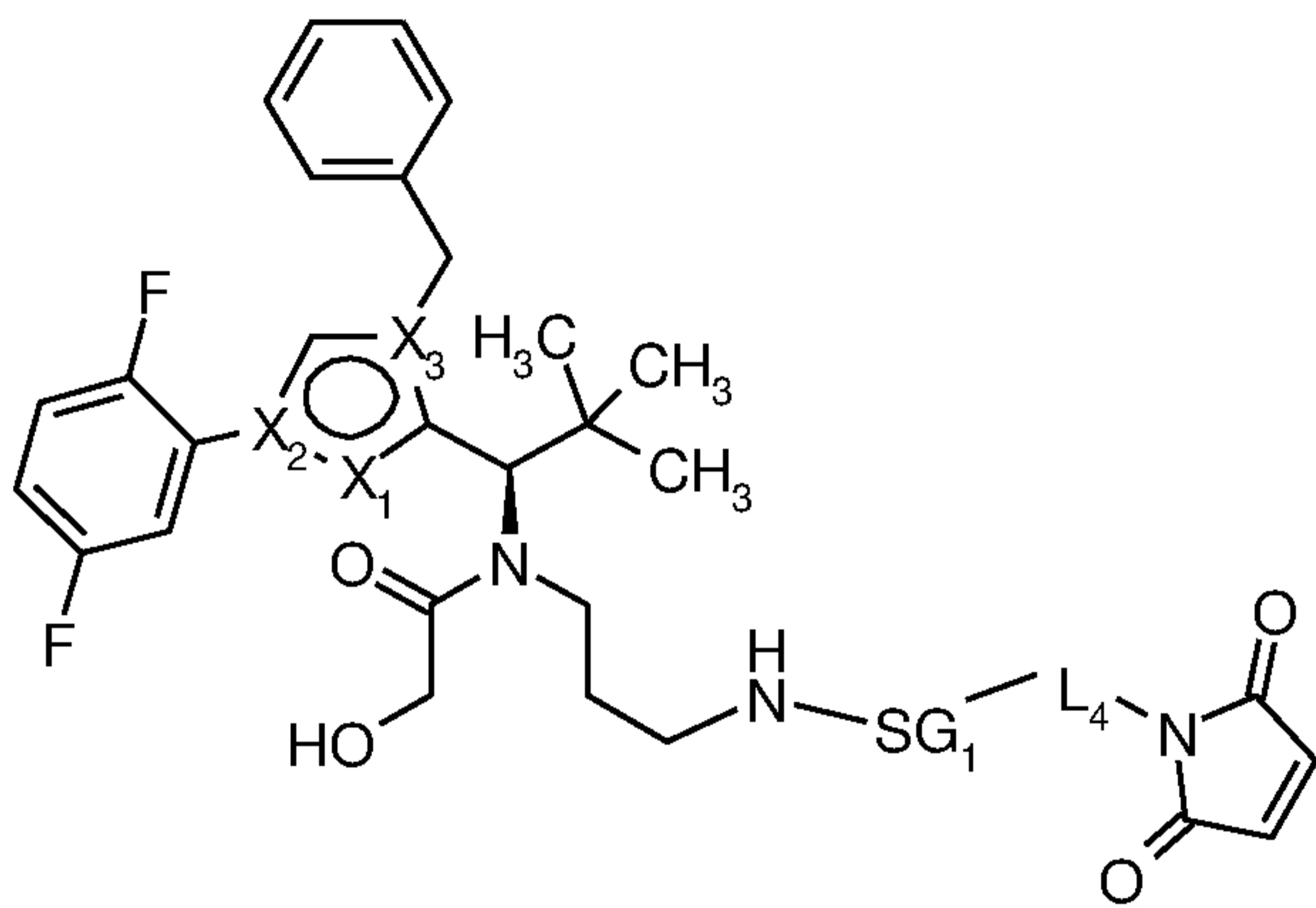
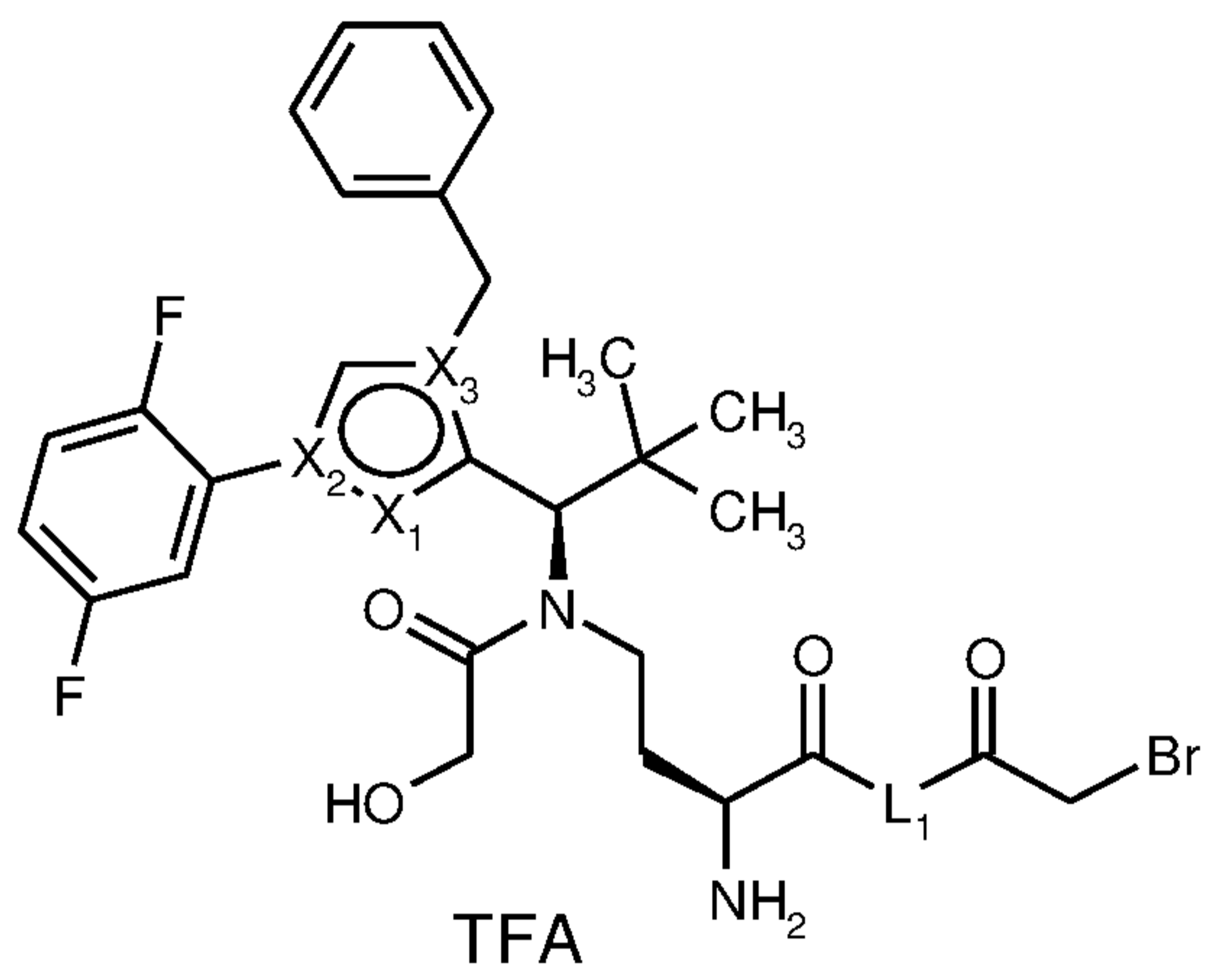
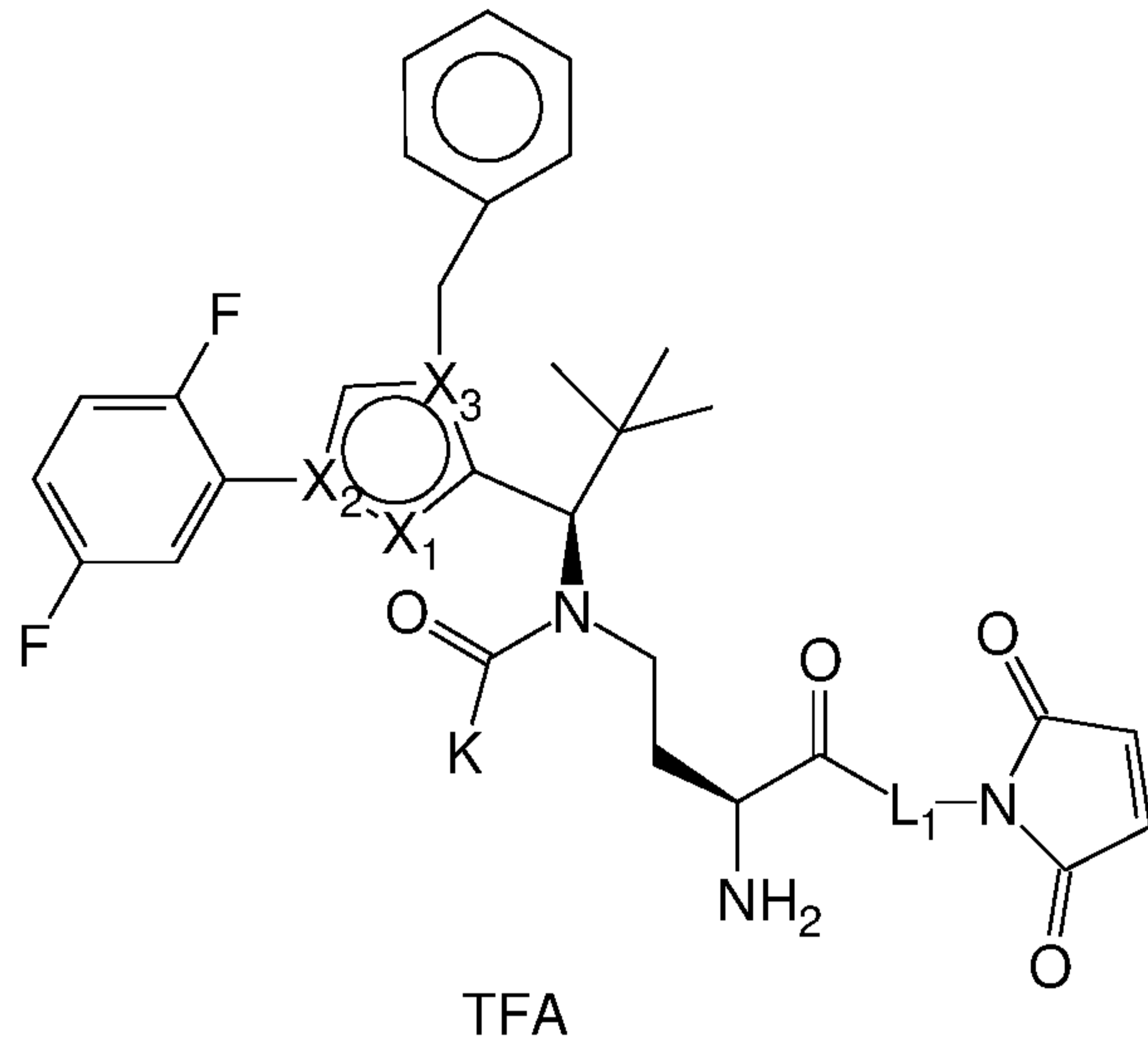


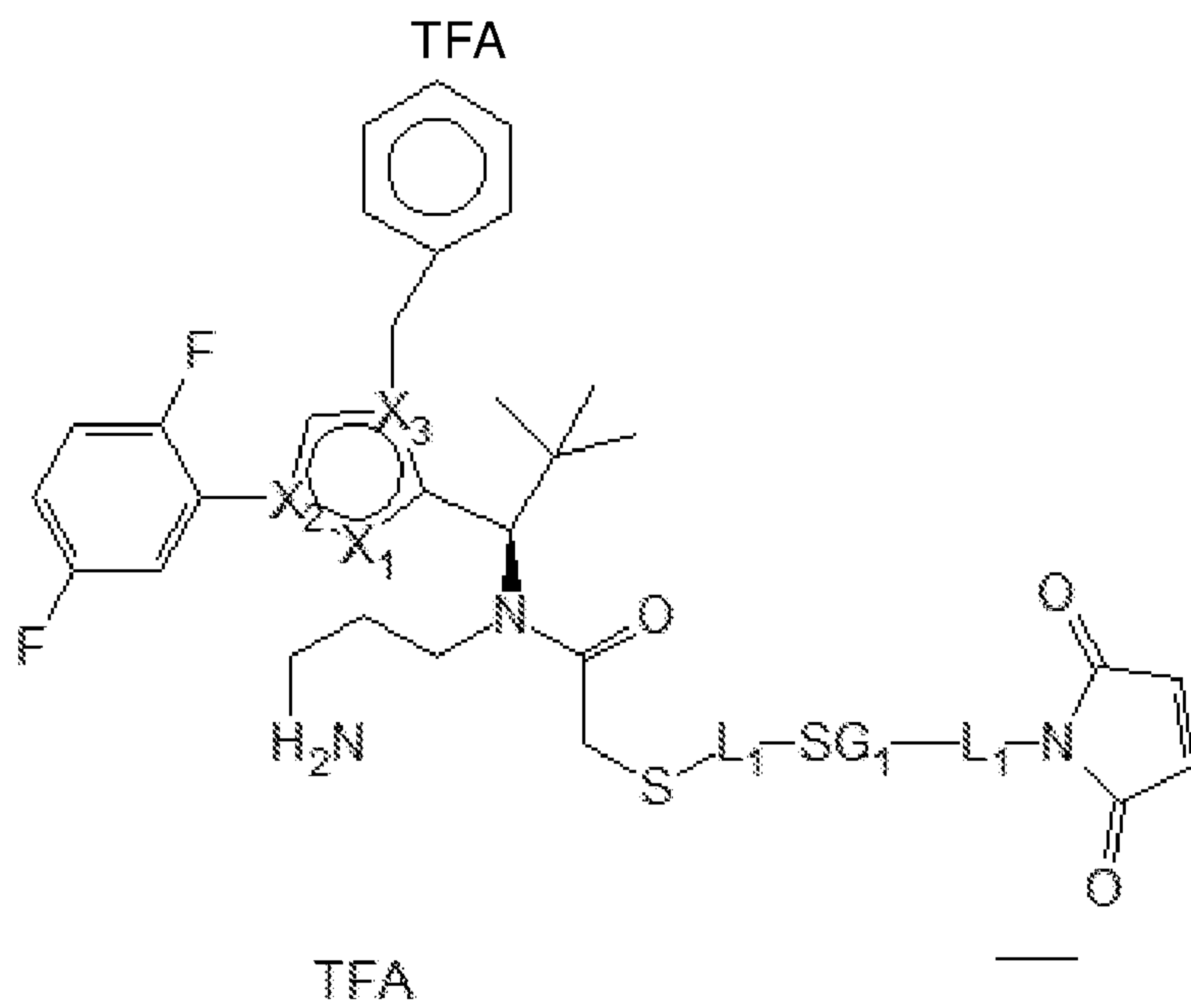
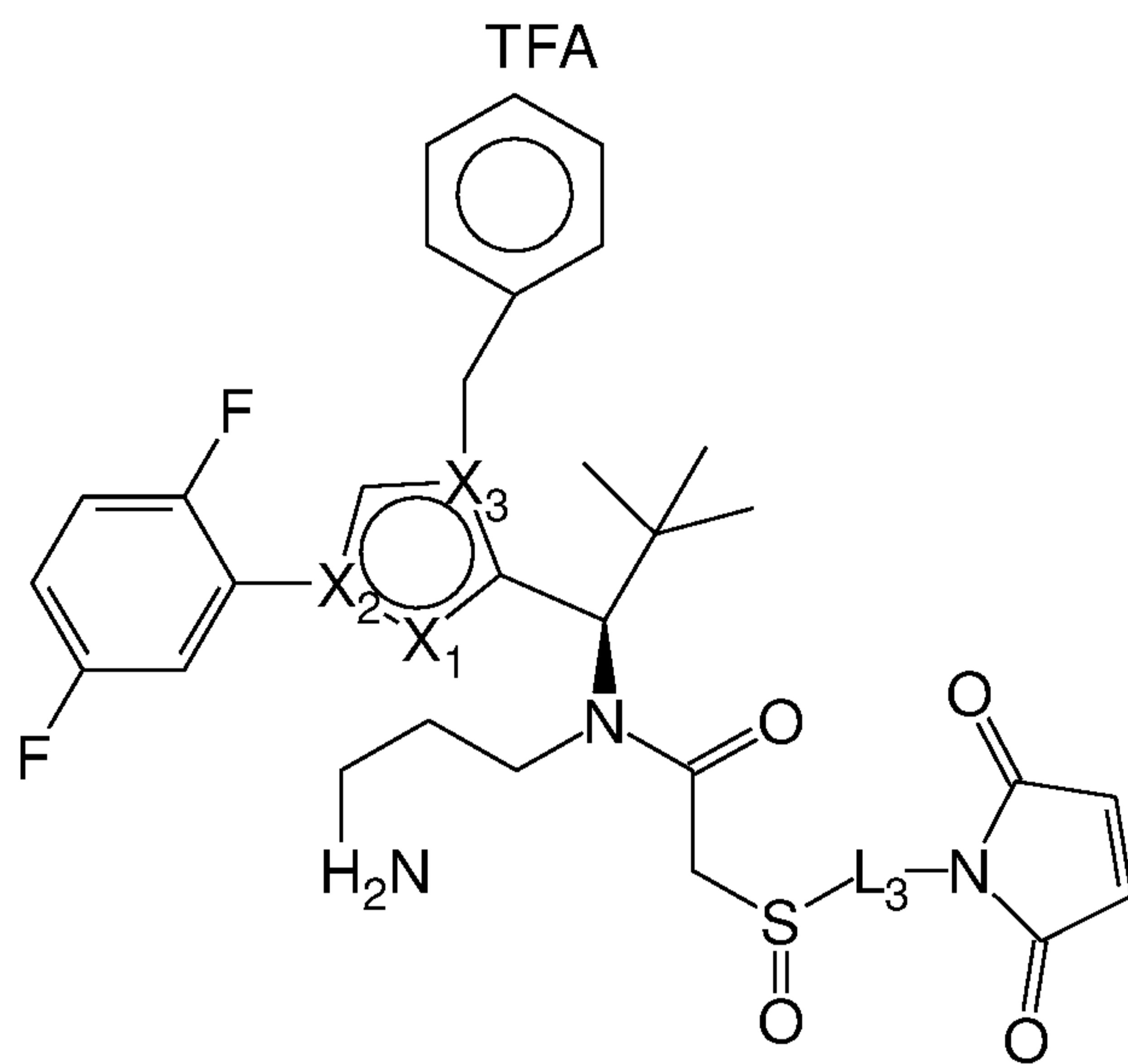
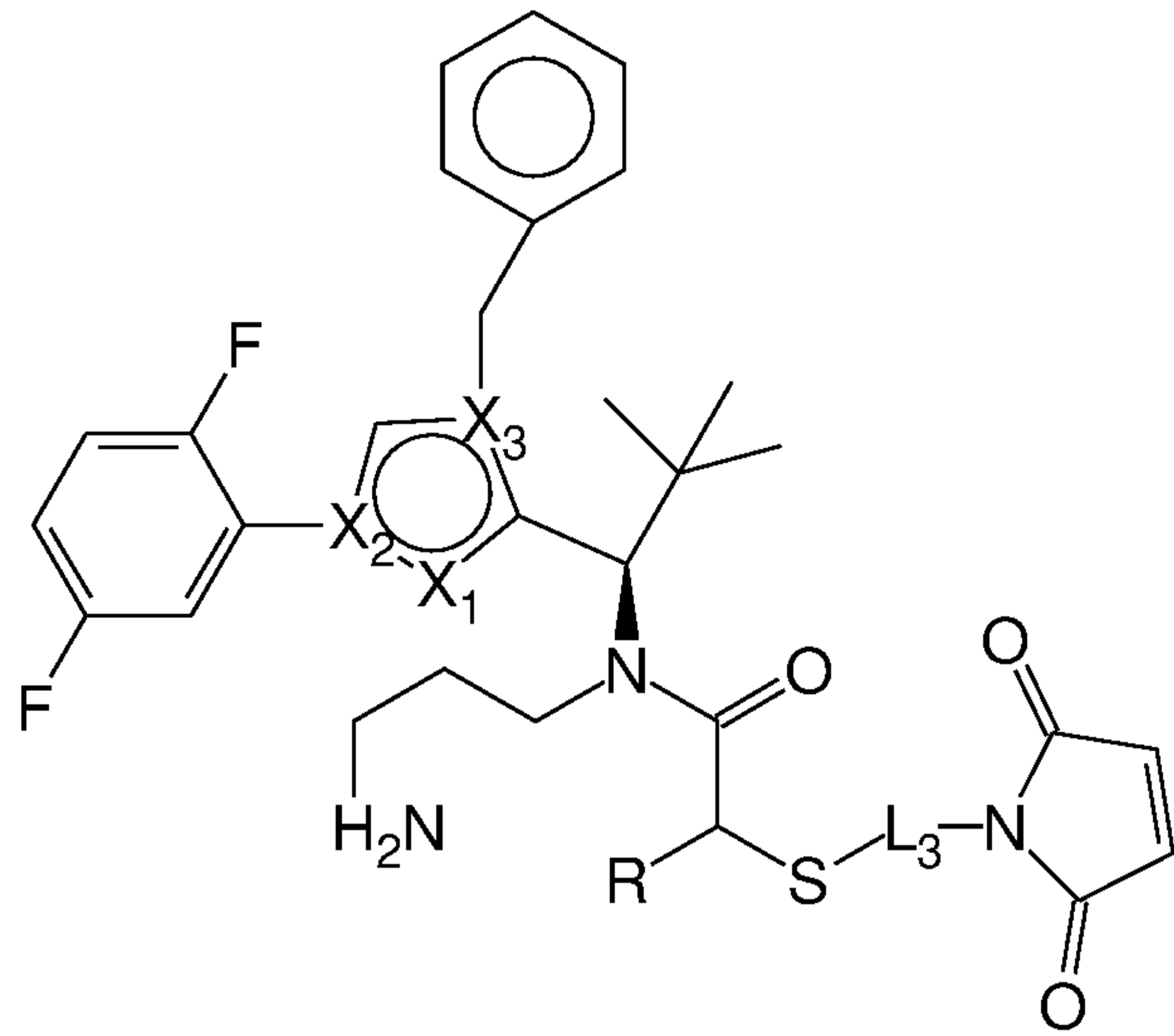
10

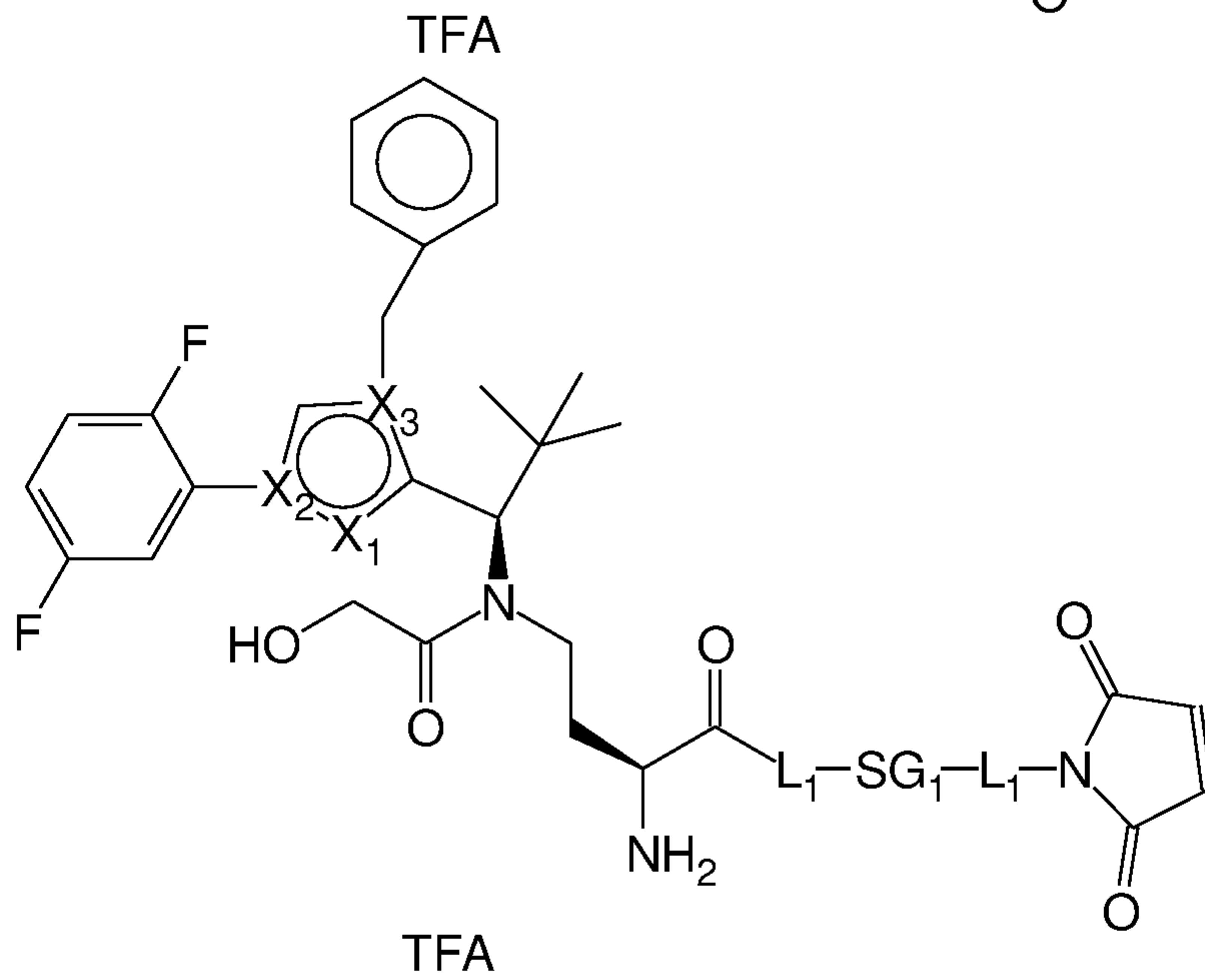
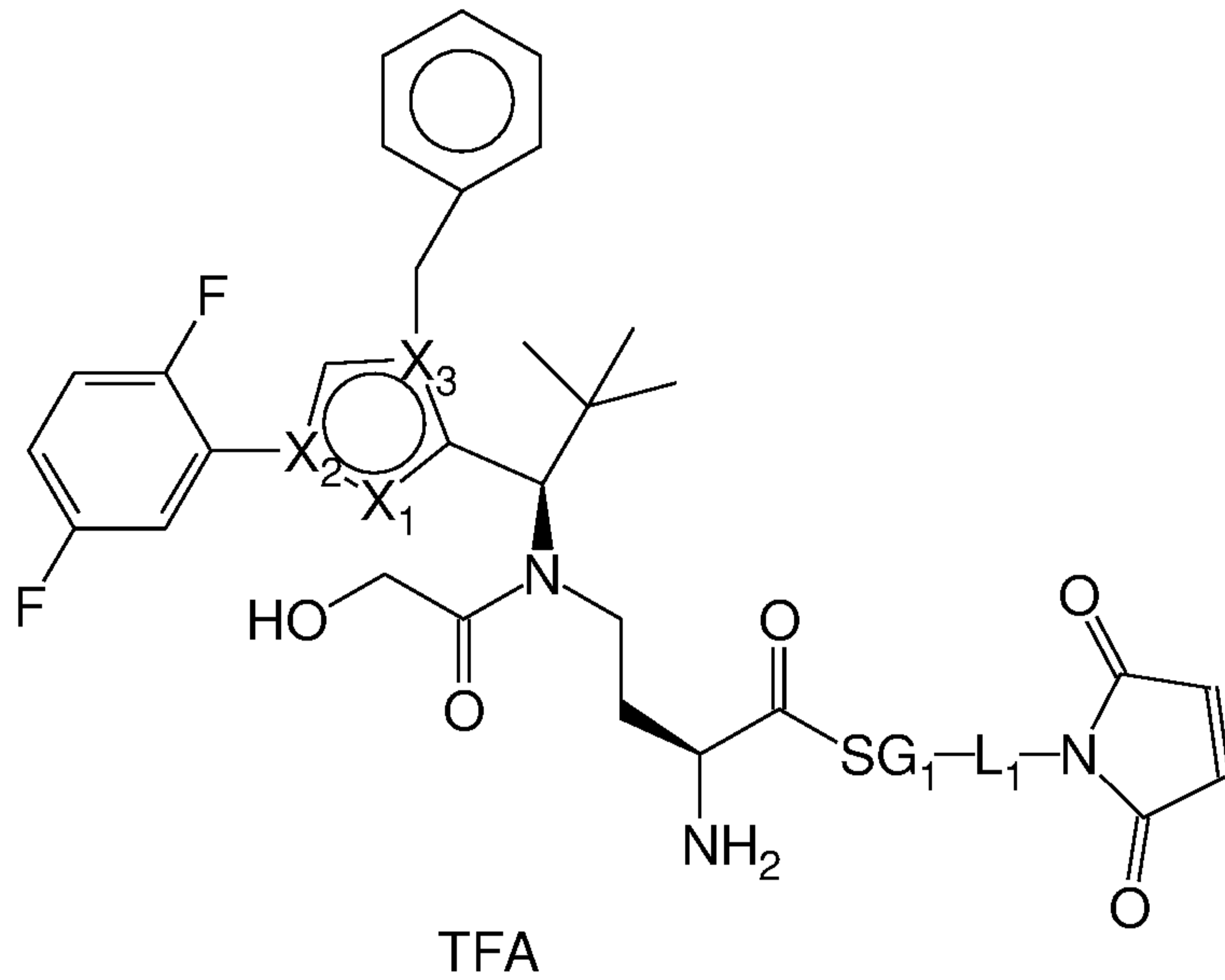
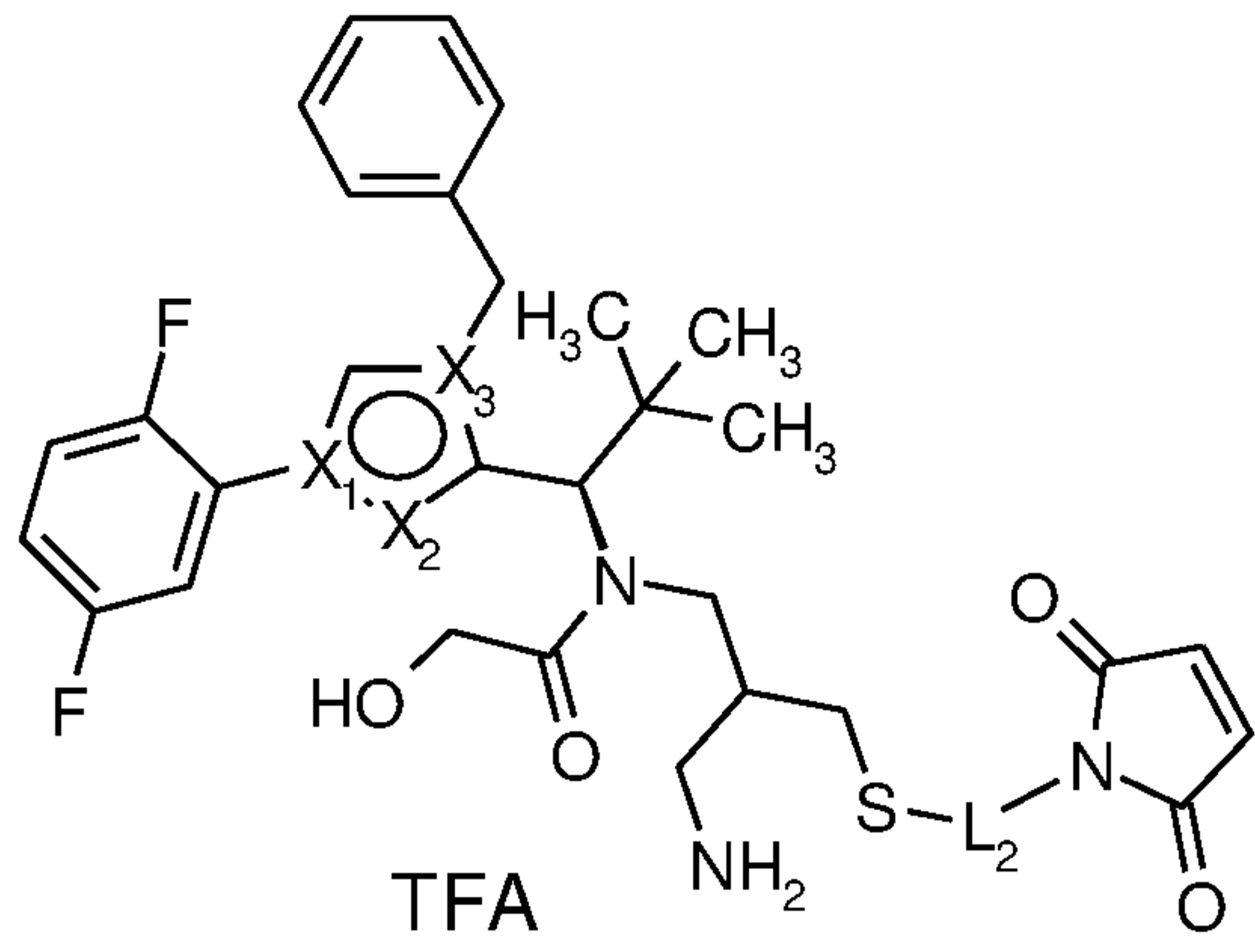
***KSP inhibitor - linker-intermediates and preparation of the conjugates***

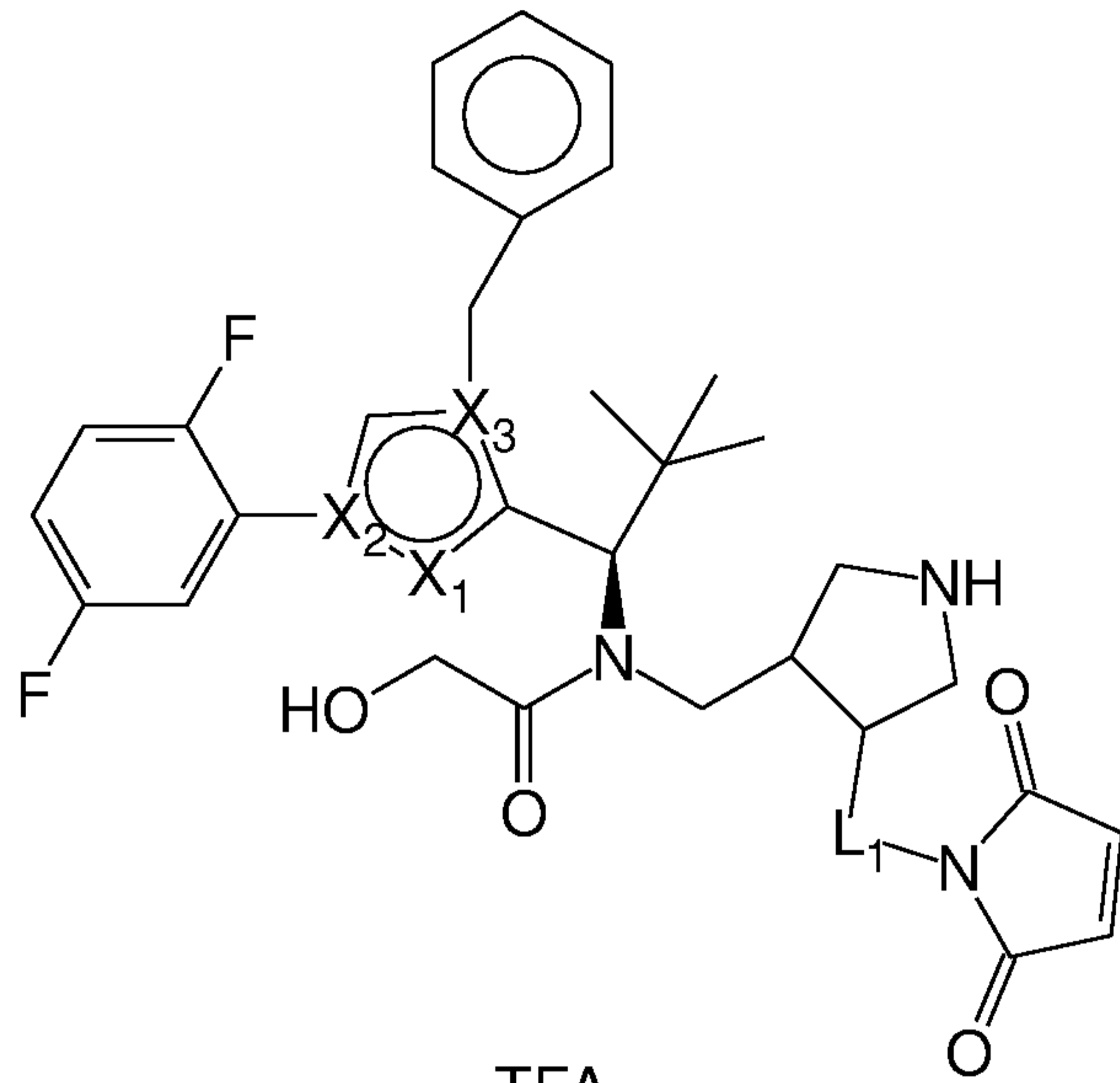
The conjugates according to the invention are prepared by initially providing the low-molecular weight KSP inhibitor with a linker. The intermediate obtained in this manner is then reacted with the binder (preferably antibody).

Preferably, for coupling to a cysteine residue, one of the compounds below is reacted with the cysteine-containing binder such as an antibody, which is optionally partially reduced for this purpose:

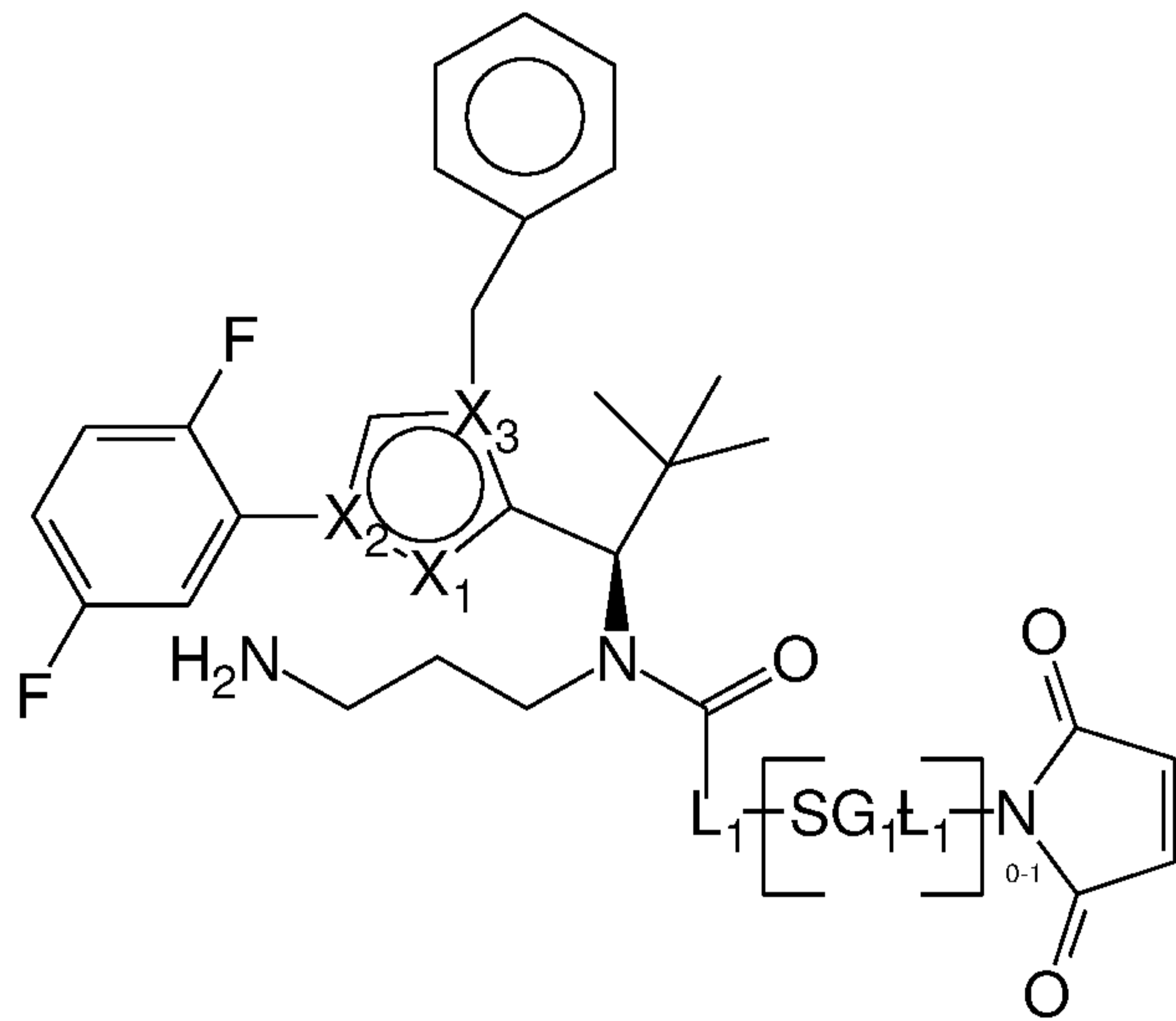




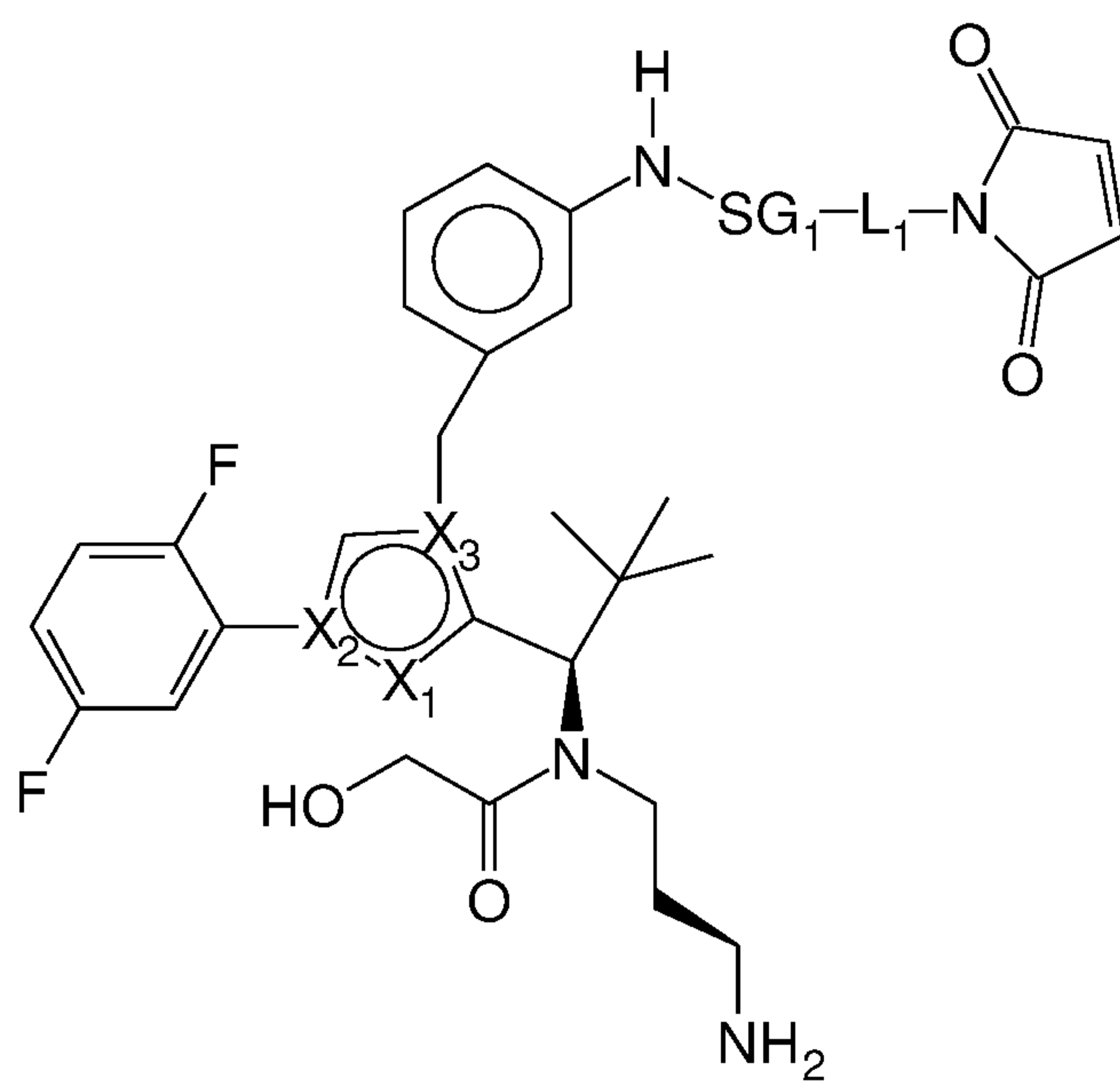




TFA



TFA



TFA

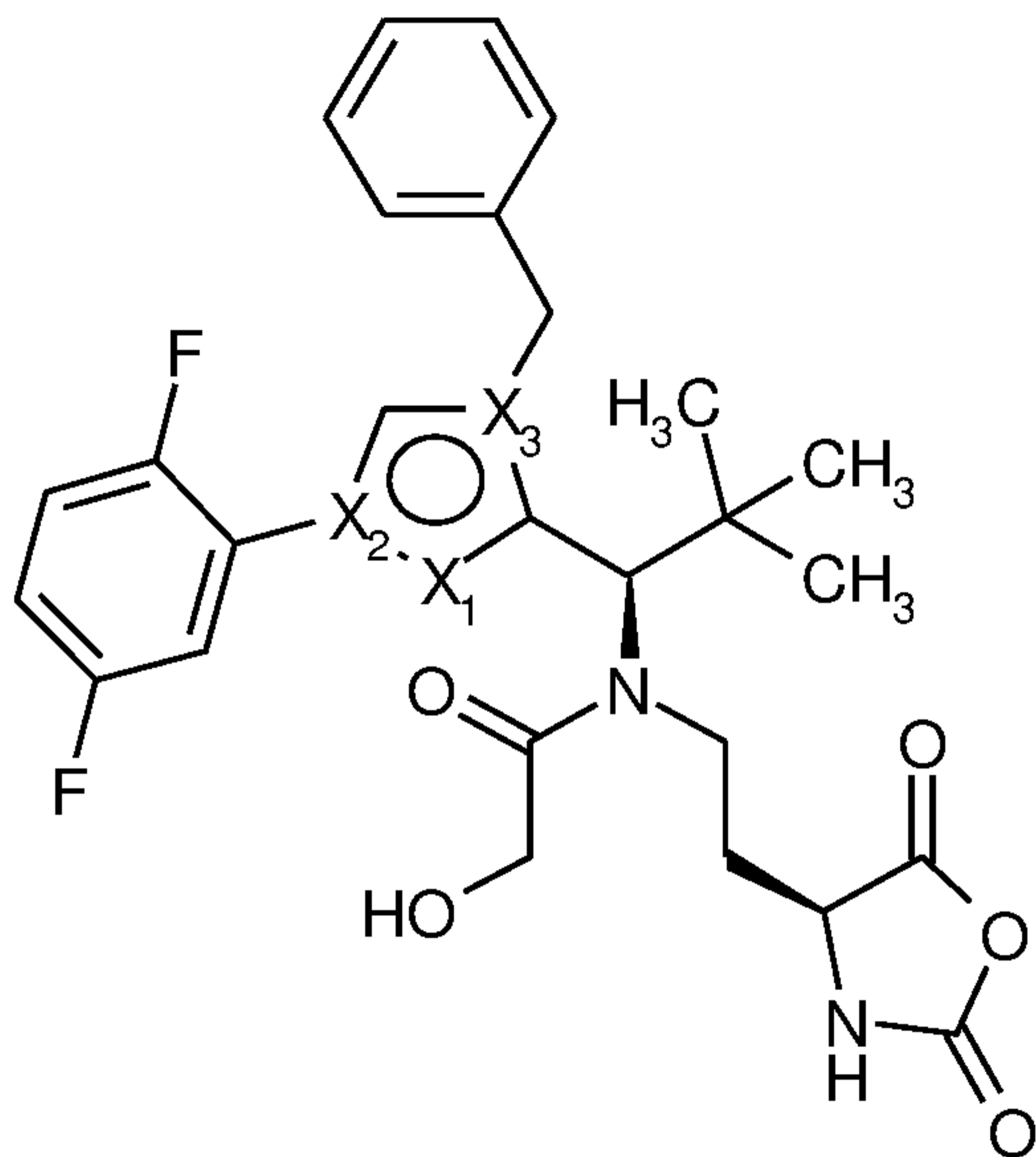
where R represents -H or -COOH,

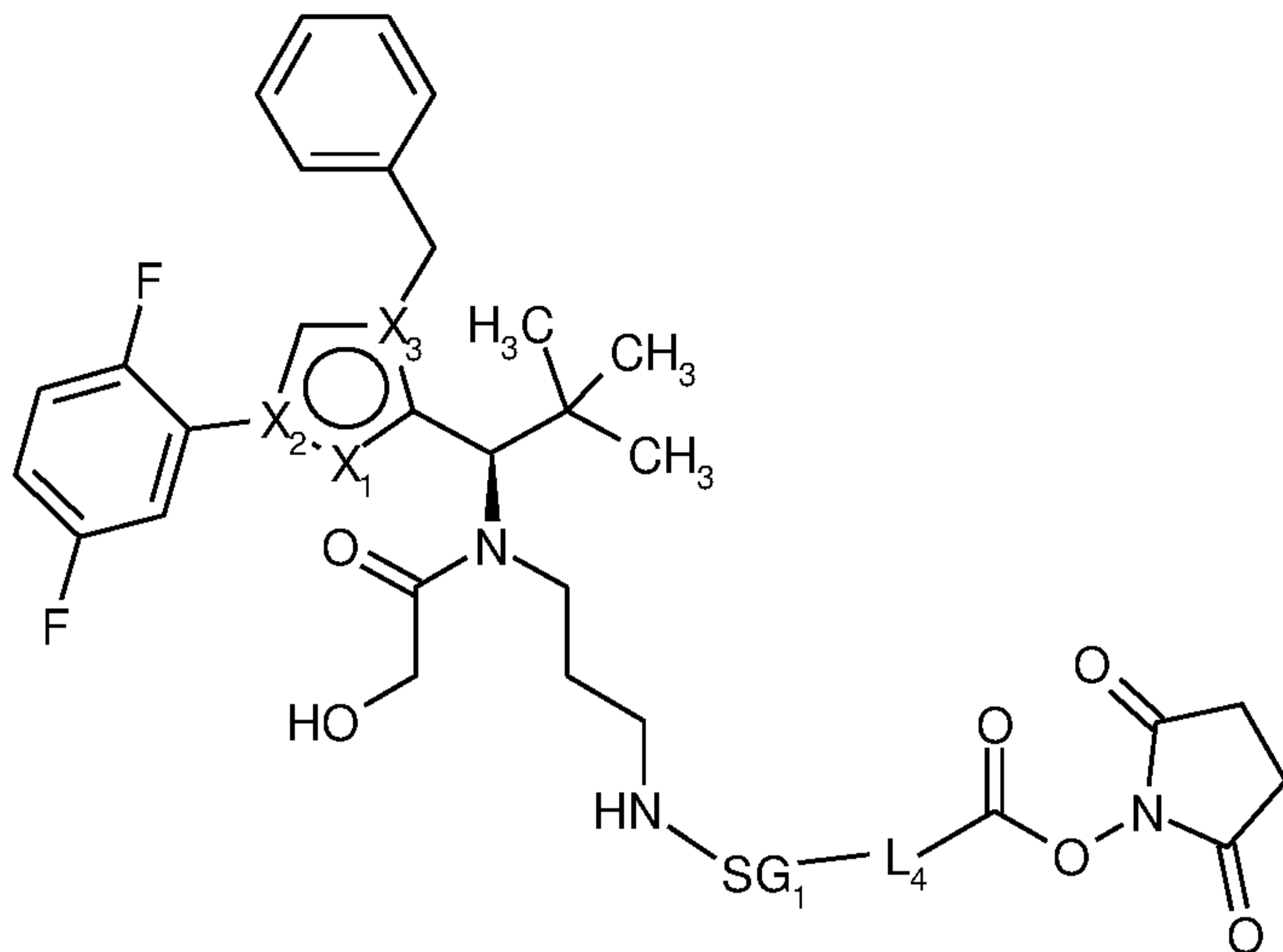
where K represents straight-chain or branched C<sub>1</sub>-C<sub>6</sub> alkyl which  
5 is optionally substituted by C<sub>1</sub>-C<sub>6</sub>-alkoxy or -OH, and

where X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub>, SG1, L1, L2, L3 and L4 have the same meaning  
as described above.

10 The compound may be employed, for example, in the form of its  
trifluoroacetic acid salt. For the reaction with the binder such  
as, for example, the antibody, the compound is preferably used  
in a 2- to 12-fold molar excess with respect to the binder.

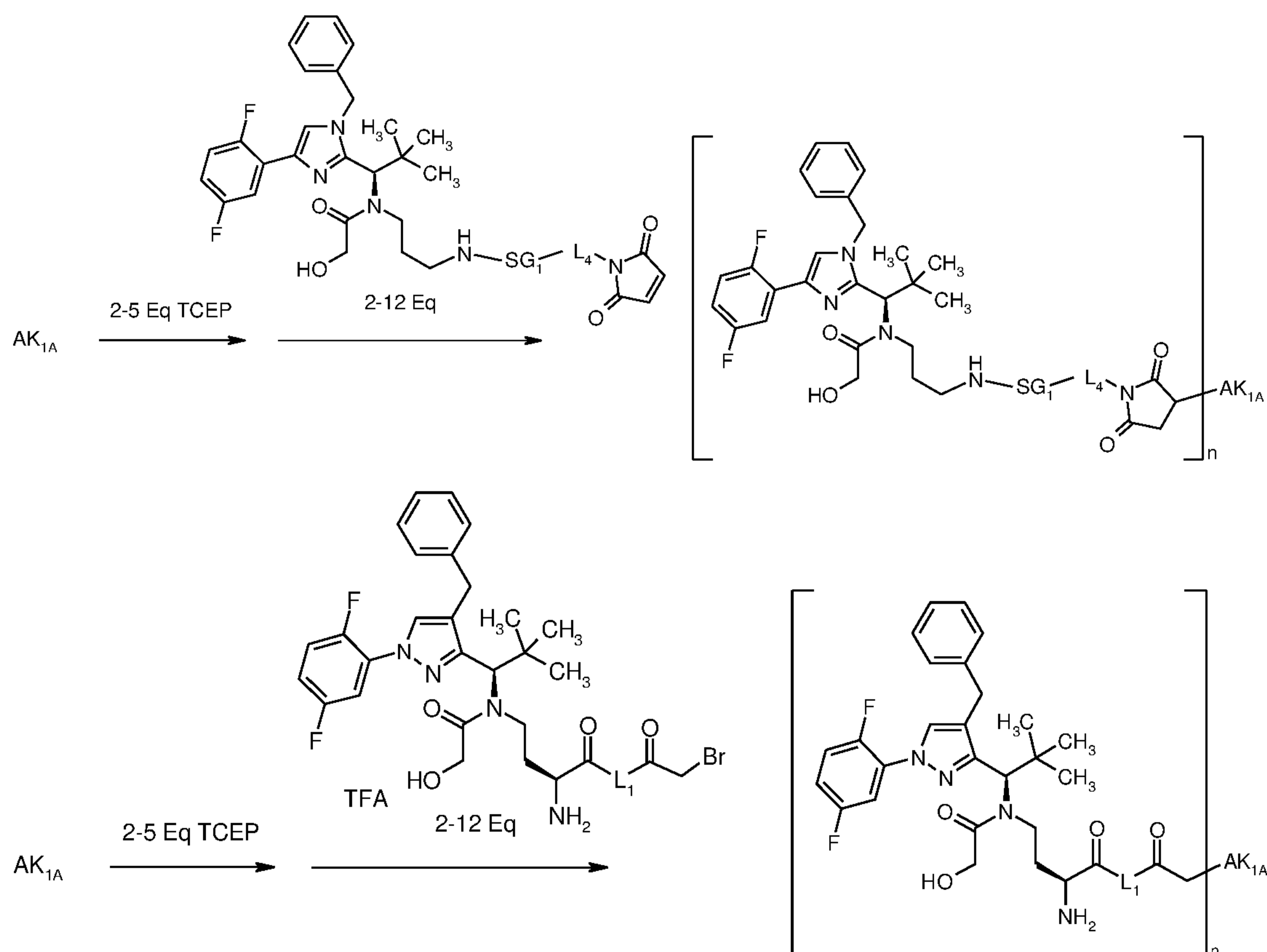
15 Preferably, for coupling to a lysine residue, one of the  
compounds below is reacted with the lysine-containing binder  
such as an antibody:





where  $X_1$ ,  $X_2$ ,  $X_3$  have the same meaning as in formula (II) and  $L_4$  has the same meaning as  $L_1$  and  $L_1$  has the same meaning as described above.

For an intermediate coupling to a cysteine residue, the reactions can be illustrated as follows:

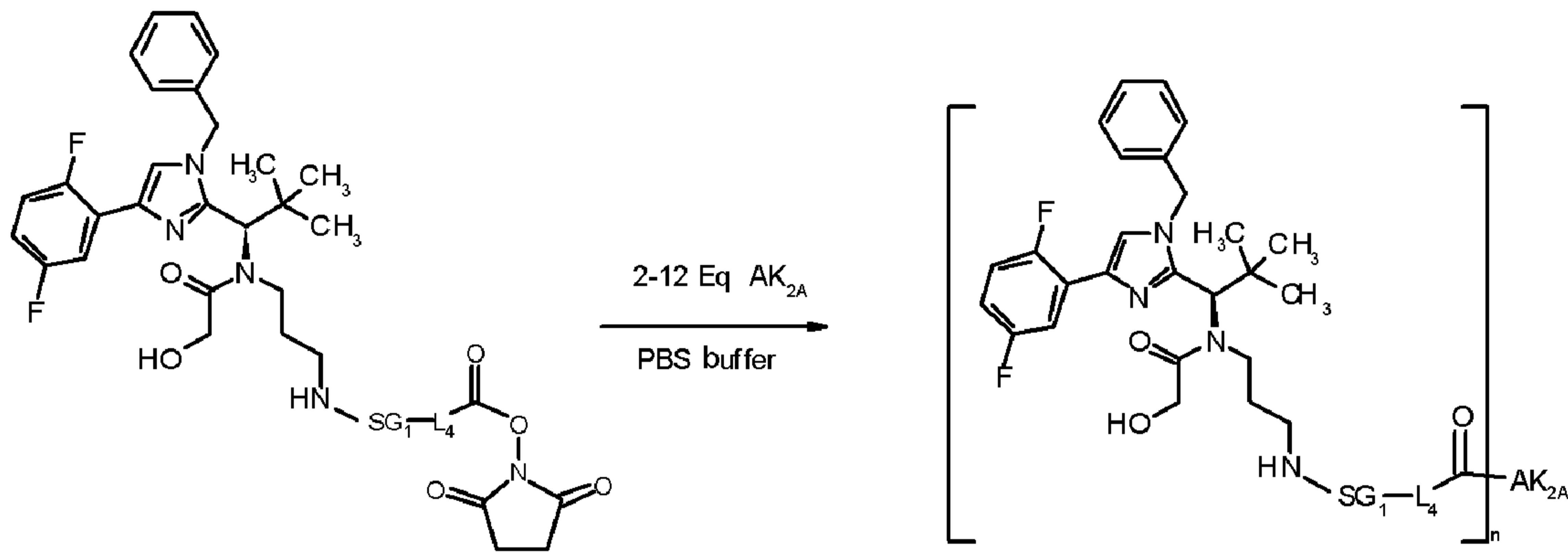


The other intermediates and other antibodies can be reacted

correspondingly.

For an intermediate coupling to a lysine radical, the reaction can be illustrated as follows:

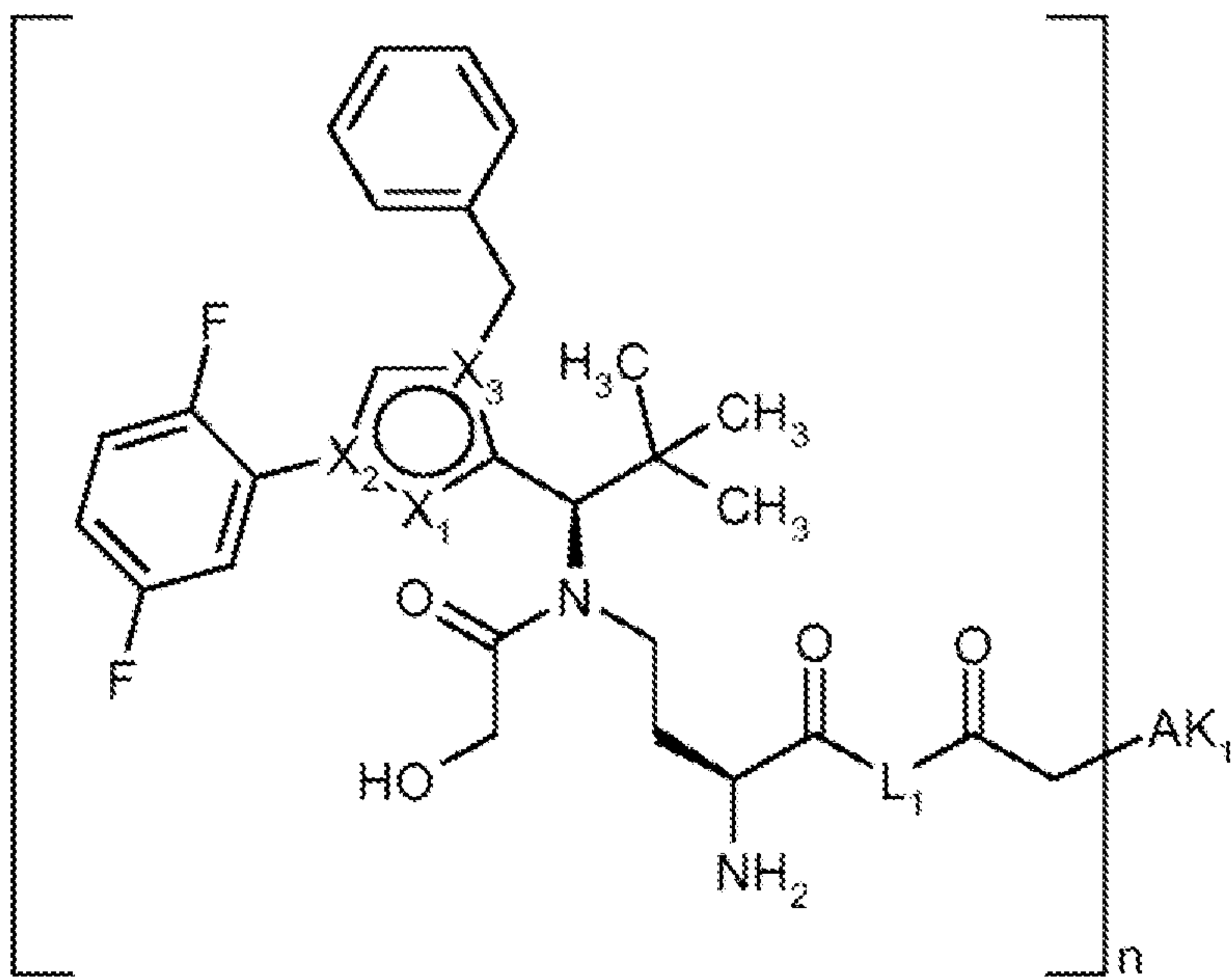
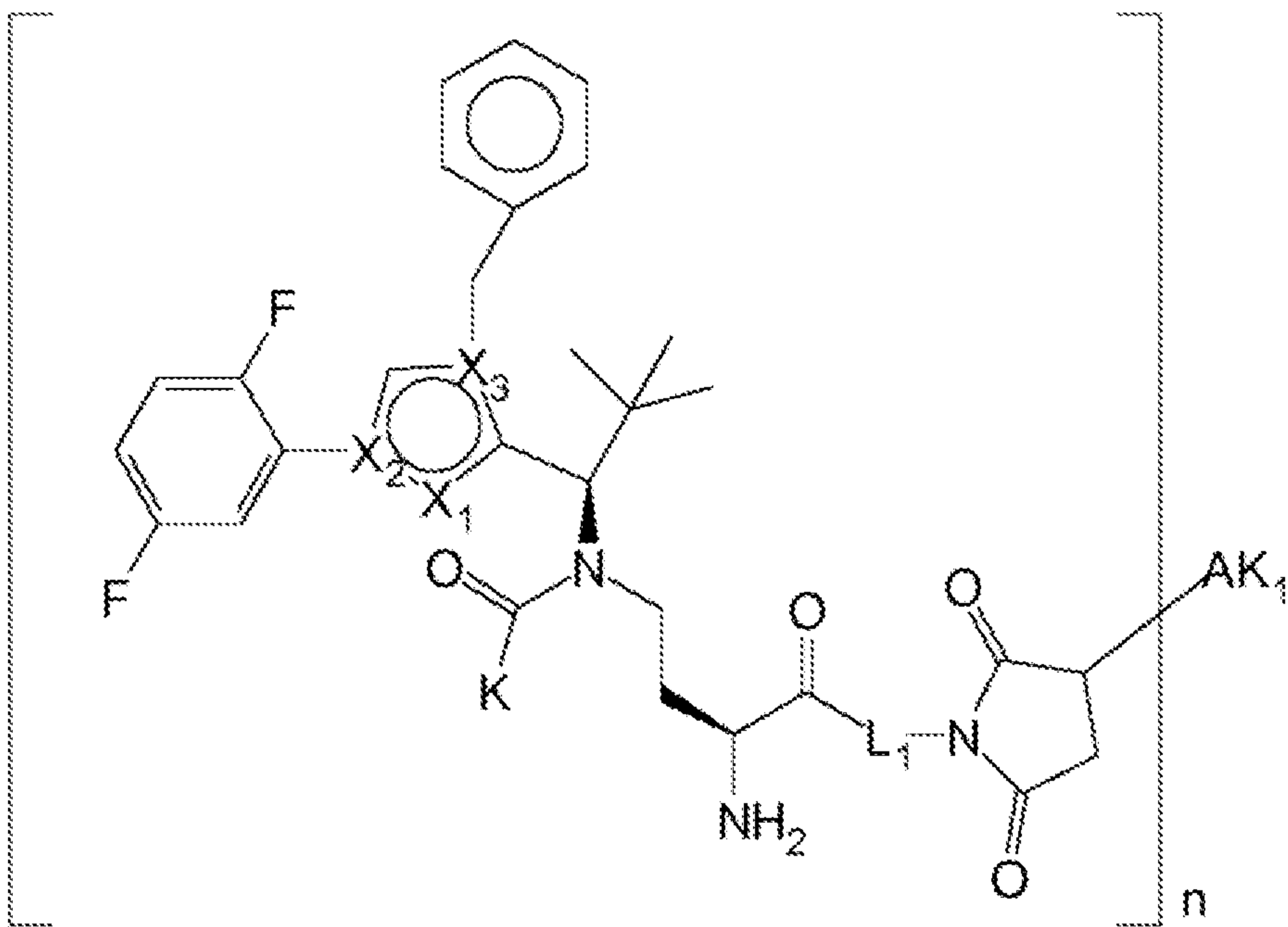
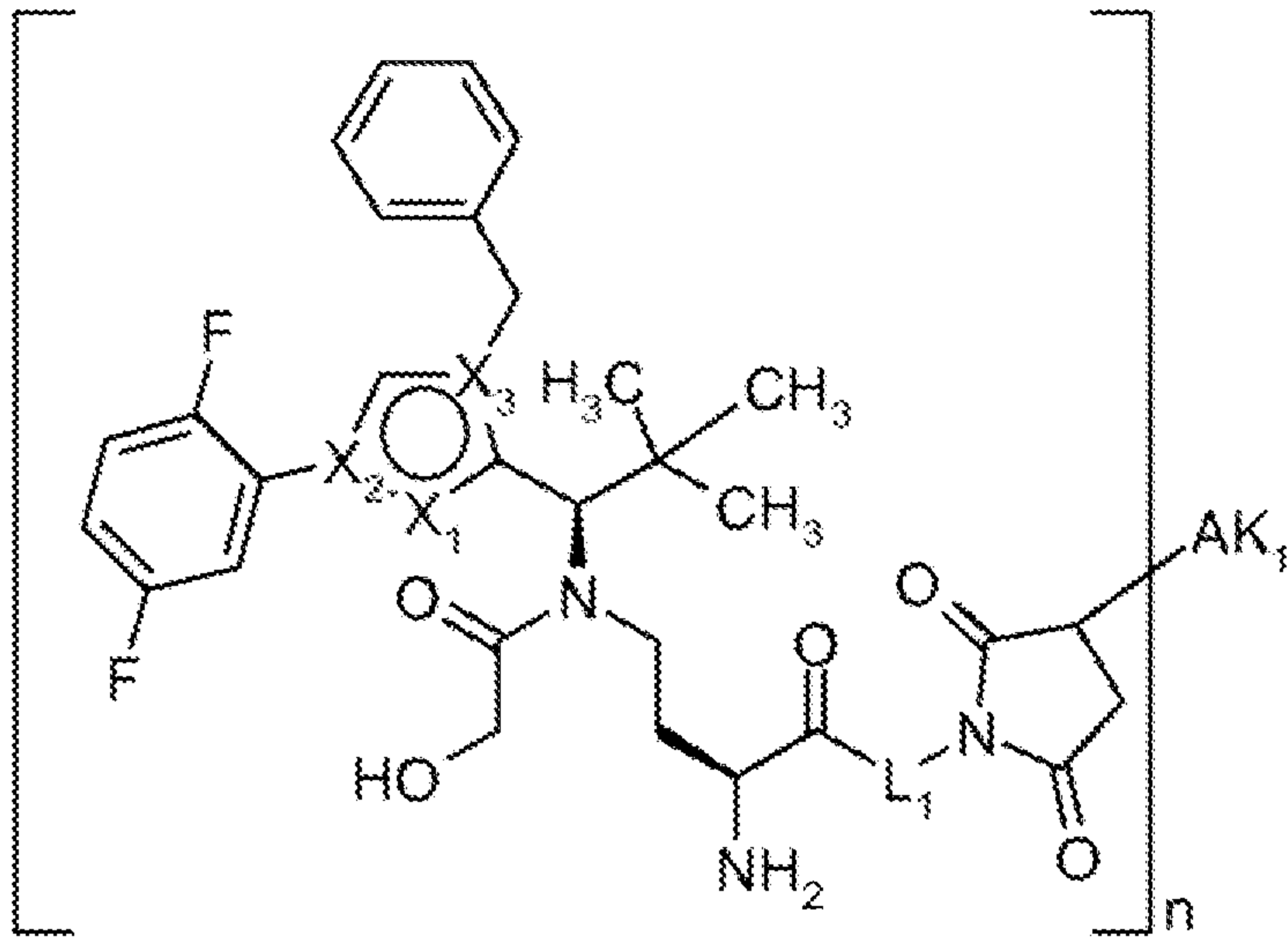
5

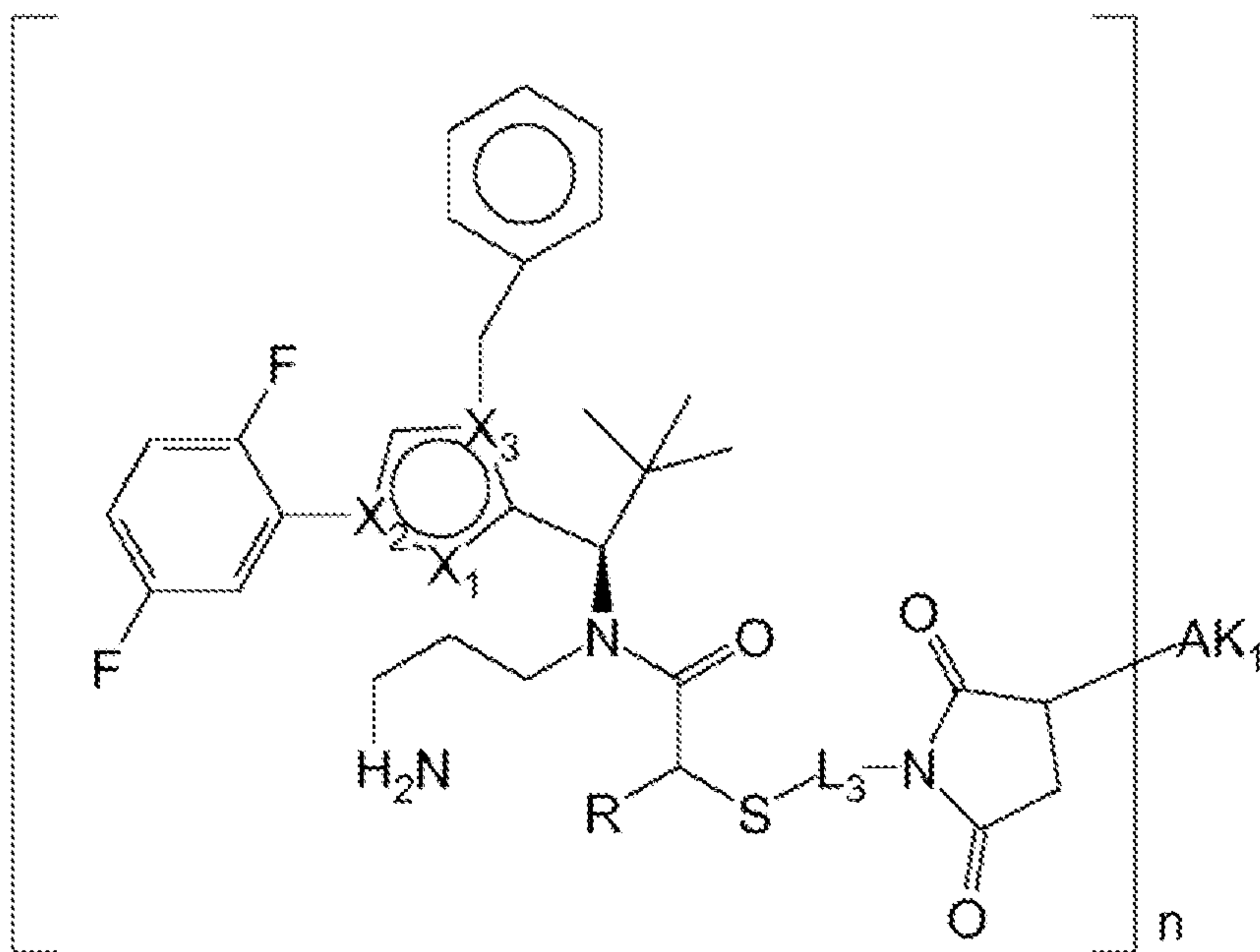
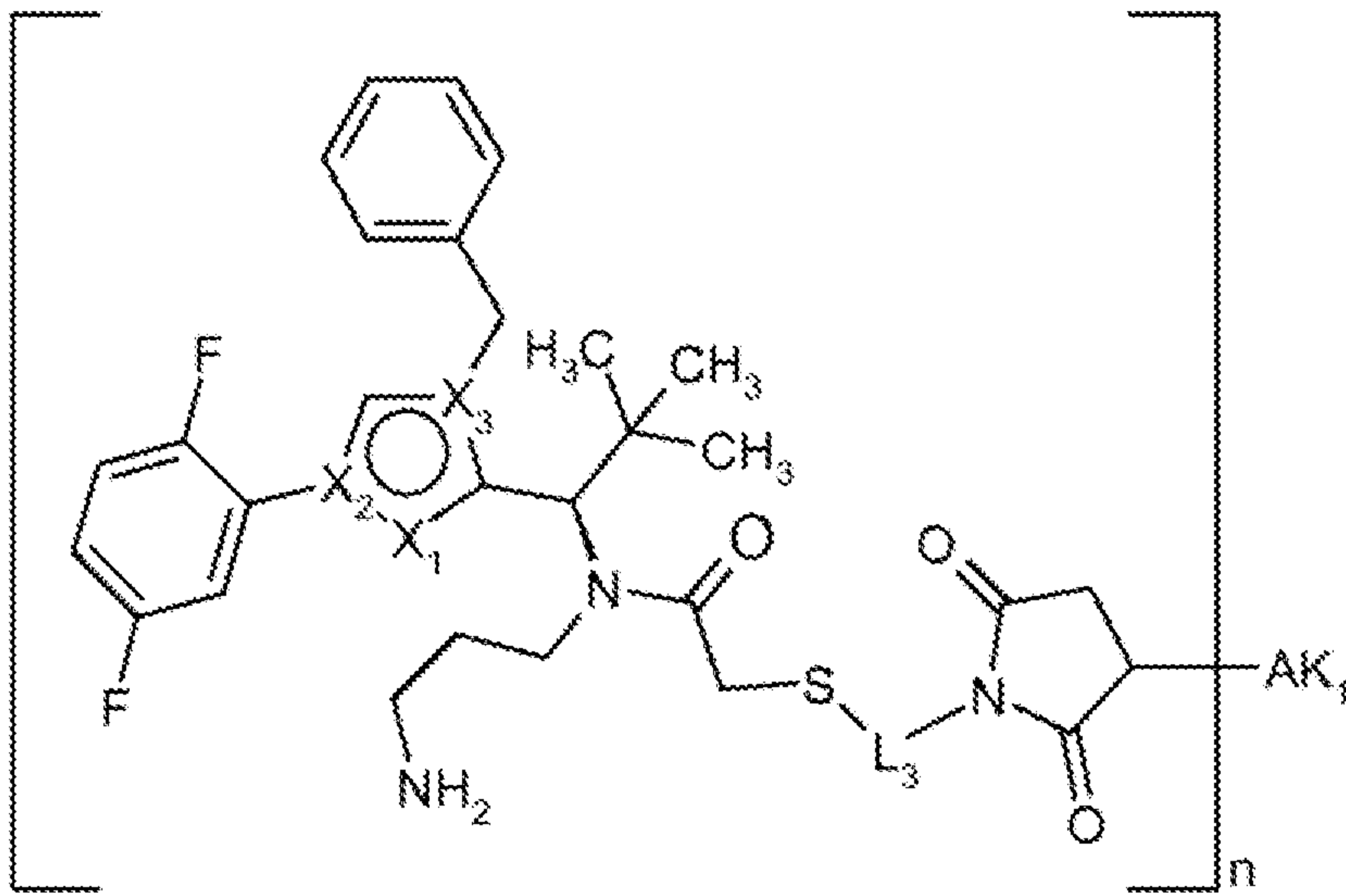
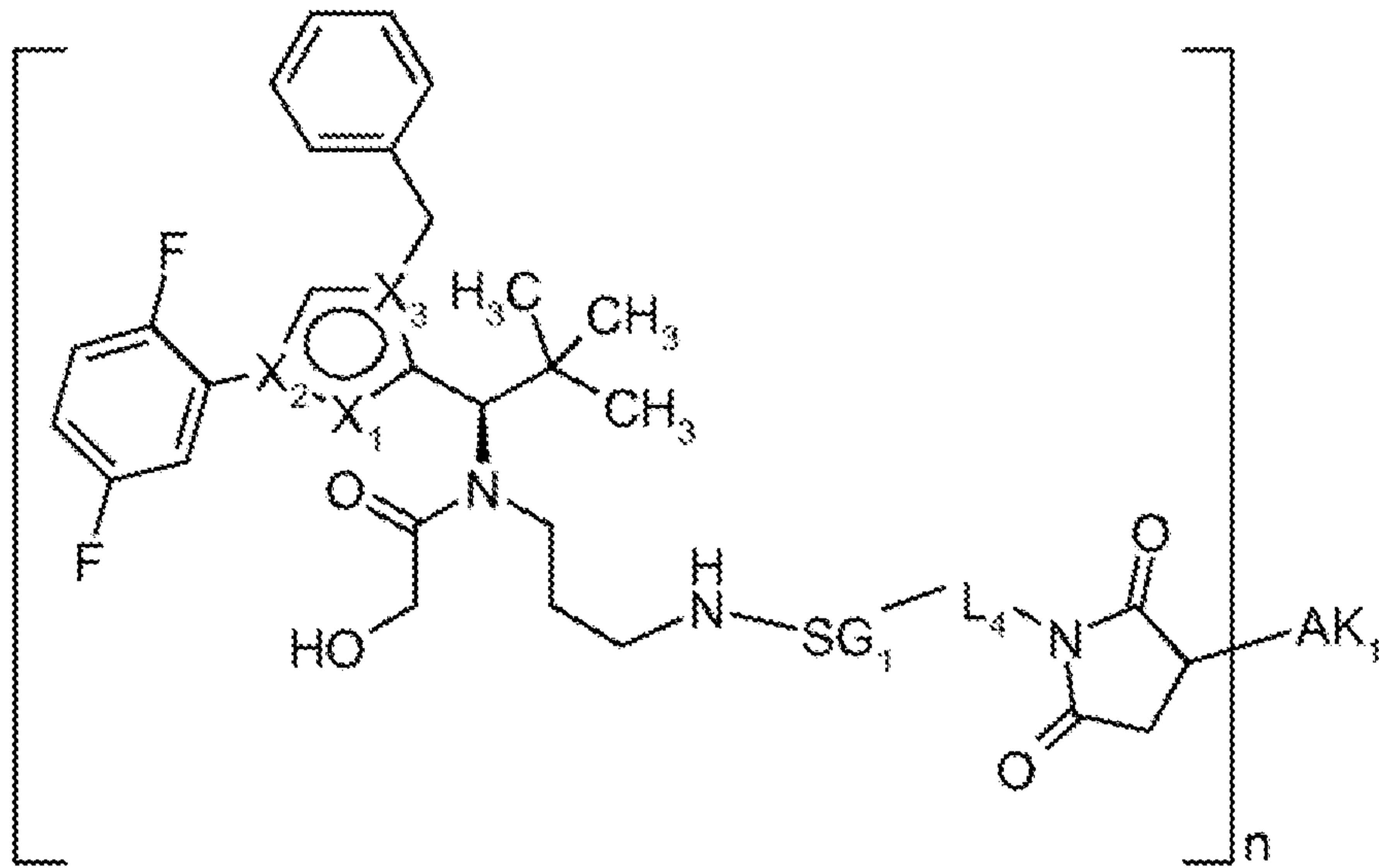


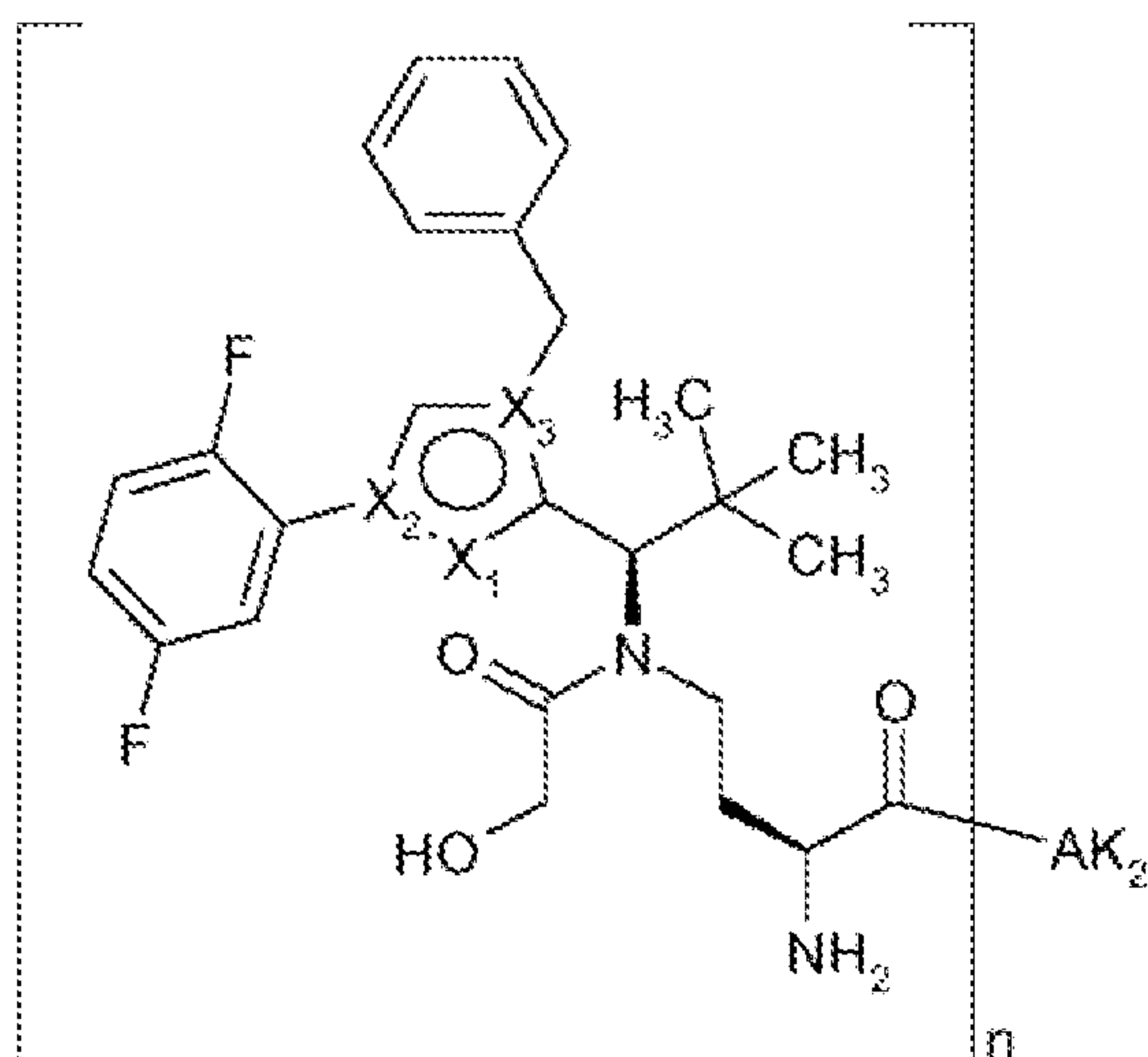
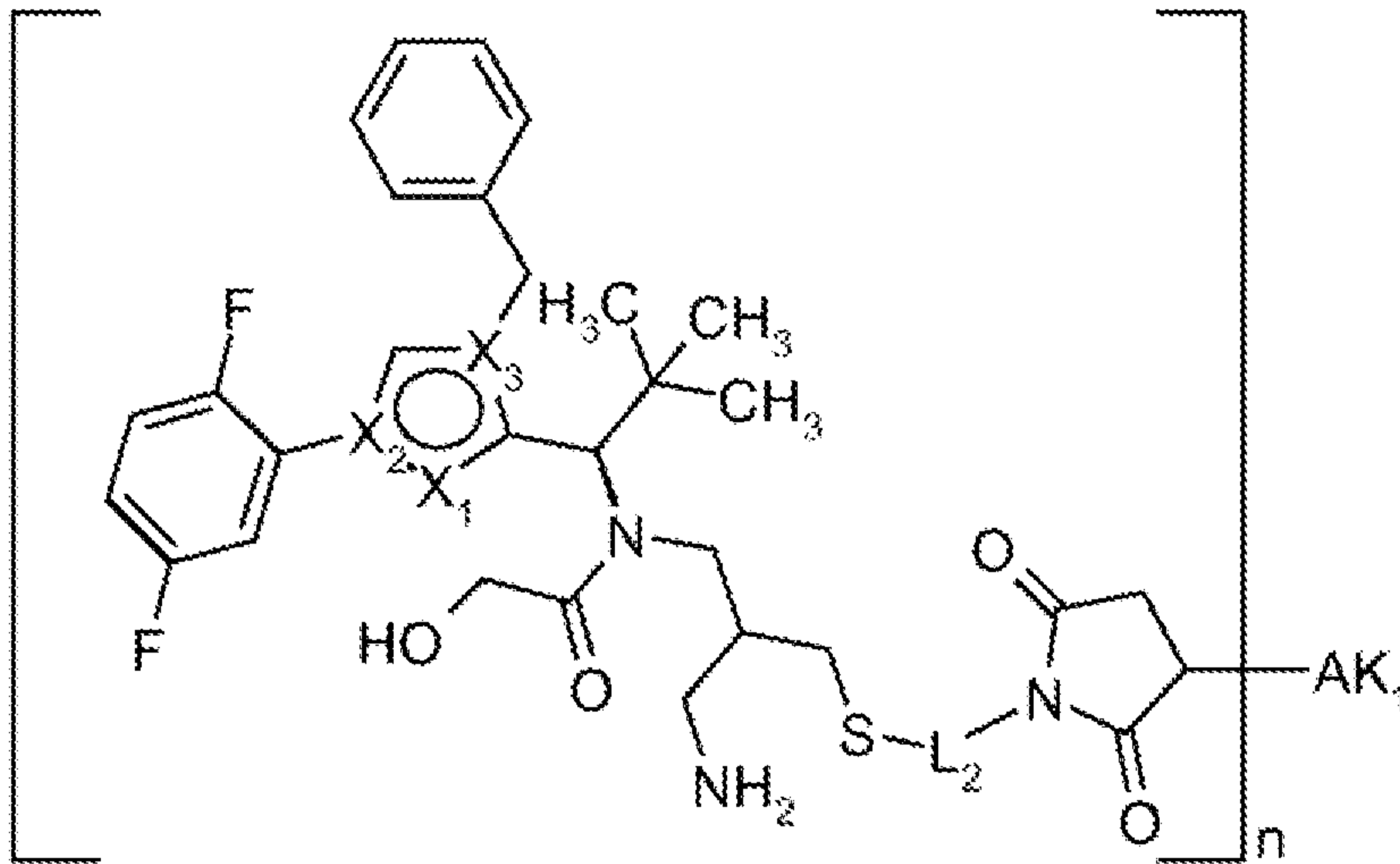
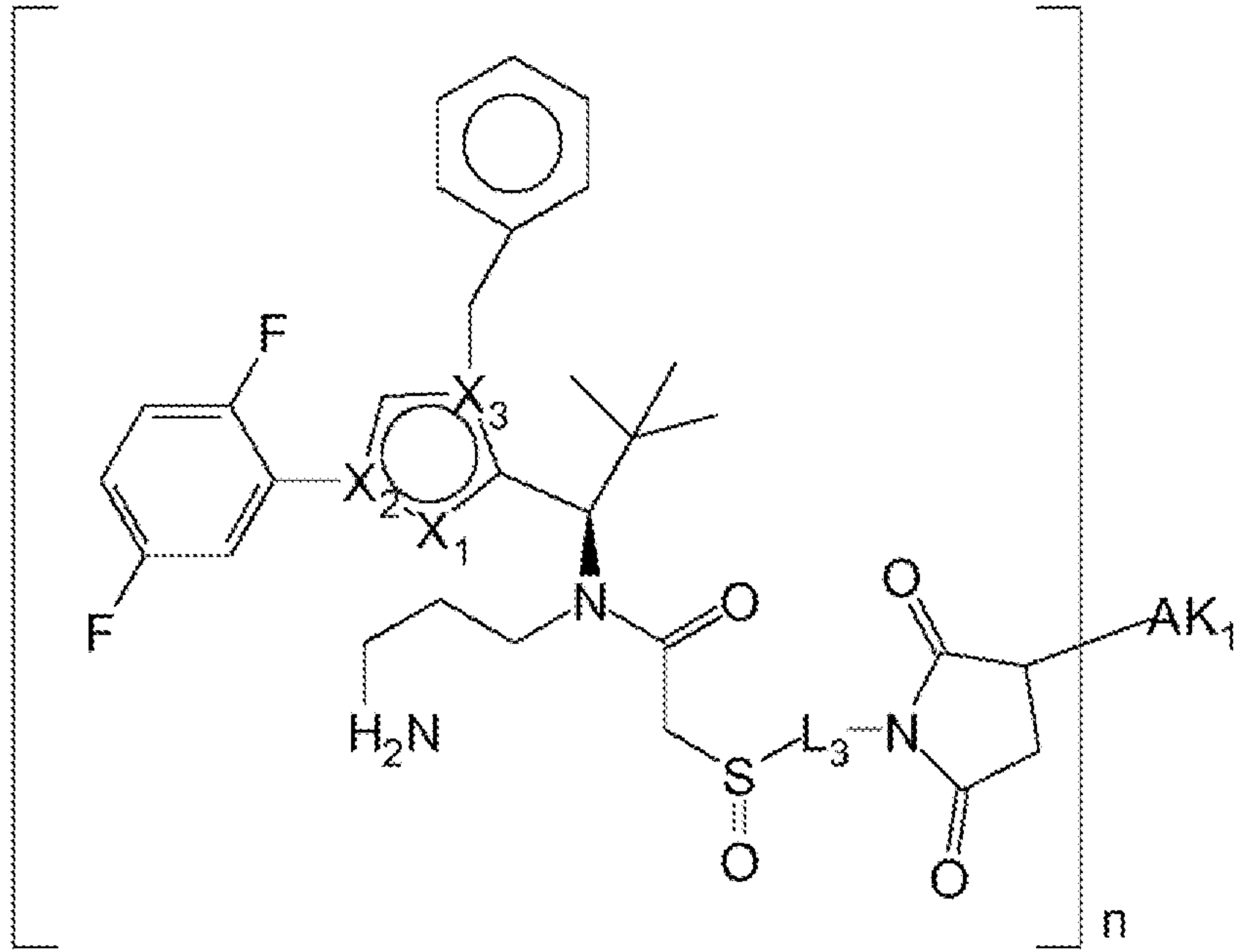
In accordance with the invention, this gives the following conjugates:

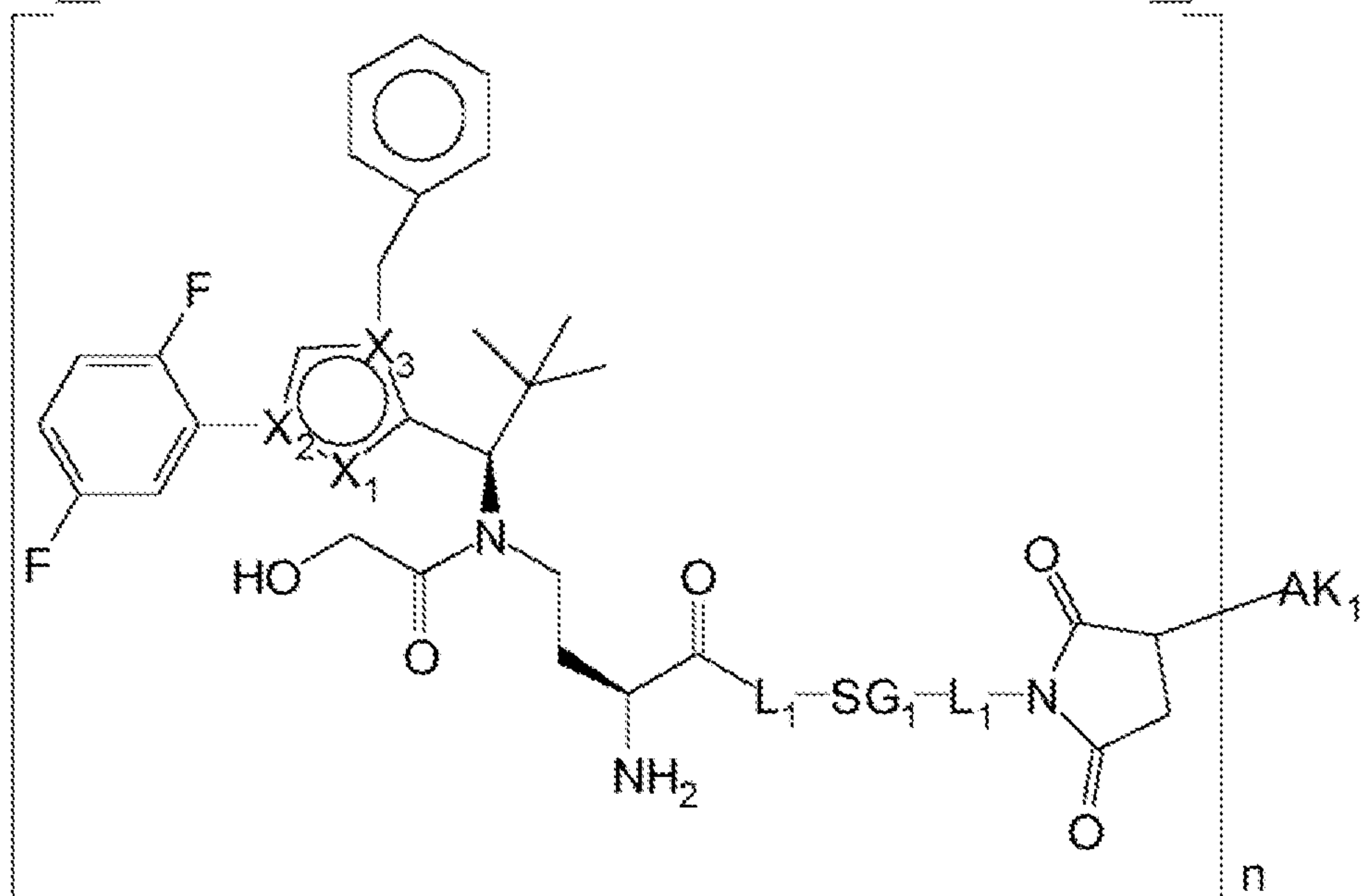
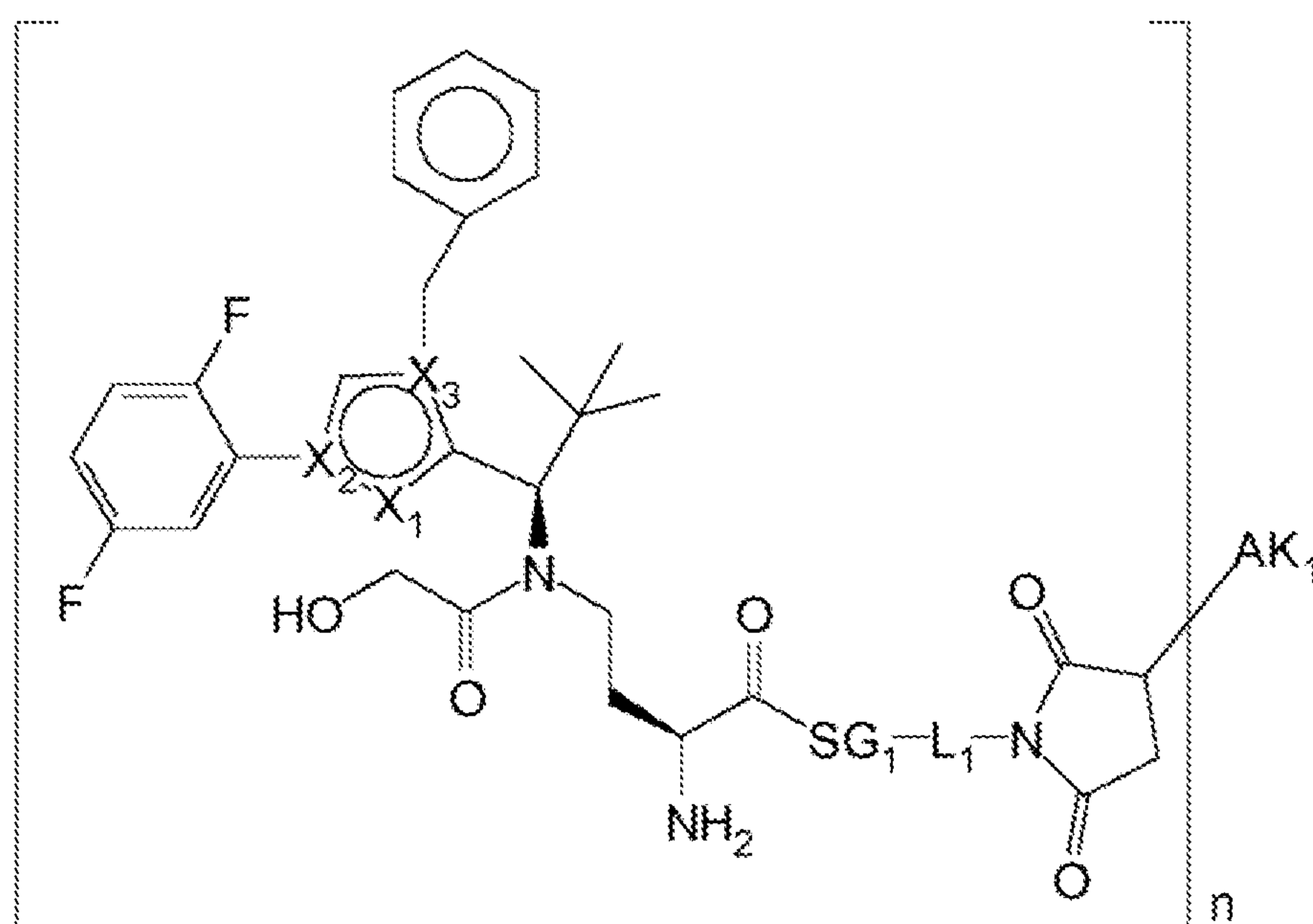
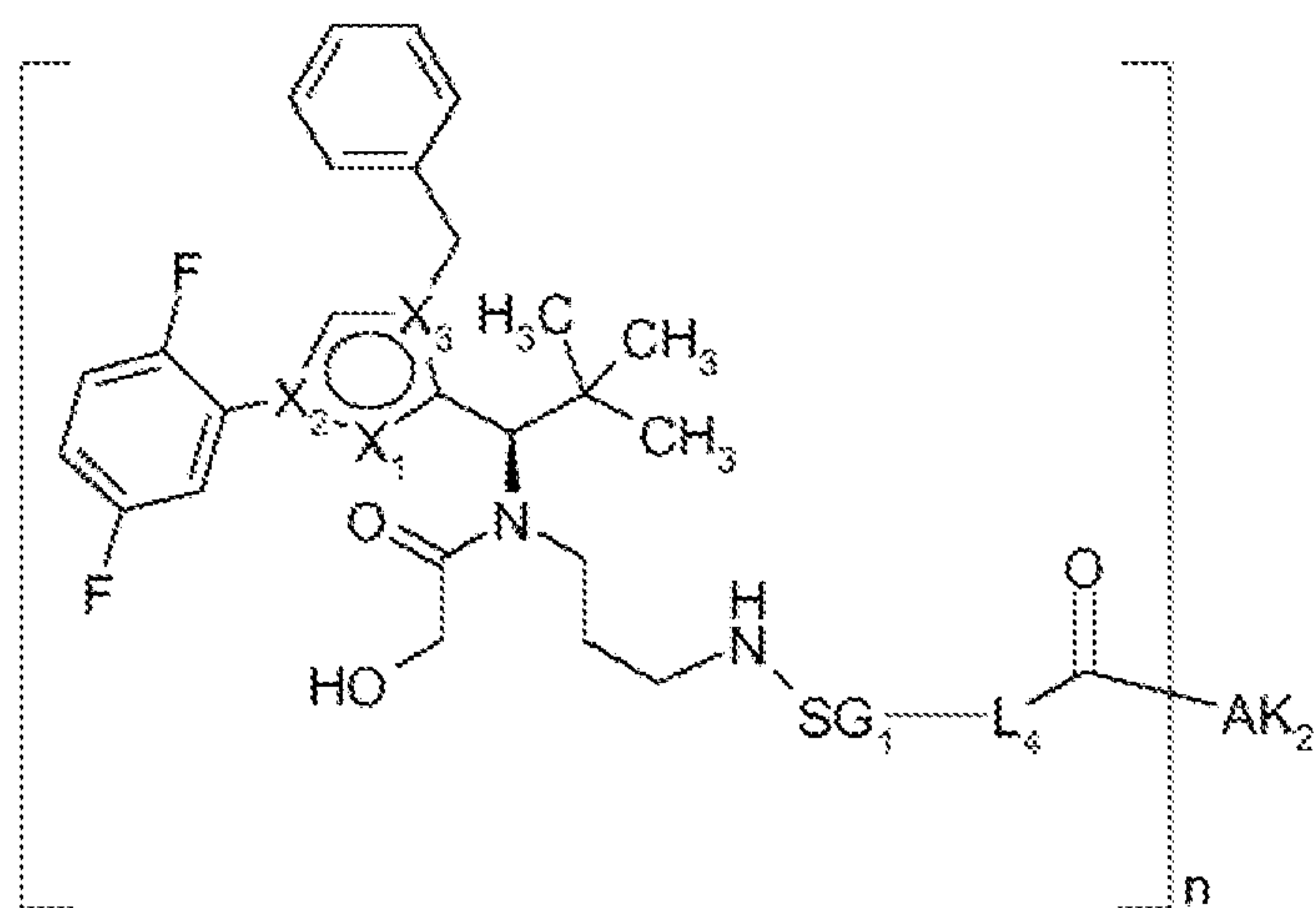
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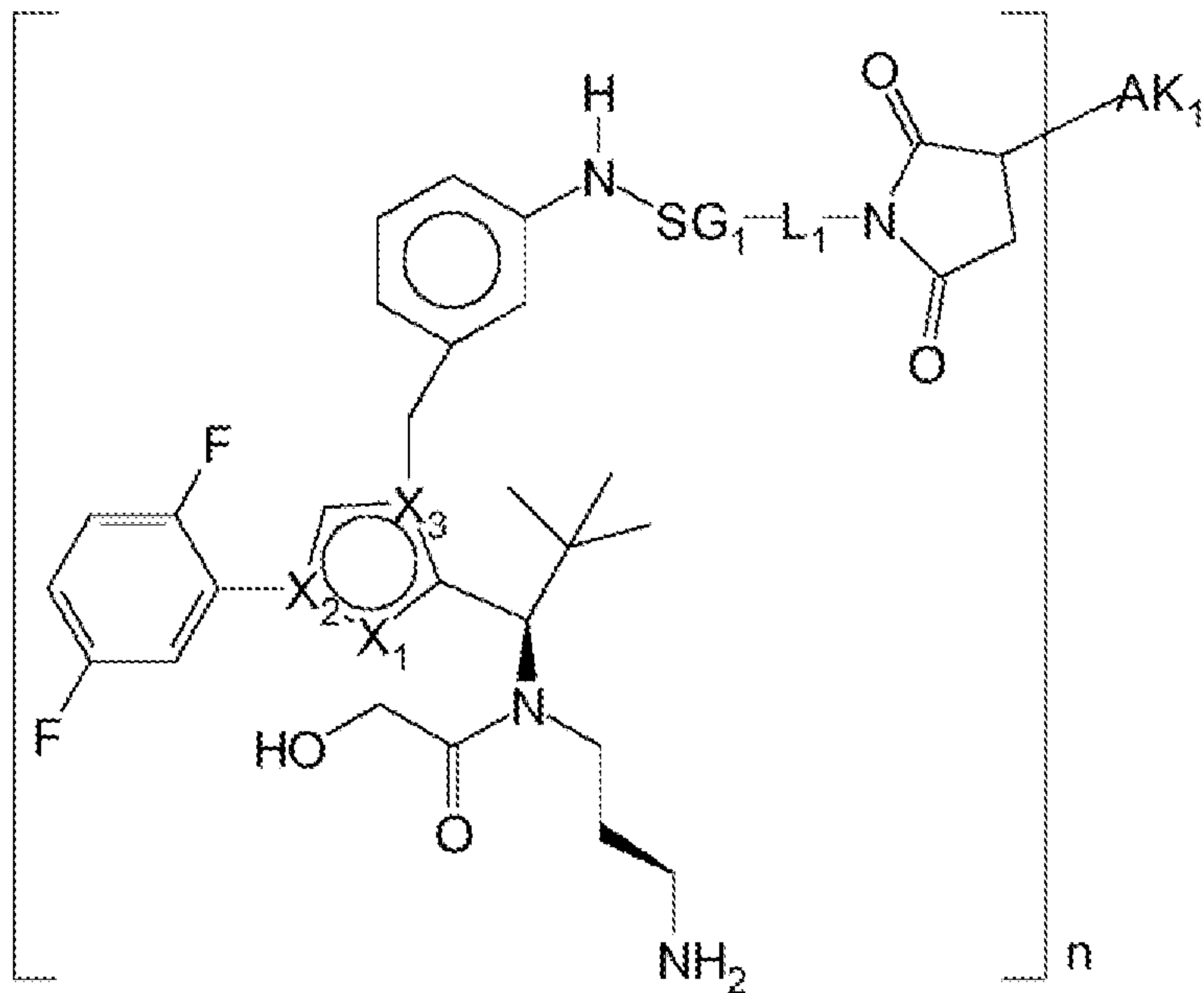




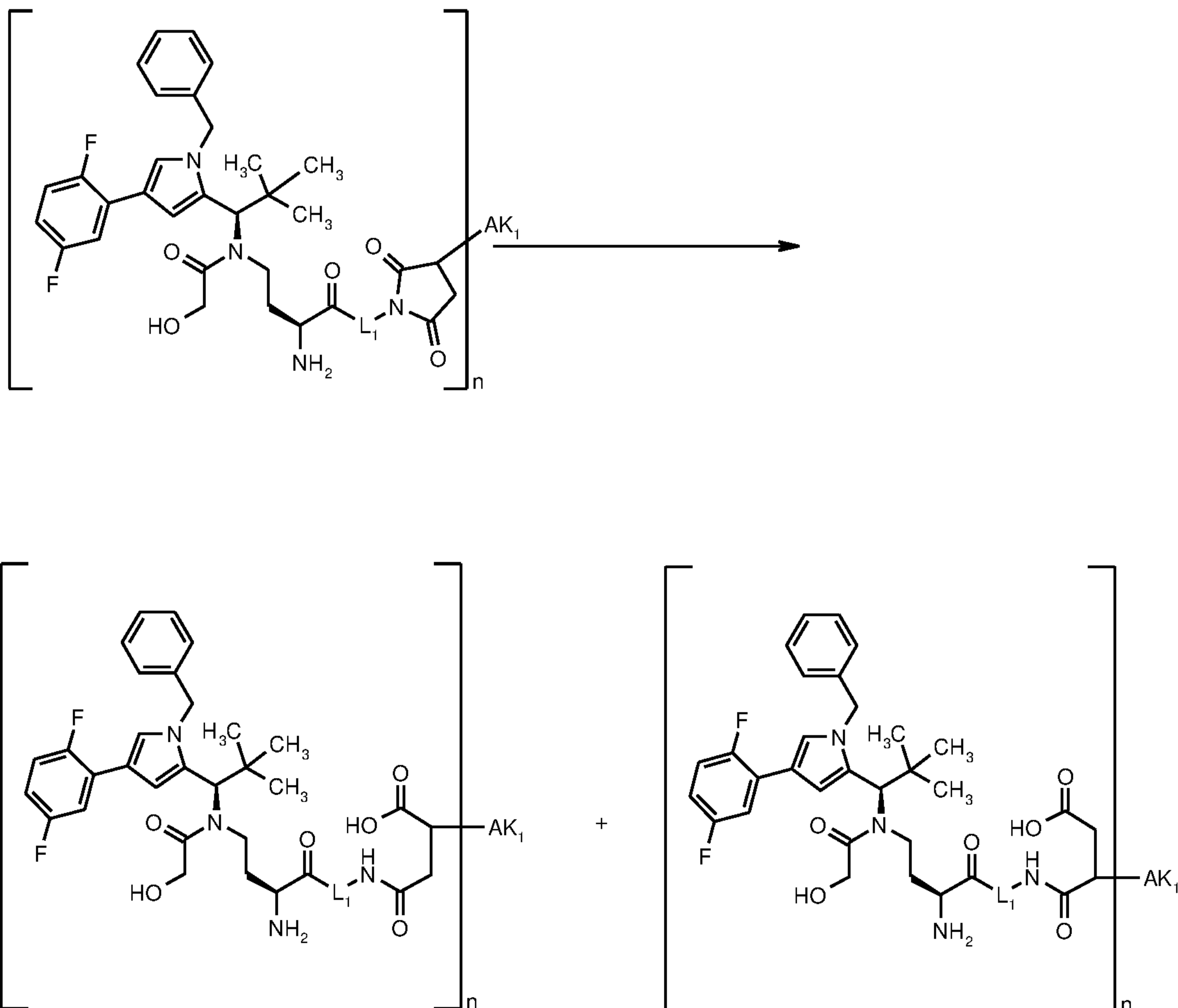








Depending on the linker, succinimide-linked ADCs may, after conjugation, be converted into the open-chain succinamides, which have an advantageous stability profile.



This reaction (ring opening) can be carried out at pH 7.5 to 9, preferably at pH 8, at a temperature of from 25°C to 37°C, for example by stirring. The preferred stirring time is 8 to 30  
5 hours.

In the above formulae, X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub> have the same meaning as in formula (II), SG1 and L1 have the same meaning as described above and L2, L3 and L4 have the same meaning as L1; R and K  
10 have the same meaning as described above. AK1 is an antibody coupled via a cysteine residue, and AK2 is an antibody coupled via a lysine residue. With particular preference, AK1 and AK2 are anti-TWEAKR antibodies, in particular antibodies which bind specifically to amino acid D in position 47 (D47) of TWEAKR (SEQ  
15 ID NO:169), in particular the anti-TWEAKR antibody TPP-2090.

### ***Binders***

In the broadest sense, the term "binder" is understood to mean  
20 a molecule which binds to a target molecule present at a certain target cell population to be addressed by the binder/active compound conjugate. The term binder is to be understood in its broadest meaning and also comprises, for example, lectins, proteins capable of binding to certain sugar chains, and  
25 phospholipid-binding proteins. Such binders include, for example, high-molecular weight proteins (binding proteins), polypeptides or peptides (binding peptides), non-peptidic (e.g. aptamers (US5,270,163) review by Keefe AD., et al., Nat. Rev. Drug Discov. 2010; 9:537-550), or vitamins) and all other cell-  
30 binding molecules or substances. Binding proteins are, for example, antibodies and antibody fragments or antibody mimetics such as, for example, affibodies, adnectins, anticalins, DARPins, avimers, nanobodies (review by Gebauer M. et al., Curr. Opinion in Chem. Biol. 2009; 13:245-255; Nuttall S.D. et al.,  
35 Curr. Opinion in Pharmacology 2008; 8:608-617). Binding peptides are, for example, ligands of a ligand/receptor pair such as, for example, VEGF of the ligand/receptor pair VEGF/KDR, such as transferrin of the ligand/receptor pair transferrin/transferrin

receptor or cytokine/cytokine receptor, such as TNFalpha of the ligand/receptor pair TNFalpha/TNFalpha receptor.

The literature also discloses various options of covalent  
5 coupling (conjugation) of organic molecules to antibodies. Preference according to the invention is given to the conjugation of the toxophores to the antibody via one or more sulphur atoms of cysteine residues of the antibody and/or via one or more NH groups of lysine residues of the antibody.  
10 However, it is also possible to bind the toxophor to the antibody via free carboxyl groups or via sugar residues of the antibody.

A "target molecule" in the broadest sense is understood to mean a molecule which is present in the target cell population and  
15 which may be a protein (for example a receptor of a growth factor) or a non-peptidic molecule (for example a sugar or phospholipid). It is preferably a receptor or an antigen.

The term "extracellular" target molecule describes a target  
20 molecule, attached to the cell, which is located at the outside of a cell, or the part of a target molecule which is located at the outside of a cell, i.e. a binder may bind on an intact cell to its extracellular target molecule. An extracellular target molecule may be anchored in the cell membrane or be a component  
25 of the cell membrane. The person skilled in the art is aware of methods for identifying extracellular target molecules. For proteins, this may be by determining the transmembrane domain(s) and the orientation of the protein in the membrane. These data are usually deposited in protein databases (e.g. SwissProt).

30 The term "cancer target molecule" describes a target molecule which is more abundantly present on one or more cancer cell species than on non-cancer cells of the same tissue type. Preferably, the cancer target molecule is selectively present  
35 on one or more cancer cell species compared with non-cancer cells of the same tissue type, where selectively describes an at least two-fold enrichment on cancer cells compared to non-cancer cells of the same tissue type (a "selective cancer target



molecule"). The use of cancer target molecules allows the selective therapy of cancer cells using the conjugates according to the invention.

5 The binder can be attached to the linker via a bond. Attachment of the binder can be via a heteroatom of the binder. Heteroatoms according to the invention of the binder which can be used for attachment are sulphur (in one embodiment via a sulphhydryl group of the binder), oxygen (according to the invention by  
10 means of a carboxyl or hydroxyl group of the binder) and nitrogen (in one embodiment via a primary or secondary amine group or amide group of the binder). These heteroatoms may be present in the natural binder or are introduced by chemical methods or methods of molecular biology. According to the invention, the  
15 attachment of the binder to the toxophor has only a minor effect on the binding activity of the binder with respect to the target molecule. In a preferred embodiment, the attachment has no effect on the binding activity of the binder with respect to the target molecule.

20

In accordance with the present invention, the term "antibody" is to be understood in its broadest meaning and comprises immunoglobulin molecules, for example intact or modified monoclonal antibodies, polyclonal antibodies or multispecific  
25 antibodies (e.g. bispecific antibodies). An immunoglobulin molecule preferably comprises a molecule having four polypeptide chains, two heavy chains (H chains) and two light chains (L chains) which are typically linked by disulphide bridges. Each heavy chain comprises a variable domain of the heavy chain  
30 (abbreviated VH) and a constant domain of the heavy chain. The constant domain of the heavy chain may, for example, comprise three domains CH1, CH2 and CH3. Each light chain comprises a variable domain (abbreviated VL) and a constant domain. The constant domain of the light chain comprises a domain  
35 (abbreviated CL). The VH and VL domains may be subdivided further into regions having hypervariability, also referred to as complementarity determining regions (abbreviated CDR) and regions having low sequence variability (framework region,

abbreviated FR). Typically, each VH and VL region is composed of three CDRs and up to four FRs. For example from the amino terminus to the carboxy terminus in the following order: FR1, CDR1, FR2, CDR2, FR3, CDR3, FR4. An antibody may be obtained  
5 from any suitable species, e.g. rabbit, llama, camel, mouse or rat. In one embodiment, the antibody is of human or murine origin. An antibody may, for example, be human, humanized or chimeric.

10 The term "monoclonal" antibody refers to antibodies obtained from a population of substantially homogeneous antibodies, i.e. individual antibodies of the population are identical except for naturally occurring mutations, of which there may be a small number. Monoclonal antibodies recognize a single antigenic  
15 binding site with high specificity. The term monoclonal antibody does not refer to a particular preparation process.

The term "intact" antibody refers to antibodies comprising both an antigen-binding domain and the constant domain of the light  
20 and heavy chain. The constant domain may be a naturally occurring domain or a variant thereof having a number of modified amino acid positions.

The term "modified intact" antibody refers to intact antibodies  
25 fused via their amino terminus or carboxy terminus by means of a covalent bond (e.g. a peptide bond) with a further polypeptide or protein not originating from an antibody. Furthermore, antibodies may be modified such that, at defined positions, reactive cysteines are introduced to facilitate coupling to a  
30 toxophor (see Junutula et al. Nat Biotechnol. 2008 Aug;26(8):925-32).

The term "human" antibody refers to antibodies which can be obtained from a human or which are synthetic human antibodies.  
35 A "synthetic" human antibody is an antibody which is partially or entirely obtainable *in silico* from synthetic sequences based on the analysis of human antibody sequences. A human antibody can be encoded, for example, by a nucleic acid isolated from a

library of antibody sequences of human origin. An example of such an antibody can be found in Söderlind et al., Nature Biotech. 2000, 18:853-856.

5 The term "humanized" or "chimeric" antibody describes antibodies consisting of a non-human and a human portion of the sequence. In these antibodies, part of the sequences of the human immunoglobulin (recipient) are replaced by sequence portions of a non-human immunoglobulin (donor). In many cases, the donor is  
10 a murine immunoglobulin. In the case of humanized antibodies, amino acids of the CDR of the recipient are replaced by amino acids of the donor. Sometimes, amino acids of the framework, too, are replaced by corresponding amino acids of the donor. In some cases the humanized antibody contains amino acids present  
15 neither in the recipient nor in the donor, which were introduced during the optimization of the antibody. In the case of chimeric antibodies, the variable domains of the donor immunoglobulin are fused with the constant regions of a human antibody.

20 The term complementarity determining region (CDR) as used herein refers to those amino acids of a variable antibody domain which are required for binding to the antigen. Typically, each variable region has three CDR regions referred to as CDR1, CDR2 and CDR3. Each CDR region may embrace amino acids according to  
25 the definition of Kabat and/or amino acids of a hypervariable loop defined according to Chotia. The definition according to Kabat comprises, for example, the region from about amino acid position 24 - 34 (CDR1), 50 - 56 (CDR2) and 89 - 97 (CDR3) of the variable light chain and 31 - 35 (CDR1), 50 - 65 (CDR2) and  
30 95 - 102 (CDR3) of the variable heavy chain (Kabat et al., Sequences of Proteins of Immunological Interest, 5th Ed. Public Health Service, National Institutes of Health, Bethesda, MD. (1991)). The definition according to Chotia comprises, for example, the region from about amino acid position 26 - 32  
35 (CDR1), 50 - 52 (CDR2) and 91 - 96 (CDR3) of the variable light chain and 26 - 32 (CDR1), 53 - 55 (CDR2) and 96 - 101 (CDR3) of the variable heavy chain (Chothia and Lesk; J Mol Biol 196: 901-917 (1987)). In some cases, a CDR may comprise amino acids from

a CDR region defined according to Kabat and Chotia.

Depending on the amino acid sequence of the constant domain of the heavy chain, antibodies may be categorized into different classes. There are five main classes of intact antibodies: IgA, IgD, IgE, IgG and IgM, and several of these can be divided into further subclasses. (Isotypes), e.g. IgG1, IgG2, IgG3, IgG4, IgA1 and IgA2. The constant domains of the heavy chain, which correspond to the different classes, are referred to as [alpha/ $\alpha$ ], [delta/ $\delta$ ], [epsilon/ $\epsilon$ ], [gamma/ $\gamma$ ] and [my/ $\mu$ ]. Both the three-dimensional structure and the subunit structure of antibodies are known.

The term "functional fragment" or "antigen-binding antibody fragment" of an antibody/immunoglobulin is defined as a fragment of an antibody/immunoglobulin (e.g. the variable domains of an IgG) which still comprise the antigen binding domains of the antibody/immunoglobulin. The "antigen binding domain" of an antibody typically comprises one or more hypervariable regions of an antibody, for example the CDR, CDR2 and/or CDR3 region. However, the "framework" or "skeleton" region of an antibody may also play a role during binding of the antibody to the antigen. The framework region forms the skeleton of the CDRs. Preferably, the antigen binding domain comprises at least amino acids 4 to 103 of the variable light chain and amino acids 5 to 109 of the variable heavy chain, more preferably amino acids 3 to 107 of the variable light chain and 4 to 111 of the variable heavy chain, particularly preferably the complete variable light and heavy chains, i.e. amino acids 1 - 109 of the VL and 1 to 113 of the VH (numbering according to W097/08320).

"Functional fragments" or "antigen-binding antibody fragments" of the invention encompass, non-conclusively, Fab, Fab', F(ab')<sub>2</sub> and Fv fragments, diabodies, Single Domain Antibodies (DABs), linear antibodies, individual chains of antibodies (single-chain Fv, abbreviated to scFv); and multispecific antibodies, such as bi and tri-specific antibodies, for example, formed from antibody fragments C. A. K Borrebaeck, editor (1995) Antibody

Engineering (Breakthroughs in Molecular Biology), Oxford University Press; R. Kontermann & S. Duebel, editors (2001) Antibody Engineering (Springer Laboratory Manual), Springer Verlag. Antibodies other than "multispecific" or "multifunctional" antibodies are those having identical binding sites. Multispecific antibodies may be specific for different epitopes of an antigen or may be specific for epitopes of more than one antigen (see, for example WO 93/17715; WO 92/08802; WO 91/00360; WO 92/05793; Tutt, et al., 1991, J. Immunol. 147:60 69; U. S. Pat. Nos. 4,474,893; 4,714,681 ; 4,925,648; 5,573,920; 5,601,819; or Kostelny et al., 1992, J. Immunol. 148: 1547 1553). An F(ab')<sub>2</sub> or Fab molecule may be constructed such that the number of intermolecular disulphide interactions occurring between the Ch1 and the CL domains can be reduced or else completely prevented.

"Epitopes" refer to protein determinants capable of binding specifically to an immunoglobulin or T cell receptors. Epitopic determinants usually consist of chemically active surface groups of molecules such as amino acids or sugar side chains or combinations thereof, and usually have specific 3-dimensional structural properties and also specific charge properties.

"Functional fragments" or "antigen-binding antibody fragments" may be fused with another polypeptide or protein, not originating from an antibody, via the amino terminus or carboxyl terminus thereof, by means of a covalent bond (e.g. a peptide linkage). Furthermore, antibodies and antigen-binding fragments may be modified by introducing reactive cysteines at defined locations, in order to facilitate coupling to a toxophore (see Junutula et al. Nat Biotechnol. 2008 Aug; 26(8):925-32).

Polyclonal antibodies can be prepared by methods known to a person of ordinary skill in the art. Monoclonal antibodies may be prepared by methods known to a person of ordinary skill in the art (Köhler and Milstein, Nature, 256, 495-497, 1975). Human and humanized monoclonal antibodies may be prepared by methods known to a person of ordinary skill in the art (Olsson et al.,

Meth Enzymol. 92, 3-16 or Cabilly et al. US 4,816,567 or Boss et al. US 4,816,397).

A person of ordinary skill in the art is aware of diverse methods  
5 for preparing human antibodies and fragments thereof, such as,  
for example, by means of transgenic mice (N Lonberg and D Huszar,  
Int Rev Immunol. 1995; 13(1):65-93) or Phage Display  
Technologien (Clackson et al., Nature. 1991 Aug  
15;352(6336):624-8). Antibodies of the invention may be obtained  
10 from recombinant antibody libraries consisting for example of  
the amino acid sequences of a multiplicity of antibodies  
compiled from a large number of healthy volunteers. Antibodies  
may also be produced by means of known recombinant DNA  
technologies. The nucleic acid sequence of an antibody can be  
15 obtained by routine sequencing or is available from publically  
accessible databases.

An "isolated" antibody or binder has been purified to remove  
other constituents of the cell. Contaminating constituents of a  
20 cell which may interfere with a diagnostic or therapeutic use  
are, for example, enzymes, hormones, or other peptidic or non-  
peptidic constituents of a cell. A preferred antibody or binder  
is one which has been purified to an extent of more than 95% by  
weight, relative to the antibody or binder (determined for  
25 example by Lowry method, UV-Vis spectroscopy or by SDS capillary  
gel electrophoresis). Moreover an antibody which has been  
purified to such an extent that it is possible to determine at  
least 15 amino acids of the amino terminus or of an internal  
amino acid sequence, or which has been purified to homogeneity,  
30 the homogeneity being determined by SDS-PAGE under reducing or  
non-reducing conditions (detection may be determined by means  
of Coomassie Blau staining or preferably by silver coloration).  
However, an antibody is normally prepared by one or more  
purification steps.

35

The term "specific binding" or "binds specifically" refers to  
an antibody or binder which binds to a predetermined  
antigen/target molecule. Specific binding of an antibody or

binder typically describes an antibody or binder having an affinity of at least  $10^{-7}$  M (as Kd value; i.e. preferably those with smaller Kd values than  $10^{-7}$  M), with the antibody or binder having an at least two times higher affinity for the predetermined antigen/target molecule than for a non-specific antigen/target molecule (e.g. bovine serum albumin, or casein) which is not the predetermined antigen/target molecule or a closely related antigen/target molecule. The antibodies preferably have an affinity of at least  $10^{-7}$  M (as Kd value; in other words preferably those with smaller Kd values than  $10^{-7}$  M), preferably of at least  $10^{-8}$  M, more preferably in the range from  $10^{-9}$  M to  $10^{-11}$  M. The Kd values may be determined, for example, by means of surface plasmon resonance spectroscopy.

The antibody-drug conjugates of the invention likewise exhibit affinities in these ranges. The affinity is preferably not substantially affected by the conjugation of the drugs (in general, the affinity is reduced by less than one order of magnitude, in other words, for example, at most from  $10^{-8}$  M to  $10^{-7}$  M).

The antibodies used in accordance with the invention are also notable preferably for a high selectivity. A high selectivity exists when the antibody of the invention exhibits an affinity for the target protein which is better by a factor of at least 2, preferably by a factor of 5 or more preferably by a factor of 10, than for an independent other antigen, e.g. human serum albumin (the affinity may be determined, for example, by means of surface plasmon resonance spectroscopy).

Furthermore, the antibodies of the invention that are used are preferably cross-reactive. In order to be able to facilitate and better interpret preclinical studies, for example toxicological or activity studies (e.g. in xenograft mice), it is advantageous if the antibody used in accordance with the invention not only binds the human target protein but also binds the species target protein in the species used for the studies. In one embodiment the antibody used in accordance with the invention, in addition

to the human target protein, is cross-reactive to the target protein of at least one further species. For toxicological and activity studies it is preferred to use species of the families of rodents, dogs and non-human primates. Preferred rodent species are mouse and rat. Preferred non-human primates are  
5 rhesus monkeys, chimpanzees and long-tailed macaques.

In one embodiment the antibody used in accordance with the invention, in addition to the human target protein, is cross-  
10 reactive to the target protein of at least one further species selected from the group of species consisting of mouse, rat and long-tailed macaque (*Macaca fascicularis*). Especially preferred are antibodies used in accordance with the invention which in addition to the human target protein are at least cross-reactive  
15 to the mouse target protein. Preference is given to cross-reactive antibodies whose affinity for the target protein of the further non-human species differs by a factor of not more than 50, more particularly by a factor of not more than ten, from the affinity for the human target protein.

20

*Antibodies directed against a cancer target molecule*

The target molecule towards which the binder, for example an antibody or an antigen-binding fragment thereof, is directed is preferably a cancer target molecule. The term "cancer target molecule" describes a target molecule which is more abundantly present on one or more cancer cell species than on non-cancer cells of the same tissue type. Preferably, the cancer target molecule is selectively present on one or more cancer cell  
30 species compared with non-cancer cells of the same tissue type, where selectively describes an at least two-fold enrichment on cancer cells compared to non-cancer cells of the same tissue type (a "selective cancer target molecule"). The use of cancer target molecules allows the selective therapy of cancer cells  
35 using the conjugates according to the invention.

Antibodies which are specific against an antigen, for example cancer cell antigen, can be prepared by a person of ordinary



skill in the art by means of methods with which he or she is familiar (such as recombinant expression, for example) or may be acquired commercially (as for example from Merck KGaA, Germany). Examples of known commercially available antibodies in cancer therapy are Erbitux® (cetuximab, Merck KGaA), Avastin® (bevacizumab, Roche) and Herceptin® (trastuzumab, Genentech). Trastuzumab is a recombinant humanized monoclonal antibody of the IgG1kappa type which in a cell-based assay (Kd = 5 nM) binds the extracellular domains of the human epidermal growth receptor with high affinity. The antibody is produced recombinantly in CHO cells.

In a preferred embodiment, the target molecule is a selective cancer target molecule.

In a particularly preferred embodiment, the target molecule is a protein.

In one embodiment, the target molecule is an extracellular target molecule. In a preferred embodiment, the extracellular target molecule is a protein.

Cancer target molecules are known to those skilled in the art. Examples of these are listed below.

Examples of cancer target molecules are:

(1) EGF receptor (NCBI reference sequence NP\_005219.2), SEQ ID NO: 213 (1210 amino acids):

>gi|29725609|ref|NP\_005219.2| EGFR receptor precursor [Homo sapiens]

MRPSGTAGAALLALLAALCPASRALEEKKVCQGTSNKLTQLGTFEDHFLSLQRMFNNCEVVL  
GNLEITYVQARNYDLSFLKTIQEVAGYVLIALNTVERIPLLENLQIIRGNMYYENSYALAVLSN  
YDANKTGLKELPMRNLQEILHGAVRFSNNPALCNVESIQWRDIVSSDFLSNMSMDFQNHLS  
CQKCDPSCPNGSCWGAGEENCQKLTKIICAQQCSGRCRGKSPSDCCHNQCAAGCTGPRESDC  
LVCRKFRDEATCKDTCPPMLLYNPTTYQMDVNPEGKYSFGATCVKKCPRNYVVTDHGSCVRA

CGADSYEMEEEDGVRKCKKCEGPCRKVCNGIGIGEFKDSLSINATNIKHFKNCTSISGDLHIL  
PVAFRGDSFTHTPPLDPQELDILKTVKEITGFLLIQAWPENRTDLHAFENLEIIRGRTKQHG  
QFSLAVVSLNITSLGLRSLKEISDGDVIISGNKNLCYANTINWKKLFGTSGQTKIISNRGE  
NSCKATGQVCHALCSPEGCWGPEPRDCVSCRNVSRGRECVDKCNLLEGEPREFVENSECIOC  
5 HPECLPQAMNITCTGRGPDNCIQCAHYIDGPHCVKTCPAGVMGENNTLVWKYADAGHVCHLC  
HPNCTYGCTGPGLEGCP TNGPKI PSIATGMVGALLLLLVVALGIGL FMRRRHIVRKRTLRL  
LQERELVEPLTPSGEAPNQALLRILKETEFKKIKVLGSGAFGTVYKGLWIPEGEKVKIPVAI  
KELREATSPKANKEILDEAYVMASVDNPHVCRLLGICLTSTVQLITQLMPFGCLLDYVREHK  
DNIGSQYLLNWCVQIAKGMNYLEDRLVHRDLAARNVLVKTPQHVKITDFGLAKLLGAEKE  
10 YHAEGGKVPKWMALESILHRIYTHQSDVWSYGVTWELMTFGSKPYDGIPASEISSILEKG  
ERLPQPPICTIDVYMIMVKCWMIDADSRPKFRELIIEFSKMARDPQRYLVIQGDERMHLPS  
TDSNFYRALMDEEDMDDVVDADAYLIPOQGGFFSSPSTSRTPLLSSLSATSNNSTVACIDRNG  
LQSCPIKEDSFLQRYSSDPTGALTEDSIDDTFLPVPEYINQSVPKRPAGSVQNPVYHNQPLN  
PAPSRDPHYQDPHSTAVGNPEYLNTVQPTCVNSTFDSPAHWAQKGS HQISLDNPDYQQDFFP  
15 KEAKPNGIFKGSTAENAEYLRVAPQSSEFIGA

The extracellular domain is marked by underlining.

(2) mesothelin (SwissProt reference Q13421-3), SEQ ID NO: 214  
20 (622 amino acids):

>sp|Q13421-3|MSLN\_HUMAN isoform 2 of mesothelin OS=Homo sapiens  
GN=MSLN

25 MALPTARPLLGSCGTPALGSLFLFSLGWVQPSRTLGETGQEAAPLDGVLANPPNISS  
LSRQLLGFPCAEVSGLSTERVRELAVALAQKNVKLSTEQLRCLAHRLSEPPEDLDALPL  
DLLLFLNPDAFSGPQACTRFFSRITKANVDLLPRGAPERQRLLPAAALACWVGRGSLSEA  
30 DVRALGGLACDLPGRFVAESA EVLLPRLVSCPGPLDQDQQAARAALQGGGPPYGPSTW  
SVSTMDALRGLLPVLGQPIIRSIPQGIVAAWRQSSRDPSWRQPERTILRPRFRREVEKT  
35 ACPSGKKAREIDESLI FYKKWELEACVDAALLATQMDRVNAIPFTYEQLDVLKHKLDELY  
PQGYPESVIQHLGYLFLKMSPEDIRKWNVTSLETLKALLEVNKGHEMSPQVATLIDRFVK

GRGQLDKDTLDTLTAFYPGYLCSLSPHEELSSVPPSSIWAVRPQDLDTCDPRQLDVLYPKA

RLAFQNMNGSEYFVKIQSFLGGAPTEDLKALSQQNVSMDLATFMKLRTDAVLPLTVAEVQ

5 KLLGPHVEGLKAEERHRPVRDWILRQRQDDLDTLGLGLQGGIPNGYLVLDLSMQEALSGT

PCLLGPGPVLTVLALLLASTLA

10 where mesothelin is encoded by amino acids 296-598. Amino acids  
37-286 are coding for the megakaryocyte-potentiating factor.  
Mesothelin is anchored in the cell membrane via a GPI anchor and  
is localized extracellularly.

15 (3) carboanhydrase IX (SwissProt reference Q16790), SEQ ID NO:  
215 (459 amino acids):

>sp|Q16790|CAH9\_HUMAN carbonic anhydrase 9 OS=Homo sapiens  
GN=CA9 PE=1 SV=2

20 MAPLCPSPWLPLLI PAPAPGLTVQLLLSLLLLVPVHPQRLPRMQEDSPLGGGSSGEDDPL

GEEDLPSEEDSPREEDPPGEEDLPGEEDLPGEEDLPEVKPKSEEEGSLKLEDLPTVEAPG

DPQEPQNNNAHRDKEGDDQSHWRYGGDPPWPRVSPACAGRFQSPVDIRPQLAAFCPALRPL

25

ELLGFQLPPLPELRLRNNGHSVQLTLPPGLEMALGPGREYRALQLHLHWGAAGRPGSEHT

VEGHRFPAEIHVVHLSTAFARVDEALGRPGGLAVLAAFLEEGPEENSAYEQLLSRLEEIA

30 EEGSETQVPGLDISALLPSDFSRYFQYEGSLTTPPCAQGVIWTVFNQTVMLSAKQLHTLS

DTLWGP GDSRLQLNFRATQPLNGRVIEASFPAGVDSSPRAAEPVQLNSCLAAGDILALVF

GLLFAVTSVAFLVQMRRQHRRGTKGGVSYRPAEVAETGA

35

The extracellular domain is marked by underlining.

(4) C4.4a (NCBI reference sequence NP\_055215.2; synonym LYPD3),

SEQ ID NO: 216 (346 amino acids):

>gi|93004088|ref|NP\_055215.2| ly6/PLAUR domain-containing  
protein 3-precursor [Homo sapiens]

5

MDPARKAGAQAMIWTAGWLLLLLLLRGGAQALECYSCVQKADDGCSPNKMKTVKCAPGVDVCT  
EAVGAVETIHGQFSLAVRGCSGSLPGKNDRGLDLHGLLAFIQLQOCAQDRCNAKLNLTSRAL  
DPAGNESAYPPNGVECYSCVGLSREACQGTSPPPVSCYNASDHVYKGCFDGNVTLTAANVTV  
SLPVRGCVQDEFCTRDGVTGPGFTLSGSCCQGSRCNSDLRNKTYFSPRIPPLVRLPPPEPTT  
10 VASTTSVTTSTSAPVRPTSTTKPMPAPTSQTPRQVEHEASRDEEPRLTGGAAGHQDRSNSG  
QYPAKGGPQQPHNKGCVAPTAGLAALLLAVAAGVLL

The mature extracellular domain is marked by underlining.

15 (5) CD52 (NCBI reference sequence NP\_001794.2 ), SEQ ID NO: 217

>gi|68342030|ref|NP\_001794.2| CAMPATH-1 antigen-precursor [Homo  
sapiens]

20 MKRFLFLLLTISLLVMVQIQTGLSGQNDTSQTS SPSASSNISGGIFLFFVANAI IHLFCFS

(6) Her2 (NCBI reference sequence NP\_004439.2), SEQ ID NO: 218

>gi|54792096|ref|NP\_004439.2| receptor tyrosine-protein kinase  
25 erbB-2 isoform a [Homo sapiens]

MELAALCRWGLLLALLPPGAASTQVCTGTDMLRRLPASPETHLDMLRHLYQGCQVVQGNLEL  
TYLPTNASLSFLQDIQEVQGYVLI AHNQVRQVPLQRLRIVRG TQLFEDNYALAVLDNGDPLN  
NTTPVTGASPGGLRELQLRSLTEILKGGVLIQRNPQLCYQDTILWKDIFHKNNQLALTLIDT  
30 NRSRACHPCSPMCKGSRGWGESSEDCQSLTRTV CAGGCARCKGPLPTDCCHEQCAAGCTGPK  
HSDCLACLHFNHSGICELHCPALVTYNTDTFESMPNPEG RYTFGASCVTACPYNYLSTDVGS  
CTLVCPLHNQEVTAE DGTQRCEKCSKPCARVCYGLGMEHLREVR AVTSANIQE FAGCKKIFG  
SLAFLPESFDGDPASNTAPLQPEQLQVFETLEEITGYLYISA WPDSLPLDLSVFQNLQVIRGR  
ILHNGAYSLTLQGLGISWLGLRSLRELGSGLALIH HNTHLCFVHTVPWDQLFRNPHQALLHT  
35 ANRPEDECVGEG LACHQLCARGHCWGPPTQCVNCSQFLRGQECV EECRVLQGLPREYVNAR  
HCLPCHPECQPQNGSVTCFGPEADQCVACAHYKDPPFCVARCPSGVKPDLSYMPIWKFPDEE  
GACQPCPINCTHSCVDLDDKGC PAEQRASPLTSIISAVV GILLVVVLGVVFGILIKRRQQKI  
RKYTMRLLQETELVEPLTPSGAMPNQAQMRILKETELRKVKVLGSGAFGTVYKGIWIPDGE

NVKIPVAIKVLRENTSPKANKEILDEAYVMAGVGSPLYVSRLLGICLTSTVQLVTQLMPYGCL  
 LDHVRENRRGLGSQDLLNWCMIKAGMSYLEDVRLVHRDLAARNVLVKSPNHVKITDFGLAR  
 LLDIDETEYHADGGKVPIKWMALESILRRRFTHQSDVWSYGVTVWELMTFGAKPYDGI PARE  
 IPDLLEKGERLPQPPICTIDVYMIMVKCWMIDSECRPRFRELVSEFSRMARDPQRFVVIQNE  
 5 DLGPASPLDSTFYRSLLEDDDMGDLVDAEEYLVPQQGFFCPDPAPGAGGMVHHRHRSSSTRS  
 GGGDLTLGLEPSEEEAPRSPLAPSEGAGSDVFDGDLGMGAAKGLQSLPTHDPSPLOQRYSEDP  
 TVPLPSETDGYVAPLTCSPQPEYVNQPDVRPQPPSPREGPLPAARPAGATLERPKTLPSPGKN  
 GVVKDVFAFGGAVENPEYLTPOGGAAPQPHPPAFSPAFDNLYYWDQDPPERGAPPSTFKGT  
 PTAENPEYLGLDVPV

10

(7) CD20 (NCBI reference sequence NP\_068769.2), SEQ ID NO: 219

>gi|23110987|ref|NP\_068769.2| B-lymphocyte antigen CD20 [Homo  
 sapiens]

15

MTTPRNSVNGTFPAEPMKGP IAMQSGPKPLFRMSSLVGPTQSFFMRESKTLGAVQIMNGLF  
 HIALGGLLMIPAGIYAPICVTVWYPLWGGIMYIISGSLLAATEKNSRKCLVKGKMIMNSLSL  
 FAAISGMILSIMDILNIKISHFLKMESLNFIRAHTPYINIYNCEPANPSEKNSPSTQYCYSI  
 QSLFLGILSVMLIFAFFQELVIAGIVENEWKRTCSRPKSNIVLLSAEEKKEQTIEIKEEVVG  
 20 LTETSSQPKNEEDIEIPIQEEEEETETNFPEPPQDQESSPIENDSSP

(8) the lymphocyte activation antigen CD30 (SwissProt ID  
 P28908), SEQ ID NO: 220

25 >gi|68348711|ref|NP\_001234.2| tumor necrosis factor receptor  
 superfamily member 8 isoform 1-precursor [Homo sapiens]

MRVLLAALGLLFLGALRAFPQDRPFEDTCHGNPSHYDKAVRRCCYRCMPGLFPTQQCPQRP  
 TDCRKQCEPDYYLDEADRCTACVTCSRDDLVEKTPCAWNSSRVCECRPGMFCSTSAVNSCAR  
 30 CFFHSVCPAGMIVKFPGTAQKNTVCEPASPGVSPACASPENCKEPSSGTIPQAKPTPVSPAT  
 SSASTMPVRGGTRLAQEAASKLTRAPDSPSSVGRPSSDPGLSPTQPCPEGSGDCRKQCEPDY  
 YLDEAGRCTACVCSRDDLVEKTPCAWNSSRTCECRPGMICATSATNSRARCVPYPICAAET  
 VTKPQDMAEKDTTFFAPPLGTQPCNPTPENGEAPASTSPTQSLLVDSQASKTLPIPTSAPV  
 ALSSTGKPVLDAGPVLFVWVILVLVVVVGSSAFLLCHRRACRKRIRQKLHLCYPVQTSQPKLE  
 35 LVDSRPRRSSTQLRSGASVTEPVAEERGLMSQPLMETCHSVGAAYLESPLQDASPAGGPSS  
 PRDLPEPRVSTEHTNKKIEKIYIMKADTVIVGTVKAELPEGRGLAGPAEPELEEELEADHTP  
 HYPEQETEPPLGSCSDVMLSVEEEGKEDPLPTAASGK

(9) the lymphocyte adhesion molecule CD22 (SwissProt ID **P20273**),  
SEQ ID NO: 221

>gi|157168355|ref|NP\_001762.2| B-cell receptor CD22 isoform 1-  
5 precursor [Homo sapiens]

MHLLGPWLLLLLVLEYLAFSDSSKWVFEHPETLYAWEGACVWIPCTYRALDGDLESFILFHNP  
EYNKNTSKFDGTRLYESTKDGKVPSEQKRVQFLGDKNKNCTLSIHPVHLNDSGQLGLRMESK  
TEKWMERIHLNVSERPFPPHIQLPPEIQESQEVTLTCLLNFSYGYPIQLQWLLEGVPMRQA  
10 AVTSTSLTIKSVFTRSELKFSPQWSHHGKIVTCQLQDADGKFLSNDTVQLNVKHTPKLEIKV  
TPSDAIVREGDSVTMTCEVSSSNPEYTTVSWLKDGTSLKKQNTFTLNLREVTKDQSGKYCCQ  
VSNDVGPGRSEEVFLQVQYAPEPSTVQILHSPAVEGSQVEFLCMSLANPLPTNYTWYHNGKE  
MQGRTEEKVHIPKILPWHAGTYSCVAENILGTGQRGPGAELDVQYPPKKVTTVIQNPMPIRE  
GDTVTLSCNYNSSNPSVTRYEWKPHGAWEEPSLGVLKIQNVGWDNTTIACAACNSWCSP  
15 VALNVQYAPRDVRVRKIKPLSEIHSGNSVSLQCDFSSSHPKEVQFFWEKNGRLLGKESQLNF  
DSISPEDAGSYSCWVNSIGQTASKAWTLEVLYAPRRLRVSMSPGDQVMGKSATLTCESDA  
NPPVSHYTWFDWNNQSLPYHSQKLRLEPVKVQHSGAYWCQGTNSVGKGRSPLSTLTVYYSPE  
TIGRRVAVGLGSCLAAILILAICGLKLQRRWKRTQSQQGLQENSSGQSFVVRNKKVRRAPLSE  
GPHSLGCYNPMMEDGISYTTLRFPENIIPRTGDAESSEMQRPPDCDDTVTYSALHKRQVGD  
20 YENVIPDFPEDEGIHYSELIQFGVGERPQAQENVVDYVILKH

(10) the myloid cell surface antigen CD33 (SwissProt ID **P20138**),  
SEQ ID NO: 222

25 >gi|130979981|ref|NP\_001763.3| myeloid cell surface antigen  
CD33 isoform 1-precursor [Homo sapiens]

MPLLLLLLPLLWAGALAMDPNFWLQVQESVTVQEGLCVLPCTFFHPIPYDKNSPVHGYWFR  
EGAIISRDSPVATNKLDQEVQEEETQGRFRLLGDPSRNNCSLSIVDARRRDNGSYFFRMERGS  
30 TKYSYKSPQLSVHVTDLTHRPKILIPGTLEPGHKNLTCSVSWACEQGTPPIFSWLSAAPT  
LGPRTTHSSVLIITPRPQDHGTNLTCQVKFAGAGVTTERTIQLNVTYVPQNPTTGIFFPGDGS  
GKQETRAGVVHGAIGGAGVTALLALCLCLIFFIVKTHRRKAARTAVGRNDTHPTTGSASPKH  
QKKSKLHGPTETSSCSGAAPTVEEMDEELHYASLNFHGMNPSKDTSTEYSEVRTQ

35 (11) the transmembrane glycoprotein NMB (SwissProt ID **Q14956**),  
SEQ ID NO: 223

>gi|52694752|ref|NP\_001005340.1| transmembrane glycoprotein NMB

isoform a-precursor [Homo sapiens]

MECLYYFLGFLLLAARLPLDAAKRFHDVLGNERPSAYMREHNQLNGWSSDENDWNEKLYPVW  
 KRGD MRWKN SWK GGRVQAVLTSDSPALVGSNITFAVNLI FPRCQKEDANGNIVYEKNCRNEA  
 5 GLSADPYVYNWTAWSESDGNGTGQSHHNVFPDGKPFPHHPGWRRWNFIYVFHTLGQYFQK  
 LGRCSVRVSVNTANVT LGPQLMEVTVYRRHGRAYVPIAQVKDVYVVTDQIPVFVTMFQKNDR  
 NSSDETFLKDLPI MF DVL IHDPSHFLNYSTINYKWSFGDNTGLFVSTNHTVNHTYVLNGTFS  
 LNLT VKAAAPGPCPPPPPPRPSKPTPSLATTLKSYDSNTPGPAGDNPLELSRIPDENCQIN  
 RYGHFQATITIVEGILEVNI IQMTDVLMPVPWPESLIDFVVT CQGSIPTEVCTIISDPTCE  
 10 ITQNTVCS PVDVDEMCLLTVRRTFNGSGTYCVNLT LGDDTSLALTSTLISVPDRDPASPLRM  
 ANSALISVGCLAI FVTVISLLVYKKHKEYNPIENSPGNVVR SKGLSVFLNRAKAVFFPGNQE  
 KDPLLKNQEFKGV S

(12) the adhesion molecule CD56 (SwissProt ID P13591), SEQ ID  
 15 NO: 224

>gi|94420689|ref|NP\_000606.3| neural cell adhesion molecule 1  
 isoform 1 [Homo sapiens]

20 MLQTKDLIWTLFFLGTAVSLQVDIVPSQGEISVGESKFFLCQVAGDAKDKDISWFSPNGEKL  
 TPNQQRISVWVNDSSSTLTIYNANIDDAGIYKCVVTGEDGSESEATVNVKIFQKLMFKNAP  
 TPQEFREGEDAVIVCDVSSLPPTIIWKHKGRDVILKKDVR FIVLSNNYLQIRGIKKTDEGT  
 YRCEGRILARGEINFKDIQVIVNVPPTIQARQNI VNATANLGQSVTLVCD AEGFPEPTMSWT  
 KDGEQIEQEEDDEKYIFSDDSSQLTIKKVDKNDEAEYICIAENKAGEQDATIHLKVFAPKI  
 25 TYVENQTAMELEEQVTLTCEASGDPIPSITWRTSTRNISSEEKTLDGHMVVRSHARVSSLTL  
 KSIQYTDAGEYICTASNTIGQDSQSMYLEVQYAPKLQGPVAVYTWEGN

QVNITCEVFAYPSATISWFRDGQLLPSSNYSNIKIYNTPSASYLEVTPDSEDFGNYNCTAV  
 NRIGQESLEFILVQADTPSSPSIDQVEPYSSTAQVQFDEPEATGGVPILKYKAEWRAVGEEV  
 30 WSKWYDAKEASMEGIVTIVGLKPETTYAVRLAALNGKGLGEISAASEFKTQPVQGEPSAPK  
 LEGQMGEDGNSIKVNLIKQDDGGSPIRHVLRVYRALSSEWKPEIRLPSGSDHVMLKSLDWNA  
 EYEVYVVAENQQGKSKAAHFVFR TSAQPTAI PANGSPTSGLSTGAI VGILIVIFVLLL VVVD  
 ITCYFLNKCGLFMCI AVNLCGKAGPGAKGKDMEEGKAAFSKDESKEPIVEVRTEEERTPNHD  
 GGKHTEPNETTPLTEPEKGPVEAKPECQETETKPAPAEVKTVPNDATQTKENESKA

35

(13) the surface molecule CD70 (SwissProt ID P32970), SEQ ID NO:  
 225

>gi|4507605|ref|NP\_001243.1| CD70 antigen [Homo sapiens]

MPEEGSGCSVRRRPYGCVLRAALVPLVAGLVICLVVCIQRFAQAQQQLPLESLGWDVAELQL  
 NHTGPQQDPRLYWQGGPALGRSFLHGPELDKGQLRIHRDGIYMVHIQVTLAICSSTTASRHH  
 5 PTTLAVGICSPASRSISLLRLSFHQGCTIASQRLTPLARGDTLCTNLTGTLTLLPSRNTDETFE  
 GVQWVRP

(14) the surface molecule CD74 (SwissProt ID P04233), SEQ ID NO:  
 226

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>gi|10835071|ref|NP\_004346.1| HLA class II histocompatibility  
 antigen gamma chain isoform b [Homo sapiens]

MHRRRSRSCREDQKPVMDQRDQLISNNEQLPMLGRRPGAPESKCSR GALYTGFSILVTL LLA  
 15 GQATTAYFLYQQQGR LDKLTVTSQNLQLENLRMKLPKPPKPVSKMRMATPLLMQALPMGALP  
 QGPMQNATKYGNMTE DHVMHLLQNADPLKVYPPLKGSFPENLRHLKNTMETIDWKVFESWMH  
 HWLLFEMSRHSLEQKPTDAPPKESLELEDPS SGLGVTKQDLGPVPM

(15) the B-lymphocyte antigen CD19 (SwissProt ID P15391), SEQ  
 20 ID NO: 227

>gi|296010921|ref|NP\_001171569.1| B-lymphocyte antigen CD19  
 isoform 1-precursor [Homo sapiens]

MPPPRLLFFLLFLTPMEVRPEEPLVVKVEEGDNAVLQCLKGTSDGPTQQLTWSRESPLKPFLL  
 KLSLGLPGLGIHMRPLAIWLFIFNVSQQMGGFYLCQPGPPSEKAWQPGWTVNVEGSGELFRW  
 NVSDLGGLGCGLKNRSSEGPSSPSGKLMSPKLYVWAKDRPEIWEGEPPCLPPRDSL NQSL SQ  
 DLTMAPGSTLWLS CGVPPDSVSRGPLSWTHVHPKGP KSLLSLELKDDRPARDMWV METGLLL  
 PRATAQDAGKYYCHRG NLTMSFHLEITARPVLWHWLLRTGGWKVSAVTLAYLIFCLCSLVGI  
 30 LHLQRALVLRKRKRMTDPTRRFFKVT PPPGSGPQNQYGNVLSLPTPT SGLGRAQRWAAGLG  
 GTAPSYGNPSSDVQADGALGSRSPPGVGP EEEEEGEGYE EEPDSEEDSEFYENDSNL GQDQL SQ  
 DGSGYENPEDEPLGPEDEDSFSNAESYENEDEELTQP VARTMDFLSPHGSAWDPSREATSLA  
 GSQSYEDMRGILYAAPQLRSIRGQPGPNHEEDADSYENMDNPDGPDPAWGGGGGRMG TWSTR

(16) the surface protein mucin-1 (SwissProt ID P15941), SEQ ID  
 35 NO: 228

>gi|65301117|ref|NP\_002447.4| mucin-1 isoform 1-precursor [Homo



sapiens]

MTPGTQSPFFLLLLLTVLTVVTGSGHASSTPGGEKETSATQRSSVPSSTEKNALSTGVSFFF  
 LSFHISNLQFNSSLEDPSTDYYQELQRDISEMFLQIYKQGGFLGLSNIKFRPGSVVVQLTLA  
 5 FREGTINVHDTVETQFNQYKTEAASRYNLTISDVSVDVPPFSAQSGAGVPGWGIALLVLC  
 VLVALAIVYLIALAVCQCRRKNYGQLDIFPARDTYHPMSEYPTYHTHGRYVPPSSTDRSPYE  
 KVSAGNGGSSLSYTNPAVAATSANL

(17) the surface protein CD138 (SwissProt ID P18827), SEQ ID NO:  
 10 229

>gi|29568086|ref|NP\_002988.3| syndecan-1-precursor [Homo  
 sapiens]

15 MRRAALWLWLCALALSLOPALPQIVATNLPPEDQDGSDDSDNFSGSGAGALQDITLSQQTP  
 STWKDTQLLTAIPTSPEPTGLEATAASTSTLPAGEGPKGEAVVLPEVEPGLTAREQEATPR  
 PRETTQLPTTHQASTTTATTAQEPATSHPHRDMQPGHHETSTPAGPSQADLHTPHTEDGGPS  
 ATERAAEDGASSQLPAAEGSGEQDFTFETSGENTAVVAVEPDRRNQSPVDQGATGASQGLLD  
 RKEVLGGVIAGGLVGLIFAVCLVGFMLYRMKKKDEGSYSLEEPKQANGGAYQKPTKQEEFYA

20

(18) the integrin alphaV (Genbank Accession No.: NP\_002201.1),  
 SEQ ID NO: 230

>gi|4504763|ref|NP\_002201.1| integrin alpha-V isoform 1-  
 25 precursor [Homo sapiens]

MAFPPRRRLRLGPRGLPLLLSGLLLPLCRAFNLVDSPA EYSGPEGSYFGFAVDFVPSASS  
 RMFLLVGAPKANTTQPGIVEGGQVLKCDWSSTRRCQPIEFDATGNRDYAKDDPLEFKSHQWF  
 GASVRSKQDKILACAPLYHWRTEMKQEREPVGT CFLQDGTKTVEYAPCRSQDIDADGQGFCQ  
 30 GGFSIDFTKADRVLLGGPGSFYWQGQLISDQVAEIVSKYDPNVYSIKYNNQLATRQAIFD  
 DSYLGYSVAVGDFNGDGIDDFVSGVPRAARTLGMVYIYDGKNMSSLYNFTGEQMAAYFGFSV  
 AATDINGDDYADVFIGAPLFMDRGS DGKLQEVGQVSVSLQRASGDFQTTKLNGFEVFARFGS  
 AIAPLGDLQDGFNDIAIAAPYGGEDKKGIVYIFNGRSTGLNAVPSQILEGQWAARSMPPSF  
 GYSMKGATDIDKNGYPDLIVGAFGVDRAILYRARPVITVNAGLEVYPSILNQDNKTCSLPGT  
 35 ALKVSCFNVRFLKADGKGVLPKLNQVVELLLDKLKQKGAIRRALFLYSRSPSHSKNMTIS  
 RGGLMQCEELIAYLRDESEFRDKLTPITIFMEYRLDYRTAADTTGLQPILNQFTPANISRQA  
 HILLDCGEDNVCKPKLEVSVSDQKKIYIGDDNPLTLIVKAQNQGE GAYEAELIVSIPLQAD  
 FIGVVRNNEALARLSCAFKTENQTRQVVC DLGNPMKAGTQLLAGLRF SVHQQSEMDTSVKFD

LQIQSSNLFDKVSPPVSHKVDLAVLAAVEIRGVSSPDHIFLPIPNWEHKENPETEEDVGPVV  
 QHIYELRNNGPSSFSKAMLHLQWPYKYNNNTLLYILHYDIDGPMNCTSDMEINPLRIKISSL  
 QTTEKNDTVAGQGERDHLITKRDLALSEGDIHTLGCGVAQCLKIVCQVGRDRGKSAILYVK  
 SLLWTETFMNKENQNHSYSLKSSASFNVIEFPYKNLPIEDITNSTLVTTNVTWGIQPAPMPV  
 5 PVWVIILAVLAGLLLLLAVLVFVMYRMGFFKRVRPPQEEQEREQLQPHENGEENSET

(19) the teratocarcinoma-derived growth factor 1 protein TDGF1  
 (Genbank Accession No.: NP\_003203.1), SEQ ID NO: 231

10 >gi|4507425|ref|NP\_003203.1| teratocarcinoma-derived growth  
 factor 1 isoform 1-precursor [Homo sapiens]

MDCRKMARFSYSVIWIMAIKVFELGLVAGLGHQEFARPSRGYLAFRDDSIWPQEEPAIRPR  
 SSQRVPPMGIQHSKELNRTCCLNGGTCMLGSFCACPPSFYGRNCEHDVRKENCGSVPHTWL  
 15 PKKCSLCKCWHGQLRCFPQAFLPGCDGLVMDEHLVASRTPPELPPSARTTTFMLVGICLSIQS  
 YY

(20) the prostate-specific membrane antigen PSMA (Swiss Prot ID:  
 Q04609), SEQ ID NO: 232

20 >gi|4758398|ref|NP\_004467.1| glutamate carboxypeptidase 2  
 isoform 1 [Homo sapiens]

MWNLLHETDSAVATARRPRWLCAGALVLAGGFLLGFLFGWFIKSSNEATNITPKHNMKAFL  
 25 DELKAENIKKFLYNFTQIPHLAGTEQNFQLAKQIQSQWKEFGLDSVELAHYDVLLSYPNKTH  
 PNYISIINEDGNEIFNTSLFEPPIPGYENVSDIVPPFSAFSPQGMPEGDLVYVNYARTEDFF  
 KLERDMKINCSGKIVIARYGKVFRGNKVKNAQLAGAKGVILYSDPADYFAPGVKSYPDGWNL  
 PGGGVQRGNILNLNGAGDPLTPGYPANEYAYRRGIAEAVGLPSIPVHPIGYYDAQKLLKMG  
 GSAPPDSSWRGSLKVPYNVGPFTGNFSTQKVKMHIHSTNEVTRIYNVIGTLRGAVEPDRYV  
 30 ILGGHRDSWVFGGIDPQSGAAVVHEIVRSFGTLKKEGWRPRRTILFASWDAEEFGLLGSTEW  
 AEENSRLQERGVAYINADSSIEGNYTLRVDCTPLMYSLVHNLTKELKSPDEGFEGKSLYES  
 WTKKSPSPEFSGMPRISKLGSGNDFEVFFQRLGIASGRARYTKNWETNKFSGYPLYHSVYET  
 YELVEKFYDPMFKYHLTVAQVRGGMVFELANSIVLPFDCRDYAVVLRKYADKIYSISMKHPQ  
 EMKTYSVSFDLSLFAVKNFTEIASKFSERLQDFDKSNPIVLRMMNDQLMFLERAFIDPLGLP  
 35 DRPFYRHVIYAPSSHNKYAGESFPGIYDALFDIESKVDPSKAWGEVQRQIYVAAFTVQAAAE  
 TLSEVA

(21) the tyrosine protein kinase EPHA2 (Swiss Prot ID: P29317),

SEQ ID NO: 233

>gi|32967311|ref|NP\_004422.2| ephrin type-A receptor 2-precursor [Homo sapiens]

5

MELQAARACFALLWGCALAAAAAQQGKEVLLDFAAAGGELGWLTHPYGKGWDLMQNIMNDM  
 PIYMYSVCNVMSGDQDNWLRTNWVYRGEAERIFIELKFTVRDCNSFPGGASSCKETFNLYYA  
 ESDLDYGTNFQKRLFTKIDTIAPDEITVSSDFEARHVKLNVEERSVGPLTRKGFYLAQDIG  
 ACVALLSVRVYYKKCPPELLQGLAHFPETIAGSDAPSLATVAGTCVDHAVVPPGGEEPMMHCA  
 10 VDGEWLVPVIGQCLCQAGYEKVEDACQACSPGFFKFEASESPCLECPEHTLPSPEGATSCECE  
 EGFFRAPQDPASMPCTRPPSAPHYLTAVGMGAKVELRWTPPQDSGGREDIVYSVTCEQCWPE  
 SGECGPCEASVRYSEPPHGLTRTSVTVSDLEPHMNYTFTVEARNGVSGLVTSRSFRTASVSI  
 NQTEPPKVRLEGRSTTSLSVSWSIPPPQQSRVWKYEVTYRKKGDSNSYNVRRTEGFSVTLDD  
 LAPDTTYLVQVQALTQEGQGAGSKVHEFQTLSPEGSGNLAVIGGVAVGVVLLLVLAVGVGFFI  
 15 HRRRKNQRARQSPEDVYFSKSEQLKPLKTYVDPHTYEDPNQAVLKFTTEIHPSCVTRQKVI  
 AGEFGEVYKGMKLTSSGKKEVPVAIKTLKAGYTEKQRVDFLGEAGIMGQFSHHNIIRLEGVI  
 SKYKPMMIITEYMENGALDKFLREKDGESVLQLVGMLRGIAAGMKYLANMNYVHRDLAARN  
 ILVNSNLVCKVSDFGLSRVLEDDPEATYTTSGGKIPIRWTAPEAISYRKFTSASDVWSFGIV  
 MWEVMTYGERPYWELSNHEVMKAINDFRLPTPMDCPSAIYQLMMQCWQQRARRPKFADIV  
 20 SILDKLIRAPDSLKTLADFDPRVSIRLPSTSGSEGVPFRTVSEWLESIKMQQYTEHFMAAGY  
 TAIEKVVQMTNDDIKRIGVRLPGHQKRIAYSLLGLKDQVNTVGIPI

(22) the surface protein SLC44A4 (Genbank Accession No: NP\_001171515), SEQ ID NO: 234

25

>gi|295849282|ref|NP\_001171515.1| choline transporter-like protein 4 isoform 2 [Homo sapiens]

MGGKQRDEDDEAYGKPKYDPSFRGPIKNRSCTDVICCVLFLLFILGYIVVGIVAWLYGDPR  
 30 QVLYPRNSTGAYCGMGENKDKPYLLYFNIFSCILSSNIISVAENGLQCPTPQTVITSLQQEL  
 CPSFLLPSAPALGRCFPWTVNTPPALPGITNDTTIQQGISGLIDSLNARDISVKIFEDFAQS  
 WYWILVALGVALVLSLLFILLRLVAGPLVLVLIIGVLGVLAYGIYYCWEEYRVLDRDKGAS I  
 SQLGFTTNLSAYQSVQETWLAALIVLAVLEAILLLMLIFLRQRIRIAIALKEASKAVGQMM  
 STMFYPLVTFVLLLICIAYWAMTALYLATSGQPQYVLWASNISSPGCEKVPINTSCNPTAHL  
 35 VNSSCPGLMCFVQGYSSKGLIQRSVFNLQIYGVVGLFWTLNWWLALGQCVLAGAFASFYWAF  
 HKPQDIPTFPLISAFIRTLRYHTGSLAFGALILTLVQIARVILEYIDHKLRGVQNPVARCIM  
 CCFKCCLWCLEKFIKFLNRNAYIMIAIYGKNFCVSAKNAFMLLMRNIVRVVLDKVTDLLEF  
 FGKLLVGGVGVLSFFFFSGRIPGLGKDFKSPHLNYYWLPIMTSILGAYVIASGFFSVFGMC

VDTLFLCFLEDLERNNGSLDRPYYSKSLKILGKKNEAPPDNKKRKK

- (23) the surface protein BMPR1B (SwissProt: O00238)
- 5 (24) the transport protein SLC7A5 (SwissProt: Q01650)
- (25) the epithelial prostate antigen STEAP1 (SwissProt: Q9UHE8)
- (26) the ovarian carcinoma antigen MUC16 (SwissProt: Q8WXI7)
- 10 (27) the transport protein SLC34A2 (SwissProt: O95436)
- (28) the surface protein SEMA5b (SwissProt: Q9P283)
- 15 (29) the surface protein LYPD1 (SwissProt: Q8N2G4)
- (30) the endothelin receptor type B EDNRB (SwissProt: P24530)
- (31) the ring finger protein RNF43 (SwissProt: Q68DV7)
- 20 (32) the prostate carcinoma-associated protein STEAP2  
(SwissProt: Q8NFT2)
- (33) the cation channel TRPM4 (SwissProt: Q8TD43)
- 25 (34) the complement receptor CD21 (SwissProt: P20023)
- (35) the B-cell antigen receptor complex-associated protein  
CD79b (SwissProt: P40259)
- 30 (36) the cell adhesion antigen CEACAM6 (SwissProt: P40199)
- (37) the dipeptidase DPEP1 (SwissProt: P16444)
- 35 (38) the interleukin receptor IL20Ralpha (SwissProt: Q9UHF4)
- (39) the proteoglycan BCAN (SwissProt: Q96GW7)

- (40) the ephrin receptor EPHB2 (SwissProt: P29323)
- (41) the prostate stem cell-associated protein PSCA (Genbank Accession No: NP\_005663.2 )
- 5 (42) the surface protein LHFPL3 (SwissProt: Q86UP9)
- (43) the receptor protein TNFRSF13C (SwissProt: Q96RJ3)
- 10 (44) the B-cell antigen receptor complex-associated protein CD79a (SwissProt: P11912)
- (45) the receptor protein CXCR5 (SwissProt: P32302)
- 15 (46) the ion channel P2X5 (SwissProt: Q93086)
- (47) the lymphocyte antigen CD180 (SwissProt: Q99467)
- (48) the receptor protein FCRL1 (SwissProt: Q96LA6)
- 20 (49) the receptor protein FCRL5 (SwissProt: Q96RD9)
- (50) the MHC class II molecule Ia antigen HLA-DOB (Genbank Accession No: NP\_002111.1)
- 25 (51) the T-cell protein VTCN1 (SwissProt: Q7Z7D3)
- (52) TWEAKR (SEQ ID NO:169 (protein); SEQ ID NO:170 (DNA) .
- 30 (53) the lymphocyte antigen CD37 (Swiss Prot: P11049)
- (54) the FGF receptor 2; FGFR2 (Gene ID: 2263; official symbol: FGFR2). The FGFR2 receptor occurs in different splice variants (alpha, beta, IIIb, IIIc). All splice variants may act as target
- 35 molecule.
- (55) the transmembrane glycoprotein B7H3 (CD276; Gene ID: 80381.

(56) the B cell receptor BAFFR (CD268; Gene ID: 115650)

(57) the receptor protein ROR 1 (Gene ID: 4919)

5 (58) the surface receptor IL3RA (CD123; Gene ID: 3561)

(59) the CXC chemokine receptor CXCR5 (CD185; Gene ID 643)

(60) the receptor protein syncytin ( Gene ID 30816)

10

In a preferred subject matter of the invention, the cancer target molecule is selected from the group consisting of the cancer target molecules (1) - (60), in particular (1), (6) and (52).

15 In a further particularly preferred subject matter of the invention, the binder binds to an extracellular cancer target molecule which is selected from the group consisting of the cancer target molecules (1) - (60), in particular (1), (6) and (52).

20

In a further particularly preferred subject matter of the invention, the binder binds specifically to an extracellular cancer target molecule which is selected from the group consisting of the cancer target molecules (1) - (60), in particular (1), (6) and (52). In a preferred embodiment the binder is, after binding to its extracellular target molecule on the target cell, internalized by the target cell as a result of the binding. This causes the binder/active compound conjugate, which may be an immunoconjugate or an ADC, to be taken up by the target cell. The binder is then processed, preferably intracellularly, with preference lysosomally.

25

In one embodiment the binder is a binding protein. In a preferred embodiment the binder is an antibody, an antigen-binding antibody fragment, a multispecific antibody or an antibody mimetic.

35

Preferred antibody mimetics are affibodies, adnectins,

anticalins, DARPins, avimers, or nanobodies. Preferred multispecific antibodies are bispecific and trispecific antibodies.

5 In a preferred embodiment the binder is an antibody or an antigen-binding antibody fragment, more preferably an isolated antibody or an isolated antigen-binding antibody fragment.

Preferred antigen-binding antibody fragments are Fab, Fab',  
10 F(ab')<sub>2</sub> and Fv fragments, diabodies, DABs, linear antibodies and scFv. Particularly preferred are Fab, diabodies and scFv.

In a particularly preferred embodiment the binder is an antibody. Particularly preferred are monoclonal antibodies or  
15 antigen-binding antibody fragments thereof. Further particularly preferred are human, humanized or chimeric antibodies or antigen-binding antibody fragments thereof.

Antibodies or antigen-binding antibody fragments which bind  
20 cancer target molecules may be prepared by a person of ordinary skill in the art using known processes, such as, for example, chemical synthesis or recombinant expression. Binders for cancer target molecules may be acquired commercially or may be prepared by a person of ordinary skill in the art using known processes,  
25 such as, for example, chemical synthesis or recombinant expression. Further processes for preparing antibodies or antigen-binding antibody fragments are described in WO 2007/070538 (see page 22 "Antibodies"). The person skilled in the art knows how processes such as phage display libraries  
30 (e.g. Morphosys HuCAL Gold) can be compiled and used for discovering antibodies or antigen-binding antibody fragments (see WO 2007/070538, page 24 ff and AK Example 1 on page 70, AK Example 2 on page 72). Further processes for preparing antibodies that use DNA libraries from B cells are described for  
35 example on page 26 (WO 2007/070538). Processes for humanizing antibodies are described on page 30-32 of WO2007070538 and in detail in Queen, et al., Pros. Natl. Acad. Sci. USA 86:10029-10033, 1989 or in WO 90/0786. Furthermore, processes for the

recombinant expression of proteins in general and of antibodies in particular are known to the person skilled in the art (see, for example, in Berger and Kimrnel (Guide to Molecular Cloning Techniques, Methods in Enzymology, Vol. 152, Academic Press, Inc.); Sambrook, et al., (Molecular Cloning: A Laboratory Manual, (Second Edition, Cold Spring Harbor Laboratory Press; Cold Spring Harbor, N.Y.; 1989) Vol. 1-3); Current Protocols in Molecular Biology, (F. M. Ausabel et al. [Eds.], Current Protocols, Green Publishing Associates, Inc. / John Wiley & Sons, Inc.); Harlow et al., (Monoclonal Antibodies: A Laboratory Manual, Cold Spring Harbor Laboratory Press (1988, Paul [Ed.])); Fundamental Immunology, (Lippincott Williams & Wilkins (1998)); and Harlow, et al., (Using Antibodies: A Laboratory Manual, Cold Spring Harbor Laboratory Press (1998)). The person skilled in the art knows the corresponding vectors, promoters and signal peptides which are necessary for the expression of a protein/antibody. Commonplace processes are also described in WO 2007/070538 on pages 41-45. Processes for preparing an IgG1 antibody are described for example in WO 2007/070538 in Example 6 on page 74 ff. Processes which allow the determination of the internalization of an antibody after binding to its antigen are known to the skilled person and are described for example in WO 2007/070538 on page 80. The person skilled in the art is able to use the processes described in WO 2007/070538 that have been used for preparing carboanhydrase IX (Mn) antibodies in analogy for the preparation of antibodies with different target molecule specificity.

#### anti-EGFR antibodies

Examples of antibodies which bind the cancer target molecules EGFR are cetuximab (INN number 7906), panitumumab (INN number 8499) and nimotuzumab (INN number 8545). Cetuximab (Drug Bank Accession Number DB00002) is a chimeric anti-EGFR1 antibody which is produced in SP2/0 mouse myeloma cells and is sold by ImClone Systems Inc/Merck KgaA/Bristol-Myers Squibb Co. Cetuximab is indicated for the treatment of metastasizing, EGFR expressing, colorectal carcinoma with wild type K-Ras gene. It



has an affinity of  $10^{-10}$ M.

Sequence:

5 Cetuximab Light Chain (kappa), SEQ ID NO: 235:

DILLTQSPVILSVSPGERVSFSCRASQSIGTNIHWYQQRTNGSPRLLIKYASESISGIPSRF  
 SGSGSGTDFTLSINSVESEDIADYYCQQNNNWPTTFGAGTKLELKRTVAAPSVFIFPPSDEQ  
 LKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDESTYSLSSSTLTLSKADY  
 10 EKHKVYACEVTHQGLSSPVTKSFNRGEC

Cetuximab Heavy Chain, SEQ ID NO: 236:

QVQLKQSGPGLVQPSQSLITCTVSGFSLTNYGVHWVRQSPGKGLEWLGVIWSSGGNTDYNTF  
 15 FTSRLSINKDNSKSQVFFKMNSLQSNDAIYYCARALTYDYEFAYWGQGLVTVSAASTKG  
 PSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYSLSS  
 VVTVPSSSLGTQTYICNVNHKPSNTKVDKRVKPKSCDKTHTCPPCPAPELLGGPSVFLFPPK  
 PKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTV  
 LHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVK  
 20 GFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVDFSCSVMHEAL  
 HNHYTQKSLSLSPGK

Panitumumab (INN number 8499) (Drug Bank Accession Number  
 DB01269) is a recombinant monoclonal human IgG2 antibody which  
 25 binds specifically to the human EGF receptor 1 and is sold by  
 Abgenix/Amgen. Panitumumab originates from the immunization of  
 transgenic mice (XenoMouse). These mice are capable of producing  
 human immunoglobulin (light and heavy chains). A specific B-cell  
 clone was selected which produces antibodies against EGFR, and  
 30 this clone was immortalized with CHO cells (Chinese hamster  
 ovary cells). These cells are now used for the production of a  
 100% human antibody. Panitumumab is indicated for the treatment  
 of EGFR-expressing, metastasizing colorectal carcinoma, which  
 is resistant to chemotherapeutic treatment with  
 35 fluoropyrimidine, oxaliplatin and irinotecan. It has an affinity  
 of  $10^{-11}$ M.

Sequence:

**Panitumumab** Light Chain (kappa), SEQ ID NO: 237:

DIQMTQSPSSLSASVGDRVTITCQASQDISNYLNWYQQKPGKAPKLLIYDASNLETGVPSRF  
 5 SGSGSGTDFTFTISSLQPEDIATYFCQHFDHLPLAFGGGKVEIKRTVAAPSVFIFPPSDEQ  
 LKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADY  
 EKHKVYACEVTHQGLSSPVTKSFNRGEC

**Panitumumab** Heavy Chain, SEQ ID NO: 238:

10 QVQLQESGPGLVKPSETLSLTCTVSGGSVSSGDYYWTWIRQSPGKGLEWIGHIYYSGNTNYN  
 PSLKSRLTISIDTSKTQFSLKLSSVTAADTAIYYCVRDRVTGAFDIWGQGMVTVSSASTKG  
 PSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYSLSS  
 VVTVPSNFGTQTYTCNVDPKPKTSTKVDKTKVERKCCVECPPEPPVAGPSVFLFPPKPKDT  
 15 LMISRTPEVTCVVVDVSHEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTFRVVSVLTVVHQD  
 WLNGKEYKCKVSNKGLPAPIEKTISKTKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFYP  
 SDIAVEWESNGQPENNYKTTPMLDSDGSFFLYSKLTVDKSRWQQGNVVFSCSVMHEALHNHY  
 TQKSLSLSPG

20 Nimotuzumab (INN number 8545) (EP 00586002, EP 00712863) is a  
 humanized monoclonal IgG1 antibody which binds specifically to  
 the human EGF receptor 1 and is sold by YM BioScienecs Inc.  
 (Mississauga Canada). It is produced in non-secreting NSO cells  
 (mammalian cell line). Nimotuzumab is approved for the treatment  
 25 of head-and-neck tumours, highly malignant astrocytoma and  
 glioblastoma multiforms (not in EU and US) and pancreatic  
 carcinoma (Orphan drug, EMA). It has an affinity of  $10^{-8}$  M.

**Nimotuzumab** Light Chain, SEQ ID NO: 239:

30 DIQMTQSPSSLSASVGDRVTITCRSSQNIVHSNGNTYLDWYQQTPGKAPKLLIYKVSNRFSG  
 VPSRFSGSGSGTDFTFTISSLQPEDIATYYCFQYSHVPWTFGQGTKLQITRTVAAPSVFIFP  
 PSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTLT  
 SKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC

35

**Nimotuzumab** Heavy Chain, SEQ ID NO: 240:

QVQLQQSGAEVKKPGSSVKVSCKASGYTFTNYYIYWVRQAPGQGLEWIGGINPTS GGSNFNE

KFKTRVTITADESSTTAYMELSSLRSEDYAFYFCTRQGLWFDS DGRGFDFWGQTTVTVSSA  
 STKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLY  
 SLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVFL  
 FPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVS  
 5 VLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQVSLT  
 CLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVDFSCSVM  
 HEALHNHYTQKSLSLSPGK

Further embodiments of EGFR antibodies are as follows:

10

- Zalutumumab / 2F8 / HuMax-EGFr, from Genmab A/S (WO 02/100348, WO 2004/056847, INN number 8605)

15

- Necitumumab / 11F8, ImClone / IMC-11F8, from ImClone Systems Inc. [Eli Lilly & Co] (WO 2005/090407 (EP 01735348-A1, US 2007/0264253-A1, US 7,598,350, WO 2005/090407-A1), INN number 9083)

20

- Matuzumab / anti-EGFR MAb, Merck KGaA / anti-EGFR MAb, Takeda / EMD 72000 / EMD-6200 / EMD-72000 and EMD-55900 / MAb 425 / monoclonal antibody 425, from Merck KGaA / Takeda ( WO 92/15683, INN number 8103 (Matuzumab))

25

- RG-7160 / GA-201 / GA201 / R-7160 / R7160 / RG7160 / RO-4858696 / RO-5083945 / RO4858696 / RO5083945, from Glycart Biotechnology AG (Roche Holding AG) (WO 2010/112413-A1, WO 2010/115554)

30

- GT-MAB 5.2-GEX / CetuGEX, from Glycotope GmbH (WO 2008/028686-A2 (EP 01900750-A1, EP 01911766-A1, EP 02073842-A2, US 2010/0028947-A1)

35

- ABT-806 / mAb-806 / ch-806 / anti-EGFR monoclonal antibody 806, from Ludwig Institute for Cancer Research / Abbott / Life Science Pharmaceuticals (WO 02/092771, WO 2005/081854 and WO

2009/023265)

- SYM-004 (consists of two chimeric IgG1 antibodies (992 and 1024)), from Symphogen A/S (WO 2010/022736-A2)
- 5
- MR1-1 /MR1-1KDEL, from IVAX Corp (Teva Pharmaceutical Industries Ltd) (Duke University), (patent: WO2001/062931-A2)
  - Antibody against the deletion mutant, EGFRvIII, from Amgen/Abgenix (WO 2005/010151, US 7,628,986)
- 10
- SC-100, from Scancell Ltd (WO 01/088138-A1)
  - MDX-447 / EMD 82633 / BAB-447 / H 447 / MAb, EGFR, Medarex/Merck KGaA, from Bristol-Myers Squibb (US) / Merck KGaA (DE) / Takeda (JP), (WO 91/05871, WO 92/15683)
  - anti-EGFR-Mab, from Xencor (WO 2005/056606)
- 15
- DXL-1218 / anti-EGFR monoclonal antibody (cancer), InNexus, from InNexus Biotechnology Inc, Pharmaprojects PH048638
- 20

In a preferred embodiment, the anti-EGFR antibodies are selected from the group consisting of cetuximab, panitumumab, nimotuzumab, zalutumumab, necitumumab, matuzumab, RG-716, GT-MAB 5.2-GEX, ISU-101, ABT-806, SYM-004, MR1-1, SC-100, MDX-447 and DXL-1218.

25

In a particularly preferred embodiment the anti-EGFR antibodies are selected from the group consisting of cetuximab, panitumumab, nimotuzumab, zalutumumab, necitumumab and matuzumab.

30

The person skilled in the art knows of processes which can be used to prepare further antibodies, from the CDR regions of the abovementioned antibodies by means of sequence variations, these further antibodies having a similar or better affinity and/or specificity for the target molecule.

35

In a further embodiment, the anti-EGFR antibodies or antigen-binding antibody fragments are selected from the group consisting of antibodies or antigen-binding antibody fragments comprising the three CDR regions of the light chain and the three CDR regions of the heavy chain of one of the following antibodies: cetuximab, panitumumab, nimotuzumab, zalutumumab, necitumumab, matuzumab, RG-716, GT-MAB 5.2-GEX, ISU-101, ABT-806, SYM-004, MR1-1, SC-100, MDX-447 and DXL-1218.

10

In a further embodiment, the anti-EGFR antibodies or antigen-binding antibody fragments are selected from the group consisting of antibodies or antigen-binding antibody fragments comprising three CDR regions of the light chain and the three CDR regions of the heavy chain of one of the following antibodies: cetuximab, panitumumab, nimotuzumab, zalutumumab, necitumumab, matuzumab. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

20

anti-Carboanhydrase IX antibodies

Examples of antibodies which bind the cancer target molecule carbonahydrase IX are described in WO 2007/070538-A2 (e.g. Claims 1 - 16).

25

In a preferred embodiment the anti-carboanhydrase IX antibodies or antigen-binding antibody fragments are selected from the group consisting of anti-carboanhydrase IX antibodies or antigen-binding antibody fragments 3ee9 (Claim 4 (a) in WO 2007/070538-A2), 3ef2 (Claim 4 (b) in WO2007/070538-A2), 1e4 (Claim 4 (c) in WO 2007/070538-A2), 3a4 (Claim 4 (d) in WO 2007/070538-A2), 3ab4 (Claim 4 (e) in WO 2007/070538-A2), 3ah10 (Claim 4 (f) in WO 2007/070538-A2), 3bb2 (Claim 4 (g) in WO 2007/070538-A2), 1aa1 (Claim 4 (h) in WO 2007/070538-A2), 5a6 (Claim 4 (i) in WO 2007/070538-A2) and 5aa3 (Claim 4 (j) in WO 2007/070538-A2).

35

anti-C4.4a antibodies:

According to the invention, use may be made of C4.4a antibodies.

5 Examples of C4.4a antibodies and antigen-binding fragments are described in WO 2012/143499 A2. By reference, all antibodies of WO 2012/143499 A2 are hereby incorporated into the description of the present invention, and they can be used in the present invention. The sequences of the antibodies are given in Table 1  
10 of WO 2012/143499 A2, where each row shows the respective CDR amino acid sequences of the variable light chain or the variable heavy chain of the antibody listed in column 1.

In one embodiment, the anti-C4.4a antibodies or antigen-binding  
15 antibody fragments thereof are, after binding to a cell expressing C4.4a, internalized by the cell.

In a further embodiment, the anti-C4.4a antibodies or antigen-binding antibody fragments comprise at least one, two or three  
20 CDR amino acid sequences of an antibody listed in Table 1 of WO 2012/143499 A2 or Table 2 of WO 2012/143499 A2. Preferred embodiments of such antibodies are likewise listed in WO 2012/143499 A2 and incorporated herein by reference.

25 anti-HER2 antibodies:

An example of an antibody binding to the cancer target molecule Her2 is trastuzumab (Genentech). Trastuzumab is a humanized antibody used *inter alia* for the treatment of breast cancer.

30 Further examples of antibodies binding to HER2 are, in addition to trastuzumab (INN 7637, CAS No.: RN: 180288-69-1) and Pertuzumab (CAS No.: 380610-27-5), the antibodies disclosed in WO 2009/123894-A2, WO 200/8140603-A2 or in WO 2011/044368-A2.  
35 An example of an anti-HER2 conjugate is trastuzumab-emtansine (INN-No. 9295). By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

anti-CD20 antibodies:

An example of an antibody binding to the cancer target molecule  
5 CD20 is rituximab (Genentech). Rituximab (CAS Number: 174722-  
31-7) is a chimeric antibody used for the treatment of non-  
Hodgkin lymphoma. By reference, these antibodies and antigen-  
binding fragments thereof are incorporated herein, and they can  
be used in the context of the present invention.

10

anti-CD52 antibodies:

An example of an antibody binding to the cancer target molecule  
CD52 is alemtuzumab (Genzyme). Alemtuzumab (CAS Number: 216503-  
15 57-0) is a humanized antibody used for the treatment of chronic  
lymphocytic leukaemia. By reference, these antibodies and  
antigen-binding fragments thereof are incorporated herein, and  
they can be used in the context of the present invention.

20 anti-Mesothelin antibodies:

Examples of anti-mesothelin antibodies are described, for  
example, in WO 2009/068204. By reference, all antibodies  
described in WO 2009/068204 are hereby incorporated into the  
25 present description, such that these antibodies can be used in  
the context of the invention disclosed herein.

The anti-mesothelin antibodies used in accordance with the  
invention are also notable preferably for an invariant binding  
30 to mesothelin. Invariant binding is characterized, for example,  
in that the antibody used in accordance with the invention binds  
to an epitope of mesothelin which cannot be masked by a further  
extracellular protein. Such a further extracellular protein is,  
for example, the protein ovarian cancer antigen 125 (CA125).  
35 Antibodies which are used with preference are characterized in  
that their binding to mesothelin is not blocked by CA125.

anti-CD30 antibodies

Examples of antibodies which bind the cancer target molecule CD30 and can be used for the treatment of cancer, for example Hodgkin lymphoma, are brentuximab, iratumumab and antibodies disclosed in WO 2008/092117, WO 2008/036688 or WO 2006/089232. An example of an anti-CD30 conjugate is brentuximab vedotin (INN No. 9144). By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

10

anti-CD22 antibodies

Examples of antibodies which bind the cancer target molecule CD22 and can be used for the treatment of cancer, for example lymphoma, are inotuzumab and epratuzumab. Examples of anti-CD22 conjugates are inotuzumab ozagamycin (INN No. 8574) or anti-CD22-MMAE and anti-CD22-MC-MMAE (CAS RN: 139504-50-0 and 474645-27-7, respectively). By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

15  
20

anti-CD33 antibodies

Examples of antibodies which bind the cancer target molecule CD33 and can be used for the treatment of cancer, for example leukaemia, are gemtuzumab and lintuzumab (INN 7580). An example of an anti-CD33 conjugate is gemtuzumab-ozagamycin. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

25  
30

anti-NMB antibodies

An example of an antibody which binds the cancer target molecule NMB and can be used for the treatment of cancer, for example melanoma or breast cancer, is glembatumumab (INN 9199). An example of an anti-NMB conjugate is glembatumumab vedotin (CAS RN: 474645-27-7). By reference, these antibodies and antigen-

35



binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

Anti-CD56 antibodies

5

An example of an antibody which binds the cancer target molecule CD56 and can be used for the treatment of cancer, for example multiple myeloma, small-cell lung carcinoma, MCC or ovarial carcinoma is lorvotuzumab. An example of an anti-CD56 conjugate  
10 is lorvotuzumab mertansine (CAS RN: 139504-50-0). By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

15 anti-CD70 antibodies

Examples of antibodies which bind the cancer target molecule CD70 and can be used for the treatment of cancer, for example non-Hodgkin lymphoma or renal cell cancer, are disclosed in WO  
20 2007/038637-A2 and WO 2008/070593-A2. An example of an anti-CD70 conjugate is SGN-75 (CD70 MMAF). By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

25 anti-CD74 antibodies

An example of an antibody which binds the cancer target molecule CD74 and can be used for the treatment of cancer, for example multiple myeloma, is milatuzumab. An example of an anti-CD74  
30 conjugate is milatuzumab-doxorubicin (CAS RN: 23214-92-8). By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

35 anti-CD19 antibodies

An example of an antibody which binds the cancer target molecule CD19 and can be used for the treatment of cancer, for example

non-Hodgkin lymphoma, is disclosed in WO 2008/031056-A2. Further antibodies and examples of an anti-CD19 conjugate (SAR3419) are disclosed in WO 2008/047242-A2. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein,  
5 and they can be used in the context of the present invention.

anti-Mucin antibodies

Examples of antibodies which bind the cancer target molecule  
10 mucin-1 and can be used for the treatment of cancer, for example non-Hodgkin lymphoma, are clivatuzumab and the antibodies disclosed in WO 2003/106495-A2, WO 2008/028686-A2. Examples of anti-mucin conjugates are disclosed in WO 2005/009369-A2. By reference, these antibodies and antigen-binding fragments  
15 thereof are incorporated herein, and they can be used in the context of the present invention.

anti-CD138 antibodies

20 Examples of antibodies which bind the cancer target molecule CD138 and conjugates thereof, which can be used for the treatment of cancer, for example multiple myeloma, are disclosed in WO 2009/080829-A1, WO 2009/080830-A1. By reference, these antibodies and antigen-binding fragments thereof are  
25 incorporated herein, and they can be used in the context of the present invention.

anti-Integrin-alphaV antibodies

30 Examples of antibodies which bind the cancer target molecule integrin alphaV and can be used for the treatment of cancer, for example melanoma, sarcoma or carcinoma, are intetumumab (CAS RN: 725735-28-4), abciximab (CAS RN: 143653-53-6), etaracizumab (CAS RN: 892553-42-3) and the antibodies disclosed in US 7,465,449,  
35 EP 719859-A1, WO 2002/012501-A1 and WO2006/062779-A2. Examples of anti-integrin AlphaV conjugates are intetumumab-DM4 and other ADCs disclosed in WO 2007/024536-A2. By reference, these antibodies and antigen-binding fragments thereof are

incorporated herein, and they can be used in the context of the present invention.

anti-TDGF1 antibodies

5

Examples of antibodies which bind the cancer target molecule TDGF1 and can be used for the treatment of cancer are the antibodies disclosed in WO 02/077033-A1, US 7,318,924, WO 2003/083041-A2 and WO 2002/088170-A2. Examples of anti-TDGF1  
10 conjugates are disclosed in WO 2002/088170-A2. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

15 anti-PSMA antibodies

Examples of antibodies which bind the cancer target molecule PSMA and can be used for the treatment of cancer, for example prostate carcinoma, are the antibodies disclosed in WO 97/35616-  
20 A1, WO 99/47554-A1, WO 01/009192-A1 and WO2003/034903. Examples of anti-PSMA conjugates are disclosed in WO 2009/026274-A1 and WO 2007/002222. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

25

anti-EPHA2 antibodies

Examples of antibodies which bind the cancer target molecule EPHA2 and can be used for preparing a conjugate and for the  
30 treatment of cancer are disclosed in WO 2004/091375-A2. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the context of the present invention.

35 anti-SLC44A4 antibodies

Examples of antibodies which bind the cancer target molecule SLC44A4 and can be used for preparing a conjugate and for the

treatment of cancer, for example pancreas or prostate carcinoma, are disclosed in WO2009/033094-A2 and US2009/0175796-A1. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the  
5 context of the present invention.

anti-HLA-DOB antibodies

An example of an antibody binding to the cancer target molecule  
10 HLA-DOB is the antibody Lym-1 (CAS RN: 301344-99-0) which can be used for the treatment of cancer, for example non-Hodgkin lymphoma. Examples of anti-HLA-DOB conjugates are disclosed, for example, in WO 2005/081711-A2. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein,  
15 and they can be used in the context of the present invention.

anti-VTCN1 antibodies

Examples of antibodies which bind the cancer target molecule  
20 VTCN1 and can be used for preparing a conjugate and for the treatment of cancer, for example ovarian carcinoma, pancreas, lung or breast cancer, are disclosed in WO 2006/074418-A2. By reference, these antibodies and antigen-binding fragments thereof are incorporated herein, and they can be used in the  
25 context of the present invention.

anti-FGFR2 antibodies

According to the invention, use may be made of anti-FGFR2  
30 antibodies.

Examples of anti-FGFR2 antibodies and antigen-binding fragments are described in WO2013076186. By reference, all antibodies of WO2013076186 are hereby incorporated into the description of the  
35 present invention, and they can be used in the present invention. The sequences of the antibodies are shown in Table 9 and Table 10 of WO2013076186. Preference is given to antibodies, antigen-binding fragments and variants of the antibodies derived from

the antibodies referred to as M048-D01 and M047-D08. Preferred anti-FGFR2 bind to the various splice variants known of FGFR2.

In one embodiment, the anti-FGFR2 antibodies or antigen-binding antibody fragments thereof are, after binding to a cell  
5 expressing FGFR2, internalized by the cell.

In a further embodiment, the anti-FGFR2 antibodies or antigen-binding antibody fragments comprise at least one, two or three  
10 CDR amino acid sequences of an antibody listed in Table 9 or Table 10 of WO2013076186. Preferred embodiments of such antibodies are likewise listed in WO2013076186 and incorporated herein by reference.

15 Anti-TWEAKR antibodies

In a preferred embodiment, when an anti-TWEAKR antibody or an antigen-binding fragment thereof is used in the processes according to the present invention, this antibody or fragment  
20 is selected from those described below. In addition, antibodies which bind to TWEAKR are known to the person skilled in the art, see, for example, WO2009/020933(A2) or WO2009140177 (A2).

The invention relates in particular to conjugates with  
25 antibodies or antigen-binding antibody fragments thereof or variants thereof which lead to strong activation of the TWEAKR (SEQ ID NO:169 (protein); SEQ ID NO:170 (DNA)), resulting in a strong induction of apoptosis in various cancer cells overexpressing TWEAKR.

30 The agonistic activity of TWEAKR with regard to the induction of apoptosis and inhibition of the proliferation of the anti-TWEAKR antibodies already described (e.g. PDL-192) is limited and does not reach the efficacy of the endogenous ligand TWEAK.  
35 This lack of agonistic activity is not based on reduced affinity, since these antibodies bind at the TWEAKR with affinities which, compared to the endogenous ligand TWEAK, are in a similar range (Michaelson JS et al, MAbs. 2011 Jul-Aug;3(4):362-75; Culp PA

et al, Clin Cancer Res. 2010 Jan 15;16(2):497-508), and even antibodies having a higher binding affinity do not necessarily display a more effective signalling activity (Culp PA, et al, Clin Cancer Res. 2010 Jan 15;16(2):497-508). In addition, it has  
5 been shown that the antitumour activity of the antibodies already described depends on the Fc effector function, and it was shown that ADCC plays an important role for the in-vivo efficacy in mouse models.

#### 10 Generation of the anti-TWEAKR antibodies

A complete human antibody phage library (Hoet RM et al, Nat Biotechnol 2005;23(3):344-8) was employed to isolate TWEAKR-specific human monoclonal antibodies of the present invention  
15 by protein panning (Hoogenboom H.R., Nat Biotechnol 2005;23(3):1105-16) using dimeric Fc-fused extracellular domains of human and mouse TWEAKR as immobilized target. 11 different Fab phages were identified, and the corresponding antibodies were cloned into a mammalian EgG expression vector  
20 which provides the CH2-CH3 domains missing in the soluble FAb. Following identification of preferred antibodies, these were expressed as full-length IgGs. These constructs were expressed, for example, transiently in mammalian cells as described by Tom et al., Chapter 12 in Methods Express: Expression Systems edited  
25 by Micheal R. Dyson and Yves Durocher, Scion Publishing Ltd, 2007 (see AK-Example 1). The antibodies were purified by protein-A chromatography and characterized further by their binding affinity to soluble monomeric TWEAKR using ELISA and BIAcore analysis, as described in AK-Example 2. To determine the  
30 cell binding characteristics of the anti-TWEAKR antibodies, binding was tested by flow cytometry on a number of cell lines (HT29, HS68, HS578). NFκB reporter gene assays were carried out to examine the agonistic activity of all 11 antibodies identified (human IgG1). The antibody having the highest in  
35 vitro activity (TPP-883) was selected for further activity and affinity maturation (see AK-Example 1 for details). A single substitution variant having improved agonistic activity was detected: G102T of CDR-H3. In the end, 7 variants were selected

based on increased affinity compared to the best single substitution variant G102T. The corresponding DNA thereof was cloned into a mammalian IgG expression vector and examined for functional activity in the NF-kappaB reporter gene assay mentioned above. Finally, the sequences obtained were compared with human germ line sequences, and deviations without any significant effect on the affinity and the efficacy were adapted. The following antibodies were obtained by antibody library screening and by affinity and/or activity maturation. "TPP-2090", "TPP-2149", "TPP-2093", "TPP-2148", "TPP-2084", "TPP-2077", "TPP-1538", "TPP-883", "TPP-1854", "TPP-1853", "TPP-1857" and "TPP-1858".

Antibodies of the invention can furthermore be obtained by methods known in the art such as antibody phage display screening (see, for example, Hoet RM et al., Nat Biotechnol 2005;23(3):344-8), the well-established hybridoma technology (see, for example, Köhler and Milstein Nature. 1975 Aug 7;256(5517):495-7) or immunization of mice, *inter alia* immunization of hMAb mice (e.g. VelocImmune mouse®).

Particular embodiments of anti-TWEAKR antibodies

One embodiment of the invention is the provision of antibodies or antigen-binding antibody fragments thereof or variants thereof showing strong induction of caspase 3/7 in one or more TWEAKR-expressing cell lines. In a preferred embodiment, the one or more TWEAKR-expressing cell line(s) is/are present in the group consisting of WiDr, A253, NCI-H322, HT29 and 786-O. "Induction of caspase 3/7" can be measured by customary methods known in the art, including those described herein. In one embodiment, the "induction of caspase 3/7" is determined in accordance with the present invention using the activity determination with capase 3/7 solution (Promega, #G8093) and reading the luminescence on a VICTOR V (Perkin Elmer). At the end of the incubation time, the caspase 3/7 activity was determined and the induction factor of caspase 3/7 was determined in comparison to untreated cells. An antibody is said

to show "strong induction" of caspase 3/7 when the induction factor is greater than 1.2, preferably greater than 1.5, even more preferably greater than 1.8, even more preferably greater than 2.1, even more preferably greater than 2.5. What is provided are anti-TWEAKR antibodies leading to stronger induction of caspase 3/7 in HT29 cells compared to agonistic antibodies already described [e.g. PDL-192(TPP-1104), P4A8(TPP-1324), 136.1(TPP-2194)] and also compared to 300 ng/ml recombinant human TWEAK. This strong activity of inducing caspase 3/7 in cancer cells was also observed in WiDr, A253, NIC-H322 and 786-O cells where in most experiments the antibodies of the invention examined induced higher factors of change compared to the reference antibodies [PDL-192(TPP-1104), P4A8(TPP-1324)] and to 300 ng/ml TWEAK. Some antibodies of the invention bind to the TWEAKR only with moderate affinity (>10 nM) which is clearly less than the affinity of the endogenous ligand TWEAK, and also less compared to other known agonistic antibodies. This property offers further possible advantages such as, for example, potentially deeper penetration into the tumour.

In this regard, one embodiment of the invention is the provision of antibodies or antigen-binding antibody fragments thereof binding specifically to a TWEAKR at a novel epitope characterized by selective binding to aspartate (D) at position 47 (D47) of TWEAKR (SEQ ID NO:169; see also Figure 1). The dependencies identified for certain TWEAKR amino acids for antibody interaction correlate with the agonistic activity determined for these antibodies. The native ligand TWEAK shows an effective activation of the TWEAKR and binds depending on leucine 46 in the cysteine-rich domain of TWEAKR (Pellegrini et al, FEBS 280:1818-1829). P4A8 displays a very low agonistic activity and interacts at least partially with domains outside of the cysteine-rich domain of TWEAKR. PDL-192 displays a moderate agonistic activity and binds depending on R56 to the cysteine-rich domain, but opposite the TWEAK ligand site. Antibodies of the present invention (e.g. TPP-2090) bind depending on D47, and TWEAK binds depending on L46. Thus, TWEAK binds to a similar but different binding site (Figure 7).



Accordingly, the antibodies of the present invention displaying strong agonistic activity bind to a novel epitope (D47-dependent) for antibodies associated with very high agonistic activity.

5

The amino acid at position 47 (D47) of TWEAKR (SEQ ID NO:169) is considered to be critical for binding of the antibodies according to the invention, which means that the antibody binds specifically to the D at position 47 (D47) of TWEAKR (SEQ ID  
10 NO:169) when the antibody loses more than 20%, alternatively more than 30%, alternatively more than 40%, alternatively more than 50%, alternatively more than 60%, alternatively more than 70%, alternatively more than 80%, alternatively more than 90%, alternatively 100% of its ELISA signal by modification of this  
15 residue into alanine, as described in AK-Example 2 and Figure 6. Alternatively, an antibody binds specifically to the D at position 47 (D47) of TWEAKR (SEQ ID NO:169) when the antibody loses more than 20%, alternatively more than 30%, alternatively more than 40%, alternatively more than 50%, alternatively more  
20 than 60%, alternatively more than 70%, alternatively more than 80%, alternatively more than 90%, alternatively 100% of its ELISA signal for TPP-2614 compared to TPP-2203. Preferably, an antibody binds specifically to the D at position 47 (D47) of TWEAKR (SEQ ID NO:169) when the antibody loses more than 80% of  
25 its ELISA signal for TPP-2614 compared to TPP-2203.

In the present application, reference is made to the following preferred antibodies of the invention, as shown in the table below: "TPP-2090", "TPP-2149", "TPP-2093", "TPP-2148", "TPP-  
30 2084", "TPP-2077", "TPP-1538", "TPP-883", "TPP-1854", "TPP-1853", "TPP-1857", "TPP-1858".

**Table:** Protein sequences of the antibodies:

	SEQ ID NO: IgG1 light chain	SEQ ID NO: IgG1 heavy chain	SEQ ID NO: L-CDR1	SEQ ID NO: L-CDR2	SEQ ID NO: L-CDR3	SEQ ID NO: H-CDR1	SEQ ID NO: H-CDR2	SEQ ID NO: H-CDR3	SEQ ID NO: VL protein	SEQ ID NO: VH protein
Anti-TWEAKR antibodies according to the invention:										
TPP-2090	1	2	3	4	5	6	7	8	9	10
TPP-2149	11	12	13	14	15	16	17	18	19	20
TPP-2093	21	22	23	24	25	26	27	28	29	30
TPP-2148	31	32	33	34	35	36	37	38	39	40
TPP-2084	41	42	43	44	45	46	47	48	49	50
TPP-2077	51	52	53	54	55	56	57	58	59	60
TPP-1538	61	62	63	64	65	66	67	68	69	70
TPP-883	71	72	73	74	75	76	77	78	79	80
TPP-1854	81	82	83	84	85	86	87	88	89	90
TPP-1853	91	92	93	94	95	96	97	98	99	100
TPP-1857	101	102	103	104	105	106	107	108	109	110
TPP-1858	111	112	113	114	115	116	117	118	119	120
Comparative antibodies:										
P3G5 (TPP-2195)	121	122								
136.1 (TPP-2194)	123	124								
P4A8 (TPP-1324)	125	126								
PDL-192 (TPP-1104)	127	128								
18.3.3 (TPP-2193)	129	130								
P2D3 (TPP-2196)	131	132								

TPP-2090 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 2 and a region of the light

chain corresponding to SEQ ID NO: 1.

5 TPP-2149 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 12 and a region of the light chain corresponding to SEQ ID NO: 11.

10 TPP-2093 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 22 and a region of the light chain corresponding to SEQ ID NO: 21.

TPP-2148 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 32 and a region of the light chain corresponding to SEQ ID NO: 31.

15 TPP-2084 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 42 and a region of the light chain corresponding to SEQ ID NO: 41.

20 TPP-2077 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 52 and a region of the light chain corresponding to SEQ ID NO: 51.

25 TPP-1538 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 62 and a region of the light chain corresponding to SEQ ID NO: 61.

30 TPP-883 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 72 and a region of the light chain corresponding to SEQ ID NO: 71.

TPP-1854 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 82 and a region of the light chain corresponding to SEQ ID NO: 81.

35 TPP-1853 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 92 and a region of the light chain corresponding to SEQ ID NO: 91.

TPP-1857 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 102 and a region of the light chain corresponding to SEQ ID NO: 101.

5 TPP-1858 is: an antibody which comprises a region of the heavy chain corresponding to SEQ ID NO: 112 and a region of the light chain corresponding to SEQ ID NO: 111.

10 TPP-2090 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 10 and a variable region of the light chain corresponding to SEQ ID NO: 9.

15 TPP-2149 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 20 and a variable region of the light chain corresponding to SEQ ID NO: 19.

20 TPP-2093 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 30 and a variable region of the light chain corresponding to SEQ ID NO: 29.

TPP-2148 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 40 and a variable region of the light chain corresponding to SEQ ID NO: 39.

25 TPP-2084 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 50 and a variable region of the light chain corresponding to SEQ ID NO: 49.

30 TPP-2077 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 60 and a variable region of the light chain corresponding to SEQ ID NO: 59.

35 TPP-1538 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 70 and a variable region of the light chain corresponding to SEQ ID NO: 69.

TPP-883 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 80 and a variable region

of the light chain corresponding to SEQ ID NO: 79.

5 TPP-1854 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 90 and a variable region of the light chain corresponding to SEQ ID NO: 89.

10 TPP-1853 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 100 and a variable region of the light chain corresponding to SEQ ID NO: 99.

TPP-1857 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 110 and a variable region of the light chain corresponding to SEQ ID NO: 109.

15 TPP-1858 is: an antibody which comprises a variable region of the heavy chain corresponding to SEQ ID NO: 120 and a variable region of the light chain corresponding to SEQ ID NO: 119.

**Table:** DNA sequences of antibodies according to the invention

20

Antibody	SEQ ID NO: IgG1 light chain	SEQ ID NO: IgG1 heavy chain
Antibodies according to the invention:		
TPP-2090	177	178
TPP-2149	179	180
TPP-2093	181	182
TPP-2148	183	184
TPP-2084	185	186
TPP-2077	187	188
TPP-1538	189	190
TPP-883	191	192
TPP-1854	193	194

TPP-1853	195	196
TPP-1857	197	198
TPP-1858	199	200
Comparative antibodies:		
P3G5 (TPP-2195)	201	202
136.1 (TPP-2194)	203	204
P4A8 (TPP-1324)	205	206
PDL-192 (TPP-1104)	207	208
18.3.3 (TPP-2193)	209	210
P2D3 (TPP-2196)	211	212

Preferred embodiments of the anti-TWEAKR antibody are those below:

- 5 1. An anti-TWEAKR antibody or an antigen-binding fragment thereof which binds specifically to the D at position 47 (D47) of the TWEAKR (SEQ ID NO:169).
2. The antibody or an antigen-binding fragment thereof  
10 according to embodiment 1 where the antibody is an agonistic antibody.
3. The antibody or an antigen-binding fragment thereof according to embodiment 1 or 2 which comprises:
- 15 a variable heavy chain comprising:
- (a) a CDR1 of the heavy chain encoded by an amino acid sequence comprising the formula PYPMX (SEQ ID NO: 171), where X is I or  
20 M;
- (b) a CDR2 of the heavy chain encoded by an amino acid sequence comprising the formula YISPSGGXTHYADSVKG (SEQ ID NO: 172), where X is S or K; and
- 25

(c) a CDR3 of the heavy chain encoded by an amino acid sequence comprising the formula GGDTYFDYFDY (SEQ ID NO: 173);

and a variable light chain comprising:

5

(a) a CDR1 of the light chain encoded by an amino acid sequence comprising the formula RASQ<sub>S</sub>ISXYLN (SEQ ID NO: 174), where X is G or S;

10 (b) a CDR2 of the light chain encoded by an amino acid sequence comprising the formula XASSLQS (SEQ ID NO: 175), where X is Q, A or N; and

(c) a CDR3 of the light chain encoded by an amino acid sequence  
15 comprising the formula QQSYXXPXIT (SEQ ID NO: 176), where X at position 5 is T or S, X at position 6 is T or S and X at position 8 is G or F.

4. The antibody or an antigen-binding fragment thereof  
20 according to any of the preceding embodiments, comprising:

a. a variable heavy chain comprising the variable CDR1  
sequence of the heavy chain, as shown in SEQ ID NO: 6, the  
variable CDR2 sequence of the heavy chain, as shown in SEQ ID  
25 NO: 7 and the variable CDR3 sequence of the heavy chain, as  
shown in SEQ ID NO: 8, and also

a variable light chain comprising the variable CDR1 sequence of  
the light chain shown in SEQ ID NO: 3, the variable CDR2 sequence  
30 of the light chain shown in SEQ ID NO: 4 and the variable CDR3  
sequence of the light chain shown in SEQ ID NO: 5 or

b. a variable heavy chain comprising the variable CDR1  
sequence of the heavy chain, as shown in SEQ ID NO: 16, the  
35 variable CDR2 sequence of the heavy chain, as shown in SEQ ID  
NO: 17, the variable CDR3 sequence of the heavy chain, as shown  
in SEQ ID NO:18, and also

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 13, the variable CDR2 sequence of the light chain shown in SEQ ID NO: 14 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:15  
5 or

c. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 26, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID  
10 NO: 27, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:28, and also

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 23, the variable CDR2  
15 sequence of the light chain shown in SEQ ID NO: 24 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:25  
or

d. a variable heavy chain comprising the variable CDR1  
20 sequence of the heavy chain, as shown in SEQ ID NO: 36, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 37, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:38, and also

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 33, the variable CDR2  
25 sequence of the light chain shown in SEQ ID NO: 34 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:35  
or

30 e. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 46, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 47, the variable CDR3 sequence of the heavy chain, as shown  
35 in SEQ ID NO:48, and also

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 43, the variable CDR2



sequence of the light chain shown in SEQ ID NO: 44 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:45 or

5 f. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 56, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 57, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:58, and also

10

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 53, the variable CDR2 sequence of the light chain shown in SEQ ID NO: 54 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:55

15

or

g. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 66, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 67, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:68, and also

20

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 63, the variable CDR2 sequence of the light chain shown in SEQ ID NO: 64 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:65 or

25

h. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 76, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 77, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:78, and also

30

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 73, the variable CDR2 sequence of the light chain shown in SEQ ID NO: 74 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:75

35

or

i. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 86, the  
5 variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 87, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:88, and also

a variable light chain comprising the variable CDR1 sequence of  
10 the light chain shown in SEQ ID NO: 83, the variable CDR2 sequence of the light chain shown in SEQ ID NO: 84 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:85  
or

15 j. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 96, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 97, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:98, and also

20 a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 93, the variable CDR2 sequence of the light chain shown in SEQ ID NO: 94 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:95

25 or

k. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 106, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID  
30 NO: 107, the variable CDR3 sequence of the heavy chain, as shown in SEQ ID NO:108, and also

a variable light chain comprising the variable CDR1 sequence of  
the light chain shown in SEQ ID NO: 103, the variable CDR2  
35 sequence of the light chain shown in SEQ ID NO: 104 and the variable CDR3 sequence of the light chain shown in SEQ ID NO:105  
or

1. a variable heavy chain comprising the variable CDR1 sequence of the heavy chain, as shown in SEQ ID NO: 116, the variable CDR2 sequence of the heavy chain, as shown in SEQ ID NO: 117, the variable CDR3 sequence of the heavy chain, as shown  
5 in SEQ ID NO:118, and also

a variable light chain comprising the variable CDR1 sequence of the light chain shown in SEQ ID NO: 113, the variable CDR2 sequence of the light chain shown in SEQ ID NO: 114 and the  
10 variable CDR3 sequence of the light chain shown in SEQ ID NO:115.

5. The antibody or the antigen-binding fragment thereof according to any of the preceding embodiments, comprising:

15 a. a variable sequence of the heavy chain, as shown in SEQ ID NO:10, and also a variable sequence of the light chain, as shown in SEQ ID NO:9, or

b. a variable sequence of the heavy chain, as shown in SEQ ID  
20 NO:20, and also a variable sequence of the light chain, as shown in SEQ ID NO:19, or

c. a variable sequence of the heavy chain, as shown in SEQ ID  
25 NO:30, and also a variable sequence of the light chain, as shown in SEQ ID NO:29, or

d. a variable sequence of the heavy chain, as shown in SEQ ID  
NO:40, and also a variable sequence of the light chain, as shown  
in SEQ ID NO:39, or

30 e. a variable sequence of the heavy chain, as shown in SEQ ID NO:50, and also a variable sequence of the light chain, as shown in SEQ ID NO:49, or

35 f. a variable sequence of the heavy chain, as shown in SEQ ID NO:60, and also a variable sequence of the light chain, as shown in SEQ ID NO:59, or

- g. a variable sequence of the heavy chain, as shown in SEQ ID NO:70, and also a variable sequence of the light chain, as shown in SEQ ID NO:69, or
- 5 h. a variable sequence of the heavy chain, as shown in SEQ ID NO:80, and also a variable sequence of the light chain, as shown in SEQ ID NO:79, or
- 10 i. a variable sequence of the heavy chain, as shown in SEQ ID NO:90, and also a variable sequence of the light chain, as shown in SEQ ID NO:89, or
- 15 j. a variable sequence of the heavy chain, as shown in SEQ ID NO:100, and also a variable sequence of the light chain, as shown in SEQ ID NO:99, or
- k. a variable sequence of the heavy chain, as shown in SEQ ID NO:110, and also a variable sequence of the light chain, as shown in SEQ ID NO:109, or
- 20 l. a variable sequence of the heavy chain, as shown in SEQ ID NO:120, and also a variable sequence of the light chain, as shown in SEQ ID NO:119.
- 25 6. The antibody according to any of the preceding embodiments which is an IgG antibody.
7. The antibody according to any of the preceding embodiments, comprising:
- 30 a. a sequence of the heavy chain, as shown in SEQ ID NO:2, and also a sequence of the light chain, as shown in SEQ ID NO:1, or
- 35 b. a sequence of the heavy chain, as shown in SEQ ID NO:12, and also a sequence of the light chain, as shown in SEQ ID NO:11, or
- c. a sequence of the heavy chain, as shown in SEQ ID NO:22,

and also a sequence of the light chain, as shown in SEQ ID NO:21,  
or

d. a sequence of the heavy chain, as shown in SEQ ID NO:32,  
5 and also a sequence of the light chain, as shown in SEQ ID NO:31,  
or

e. a sequence of the heavy chain, as shown in SEQ ID NO:42,  
and also a sequence of the light chain, as shown in SEQ ID NO:41,  
10 or

f. a sequence of the heavy chain, as shown in SEQ ID NO:52,  
and also a sequence of the light chain, as shown in SEQ ID NO:51,  
or

15 g. a sequence of the heavy chain, as shown in SEQ ID NO:62,  
and also a sequence of the light chain, as shown in SEQ ID NO:61,  
or

20 h. a sequence of the heavy chain, as shown in SEQ ID NO:72,  
and also a sequence of the light chain, as shown in SEQ ID NO:71,  
or

i. a sequence of the heavy chain, as shown in SEQ ID NO:82,  
25 and also a sequence of the light chain, as shown in SEQ ID NO:81,  
or

j. a sequence of the heavy chain, as shown in SEQ ID NO:92,  
and also a sequence of the light chain, as shown in SEQ ID NO:91,  
30 or

k. a sequence of the heavy chain, as shown in SEQ ID NO:102,  
and also a sequence of the light chain, as shown in SEQ ID  
NO:101, or

35 l. a sequence of the heavy chain, as shown in SEQ ID NO:112,  
and also a sequence of the light chain, as shown in SEQ ID  
NO:111.

8. The antigen-binding fragment according to any of the preceding embodiments or an antigen-binding fragment of an antibody according to any of the preceding embodiments which is  
5 an scFv, Fab, Fab ' fragment or a F(ab ' )<sub>2</sub> fragment.

9. The antibody or the antigen-binding fragment according to any of the preceding embodiments which is a monoclonal antibody or an antigen-binding fragment thereof.  
10

10. The antibody or the antigen-binding fragment according to any of the preceding embodiments which is a human, humanized or chimeric antibody or an antigen-binding fragment.

15 Particular preference is given to the anti-TWEAKR antibody TPP-2090.

***Isotopes, salts, solvates, isotopic variants***

20 The present invention also encompasses all suitable isotopic variants of the compounds according to the invention. An isotopic variant of a compound according to the invention is understood here as meaning a compound in which at least one atom within the compound according to the invention has been  
25 exchanged for another atom of the same atomic number, but with a different atomic mass than the atomic mass which usually or predominantly occurs in nature. Examples of isotopes which can be incorporated into a compound according to the invention are those of hydrogen, carbon, nitrogen, oxygen, phosphorus,  
30 sulphur, fluorine, chlorine, bromine and iodine, such as <sup>2</sup>H (deuterium), <sup>3</sup>H (tritium), <sup>13</sup>C, <sup>14</sup>C, <sup>15</sup>N, <sup>17</sup>O, <sup>18</sup>O, <sup>32</sup>P, <sup>33</sup>P, <sup>33</sup>S, <sup>34</sup>S, <sup>35</sup>S, <sup>36</sup>S, <sup>18</sup>F, <sup>36</sup>Cl, <sup>82</sup>Br, <sup>123</sup>I, <sup>124</sup>I, <sup>129</sup>I and <sup>131</sup>I. Particular isotopic variants of a compound according to the invention, especially those in which one or more radioactive isotopes have  
35 been incorporated, may be beneficial, for example, for the examination of the mechanism of action or of the active compound distribution in the body; due to comparatively easy preparability and detectability, especially compounds labelled

with  $^3\text{H}$  or  $^{14}\text{C}$  isotopes are suitable for this purpose. In addition, the incorporation of isotopes, for example of deuterium, can lead to particular therapeutic benefits as a consequence of greater metabolic stability of the compound, for example an extension of the half-life in the body or a reduction in the active dose required; such modifications of the compounds according to the invention may therefore in some cases also constitute a preferred embodiment of the present invention. Isotopic variants of the compounds according to the invention can be prepared by the processes known to those skilled in the art, for example by the methods described below and the procedures described in the working examples, by using corresponding isotopic modifications of the respective reagents and/or starting compounds.

Preferred salts in the context of the present invention are physiologically acceptable salts of the compounds according to the invention. Also encompassed are salts which are not themselves suitable for pharmaceutical applications but can be used, for example, for isolation or purification of the compounds according to the invention.

Physiologically acceptable salts of the compounds according to the invention include acid addition salts of mineral acids, carboxylic acids and sulphonic acids, for example salts of hydrochloric acid, hydrobromic acid, sulphuric acid, phosphoric acid, methanesulphonic acid, ethanesulphonic acid, benzenesulphonic acid, toluenesulphonic acid, naphthalenedisulphonic acid, acetic acid, trifluoroacetic acid, propionic acid, lactic acid, tartaric acid, malic acid, citric acid, fumaric acid, maleic acid and benzoic acid.

Physiologically acceptable salts of the inventive compounds also include salts of conventional bases, by way of example and with preference alkali metal salts (e.g. sodium and potassium salts), alkaline earth metal salts (e.g. calcium and magnesium salts) and ammonium salts derived from ammonia or organic amines having 1 to 16 carbon atoms, by way of example and with preference

ethylamine, diethylamine, triethylamine, ethyldiisopropylamine,  
 monoethanolamine, diethanolamine, triethanolamine,  
 dicyclohexylamine, dimethylaminoethanol, procaine,  
 dibenzylamine, N-methylpiperidine, N-methylmorpholine,  
 5 arginine, lysine and 1,2-ethylenediamine.

Solvates in the context of the invention are described as those  
 forms of the compounds according to the invention which form a  
 complex in the solid or liquid state by coordination with solvent  
 10 molecules. Hydrates are a specific form of the solvates in which  
 the coordination is with water. Solvates preferred in the  
 context of the present invention are hydrates.

In addition, the present invention also encompasses prodrugs of  
 15 the compounds according to the invention. The term "prodrugs"  
 here denotes compounds which may themselves be biologically  
 active or inactive, but are converted (for example by metabolic  
 or hydrolytic means) to inventive compounds during their  
 residence time in the body.

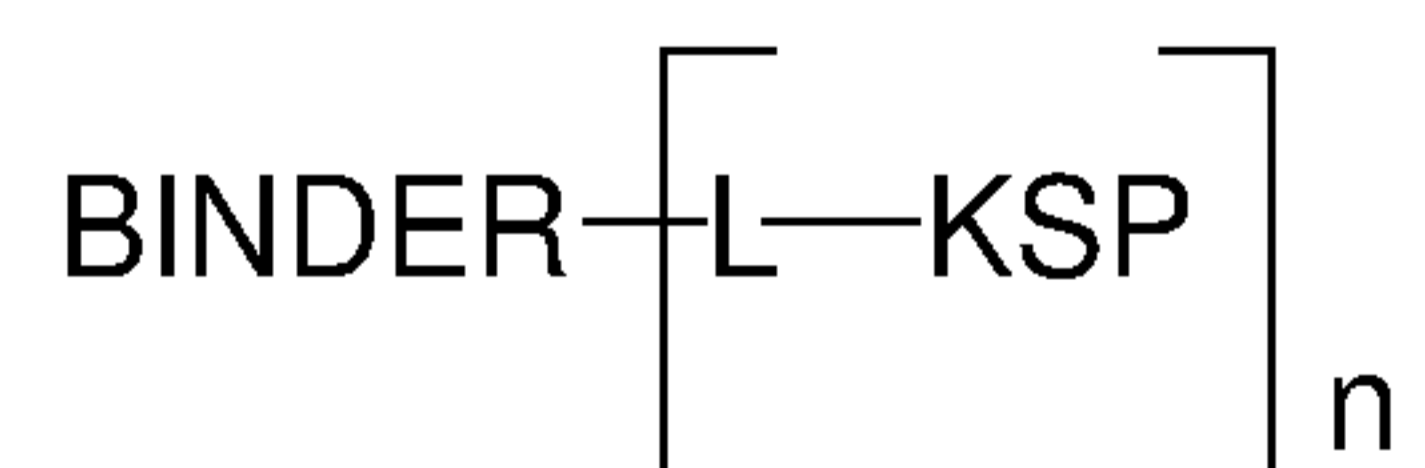
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### ***Particular embodiments***

The following embodiments are particularly preferred:

25 Embodiment A:

An ADC of the formula

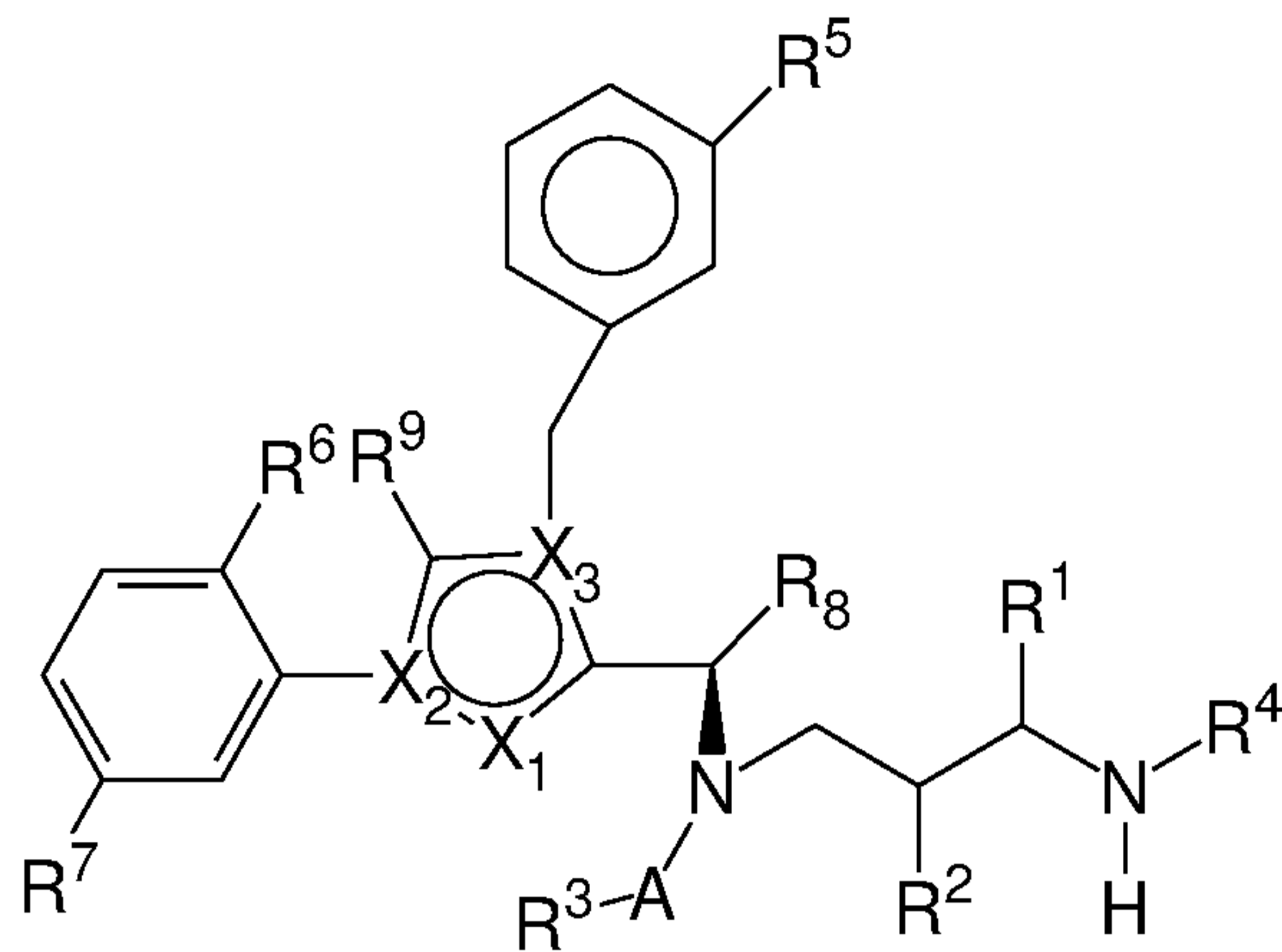


30

where KSP-L- is a compound of the formula (II), (IIa), (IIb),  
 (IIc), (IId), (IIe) below or the formula (IIf) below, the binder  
 is an anti-TWEAKR antibody (particularly preferably an anti-  
 TWEAKR antibody which binds specifically to amino acid D in  
 35 position 47 (D47) of TWEAKR (SEQ ID NO:169), especially the  
 anti-TWEAK R antibody TPP-2090), and n is a number from 1 to 10:



formula (IIIf):



(IIIf)

5 where

X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

10 X<sub>1</sub> represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represents N;

X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C; or

X<sub>1</sub> represents CH, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

15 A represents CO (carbonyl);

R<sup>1</sup> represents -L-#1, H, -COOH, -CONHNH<sub>2</sub>, -(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub>, -CONZ''(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub> and -CONZ''CH<sub>2</sub>COOH, where Z'' represents H or NH<sub>2</sub>;

20

R<sup>2</sup> and R<sup>4</sup> represent H or -L-#1, or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents H or -L-#1;

25 R<sup>3</sup> represents -L-#1 or a C<sub>1-10</sub>-alkyl-, which may optionally be substituted by -OH, O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-NH-alkyl, NH-CO-alkyl, NH-CO-NH-alkyl, S(O)<sub>n</sub>-alkyl, SO<sub>2</sub>-NH-alkyl, NH-alkyl, N(alkyl)<sub>2</sub> or NH<sub>2</sub> (where alkyl is preferably C<sub>1-3</sub>-alkyl);

R<sup>5</sup> represents -L-#1, H or F;

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, (optionally fluorinated) C<sub>1-3</sub>-alkyl, (optionally fluorinated) C<sub>2-4</sub>-alkenyl,  
5 (optionally fluorinated) C<sub>2-4</sub>-alkynyl, hydroxy or halogen;

R<sup>8</sup> represents a branched C<sub>1-5</sub>-alkyl group which may be substituted by -L-#1; and

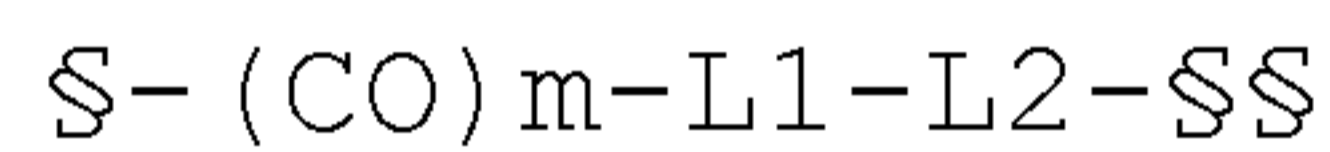
10 R<sup>9</sup> represents H or F,

where one of the substituents R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup>, R<sup>8</sup> and R<sup>10</sup> represents -L-#1, and

15 -L- represents the linker and #1 represents the bond to the antibody,

and salts, solvates and salts of the solvates of the ADC.

20 The linker is preferably a linker



where

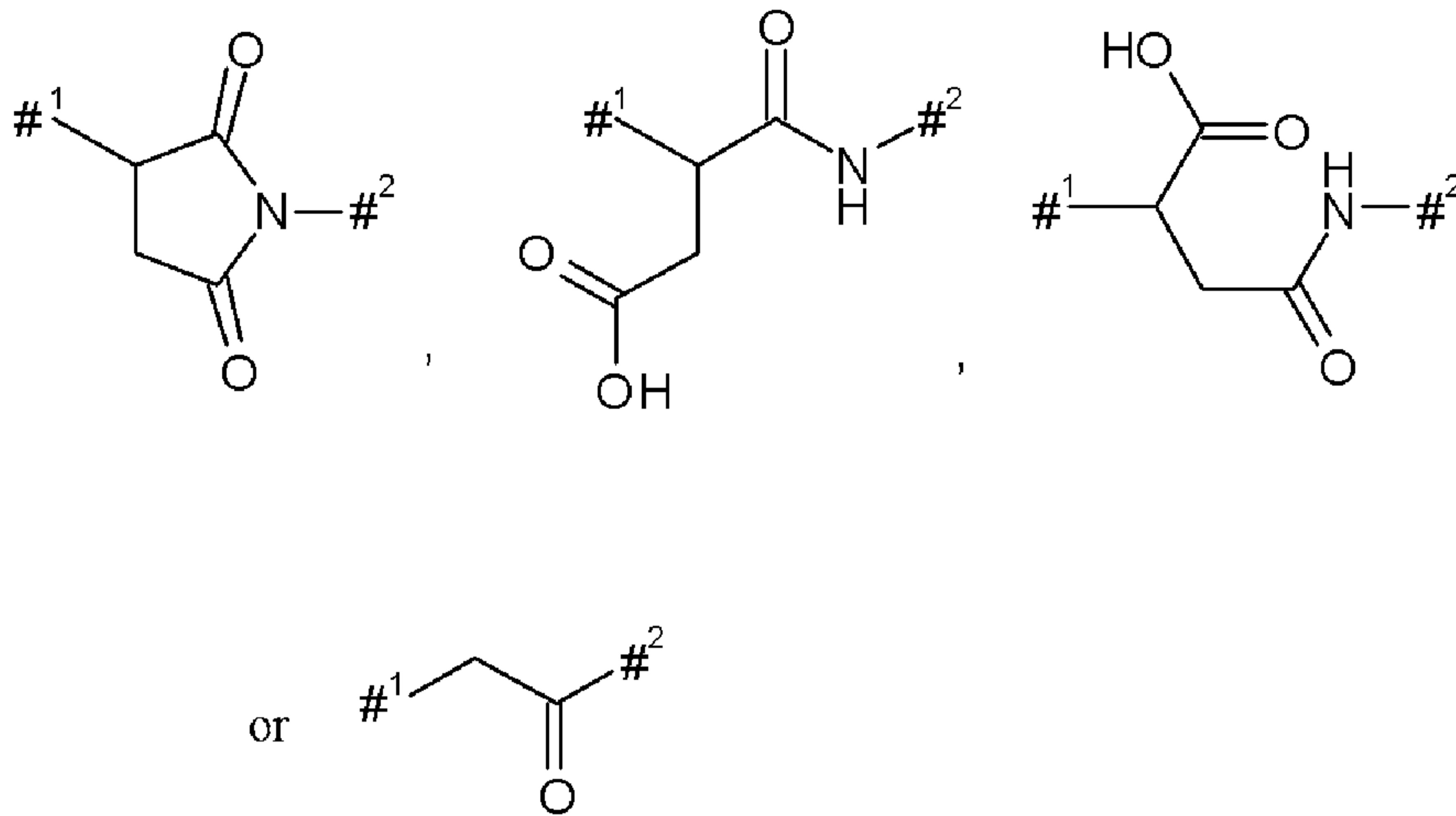
25

m is 0 or 1;

\\$ represents the bond to KSP and

30 \\$\$ represents the bond to the antibody, and

L2 represents

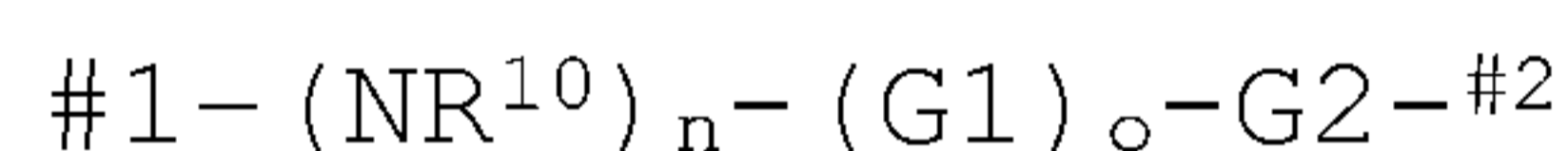


where

5 #<sup>1</sup> denotes the point of attachment to the sulphur atom of the antibody,

#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>,

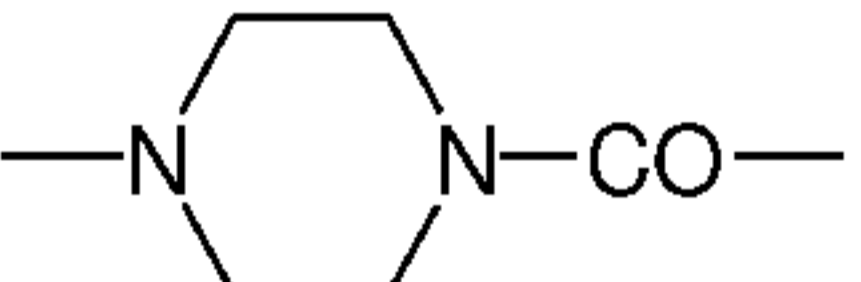
10 and L<sup>1</sup> is represented by formula



where

15

R<sup>10</sup> represents H, NH<sub>2</sub> or C<sub>1</sub>-C<sub>3</sub>-alkyl;

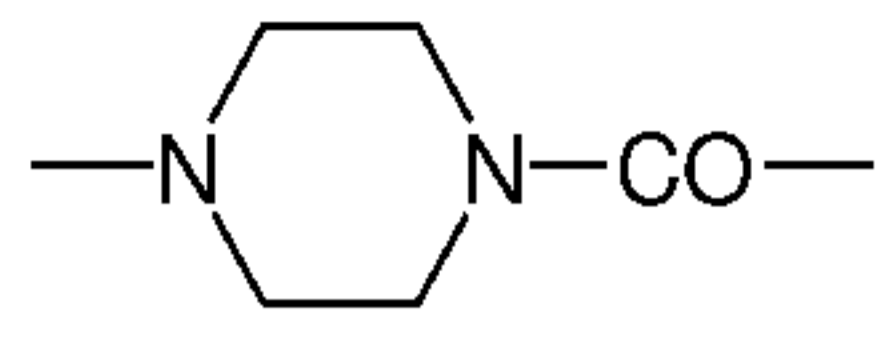
G1 represents -NHCO- or  ;

20 n is 0 or 1;

o is 0 or 1; and

25 G2 represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-,

-NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- and a 3- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, or -SO- (preferably



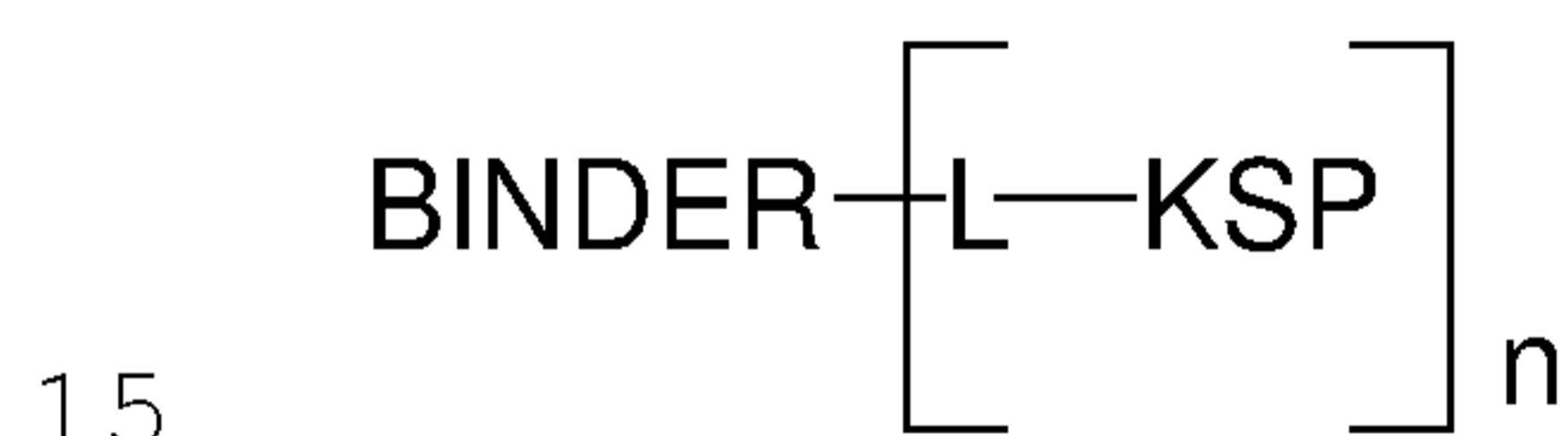
5 substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulfoxide or sulphonic acid.

Here, #<sup>1</sup> is the bond to the KSP inhibitor and #<sup>2</sup> is the bond to the coupling group to the binder (e.g. L2).

10

Embodiment B:

An ADC of the formula

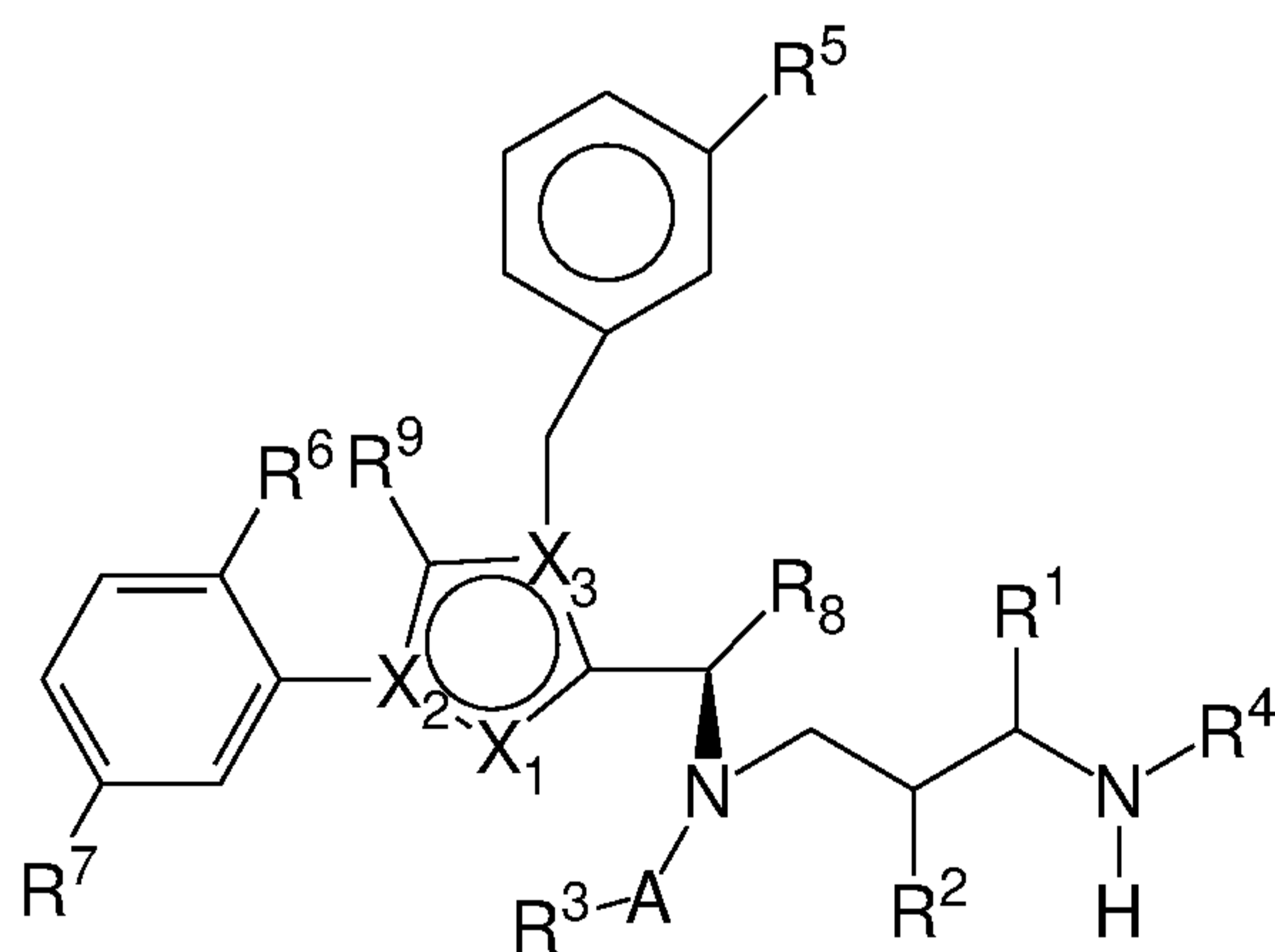


15

where KSP-L- is a compound of the formula (II), (IIa), (IIb), (IIc), (IId), (IIe), (IIf) below or of the formula (IIg) below, the binder is an antibody and n is a number from 1 to 10:

20

formula (IIg):



(IIg)

25 where

X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

X<sub>1</sub> represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represents N;

X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C; or

5

X<sub>1</sub> represents CH, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

A represents CO (carbonyl);

10 R<sup>1</sup> represents -L-#1, H, -COOH, -CONH<sub>2</sub>, -(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub>, -CONZ''(CH<sub>2</sub>)<sub>1-3</sub>NH<sub>2</sub> and -CONZ''CH<sub>2</sub>COOH, where Z'' represents H or NH<sub>2</sub>;

15 R<sup>2</sup> and R<sup>4</sup> represent H or -L-#1, or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents H or -L-#1;

20 R<sup>3</sup> represents -L-#1 or a C<sub>1-10</sub>-alkyl-, which may optionally be substituted by -OH, O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-NH-alkyl, NH-CO-alkyl, NH-CO-NH-alkyl, S(O)<sub>n</sub>-alkyl, SO<sub>2</sub>-NH-alkyl, NH-alkyl, N(alkyl)<sub>2</sub> or NH<sub>2</sub> (where alkyl is preferably C<sub>1-3</sub>-alkyl);

R<sup>5</sup> represents -L-#1, H or F;

25 R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, (optionally fluorinated) C<sub>1-3</sub>-alkyl, (optionally fluorinated) C<sub>2-4</sub>-alkenyl, (optionally fluorinated) C<sub>2-4</sub>-alkynyl, hydroxy or halogen;

R<sup>8</sup> represents a branched C<sub>1-5</sub>-alkyl group; and

30

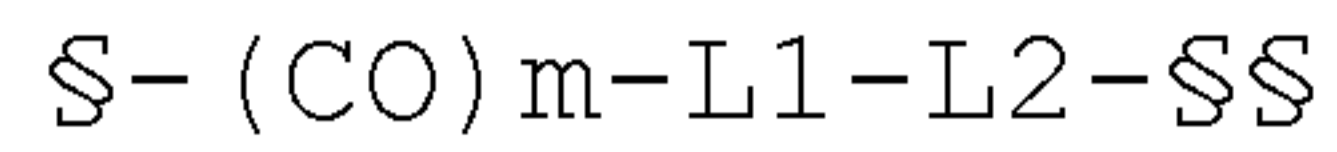
R<sup>9</sup> represents H or F,

where one of the substituents R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup>, R<sup>5</sup> and R<sup>10</sup> represents -L-#1, and

35

-L- represents the linker and #1 represents the bond to the antibody,

where -L- is represented by



5 where

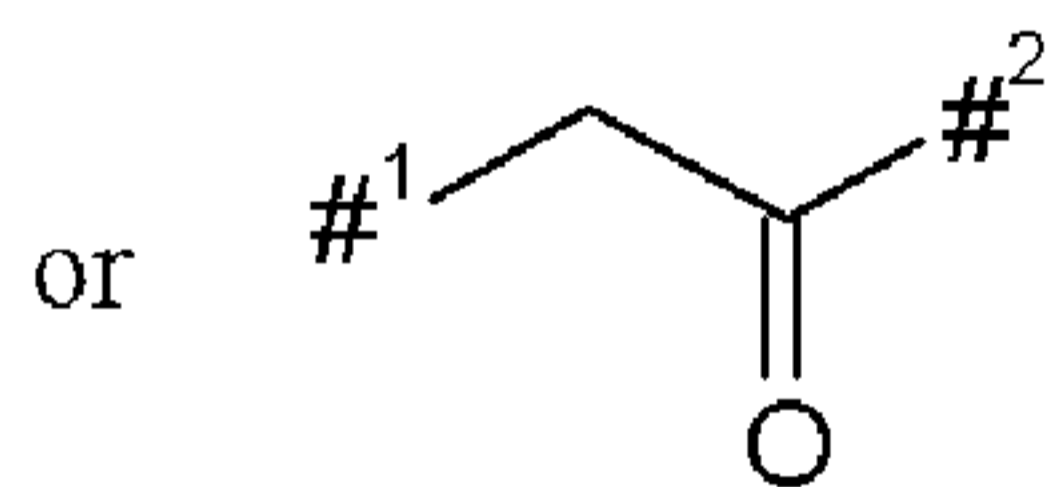
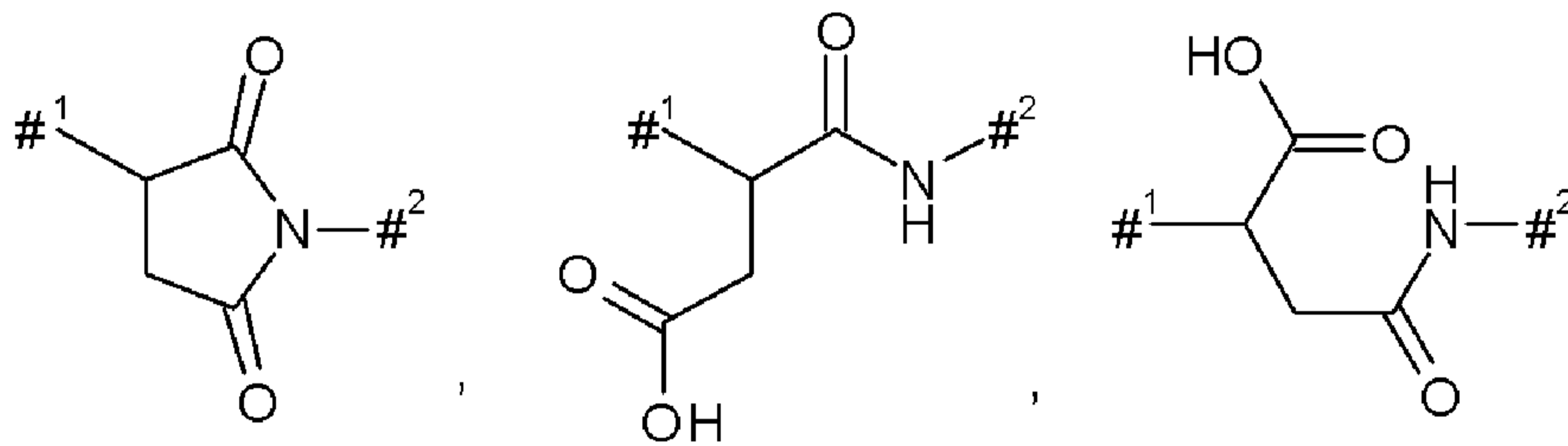
m is 0 or 1;

§ represents the bond to KSP and

10

§§ represents the bond to the antibody, and

L2 represents



15

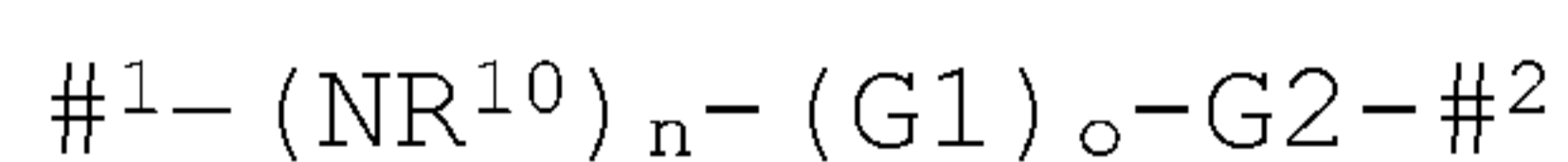
where

#<sup>1</sup> denotes the point of attachment to the sulphur atom of the  
20 antibody,

#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>,

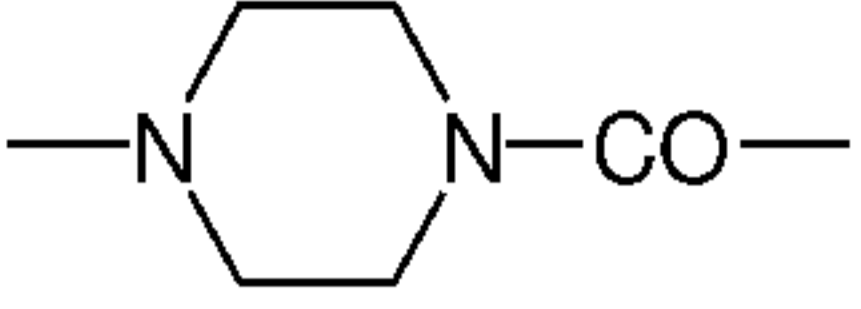
and L1 is represented by formula

25



where

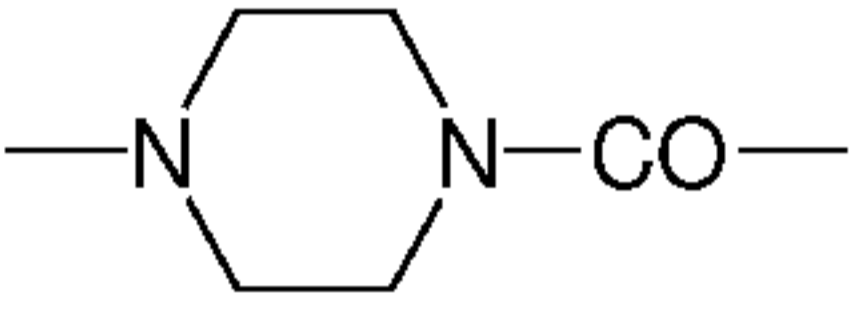
R<sup>10</sup> represents H, NH<sub>2</sub> or C<sub>1</sub>-C<sub>3</sub>-alkyl;

G1 represents -NHCO- or  ;

5 n is 0 or 1;

o is 0 or 1; and

G2 represents a straight-chain or branched hydrocarbon chain  
 10 having 1 to 100 carbon atoms from arylene groups and/or straight-  
 chain and/or branched and/or cyclic alkylene groups and which  
 may be interrupted once or more than once by one or more of the  
 groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-,  
 -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- and a 3- to 10-membered aromatic or  
 15 non-aromatic heterocycle having up to 4 heteroatoms selected  
 from the group consisting of N, O and S, or -SO- (preferably

, where the side chains, if present, may be  
 substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>,  
 sulphonamide, sulphone, sulphoxide or sulphonic acid,

20

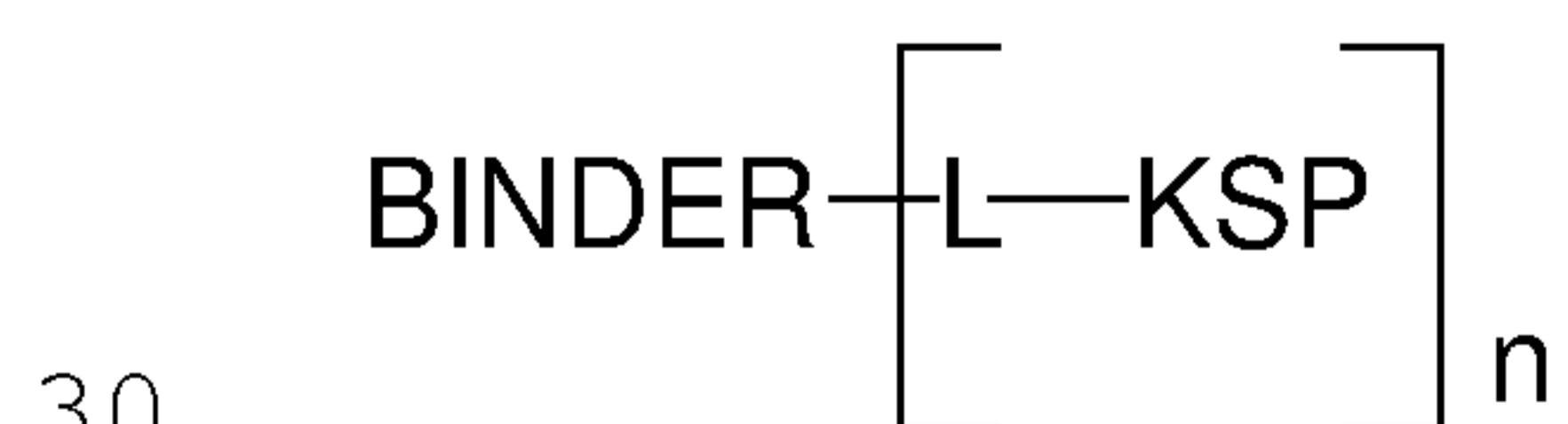
#<sup>1</sup> is the bond to the KSP inhibitor and #<sup>2</sup> is the bond to the  
 coupling group to the antibody (e.g. L2),

and salts, solvates and salts of the solvates of the ADC.

25

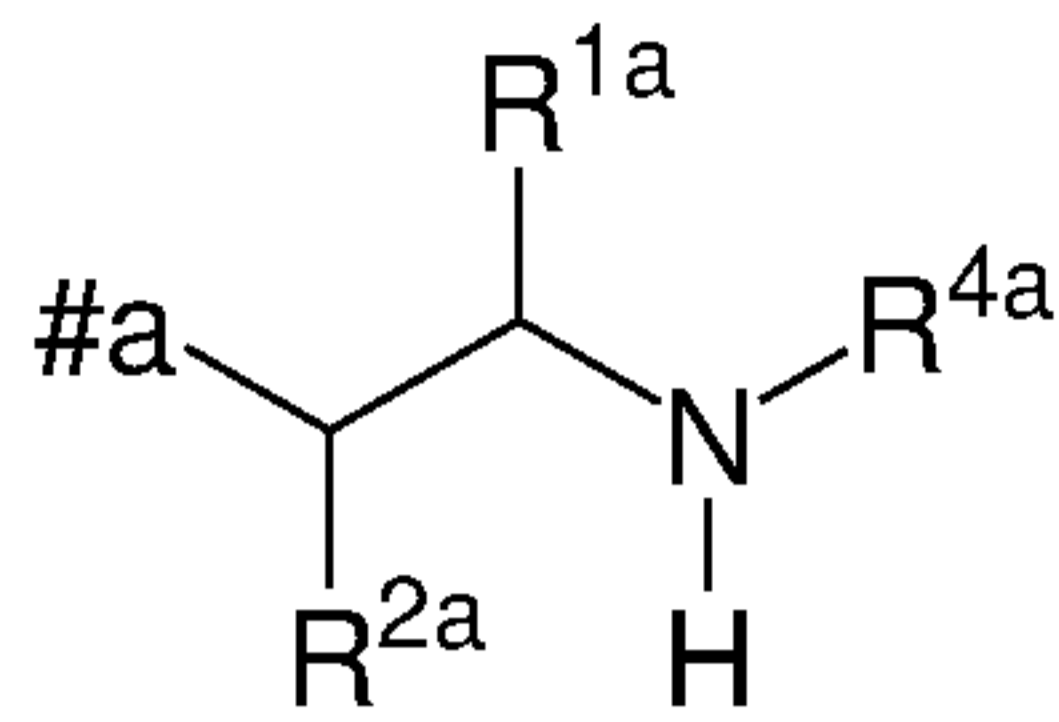
### Embodiment C:

An ADC of the formula



where KSP-L- is a compound having the substructure I(sub) below,  
 the binder is an anti-TWEAKR antibody (particularly preferably  
 an anti-TWEAKR antibody which binds specifically to amino acid  
 35 D in position 47 (D47) of TWEAKR (SEQ ID NO:169), especially the

anti-TWEAK R antibody TPP-2090), anti-HER2 antibody or anti-EGFR antibody (preferably nimotuzumab), and n is a number from 1 to 10:



5

where

\#a represents a bond to the remainder of the molecule;

10

R<sup>1a</sup> represents -L-\#1, H or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, halogen, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub>, COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted  
20 by -NH<sub>2</sub>;

R<sup>2a</sup> and R<sup>4a</sup> independently of one another represent H, -L-\#1, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or R<sup>2a</sup> and R<sup>4a</sup> together represent (with formation of a pyrrolidine ring) -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where  
25 R<sup>10</sup> represents -L-\#1, H, NH<sub>2</sub>, COOH, SO<sub>3</sub>H, SH or OH, and where Z represents -H, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z'  
30 represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH,

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where  
35



$Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

where one of the substituents  $R^{1a}$ ,  $R^{2a}$ ,  $R^{4a}$  or  $R^{10}$  represents -L-  
#1,

5

-L- represents the linker and #1 represents the bond to the  
antibody,

where -L- is represented by

10

$\S-(CO)_m-L1-L2-\S\S$

where

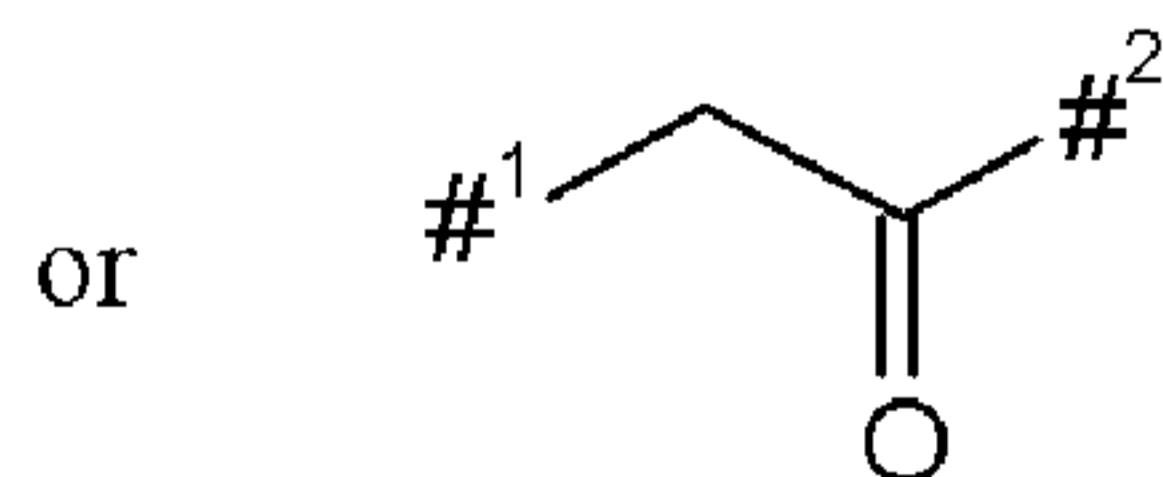
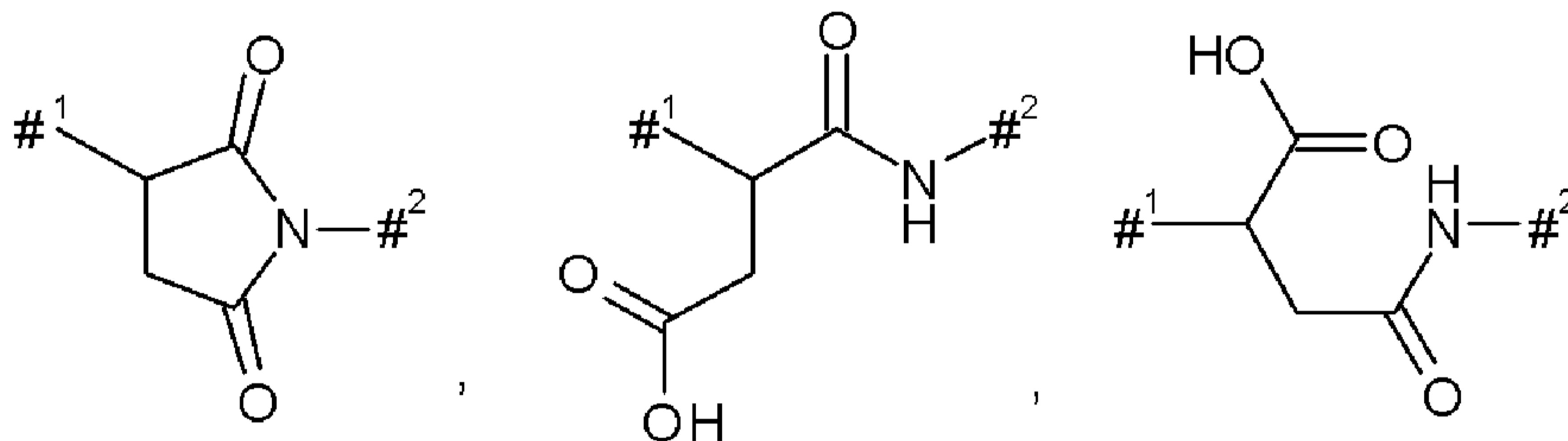
15 m is 0 or 1;

$\S$  represents the bond to KSP and

$\S\S$  represents the bond to the antibody, and

20

L2 represents

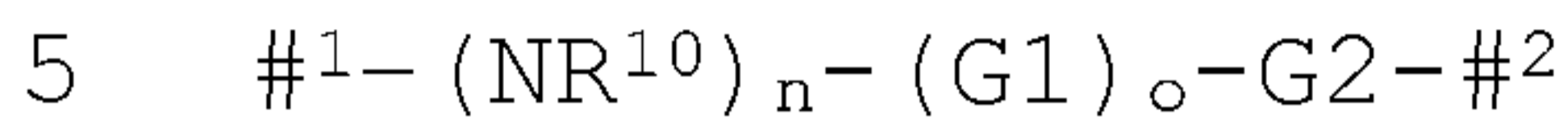


25 where

#<sup>1</sup> denotes the point of attachment to the sulphur atom of the  
antibody,

#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>,

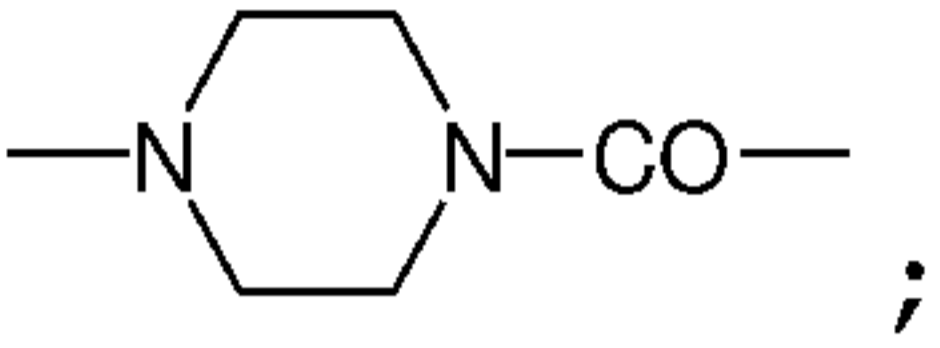
and L<sup>1</sup> is represented by formula



where

R<sup>10</sup> represents H, NH<sub>2</sub> or C<sub>1</sub>-C<sub>3</sub>-alkyl;

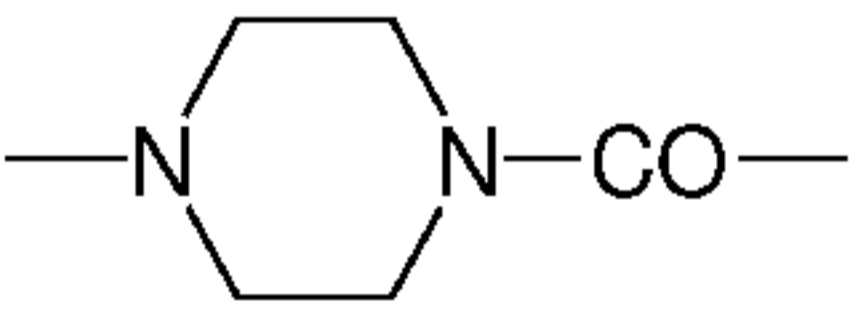
10

G<sub>1</sub> represents -NHCO- or  ;

n is 0 or 1;

15 o is 0 or 1; and

G<sub>2</sub> represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- and a 3- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, or -SO- (preferably

25 ) , where the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid,

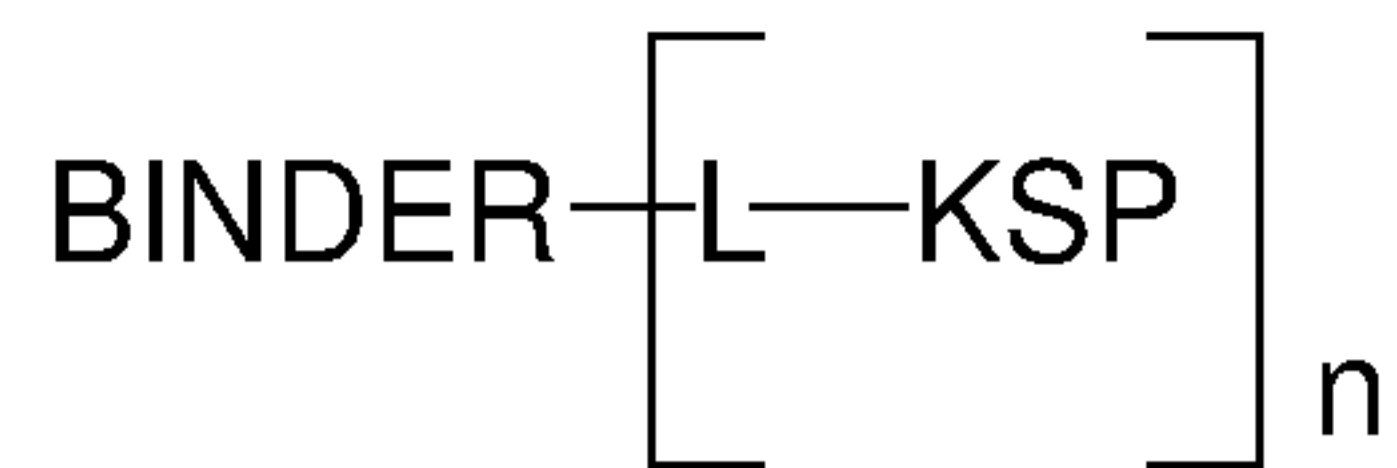
30 #<sup>1</sup> is the bond to the KSP inhibitor and #<sup>2</sup> is the bond to the coupling group to the antibody (e.g. L<sub>2</sub>),

and salts, solvates and salts of the solvates of the ADC.

#### Embodiment D:

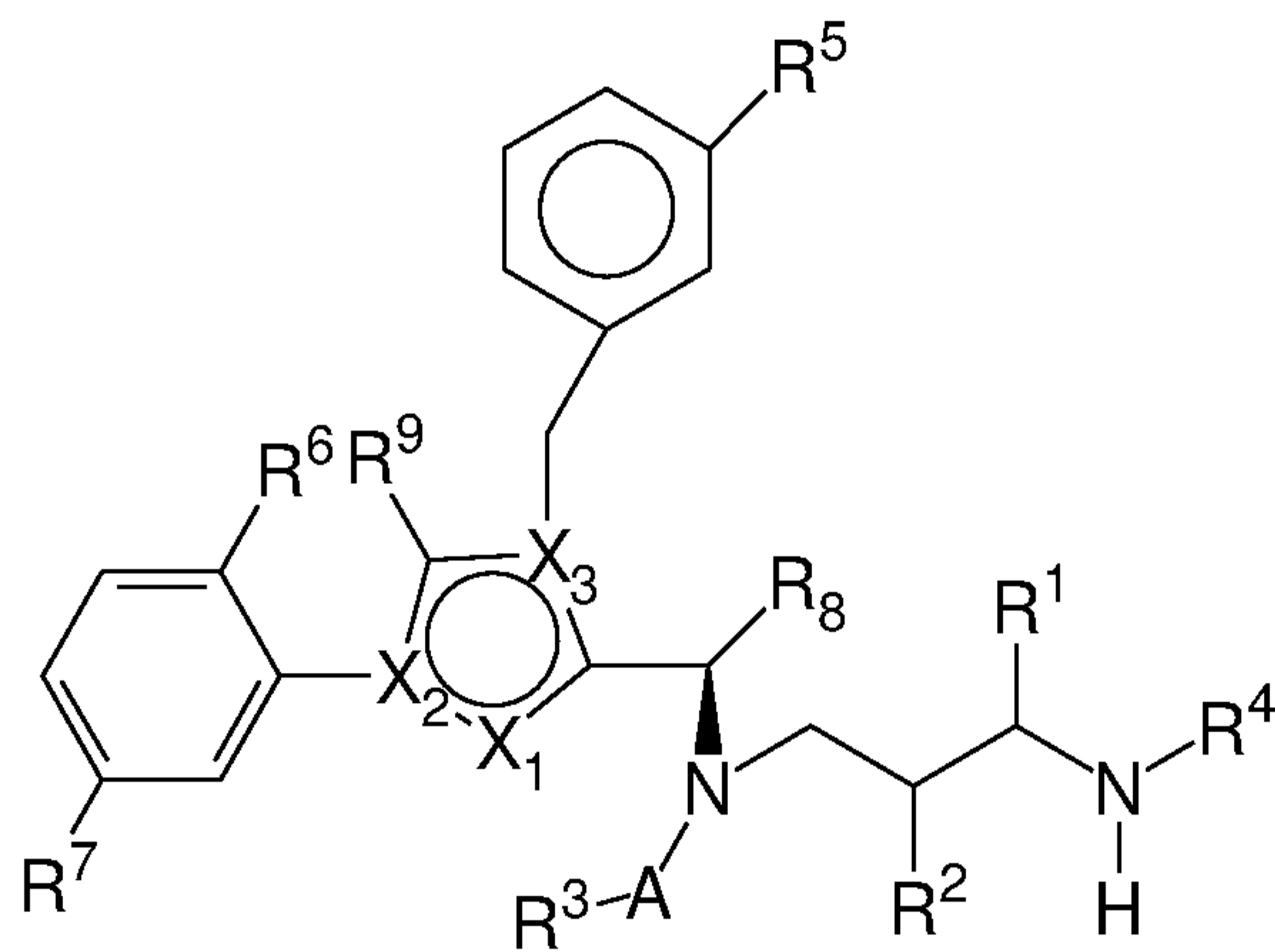
35

An ADC of the formula



where KSP-L- is a compound of the formula (II), (IIa), (IIb), (IIc), (IId), (IIe), (IIf), (IIg) below or of the formula (IIh) below, the binder is an antibody and n is a number from 1 to 10:

formula (IIh):



(II)

10

where

X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

15 X<sub>1</sub> represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represents N;

X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C; or

X<sub>1</sub> represents CH, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

20

A represents CO (carbonyl);

R<sup>1</sup> represents -L-#1;

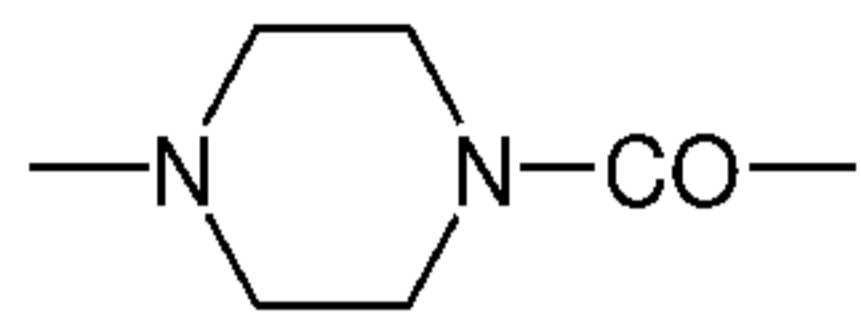
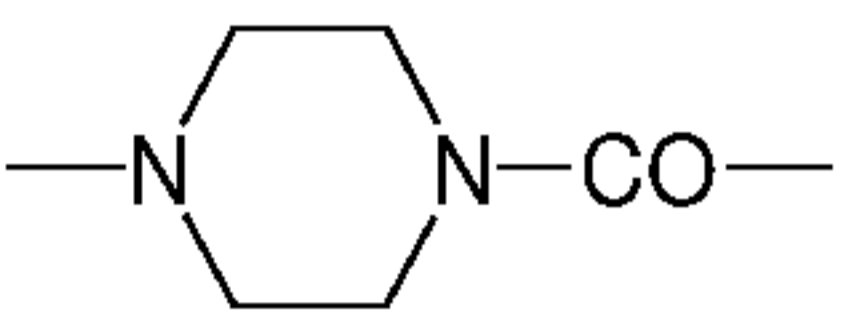
25 R<sup>2</sup> and R<sup>4</sup> represent H, or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents H;

R<sup>3</sup> represents C<sub>1-10</sub>-alkyl-, which may optionally be substituted by -OH, O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-NH-alkyl, NH-CO-alkyl, NH-CO-NH-alkyl, S(O)<sub>n</sub>-alkyl, SO<sub>2</sub>-NH-alkyl, NH-alkyl, N(alkyl)<sub>2</sub> or NH<sub>2</sub> (where alkyl is preferably C<sub>1-3</sub>-alkyl), or -MOD;

5

where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

R<sup>10</sup> represents H or C<sub>1-3</sub>-alkyl;

10 G1 represents -NHCO- , -CONH- or  (where, if G1 represents -NHCO- or  , R<sup>10</sup> does not represent NH<sub>2</sub>);

n is 0 or 1;

15 o is 0 or 1; and

G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-,  
 20 SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>-  
 (where R<sup>y</sup> represents H, phenyl, C<sub>1-10</sub>-alkyl, C<sub>2-10</sub>-alkenyl or C<sub>2-10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1-3</sub>-alkyl or phenyl), where the hydrocarbon chain including any side  
 25 chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group -MOD preferably has at least one group -COOH;

30 R<sup>5</sup> represents H or F;

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, (optionally fluorinated) C<sub>1-3</sub>-alkyl, (optionally fluorinated) C<sub>2-4</sub>-alkenyl, (optionally fluorinated) C<sub>2-4</sub>-alkynyl, hydroxy or halogen;

35

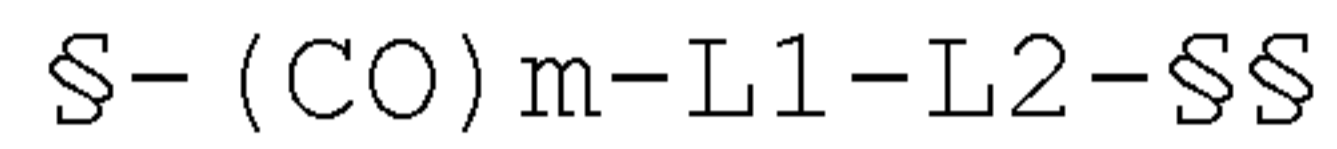
R<sup>8</sup> represents a branched C<sub>1-5</sub>-alkyl group; and

R<sup>9</sup> represents H or F,

where -L- represents the linker and #1 represents the bond to the antibody,

5

where -L- is represented by



10 where

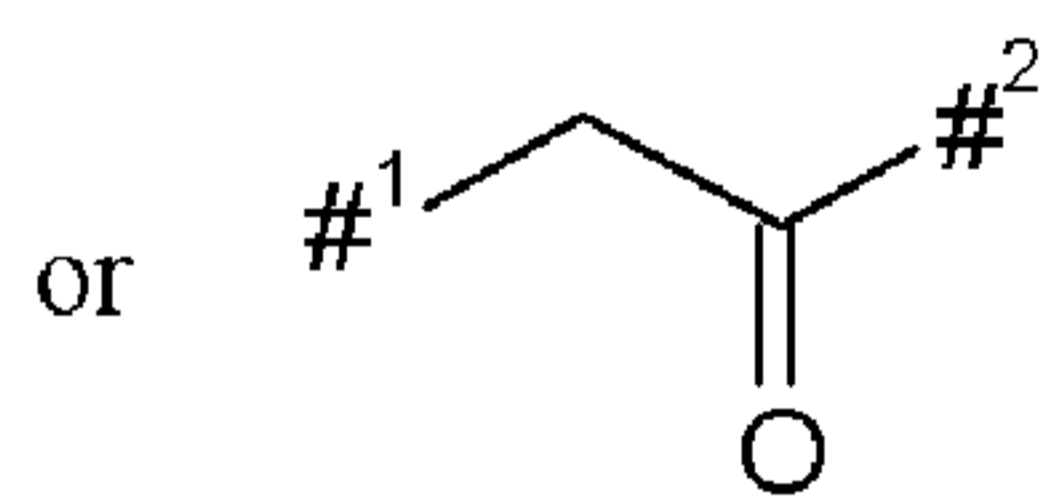
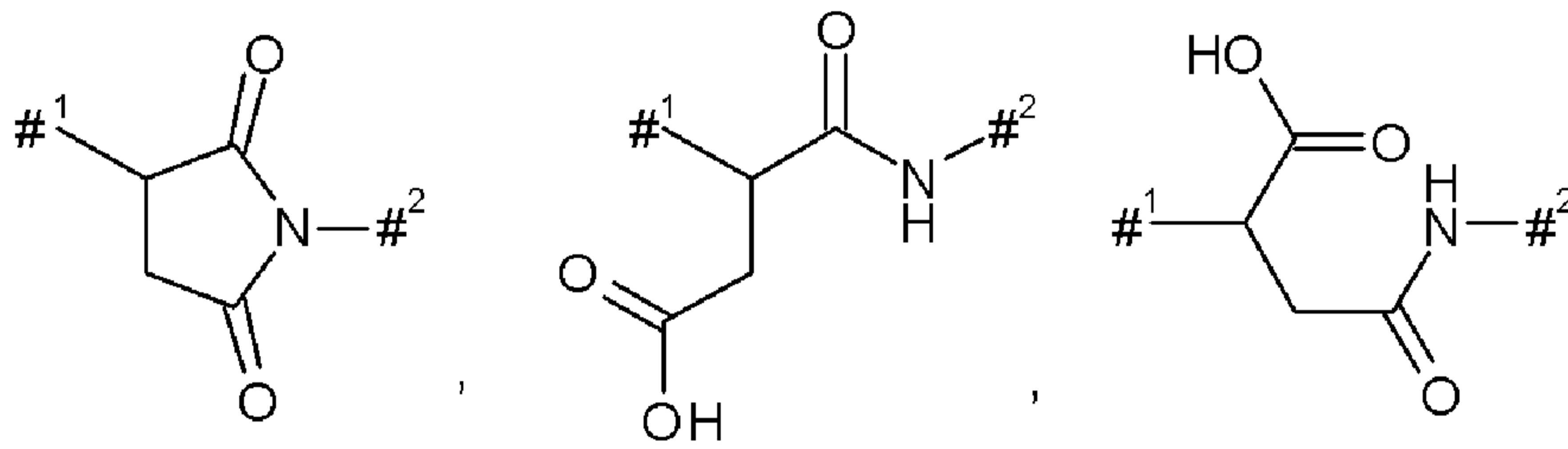
m is 0 or 1;

\S represents the bond to KSP and

15

\S\S represents the bond to the antibody, and

L2 represents



20

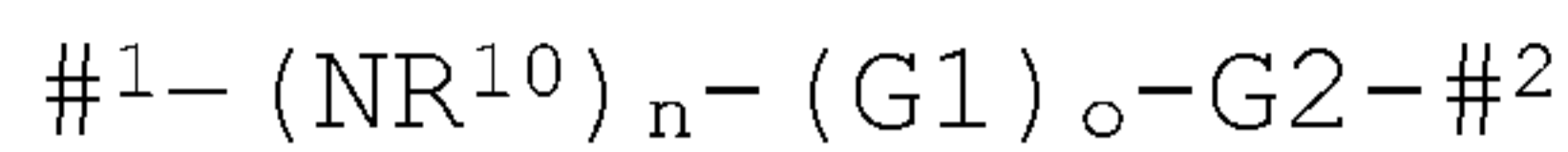
where

#<sup>1</sup> denotes the point of attachment to the sulphur atom of the antibody,

25

#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>,

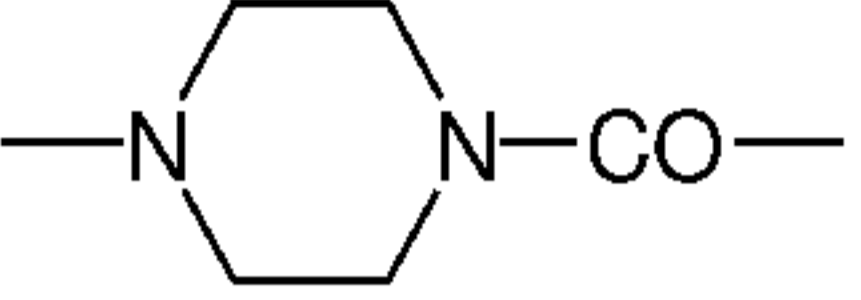
and L1 is represented by formula



where

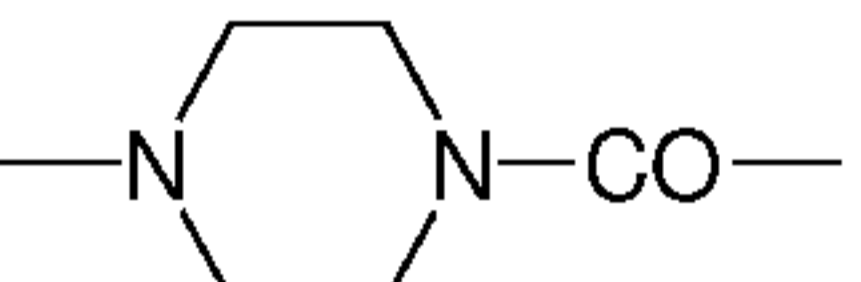
5

R<sub>10</sub> represents H, NH<sub>2</sub> or C<sub>1</sub>-C<sub>3</sub>-alkyl;

G1 represents -NHCO- or  ;

10 n is 0 or 1;

o is 0 or 1; and

G2 represents a straight-chain or branched hydrocarbon chain  
 15 having 1 to 100 carbon atoms from arylene groups and/or straight-  
 chain and/or branched and/or cyclic alkylene groups and which  
 may be interrupted once or more than once by one or more of the  
 groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-,  
 -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H,  
 20 C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl) and a 3- to 10-membered aromatic or non-  
 aromatic heterocycle having up to 4 heteroatoms selected from  
 the group consisting of N, O and S, -SO- or -SO<sub>2</sub>- (preferably  
  
 ), where the hydrocarbon chain including the side  
 chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH,  
 25 -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic  
 acid.

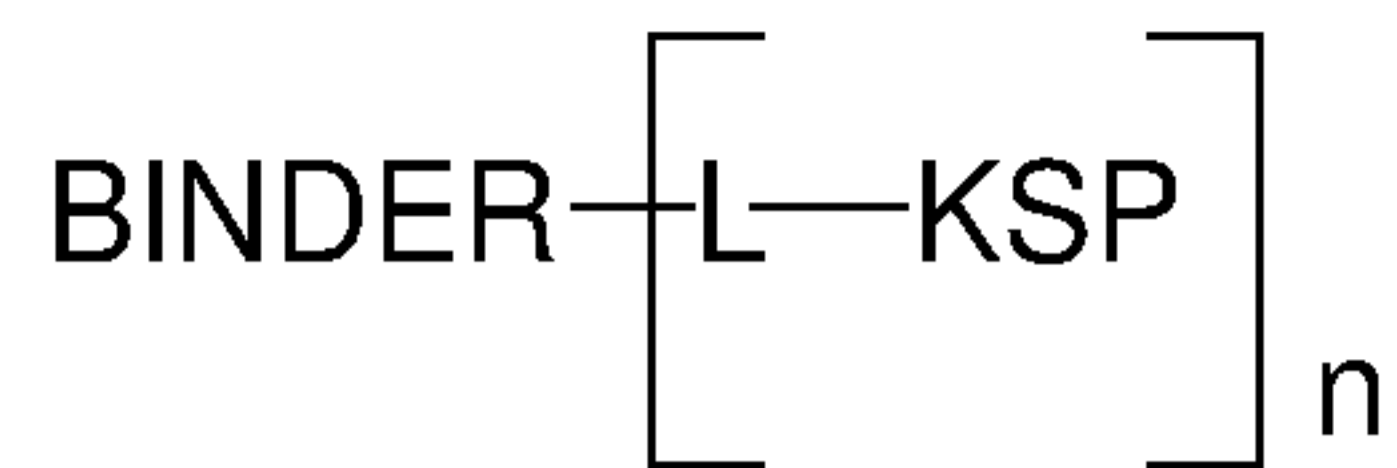
#<sup>1</sup> is the bond to the KSP inhibitor and #<sup>2</sup> is the bond to the  
 coupling group to the antibody (e.g. L2),

30

and salts, solvates and salts of the solvates of the ADC.

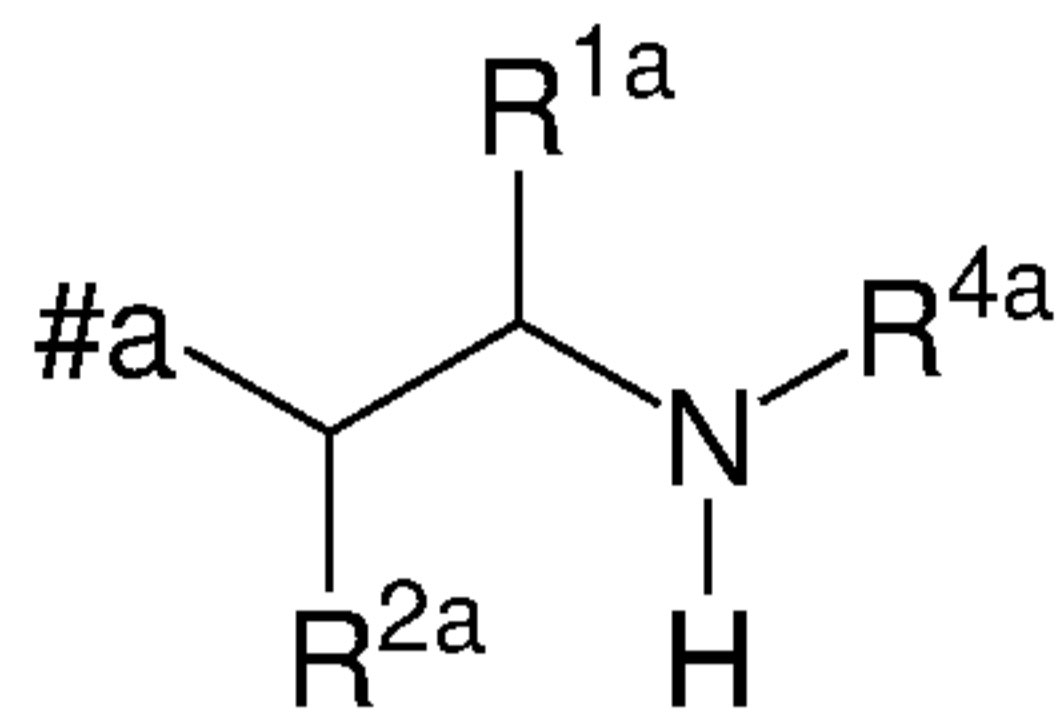
#### Embodiment E:

35 An ADC of the formula



where KSP-L- is a compound having the substructure I(sub) below,  
the binder is nimutuzumab and n is a number from 1 to 10:

5



where

10 #a represents a bond to the remainder of the molecule;

R<sup>1a</sup> represents -L-#1, H or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, halogen, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

15 where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub>, COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>,  
20 or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>;

R<sup>2a</sup> and R<sup>4a</sup> independently of one another represent H, -L-#1, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or R<sup>2a</sup> and R<sup>4a</sup> together represent (with  
25 formation of a pyrrolidine ring) -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents -L-#1, H, NH<sub>2</sub>, COOH, SO<sub>3</sub>H, SH or OH, and where Z represents -H, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  
30 -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH,

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by -

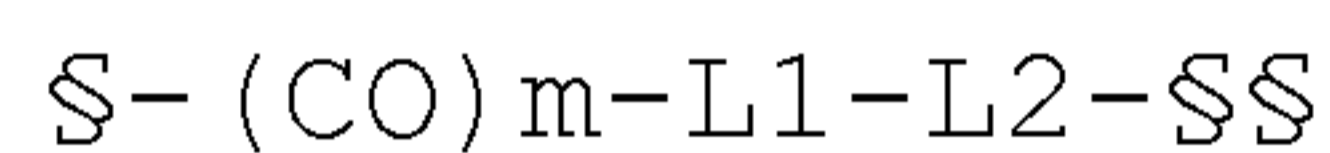
NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

5 where one of the substituents R<sup>1a</sup>, R<sup>2a</sup>, R<sup>4a</sup> or R<sup>10</sup> represents -L-  
#1,

-L- represents the linker and #1 represents the bond to the  
antibody,

10

where -L- is represented by



15 where

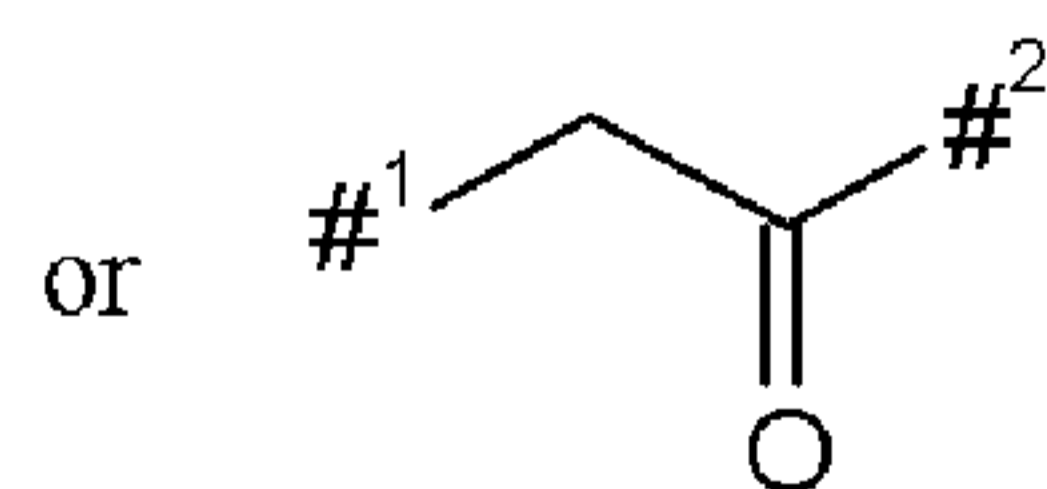
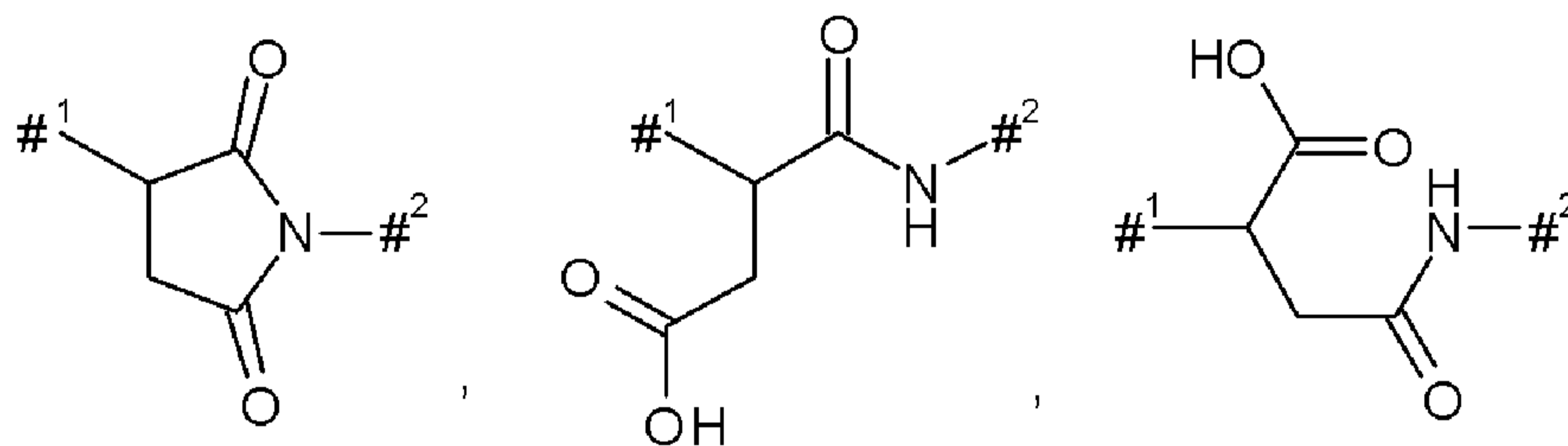
m is 0 or 1;

§ represents the bond to KSP and

20

§§ represents the bond to the antibody, and

L2 represents



25

where

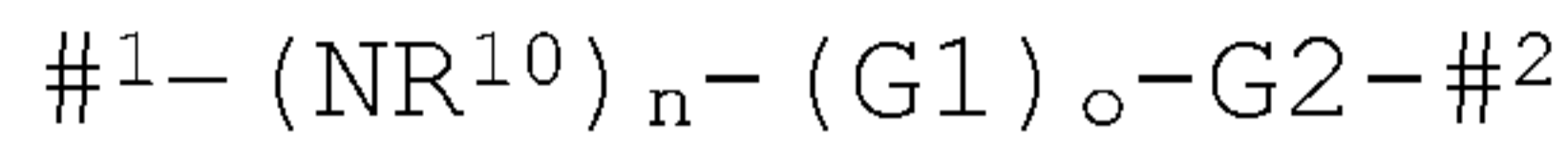
#<sup>1</sup> denotes the point of attachment to the sulphur atom of the



antibody,

#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>,

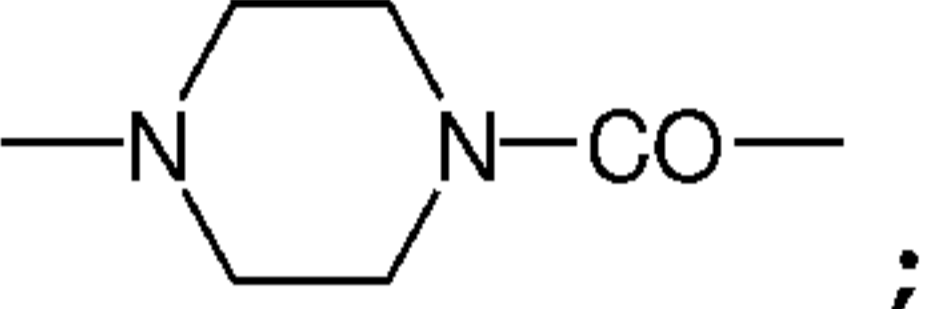
5 and L<sup>1</sup> is represented by formula



where

10

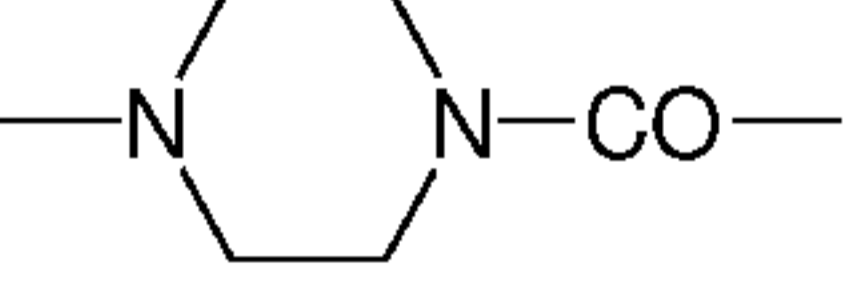
R<sup>10</sup> represents H, NH<sub>2</sub> or C<sub>1</sub>-C<sub>3</sub>-alkyl;

G1 represents -NHCO- or  ;

15 n is 0 or 1;

o is 0 or 1; and

G2 represents a straight-chain or branched hydrocarbon chain  
 20 having 1 to 100 carbon atoms from arylene groups and/or straight-  
 chain and/or branched and/or cyclic alkylene groups and which  
 may be interrupted once or more than once by one or more of the  
 groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-,  
 -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- and a 3- to 10-membered aromatic or  
 25 non-aromatic heterocycle having up to 4 heteroatoms selected  
 from the group consisting of N, O and S, or -SO- (preferably

) , where the side chains, if present, may be  
 substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>,  
 sulphonamide, sulphone, sulphoxide or sulphonic acid,

30

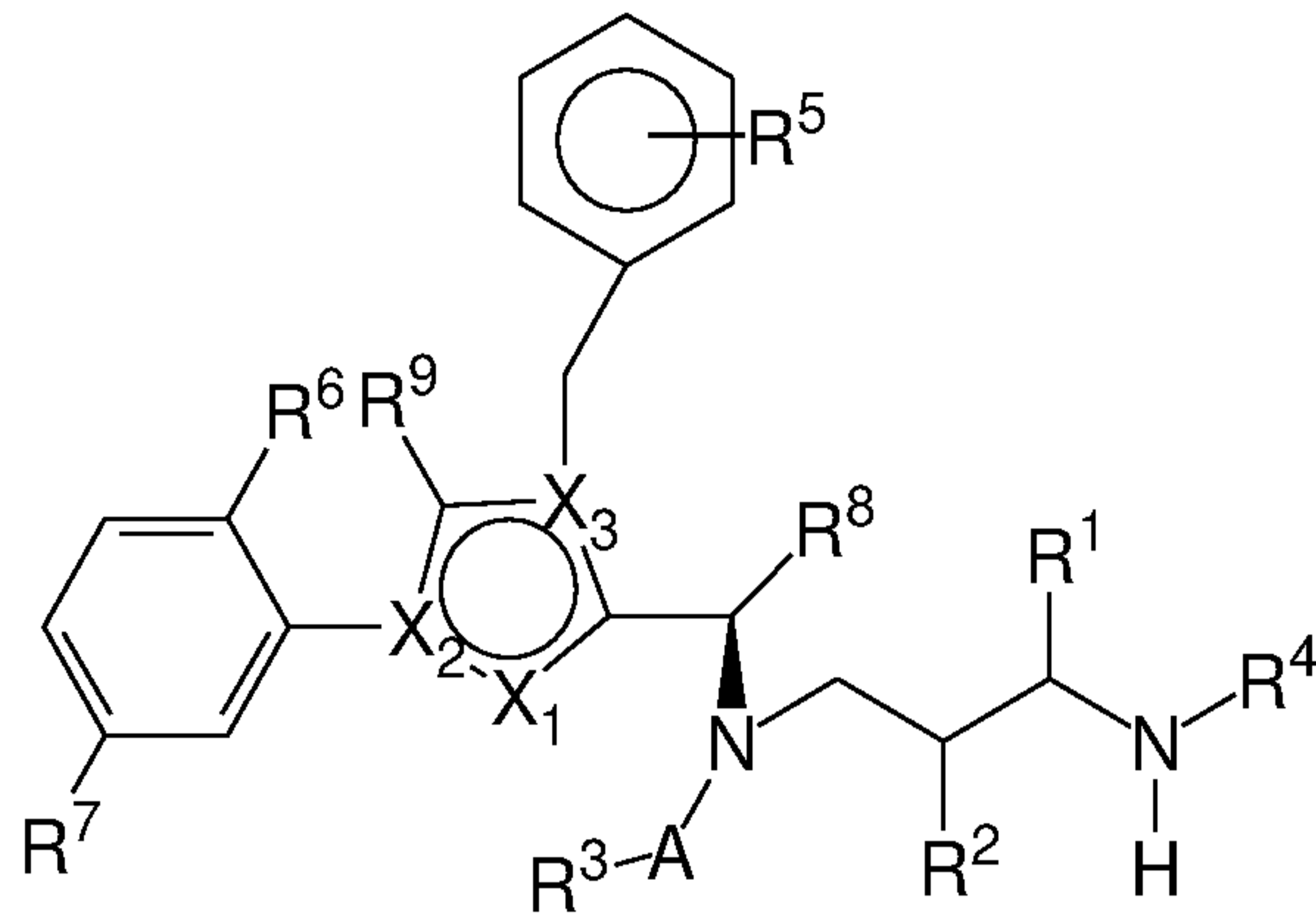
#<sup>1</sup> is the bond to the KSP inhibitor and #<sup>2</sup> is the bond to the  
 coupling group to the antibody (e.g. L<sub>2</sub>),

and salts, solvates and salts of the solvates of the ADC.

35

In this embodiment, KSP-L- particularly preferably has the  
 formula (IIIi) below:

formula (IIIi):



(IIIi)

5

where

$X_1$  represents N,  $X_2$  represents N and  $X_3$  represents C; or

10  $X_1$  represents CH or CF,  $X_2$  represents C and  $X_3$  represents N; or

$X_1$  represents NH,  $X_2$  represents C and  $X_3$  represents C; or

$X_1$  represents CH,  $X_2$  represents N and  $X_3$  represents C

15

(with  $X_1$  representing CH,  $X_2$  representing C and  $X_3$  representing N being preferred);

20  $R^1$  represents H, -MOD or  $-(CH_2)_{0-3}Z$ , where Z represents -H, -NH $Y^3$ , -O $Y^3$ , -S $Y^3$ , halogen, -CO-N $Y^1Y^2$  or -CO-O $Y^3$ ,

25 where  $Y^1$  and  $Y^2$  independently of one another represent H, NH $_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  (e.g.  $-(CH_2)_{0-3}Z'$ ) or  $-CH(CH_2W)Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H, NH $_2$ , SO $_3H$ , COOH, -NH-CO-CH $_2$ -CH $_2$ -CH(NH $_2$ )COOH or  $-(CO-NH-CHY^4)_{1-3}COOH$ , where W represents H or OH,

30 where  $Y^4$  independently of one another represents straight-chain or branched C $_1$ - $_6$ -alkyl which is optionally substituted by -NHCONH $_2$ , or represents aryl or benzyl which are optionally

substituted by  $-\text{NH}_2$ ;

$R^2$  represents H,  $-\text{MOD}$ ,  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  or  $-(\text{CH}_2)_{0-3}\text{Z}$ , or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring) represent  $-\text{CH}_2-$   
5  $\text{CHR}^{10}-$  or  $-\text{CHR}^{10}-\text{CH}_2-$ , where  $R^{10}$  represents H,  $\text{NH}_2$ ,  $\text{SO}_3\text{H}$ ,  $\text{COOH}$ ,  $\text{SH}$ ,  
or  $\text{OH}$ ;

where Z represents  $-\text{H}$ , halogen,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

10

where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$  or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , and  $\text{Y}^3$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$   
5 represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$ ;

15 where  $\text{Y}^4$  independently of one another represents straight-chain  
or branched  $\text{C}_{1-6}$  alkyl which is optionally substituted by  $-\text{NHCONH}_2$ , or represents aryl or benzyl which are optionally  
substituted by  $-\text{NH}_2$ , and  $\text{Y}^5$  represents H or  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ , where  
 $\text{Y}^6$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl;

20

$R^4$  represents H,  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  or  $-(\text{CH}_2)_{0-3}\text{Z}$ , preferably H,

where Z represents  $-\text{H}$ , halogen,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

25

where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$  or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , and  $\text{Y}^3$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$   
represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$ ;

30 where  $\text{Y}^4$  independently of one another represents straight-chain  
or branched  $\text{C}_{1-6}$  alkyl which is optionally substituted by  $-\text{NHCONH}_2$ , or represents aryl or benzyl which are optionally  
substituted by  $-\text{NH}_2$ , and  $\text{Y}^5$  represents H or  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ , where  
 $\text{Y}^6$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl;

35

or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring)  
represent  $-\text{CH}_2-\text{CHR}^{10}-$  or  $-\text{CHR}^{10}-\text{CH}_2-$ , where  $R^{10}$  represents H,  $\text{NH}_2$ ,  
 $\text{SO}_3\text{H}$ ,  $\text{COOH}$ ,  $\text{SH}$  or  $\text{OH}$ ;

A represents CO, SO, SO<sub>2</sub>, SO<sub>2</sub>NH or CNNH;

R<sup>3</sup> represents -L-#1,

5

R<sup>5</sup> represents H, NH<sub>2</sub>, NO<sub>2</sub>, halogen (in particular F, Cl, Br), -CN, CF<sub>3</sub>, -OCF<sub>3</sub>, -CH<sub>2</sub>F, -CH<sub>2</sub>F, SH or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

10 where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

15 R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano, (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy, NO<sub>2</sub>, NH<sub>2</sub>, COOH or halogen (in particular F, Cl, Br),

20 R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl or -(CH<sub>2</sub>)<sub>0-2</sub>(HZ<sup>2</sup>), where HZ<sup>2</sup> represents a 4- to 7-membered heterocycle having up to two heteroatoms selected from the group consisting of N, O and S, where each of these groups may be substituted by -OH,  
25 CO<sub>2</sub>H or NH<sub>2</sub>;

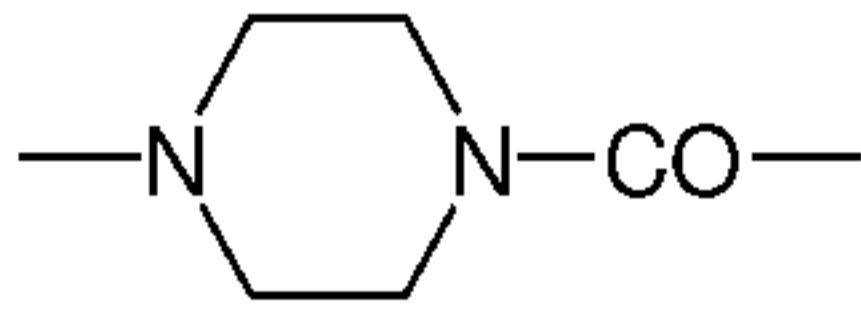
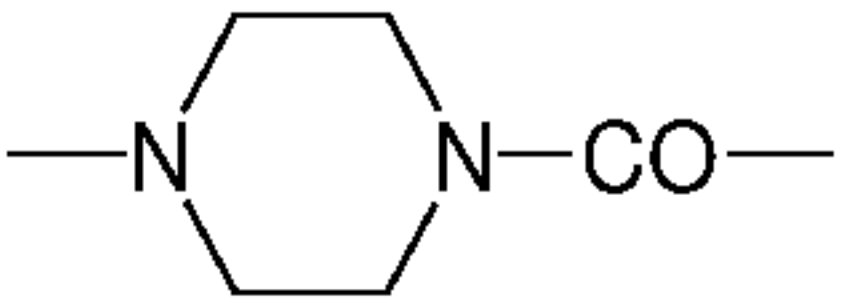
R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

30 L represents the linker and #1 represents the bond to the binder or derivative thereof,

where -MOD represents -(NR<sup>10</sup>)<sub>n</sub>-(G1)<sub>o</sub>-G2-H, where

R<sup>10</sup> represents H or C<sub>1</sub>-C<sub>3</sub>-alkyl;

35

G1 represents -NHCO- , -CONH- or  (where, if G1 represents -NHCO- or  , R<sup>10</sup> does not represent NH<sub>2</sub>);

n is 0 or 1;

o is 0 or 1; and

5

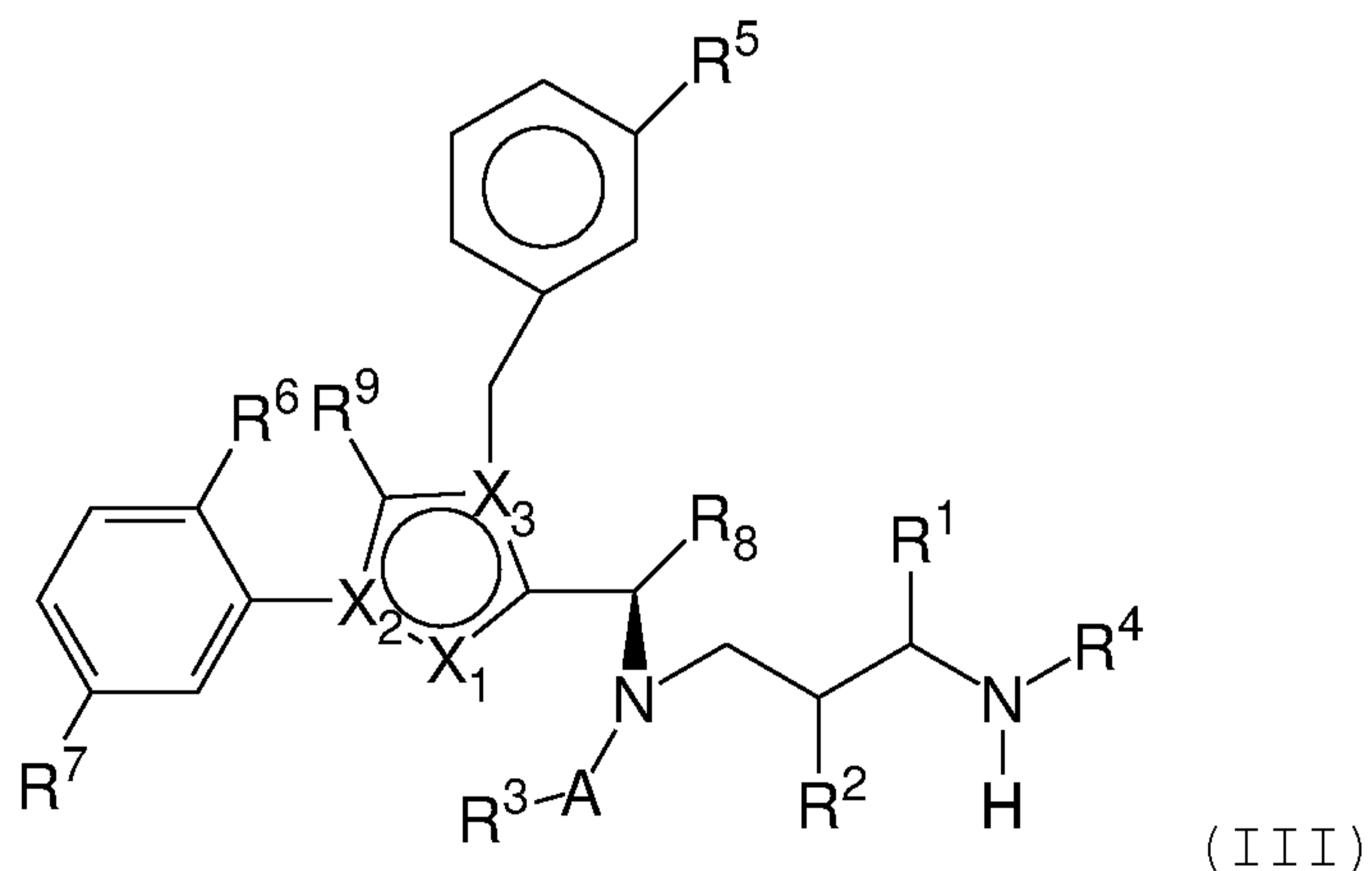
G2 represents a straight-chain and/or branched hydrocarbon group which has 1 to 10 carbon atoms and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>-  
 10 (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl), where the hydrocarbon chain including any side  
 15 chains may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, where the group -MOD preferably has at least one group -COOH;

and the salts, solvates and salts of the solvates thereof.

20

Embodiment F:

Compounds of the general formula:



25

where

X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C, or X<sub>1</sub>  
 30 represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represents N;

$R^1$  represents H or  $-(CH_2)_{0-3}Z$ , where Z represents -H,  $-NHY^3$ ,  $-OY^3$ ,  $-SY^3$ , halogen,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

5 where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$ ,  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$  or  $-CH(CH_2W)Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $NH_2$ ,  $SO_3H$ ,  $COOH$ ,  $-NH-CO-CH_2-CH_2-CH(NH_2)COOH$  or  $-(CO-NH-CHY^4)_{1-3}COOH$ ; where W represents H or OH;

10

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$ -alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ ;

15

$R^2$  and  $R^4$  independently of one another represent H,  $-CO-CHY^4-NHY^5$  or  $-(CH_2)_{0-3}Z$ , or  $R^2$  and  $R^4$  together (with formation of a pyrrolidine ring) represent  $-CH_2-CHR^{10}-$  or  $-CHR^{10}-CH_2-$ , where  $R^{10}$  represents H,  $NH_2$ ,  $SO_3H$ ,  $COOH$ , SH or OH,

20

where Z represents -H, halogen,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,

25 where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ;

30 where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ , and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

A represents CO, SO,  $SO_2$ ,  $SO_2NH$  or CNNH;

35

$R^3$  represents an optionally substituted alkyl, aryl, heteroaryl, heteroalkyl, heterocycloalkyl group, preferably a  $C_{1-10}$ -alkyl,  $C_{6-10}$ -aryl or  $C_{6-10}$ -aralkyl,  $C_{5-10}$ -heteroalkyl,  $C_{1-10}$ -alkyl-O- $C_{6-10}$ -

aryl or C<sub>5-10</sub>-heterocycloalkyl group which may be substituted by 1-3 -OH groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH,

(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl);

R<sup>5</sup> represents H, F, NH<sub>2</sub>, NO<sub>2</sub>, halogen, SH or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano, (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy or halogen,

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl or optionally substituted oxetane; and

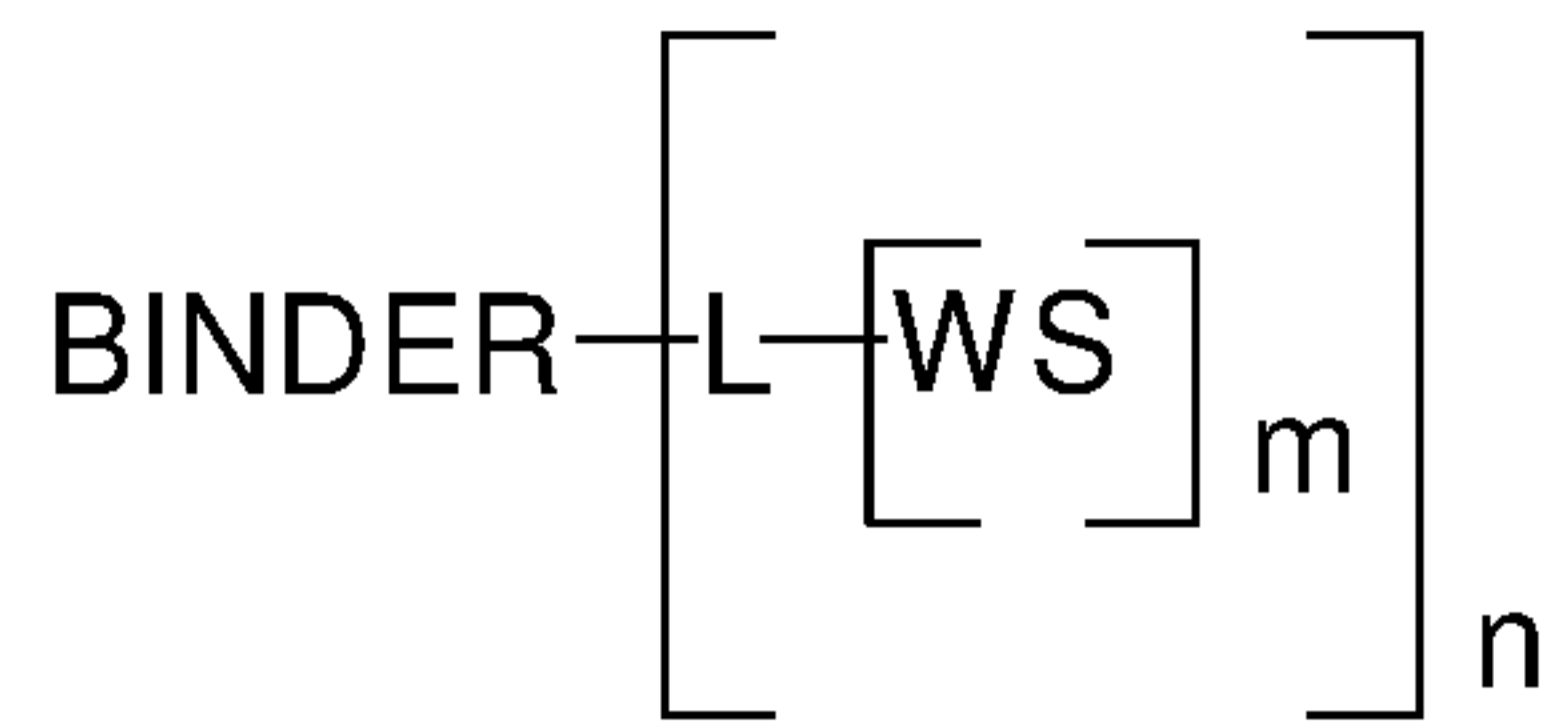
R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

and the salts, solvates and salts of the solvates thereof.

#### Embodiment G:

The invention also provides binder/active compound of the

general formula below:



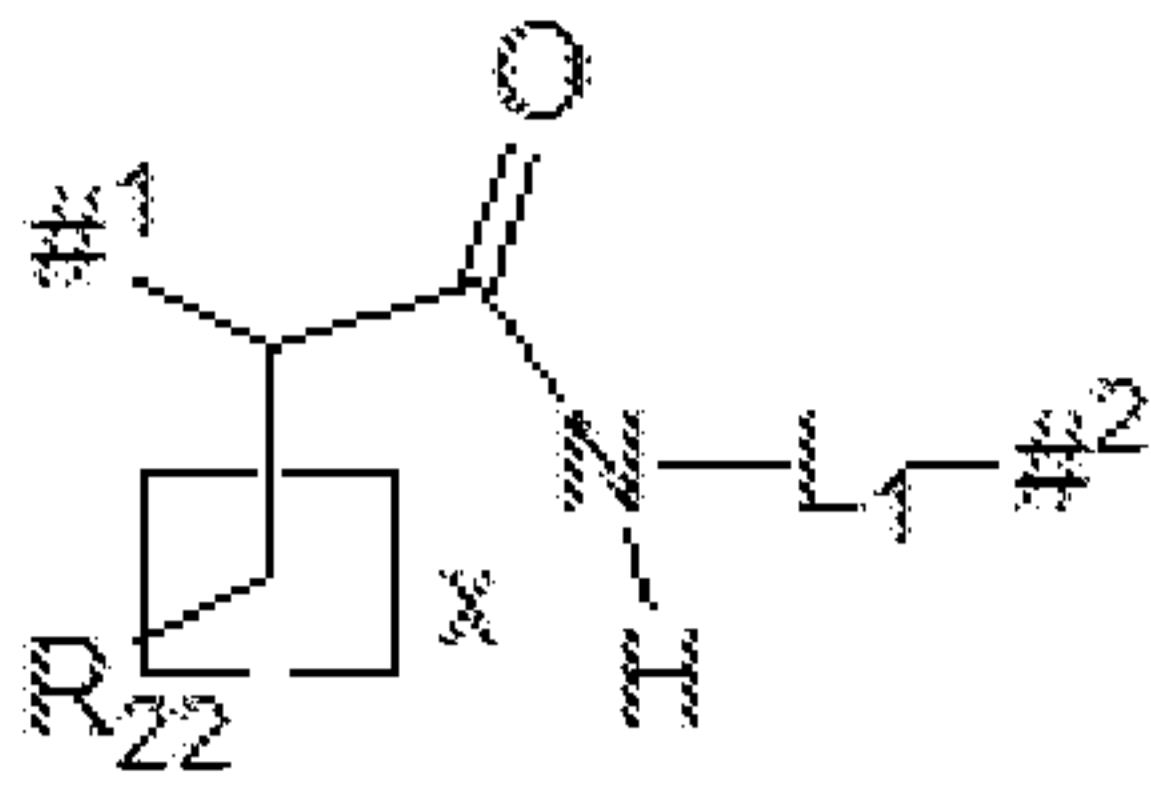
5 where BINDER represents the binder (preferably: antibody) or a derivative thereof (preferably: cysteine residue), preferably an antibody, L represents the linker, WS represents the active compound, preferably a KSP inhibitor such as, for example, a KSP inhibitor according to the invention of one of the formulae  
 10 (II), (IIa), (IIb), (IIc), (IId), (IIe), (IIf), (IIg) or (IIh), m represents a number from 1 to 2, preferably 1, and n represents a number from 1 to 50, preferably from 1.2 to 20 and particularly preferably from 2 to 8, where L has one of the structures below. Here, m is the number of active compound molecules per linker  
 15 and n a mean of the number of active compound/linker conjugates per BINDER. The sum of all WS present in a conjugate molecule is therefore the product of m and n.

WS is an active compound which has local or systemic therapeutic  
 20 action in animals, preferably in humans. These active compounds generally have a molecular weight below 5 kDa, preferably below 1.5 kDa. Preferred active compounds are antiproliferative substances, for example cytotoxic or cytostatic substances. Preferred active compounds are cytotoxic substances, inhibitors  
 25 of angiogenesis, cell cycle inhibitors, PI3 kinase or m-TOR inhibitors, inhibitors of the MAPK signalling cascade pathway, HDAC inhibitors, proteasome inhibitors, PARP inhibitors, Wnt/Hedgehog signal cascade path inhibitors and RNA polymerase inhibitors. Cytotoxic substances are, *inter alia*, DNA-binding  
 30 or intercalating substances, DNA-alkylating substances, microtubulin-stabilizing substances or -destabilizing substances, platinum compounds and topoisomerase I inhibitors. Exemplary DNA-binding substances are, for example, anthracyclins such as doxorubicin or daunorubicin). Exemplary DNA-alkylating  
 35 substances are, for example, calicheamicins, temozolomide or cyclophosphamide and derivatives. Exemplary microtubulin-

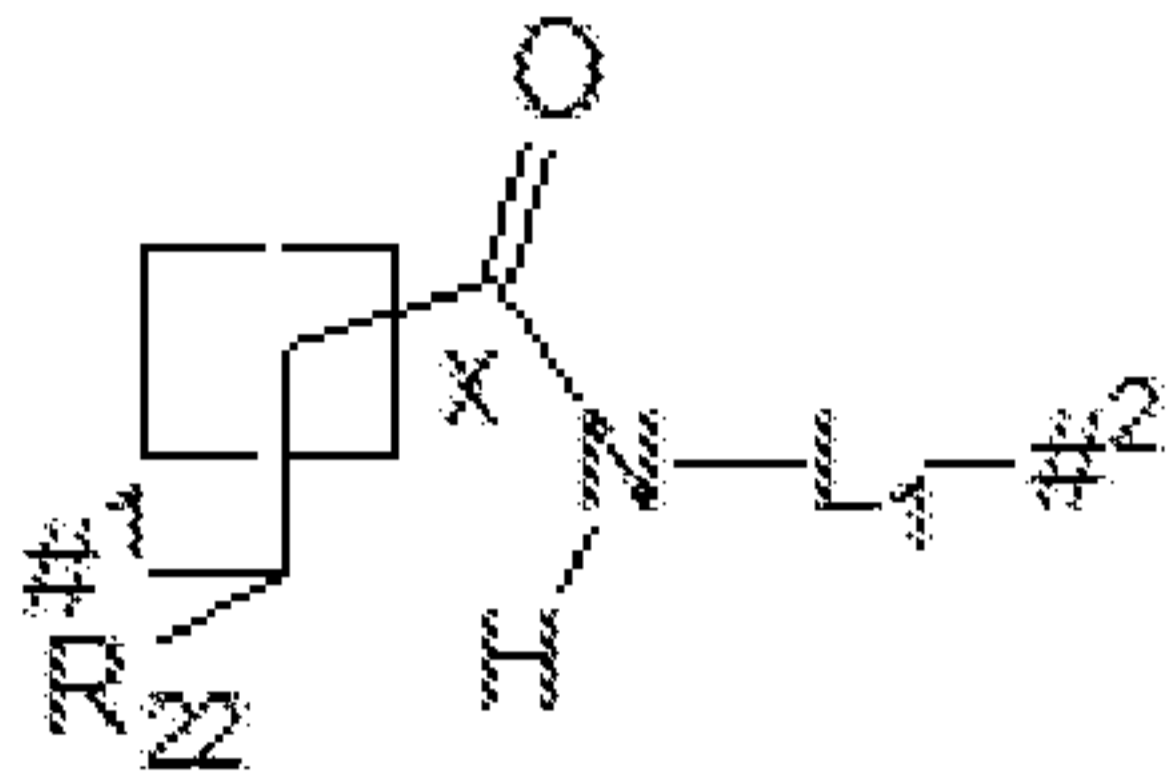


stabilizing or -destabilizing substances are, for example, taxanes such as paclitaxel, docetaxel, maytansinoides and auristatins, tubulysines, vinca alkaloids, epothilones and derivatives thereof. Examples of maytansinoides are maytansins, 5 maytansinols, DM-1 and DM-4 (see, for example, US patent US5208020). Examples of auristatins are auristatin E, monomethylauristatin E (MMAE), auristatin F, monomethylauristatin F (MMAF) and dolastin (see, *inter alia*, WO 09/117531, WO 2005/081711, WO 04/010957; WO02/088172 or 10 WO01/24763). Examples of vinca alkaloids are vincristine and vinblastine. Examples of epothilones are epothilone A, B, C, D, E or F (see, *inter alia*, WO 98/13375; WO2004/005269; WO 2008/138561; WO 2009/002993; WO 2009/055562; WO 2009/012958; WO2009/026177; WO 2009/134279; WO 2010/033733; WO2010/034724; 15 WO 2011/017249; WO2011/057805). Examples of platinum compounds are cisplatin and carboplatin. Examples of topoisomerase I inhibitors are camptothecin and derivatives. Examples of inhibitors of angiogenesis are MetAP2 inhibitors such as, for example, fumagillol. Examples of cell cycle inhibitors are CDK 20 inhibitors (e.g. BMS-387032 or PD0332991), Rho kinase inhibitors such as, for example, GSK429286, PLK inhibitors such as, for example, volasertib, aurora kinase inhibitors such as, for example, AZD1152 or MLN805Z. Examples of inhibitors of the MAPK signalling cascade pathway are *inter alia* MEK inhibitors (e.g. 25 PD0325901), Ras inhibitors, JNK inhibitors, B-Raf inhibitors (e.g. SB590885) or p38 MAPK inhibitors (e.g. SB202190). Examples of HDAC inhibitors are belinostat and givinostat. Examples of PARP inhibitors are iniparib and olaparib. Examples of RNA polymerase inhibitors are amatoxins such as, for example, alpha- 30 amantin, amanin and amanullin. Particularly preferred active compounds are vinca alkaloids, auristatins, maytansinoides, tubulysins, duocarmycins, kinase inhibitors, MEK inhibitors and KSP inhibitors.

35 Here, L represents one of the formulae A3 and A4 below



Formula A3

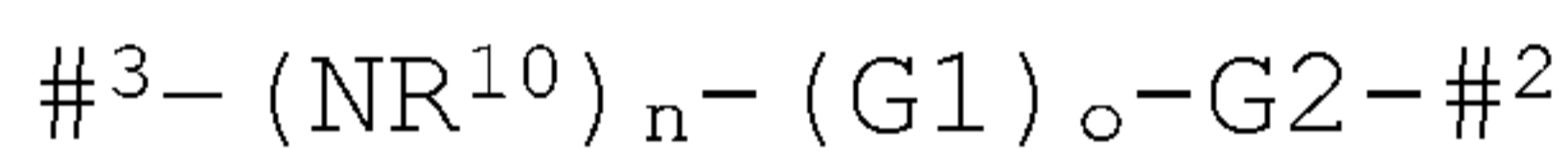


Formula A4

5 where #<sup>1</sup> denotes the point of attachment to the sulphur atom of the binder, #<sup>2</sup> denotes the point of attachment to the active compound, x represents 1 or 2, and R<sup>22</sup> represents COOH, COOR, COR (where R in each case represents C<sub>1-3</sub>-alkyl), CONH<sub>2</sub>, Br, preferably COOH.

10

L1 has the same meaning as above. Preferably, -L1-#<sup>2</sup> is represented by the formula below:

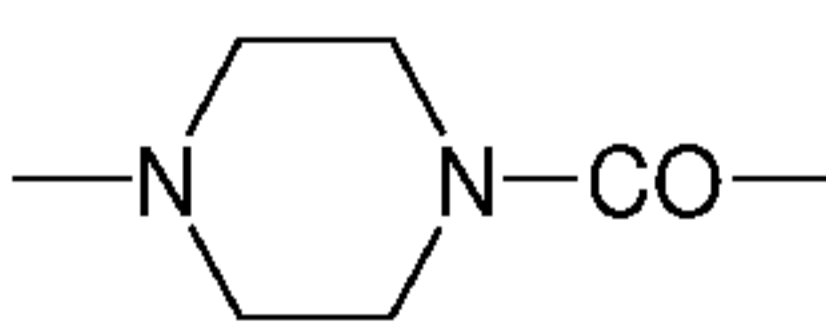
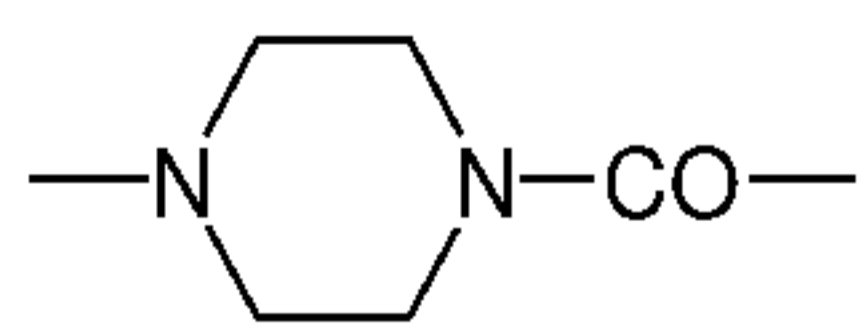


15

where

#<sup>3</sup> denotes the point of attachment to the nitrogen atom,

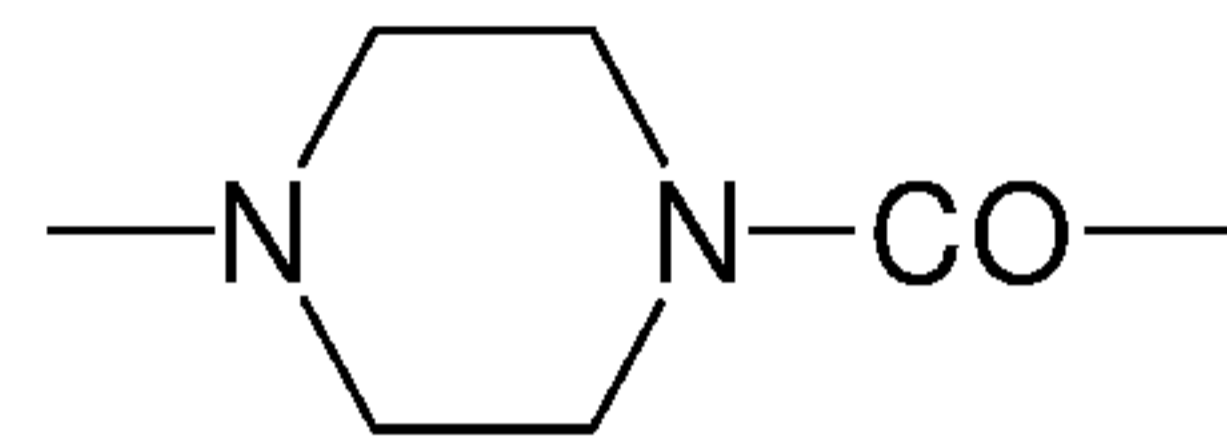
20 R<sup>10</sup> represents H, NH<sub>2</sub> or C<sub>1-3</sub>-alkyl;

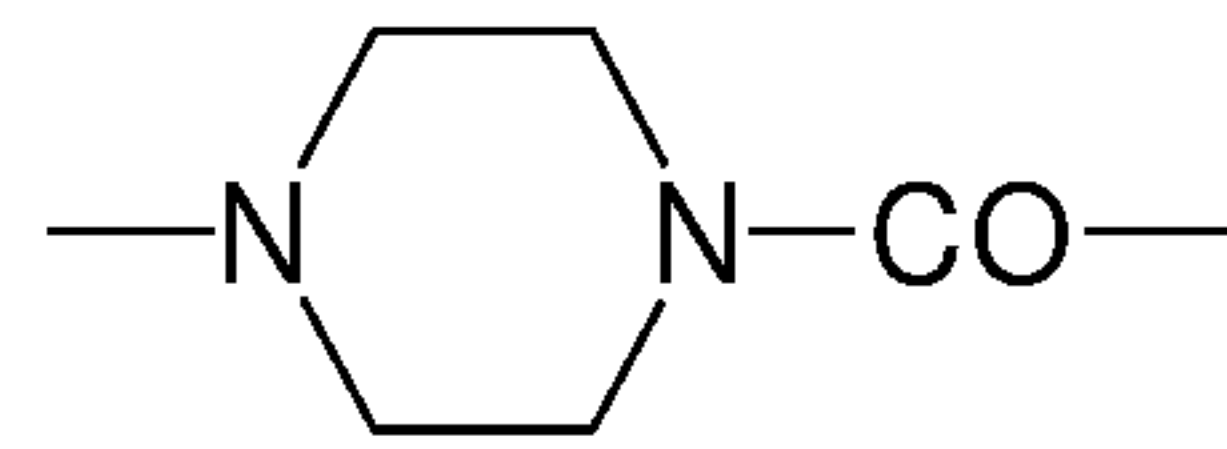
G1 represents -NHCO-, -CONH- or  (where, if G1 represents NHCO or , R<sup>10</sup> does not represent NH<sub>2</sub>),

25 n is 0 or 1;

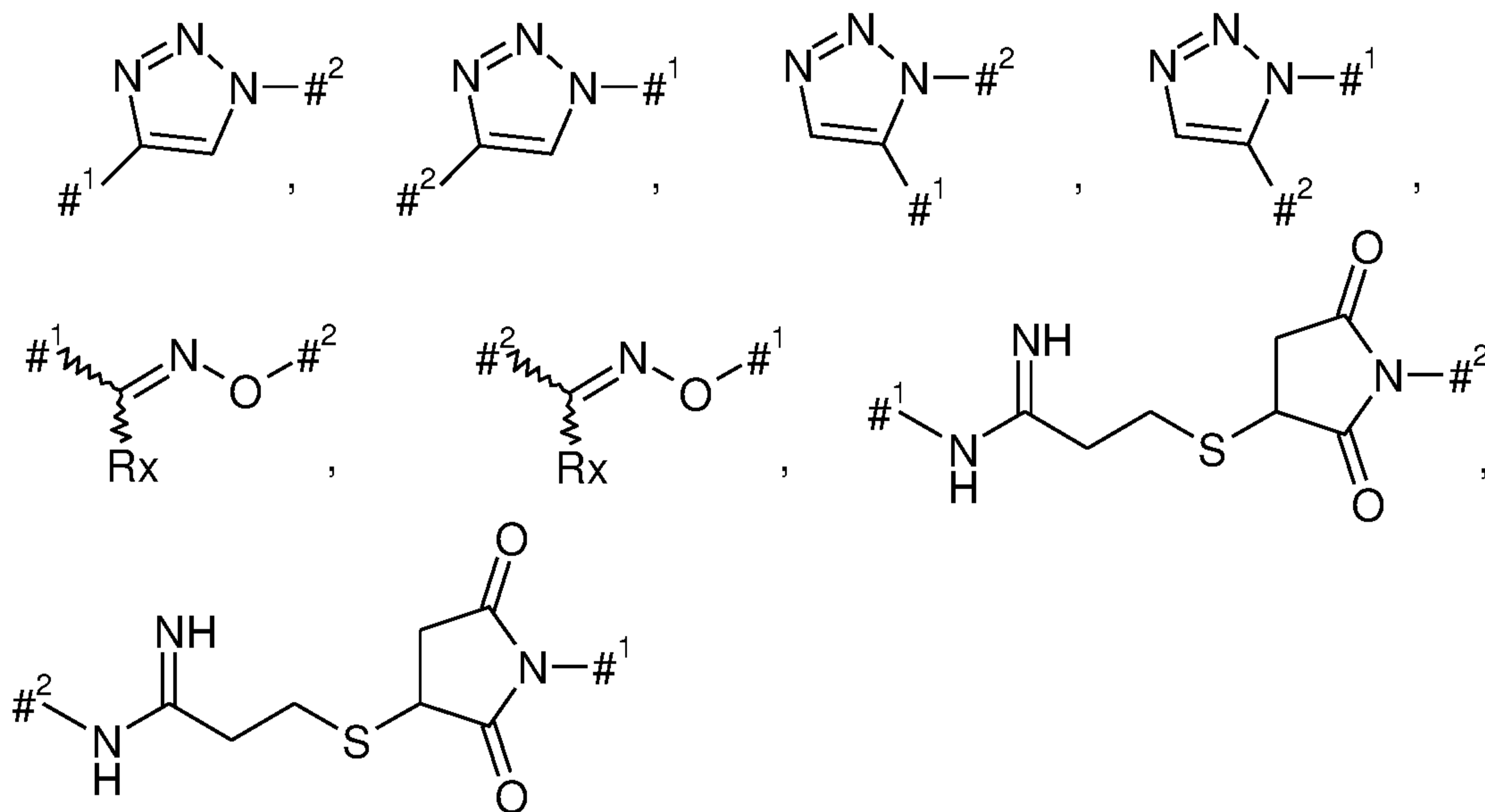
o is 0 or 1; and

G2 represents a straight-chain or branched hydrocarbon chain which has 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more  
 5 of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, -C(NH)NR<sup>y</sup>-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>- (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -  
 10 CR<sup>x</sup>=N-O- (where R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl) and/or a 3- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O



and S, -SO- or -SO<sub>2</sub>- (preferably ) , where the hydrocarbon chain including the side chains, if present, may be  
 15 substituted by NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid.

Further interrupting groups in G2 are preferably



20

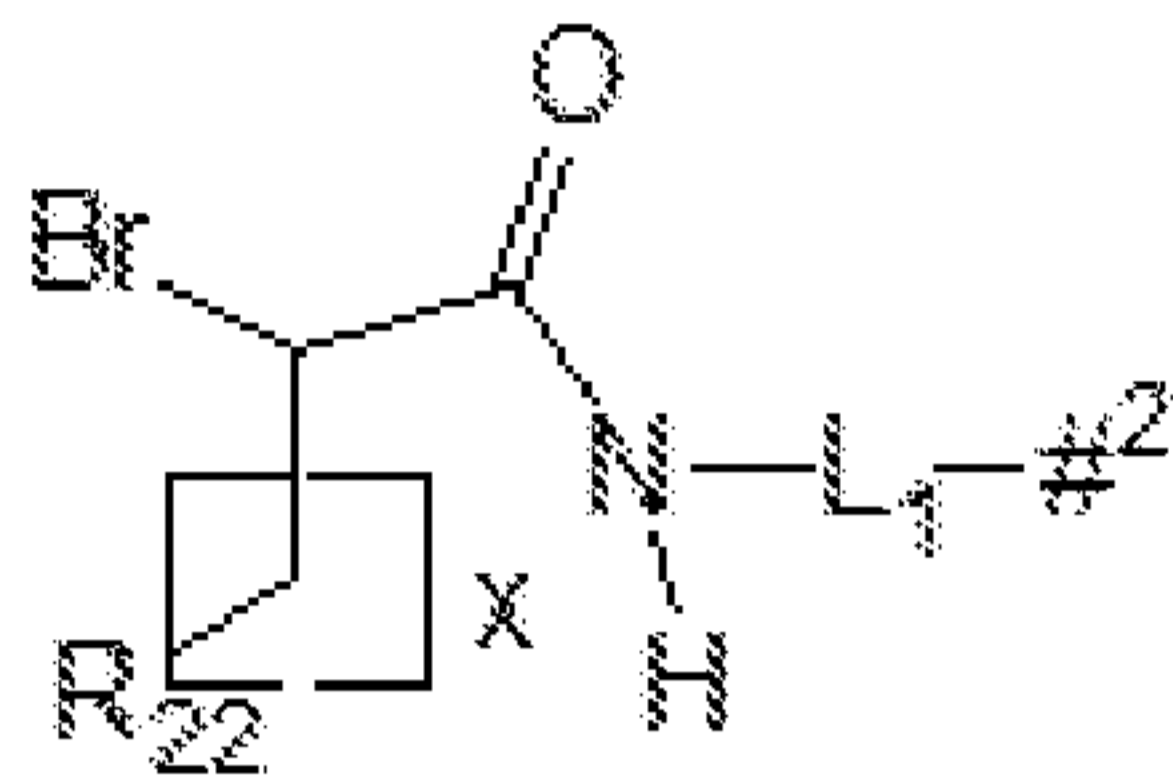
where Rx represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl.

In the conjugate according to the invention or in a mixture of  
 25 the conjugates according to the invention, the bonds to a cysteine residue of the binder are present, to an extent of

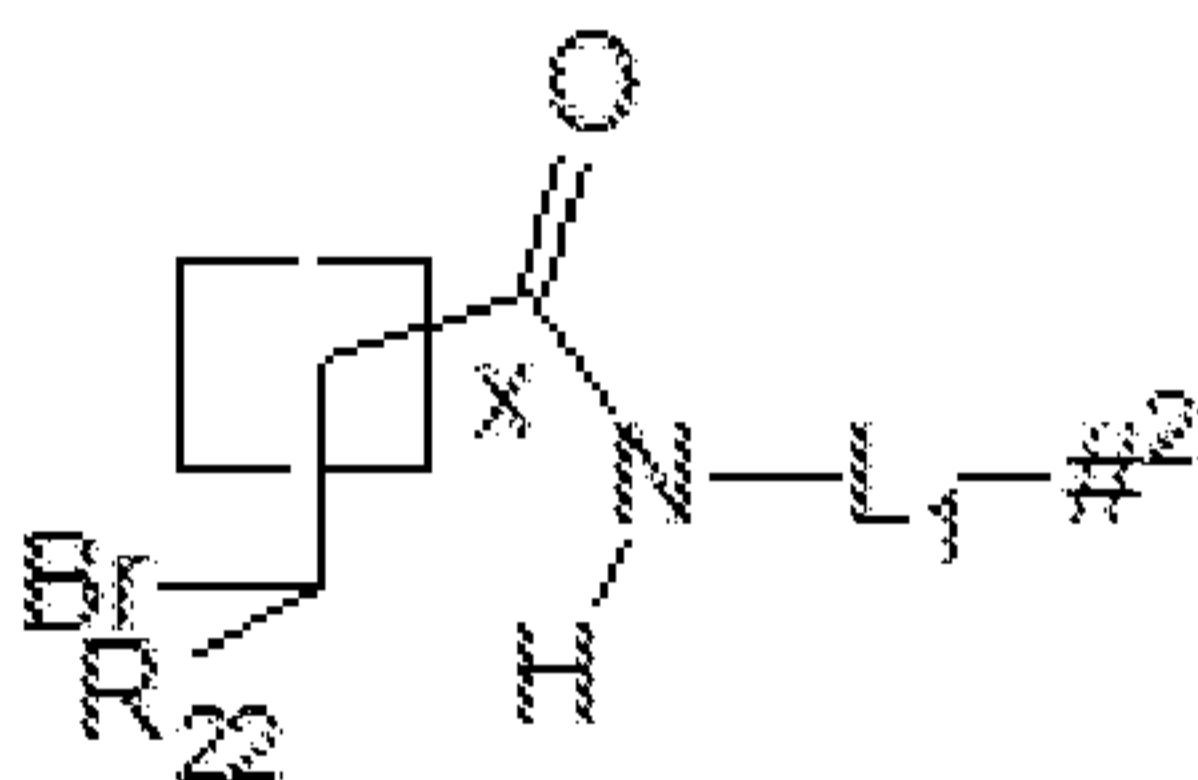
preferably more than 80%, particularly preferably more than 90% (in each case based on the total number of bonds of the linker to the binder) as one of the two structures of the formula A3 or A4.

5

The conjugates with the linkers of formula A3 or A4 can be obtained by coupling the binders to the appropriate bromine derivatives of the formulae A3' and A4', respectively, below:



10 Formula A3'

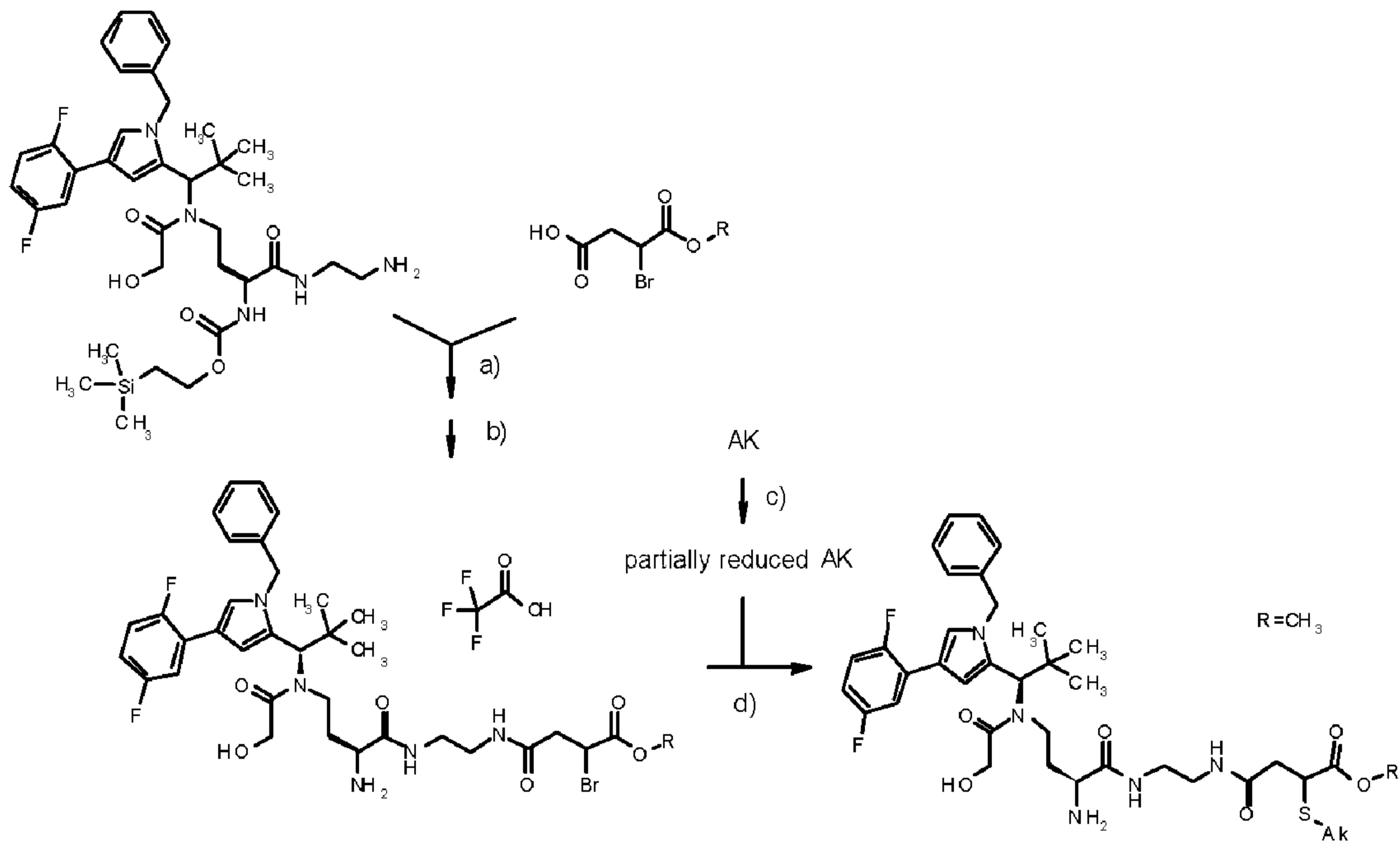


Formula A4'

These bromine derivatives of the formula A3' or A4' can be obtained by reacting HOOCCH<sub>2</sub>CHBrCOOR<sub>22</sub> or HOOCCHBrCH<sub>2</sub>COOR<sub>22</sub> with an amine group of the binder, as illustrated in an exemplary manner in Schemes 30 to 32 below.

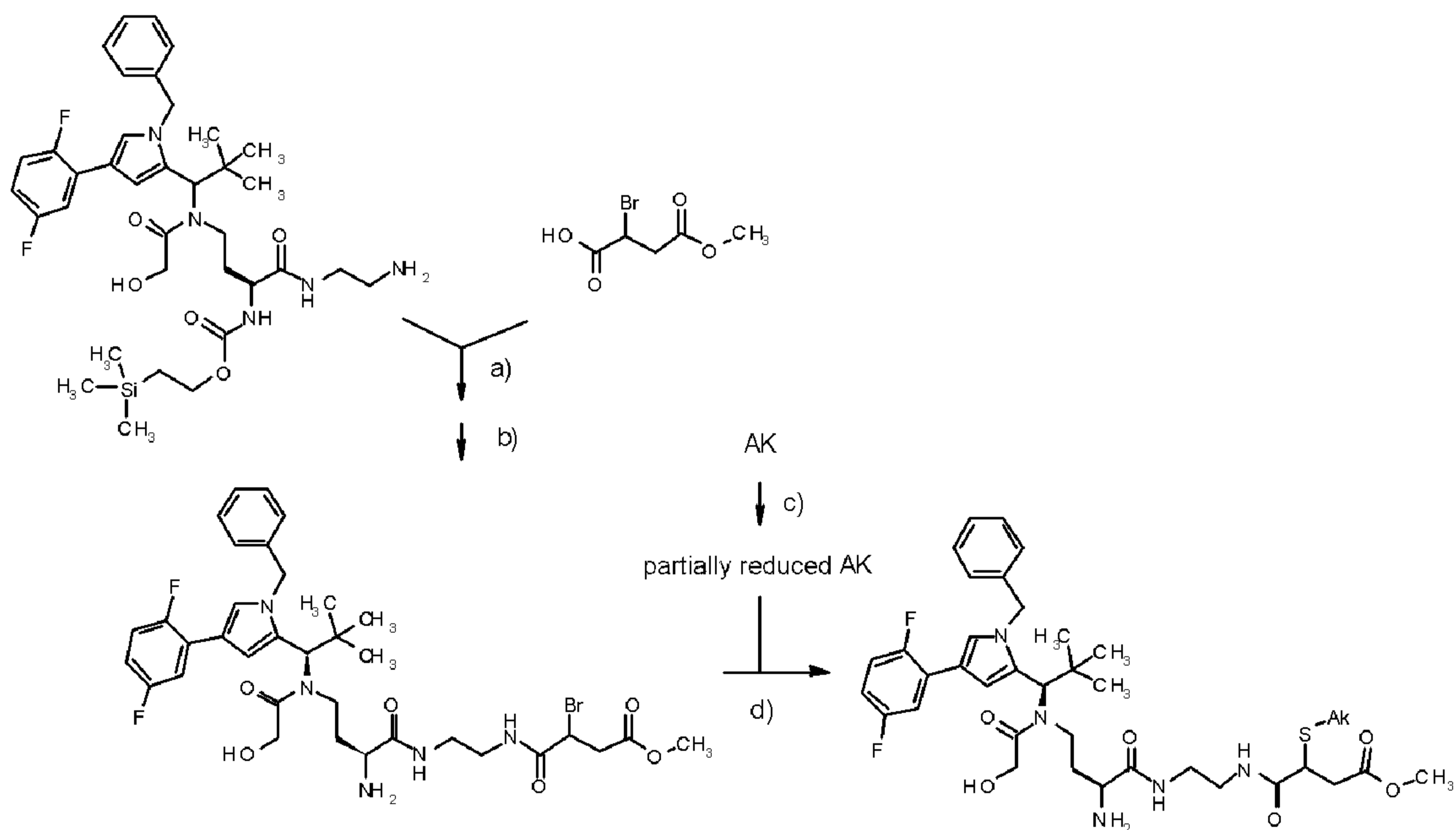
Scheme 30:

20



- [a): 2-bromo-1-ethylpyridinium tetrafluoroborate (BEP), DCM, pyridine, RT; b) zinc chloride, trifluoroethanol, 50°C, EDTA; c) 3-4 equivalents of TCEP, PBS buffer; d) PBS buffer, 20h RT.]

### Scheme 31:

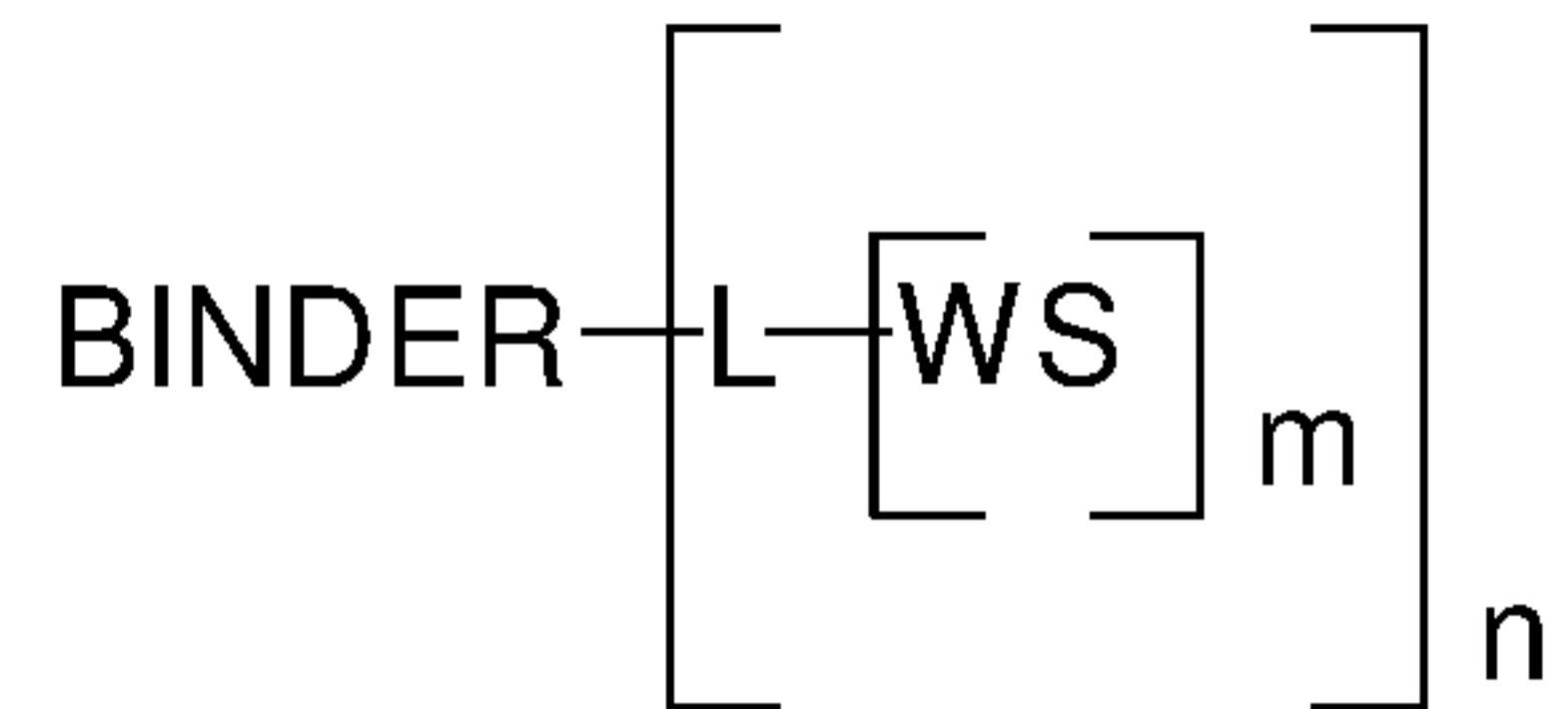


10

- [a): 2-bromo-1-ethylpyridinium tetrafluoroborate (BEP), DCM, pyridine, RT; b) zinc chloride, trifluoroethanol, 50°C, EDTA; c) 3-4 equivalents of TCEP, PBS buffer; d) PBS buffer, 20h RT.]

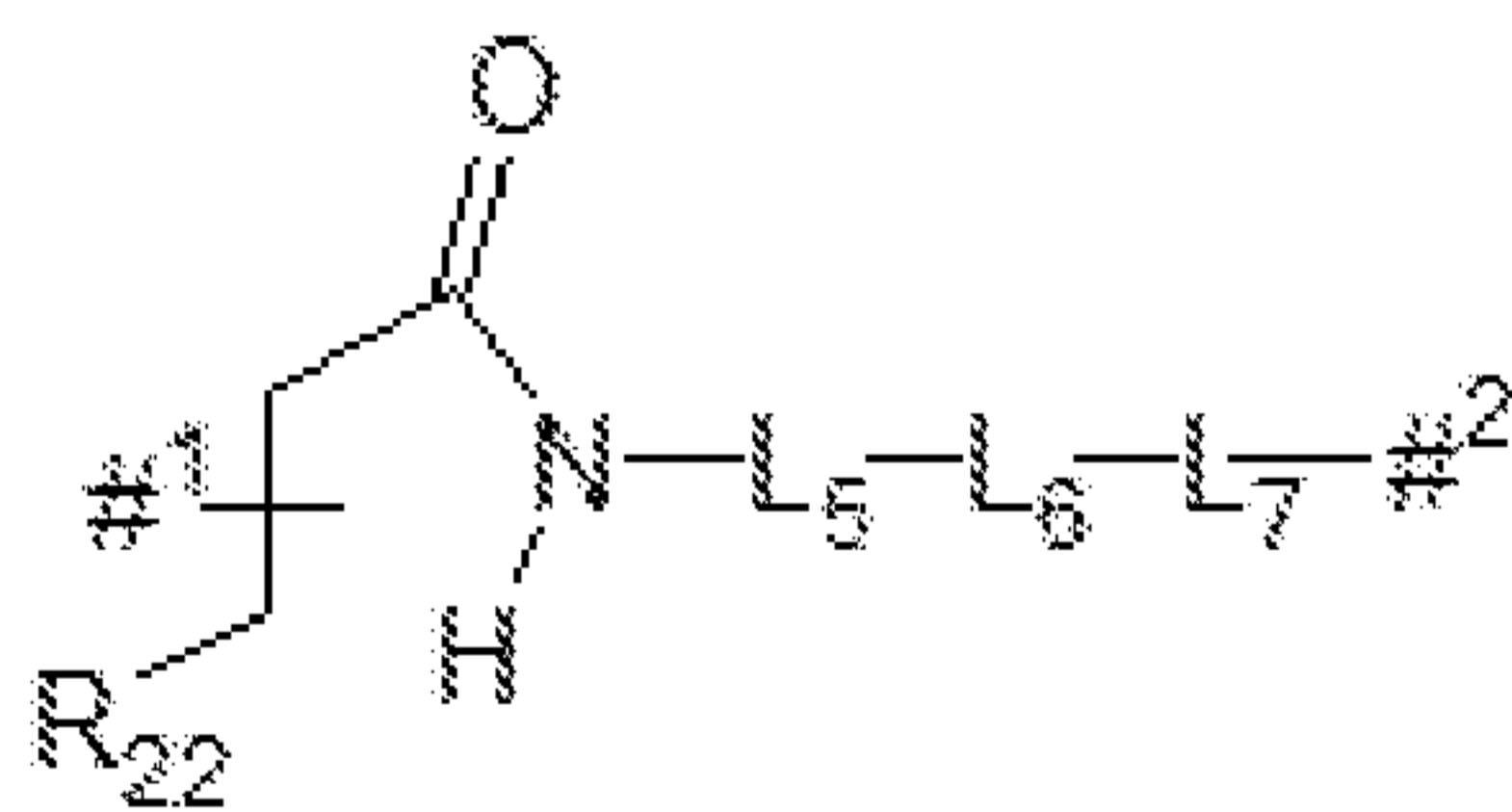
Embodiment H:

The invention also provides binder/active compound of the  
5 general formula below:



where BINDER represents the binder (preferably: antibody) or a  
10 derivative thereof (preferably: cysteine residue), preferably  
an antibody, L represents the linker, WS represents the active  
compound, preferably a KSP inhibitor such as, for example, a KSP  
inhibitor according to the invention of one of the formulae  
15 (II), (IIa), (III) or (IIIa), m represents a number from 1 to  
2, preferably 1, and n represents a number from 1 to 50,  
preferably from 1.2 to 20 and particularly preferably from 2 to  
8, where L has one of the structures below. Here, m is the number  
of active compound molecules per linker and n a mean of the  
20 number of active compound/linker conjugates per BINDER. The sum  
of all WS present in a conjugate molecule is therefore the  
product of m and n.

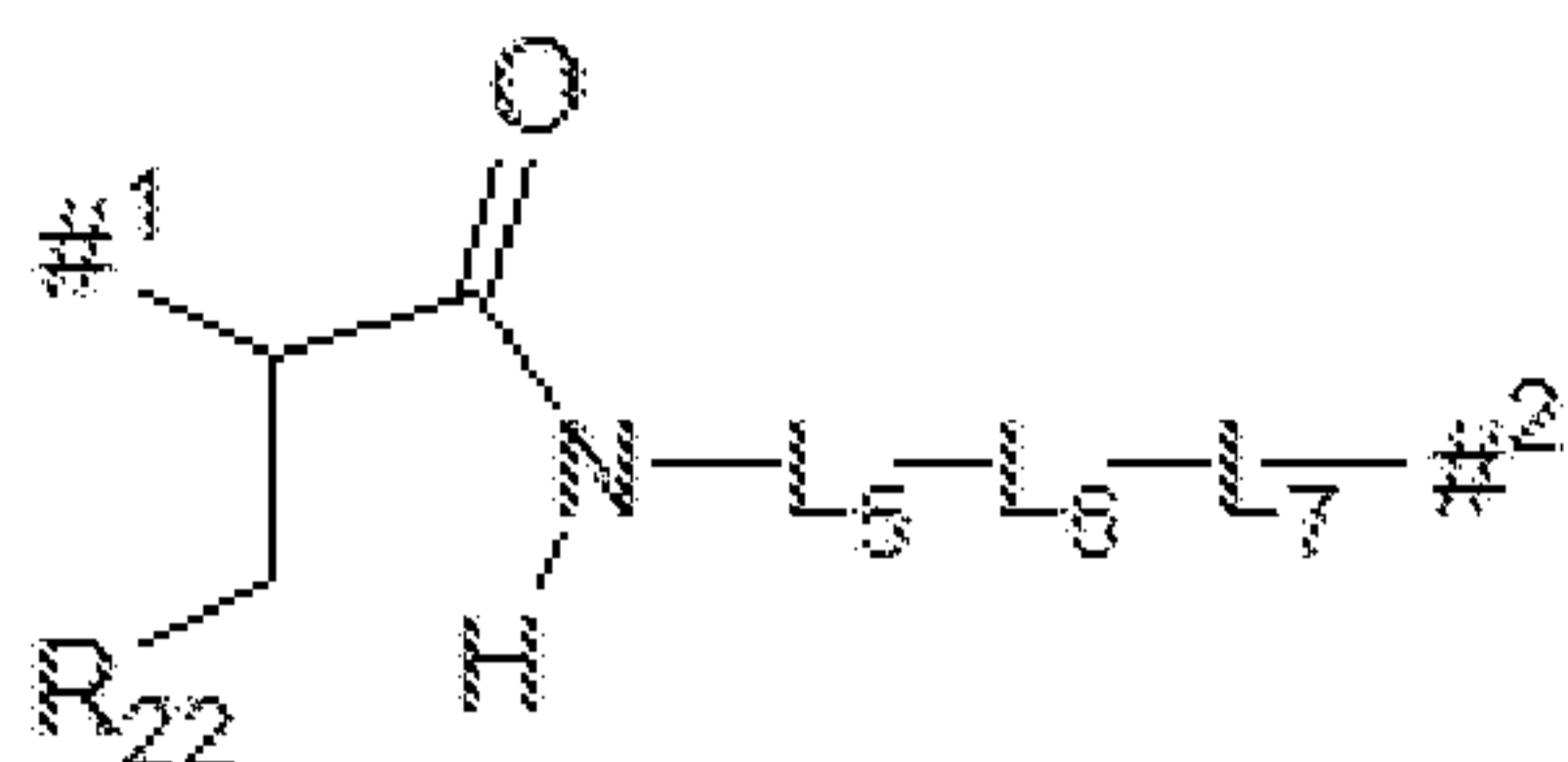
Here, L represents:



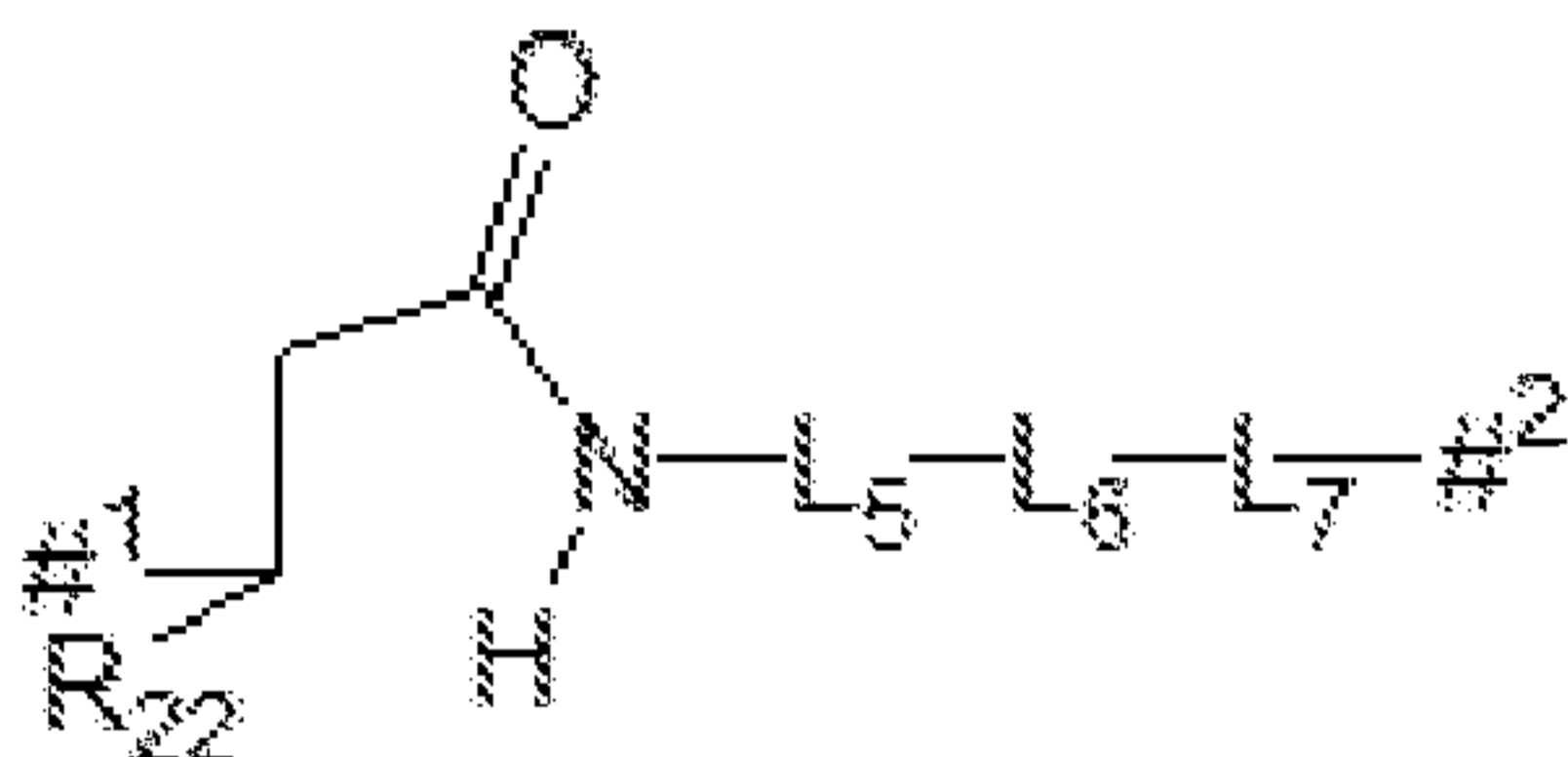
25 Formula A

where #<sup>1</sup> denotes the point of attachment to the sulphur atom of  
the binder, #<sup>2</sup> denotes the point of attachment to the active  
compound and R<sup>22</sup> represents COOH, COOR, COR (where R in each case  
30 represents C<sub>1-3</sub>-alkyl), CONH<sub>2</sub>, Br, preferably COOH. The link to  
the sulphur atom of the binder may thus have one of the

structures below:



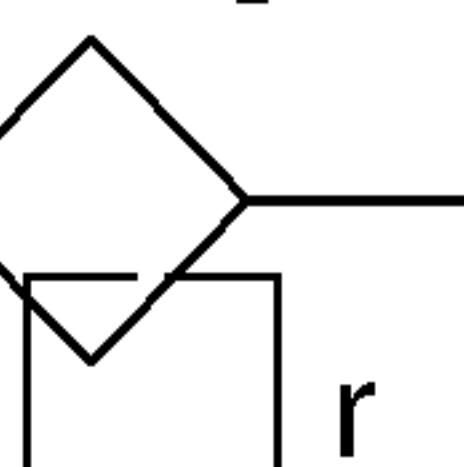
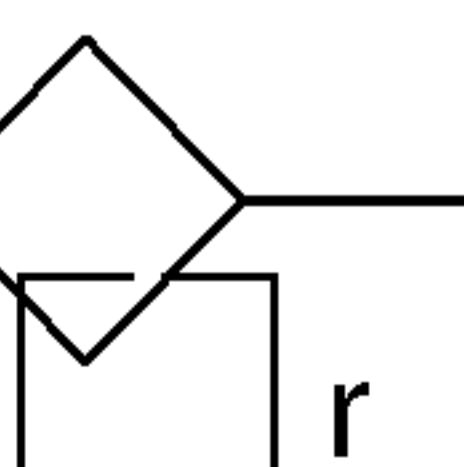
Formula A1



5 Formula A2

In the case of active compound/binder conjugates containing more than one active compound molecule WS per active compound/binder conjugate, both structures according to the formulae A1 and/or  
 10 A2 may be present in an active compound/binder conjugate. Since the active compound/binder conjugates according to the invention may be mixtures of different active compound/binder conjugates, it is also possible for this mixture to comprise both active compound/binder conjugates of formula A1 or formula A2 and those  
 15 of formula A1 and A2.

L<sub>5</sub> is a group selected from  $-(CH_2)_m-(CHR^S)_n-(OCH_2CH_2)_o-(X)_p-(CH_2)_q-$ , where m, n, o, p and q independently of one another have the following values: m=0-10; n=0 or 1; o=0-10; p=0 or 1; and q=0-  
 20 10, where m+n+o=1-15, preferably 1-6. X represents a 5- or 6-membered aromatic or nonaromatic hetero- or homocycle, preferably  $-C_6H_4-$  or  $-C_6H_{10}-$ . R<sup>S</sup> represents an acid group, preferably  $-COOH$  or  $SO_3H$ .

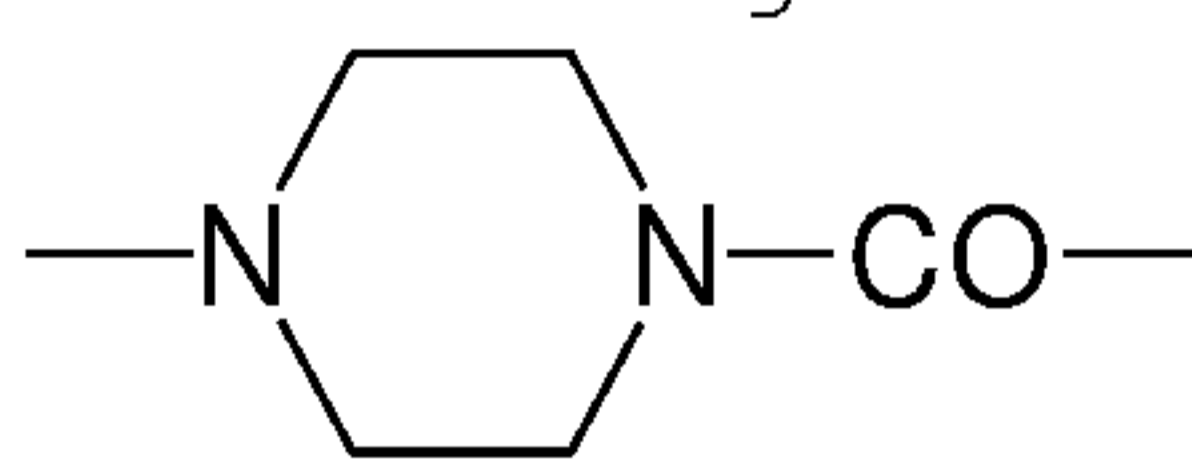
25 L<sub>6</sub> is a group selected from  $-CONH-$ ,  $-OCONH-$ ,  $-NHCO-$ ,  $-NHC(O)O-$ ,  
 $-OCO-N$   and  $-CO-N$   where r is 1, 2 or 3.

L<sub>7</sub> is a single bond or a group selected from a straight-chain or

branched hydrocarbon chain which has 1 to 100 (preferably 1 to 10) carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups

5 -O-, -S-, -SO-, SO<sub>2</sub>, -NR<sup>y</sup>-, -NR<sup>y</sup>CO-, -C(NH)NR<sup>y</sup>-, CONR<sup>y</sup>-, -NR<sup>y</sup>NR<sup>y</sup>-, -SO<sub>2</sub>NR<sup>y</sup>NR<sup>y</sup>-, -CONR<sup>y</sup>NR<sup>y</sup>- (where R<sup>y</sup> represents H, phenyl, C<sub>1</sub>-C<sub>10</sub>-alkyl, C<sub>2</sub>-C<sub>10</sub>-alkenyl or C<sub>2</sub>-C<sub>10</sub>-alkynyl, each of which may be substituted by NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid), -CO-, -CR<sup>x</sup>=N-O- (where

10 R<sup>x</sup> represents H, C<sub>1</sub>-C<sub>3</sub>-alkyl or phenyl) and/or a 3- to 10-membered, preferably 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, -SO- or -SO<sub>2</sub>- (preferably

), where the hydrocarbon chain including the

15 side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid.

L<sub>5</sub> is preferably a group -(CH<sub>2</sub>)<sub>m</sub>-(CHR<sup>s</sup>)<sub>n</sub>-(OCH<sub>2</sub>CH<sub>2</sub>)<sub>o</sub>-(X)<sub>p</sub>-(CH<sub>2</sub>)<sub>q</sub>-,

20 where m=1-3, n=0, o=0-7, p=0 and q=0 or 1. Particular preference is given to a group -(CH<sub>2</sub>)<sub>m</sub>-(CHR<sup>s</sup>)<sub>n</sub>-(OCH<sub>2</sub>CH<sub>2</sub>)<sub>o</sub>-(X)<sub>p</sub>-(CH<sub>2</sub>)<sub>q</sub>-, where m=1 or 2, n=0, o=0 or 1, p=0 and q=0 or 1.

L<sub>6</sub> is preferably a group selected from -CONH- and -NHCO-.

25

L<sub>7</sub> is preferably a single bond or -[(CH<sub>2</sub>)<sub>x</sub>-(X<sup>4</sup>)<sub>y</sub>]<sub>w</sub>-(CH<sub>2</sub>)<sub>z</sub>-,

where

30 w = 0 to 20;

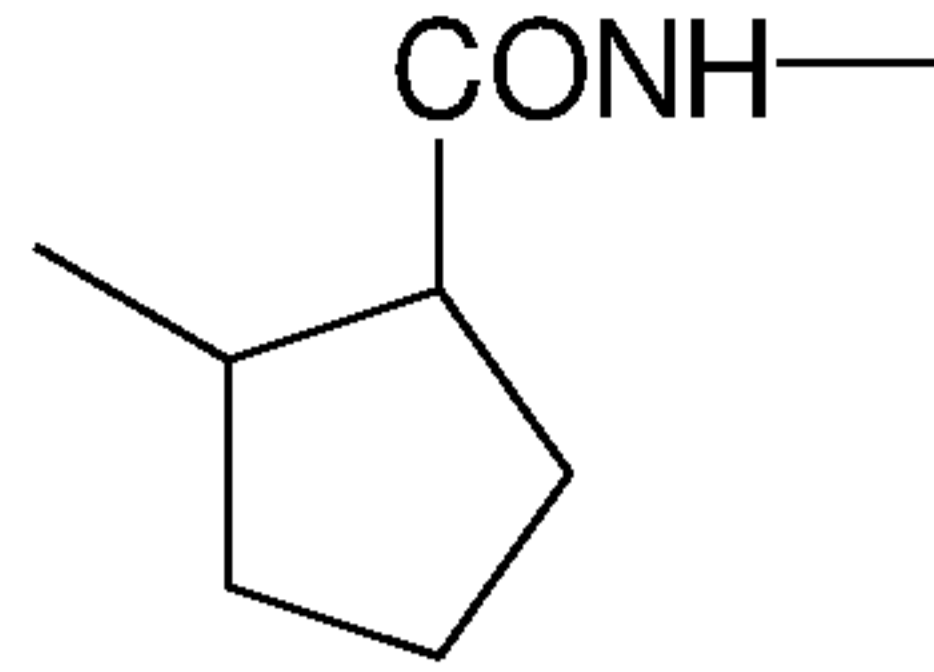
x = 0 to 5;

y = 0 or 1;

35

z = 1 to 5; and





X<sup>4</sup> represents -O-, -CONH-, -NHCO- or

Particularly preferably, L<sub>7</sub> is a single bond or a group -[(CH<sub>2</sub>)<sub>x</sub>-NHCO-], where x = 1 to 5.

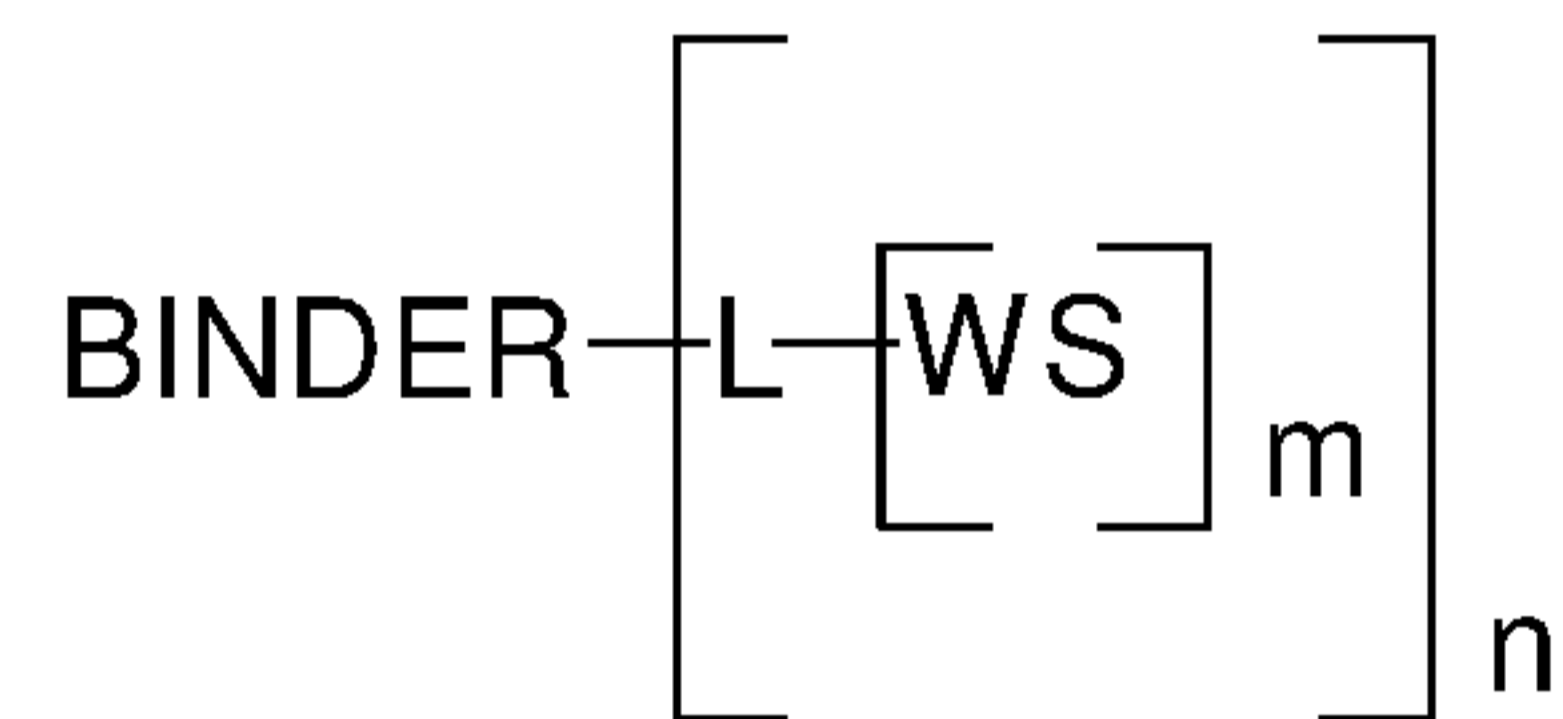
5

Particularly preferably, -L<sub>5</sub>-L<sub>6</sub>-L<sub>7</sub>- represents -(CH<sub>2</sub>)<sub>m</sub>-(CHR<sup>S</sup>)<sub>n</sub>-(OCH<sub>2</sub>CH<sub>2</sub>)<sub>o</sub>-(X)<sub>p</sub>-(CH<sub>2</sub>)<sub>q</sub>--NHCO--[(CH<sub>2</sub>)<sub>x</sub>-NHCO-], where m=1 or 2, n=0, o=0 or 1, p=0, and q=0 or 1, and x=1-5.

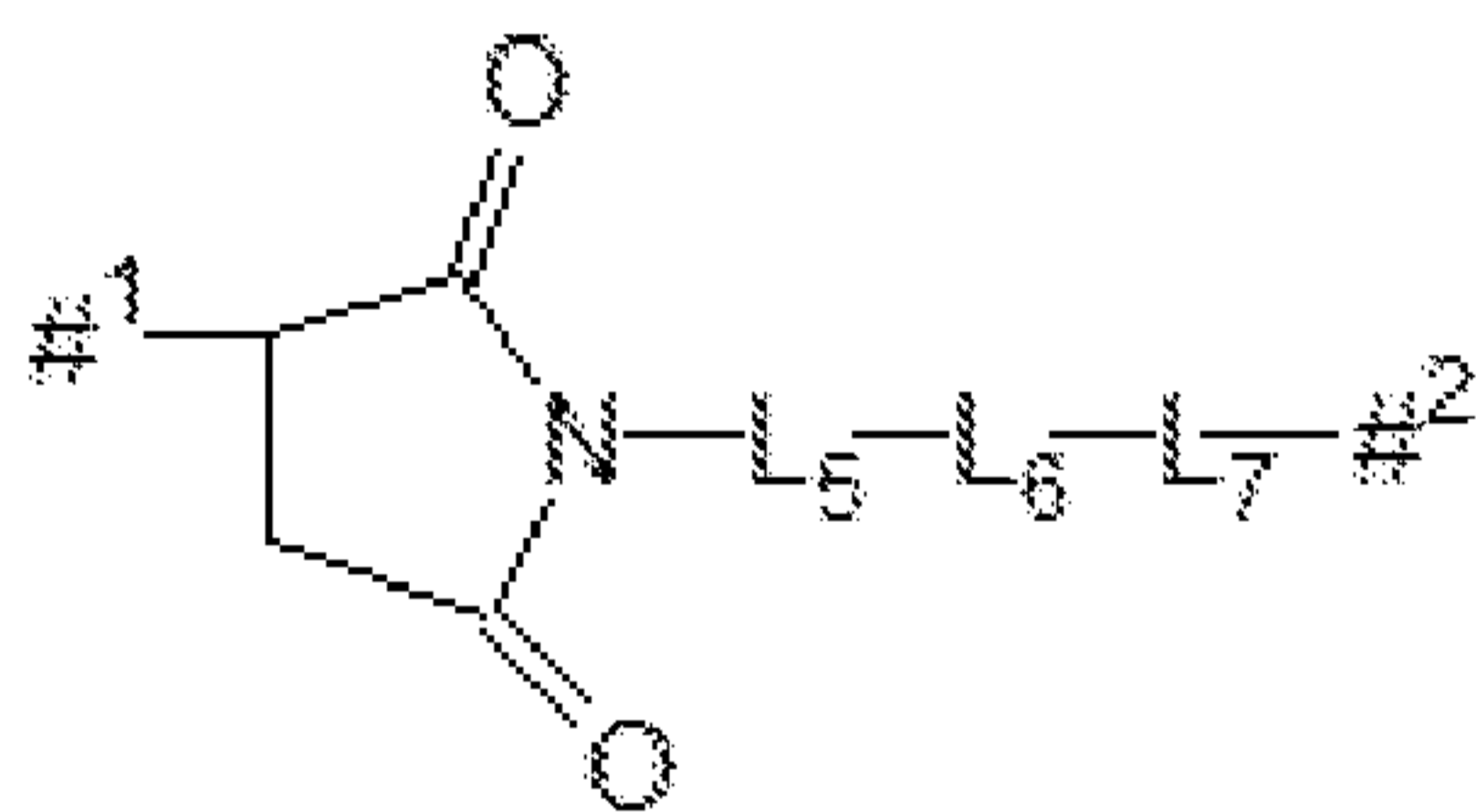
10 However, it is also possible that these two structures are jointly present in the conjugate according to the invention.

According to the invention, these binder/active compound conjugates can be prepared from the compounds of the formula

15



where L has the formula A' below:



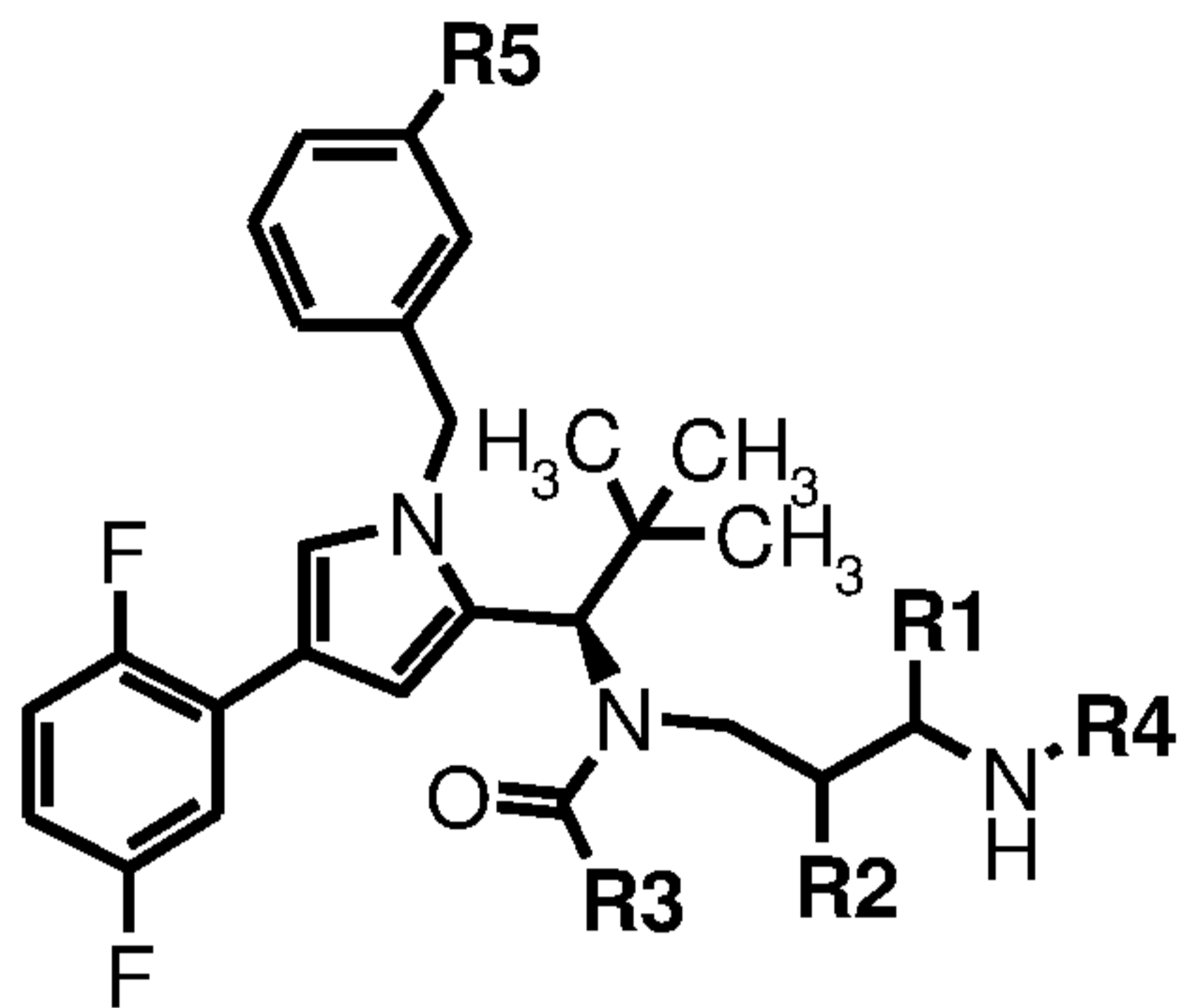
20 Formula A'

Preferably, the conversion of A' into A is carried out by stirring in a pH buffer having a pH of from 7.5 to 8.5, preferably 8, at a temperature below 37°C, preferably from 10 to 25°C, over a period of up to 40 hours, preferably 1 to 15 hours.

25

Embodiment I:

An antibody conjugate of the formula



5

where

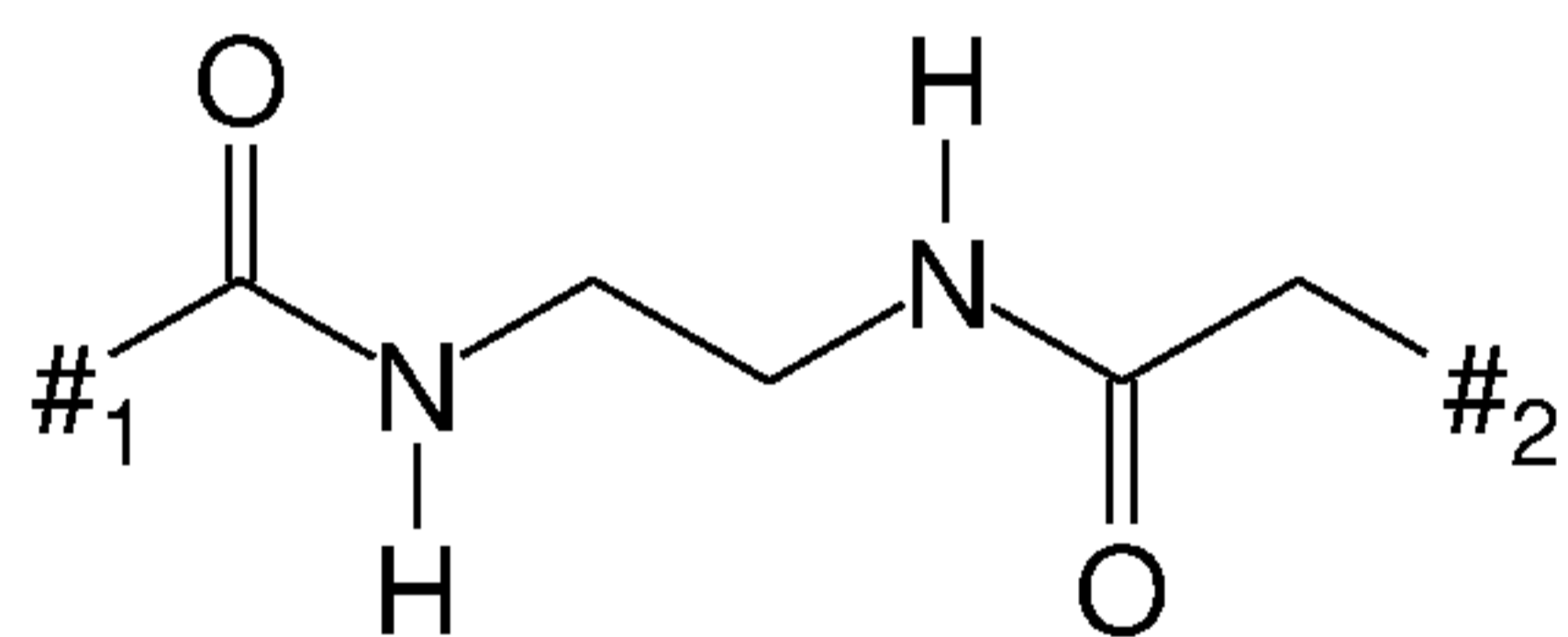
R2, R4 and R5 represent H;

10 R3 represents  $-\text{CH}_2\text{OH}$ ;

R1 represents  $-\text{L1-L2-BINDER}$ , where

L1 represents

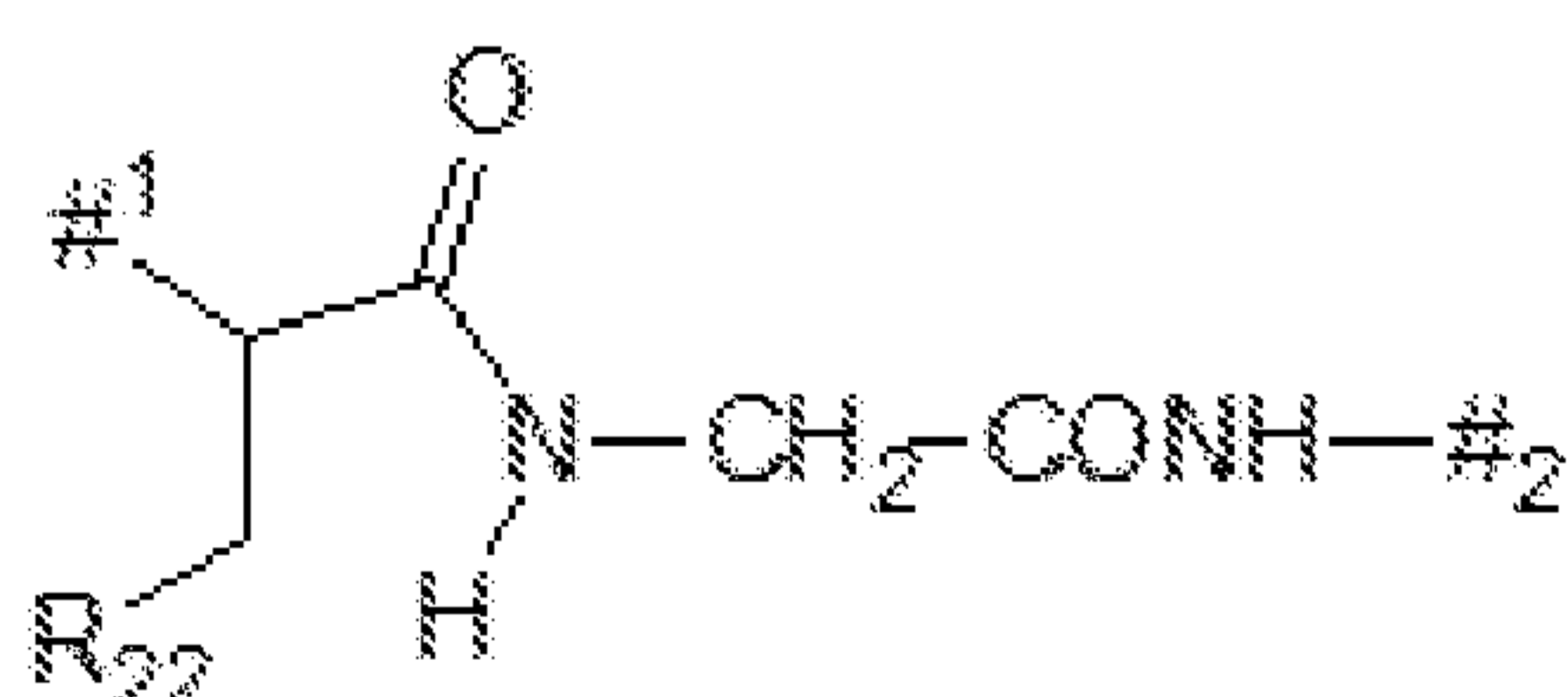
15



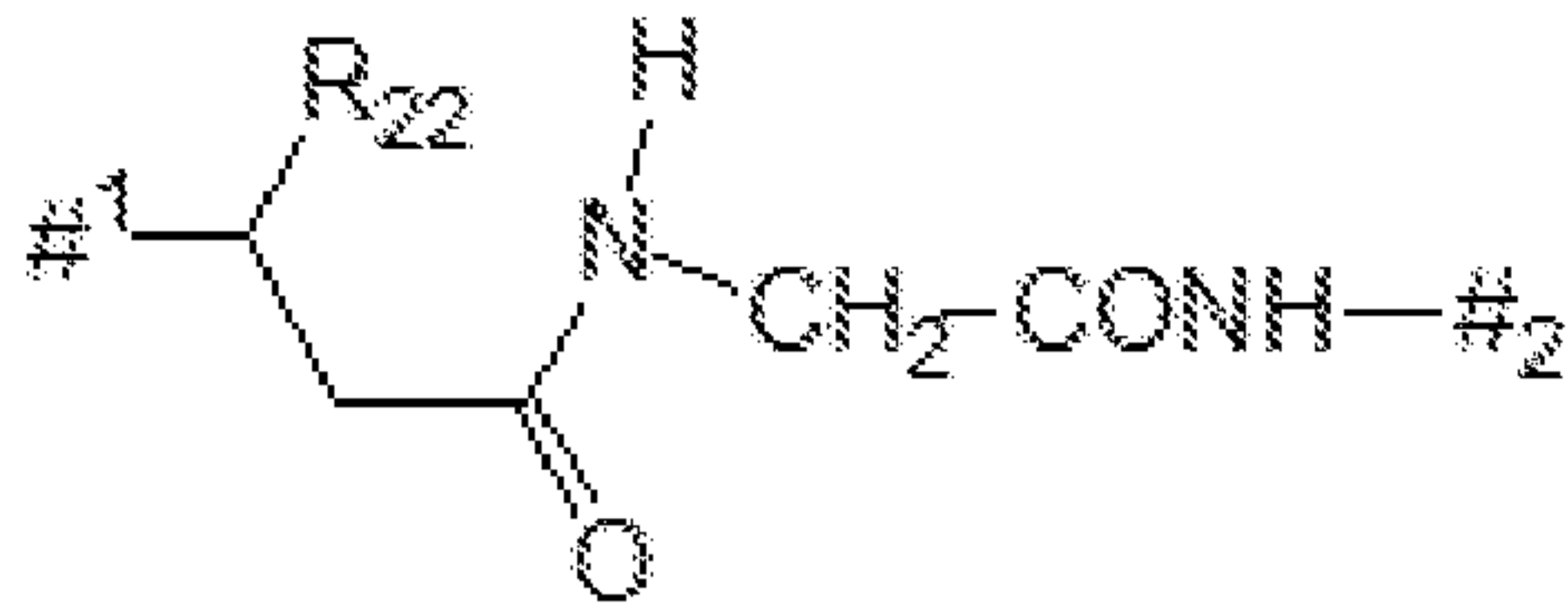
where #2 represents the attachment to L2 and #1 represents the attachment to the other attachment;

20

and L2 represents one or both of the structure of the formulae A5 and A6 below:



25 Formula A5



Formula A6

where

5

#<sup>1</sup> denotes the point of attachment to the sulphur atom of the binder,

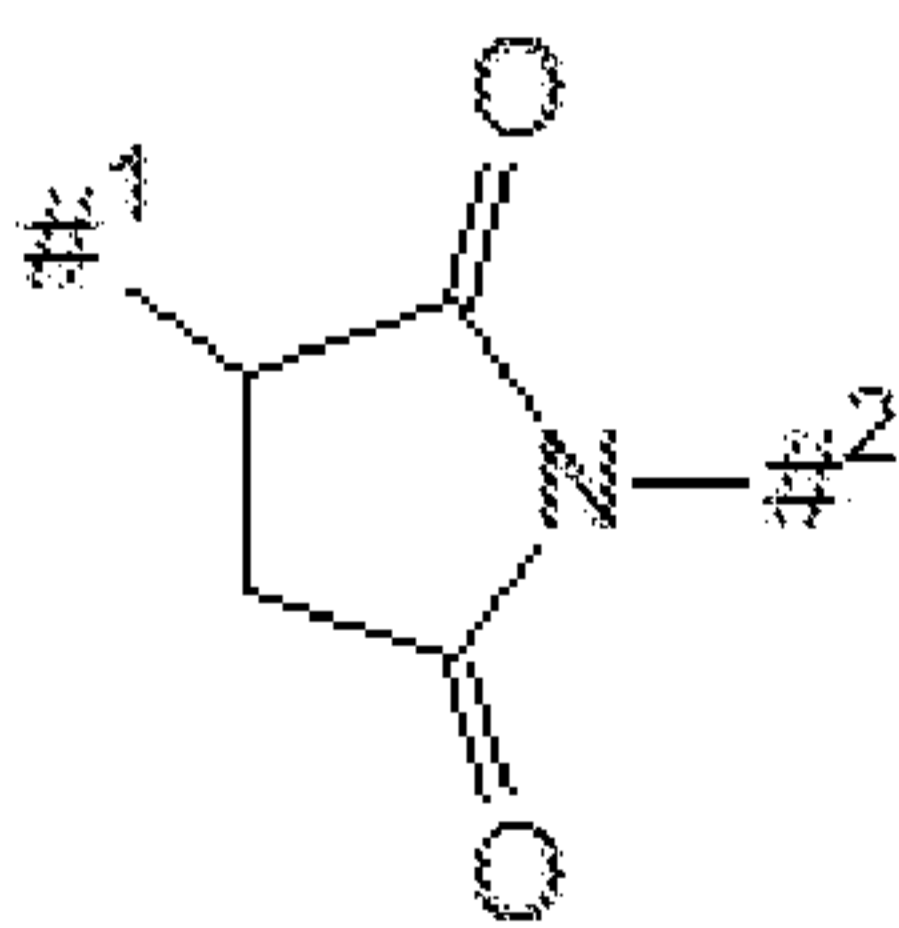
#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>, and

10

R<sup>22</sup> represents COOH, COOR, COR, CONHR (where R in each case represents C1-3-alkyl), CONH<sub>2</sub>, preferably COOH.

In a conjugate according to the invention or in a mixture of the  
 15 conjugates according to the invention, the bonds to a cysteine residue of the binder are present, to an extent of preferably more than 80%, particularly preferably more than 90% (in each case based on the total number of bonds of the linker to the binder) particularly preferably as one of the two structures of  
 20 the formula A5 or A6.

Here, the structures of the formula A5 or A6 are generally present together, preferably in a ratio of from 60:40 to 40:60, based on the number of bonds to the binder. The remaining bonds  
 25 are then present as the structure



The binder is preferably a binder protein or peptide,  
 30 particularly preferably a human, humanized or chimeric monoclonal antibody or an antigen-binding fragment thereof, in

particular an anti-TWEAKR antibody or an antigen-binding fragment thereof or an anti-EGFR antibody or an antigen-binding fragment thereof. Particular preference is given to an anti-TWEAKR antibody which binds specifically to amino acid D in position 47 (D47) of TWEAKR (SEQ ID NO:169), in particular the anti-TWEAKR antibody TPP-2090, or the anti-EGFR antibodies cetuximab or nimotuzumab. As an alternative to the binder, a cysteine residue may also be present.

## 10 ***Therapeutic use***

The hyper-proliferative diseases, for the treatment of which the compounds according to the invention may be employed, include in particular the group of cancer and tumour diseases. In the context of the present invention, these are understood to mean especially the following diseases, but without any limitation thereto: mammary carcinomas and mammary tumours (mammary carcinomas including ductal and lobular forms, also *in situ*), tumours of the respiratory tract (small-cell and non-small cell carcinoma, bronchial carcinoma), cerebral tumours (e.g. of the brain stem and of the hypothalamus, astrocytoma, ependymoma, glioblastoma, glioma, medulloblastoma, meningioma and neuroectodermal and pineal tumours), tumours of the digestive organs (carcinomas of the oesophagus, stomach, gall bladder, small intestine, large intestine, rectum and anal carcinomas), liver tumours (*inter alia* hepatocellular carcinoma, cholangiocarcinoma and mixed hepatocellular cholangiocarcinoma), tumours of the head and neck region (larynx, hypopharynx, nasopharynx, oropharynx, lips and oral cavity carcinomas, oral melanomas), skin tumours (basaliomas, spinaliomas, squamous cell carcinomas, Kaposi's sarcoma, malignant melanoma, non-melanomatous skin cancer, Merkel cell skin cancer, mast cell tumours), tumours of soft tissue (*inter alia* soft tissue sarcomas, osteosarcomas, malignant fibrous histiocyctomas, chondrosarcomas, fibrosarcomas, hemangiosarcomas, leiomyosarcomas, liposarcomas, lymphosarcomas and rhabdomyosarcomas), tumours of the eyes (*inter alia* intraocular melanoma and retinoblastoma), tumours of the

endocrine and exocrine glands (e.g. of the thyroid and parathyroid glands, pancreas and salivary gland carcinomas, adenocarcinomas), tumours of the urinary tract (tumours of the bladder, penis, kidney, renal pelvis and ureter) and tumours of the reproductive organs (carcinomas of the endometrium, cervix, ovary, vagina, vulva and uterus in women and carcinomas of the prostate and testes in men). These also include proliferative blood diseases of the blood, the lymph system and the spinal cord, in solid form and as circulating cells, such as leukaemias, lymphomas and myeloproliferative diseases, for example acute myeloid, acute lymphoblastic, chronic lymphocytic, chronic myelogenous and hairy cell leukaemia, and AIDS-correlated lymphomas, Hodgkin's lymphomas, non-Hodgkin's lymphomas, cutaneous T cell lymphomas, Burkitt's lymphomas and lymphomas in the central nervous system.

These well-characterized diseases in humans can also occur with a comparable aetiology in other mammals and can likewise be treated there with the compounds of the present invention.

The treatment of the cancer diseases mentioned above with the compounds according to the invention comprises both a treatment of the solid tumors and a treatment of metastasizing or circulating forms thereof.

In the context of this invention, the term "treatment" or "treat" is used in the conventional sense and means attending to, caring for and nursing a patient with the aim of combating, reducing, attenuating or alleviating a disease or health abnormality, and improving the living conditions impaired by this disease, as, for example, in the event of a cancer.

The present invention thus further provides for the use of the compounds according to the invention for the treatment and/or prevention of disorders, in particular the disorders mentioned above.

The present invention further provides for the use of the

compounds according to the invention for producing a medicament for the treatment and/or prevention of disorders, in particular the disorders mentioned above.

5 The present invention further provides for the compounds according to the invention for use in a method for treatment and/or prevention of disorders, in particular the disorders mentioned above.

10 The present invention further provides for compounds for the use in a method for treatment and/or prevention of disorders, in particular the disorders mentioned above, using an effective amount of at least one of the compounds according to the invention.

15

The compounds according to the invention can be used alone or, if required, in combination with one or more other pharmacologically active substances, provided that this combination does not lead to undesirable and unacceptable side effects. The present invention furthermore therefore provides medicaments containing at least one of the compounds according to the invention and one or more further active compounds, in particular for treatment and/or prevention of the abovementioned disorders.

25

For example, the compounds of the present invention can be combined with known anti-hyper-proliferative, cytostatic or cytotoxic substances for the treatment of cancer diseases. Examples of suitable combination active compounds include:

30

131I-chTNT, abarelix, abiraterone, aclarubicin, afatinib, aflibercept, aldesleukin, alemtuzumab, alisertib, alitretinoin, alphasaradin (radium-223 chloride), altretamine, aminoglutethimide, AMP-514, amrubicin, amsacrine, anastrozole, arglabin, arsenic trioxide, asparaginase, AT9283, axitinib, azacitidine, basiliximab, belotecan, bendamustin, bevacizumab, bexaroten, bicalutamide, bisantrene, bleomycin, BMS-936559, bosutinib, bortezomib, brentuximab vedotin, buserelin,

35

busulfan, cabazitaxel, cabozantinib, calcium folinate, calcium  
levofolinate, capecitabine, carboplatin, carfilzomib  
(proteasome inhibitor), carmofur, carmustine, catumaxomab,  
celecoxib, celmoleukin, cetuximab, chlorambucil, chlormadinone,  
5 chlormethine, cisplatin, cladribine, clodronic acid,  
clofarabine, copanlisib, crisantaspase, crizotinib,  
cyclophosphamide, CYC116, cyproterone, cytarabine, dacarbazine,  
dactinomycin, darbepoetin-alfa, dabrafenib, danusertib,  
dasatinib, daunorubicin, decitabine, degarelix, denileukin-  
10 diftitox, denosumab, deslorelin, dibrospidium chloride,  
docetaxel, doxifluridine, doxorubicin, doxorubicin + estrone,  
eculizumab, edrecolomab, elliptinium acetate, eltrombopag,  
endostatin, ENMD-2076, enocitabine, epirubicin, epitiostanol,  
epoetin-alfa, epoetin-beta, eptaplatin, eribulin, erlotinib,  
15 estradiol, estramustine, etoposide, everolimus, exemestane,  
fadrozole, filgrastim, fludarabine, fluorouracil, flutamide,  
formestane, fotemustine, fulvestrant, gallium nitrate,  
ganirelix, gefitinib, gemcitabine, gemtuzumab, glutoxim,  
goserelin, histamine dihydrochloride, histrelin,  
20 hydroxycarbamide, I-125 seeds, ibandronic acid, ibritumomab-  
tiuxetan, ibrutinib, idarubicin, ifosfamide, imatinib,  
imiquimod, INCB24360, improsulfan, interferon-alfa, interferon-  
beta, interferon-gamma, ipilimumab, irinotecan, ixabepilone,  
lambrolizumab, lanreotide, lapatinib, lenalidomide,  
25 lenograstim, lentinan, letrozole, leuprorelin, levamisole,  
lisuride, lobaplatin, lomustine, lonidamine, masoprocol,  
medroxyprogesterone, megestrol, melphalan, mepitiostane,  
mercaptapurine, methotrexate, methoxsalen,  
methylaminolevulinate, methyltestosterone, mifamurtide,  
30 miltefosine, miriplatin, mitobronitol, mitoguazone, mitolactol,  
mitomycin, mitotane, mitoxantrone, MLN-8054, Mps1 inhibitors  
(disclosed in WO2013/087579, in particular Example 01.01,  
WO2014/131739, in particular Example 2), nedaplatin, nelarabine,  
nemorubicin, nilotinib, nilutamide, nimotuzumab, nimustine,  
35 nitracrine, nivolumab, NMS-P715, NMS-P937, ofatumumab,  
omeprazole, oprelvekin, oxaliplatin, p53 gene therapy,  
paclitaxel, palbociclib, palifermin, palladium-103 seed,  
pamidronic acid, panitumumab, pazopanib, pegaspargase, PEG-

epoetin-beta (methoxy-PEG-epoetin-beta), pegfilgrastim, Peg-  
interferon-alfa-2b, pemetrexed, pentazocin, pentostatin,  
peplomycin, perfosfamide, picibanil, pirarubicin, plerixafor,  
plicamycin, poliglusam, polyestradiol phosphate,  
5 polysaccharide-K, ponatinib, porfimer-sodium, pralatrexate,  
prednimustine, procarbazine, quinagolide, R763, raloxifene,  
raltitrexed, ranimustine, razoxane, refametinib, regorafenib,  
risedronic acid, rituximab, romidepsin, romiplostim,  
roninciclib, ruxolitinib, sargramostim, sipuleucel-T,  
10 sizofiran, sobuzoxane, sodium glycididazole, SNS-314,  
sorafenib, streptozocin, sunitinib, talaporfin, tamibarotene,  
tamoxifen, tasonermin, teceleukin, tegafur, tegafur + gimeracil  
+ oteracil, temoporfin, temozolomide, temsirolimus, teniposide,  
testosterone, tetrofosmin, thalidomide, thiotepa, thymalfasin,  
15 TKM-PLK1, tioguanine, tocilizumab, topotecan, toremifene,  
tositumomab, tozasertib, trabectedin, trametinib, trastuzumab,  
trastuzumab emtansine, treosulfan, tretinoin, trilostane,  
triptorelin, trofosfamide, tryptophan, ubenimex, valrubicin,  
vandetanib, vapreotide, vemurafenib, vinblastine, vincristine,  
20 vindesine, vinflunine, vinorelbin, volasertib, vorinostat,  
vorozol, XL228, yttrium-90 glass microbeads, zinostatin,  
zinostatin-stimalamer, zoledronic acid, zorubicin.

In addition, the compounds of the present invention can be  
25 combined, for example, with binders which, by way of example,  
can bind to the following targets: OX-40, CD137/4-1BB, DR3,  
IDO1/IDO2, LAG-3, CD40.

In addition, the compounds according to the invention can also  
30 be used in combination with radiotherapy and/or surgical  
intervention.

Generally, the following aims can be pursued with the  
combination of compounds of the present invention with other  
35 cytostatically or cytotoxically active agents:

- improved efficacy in slowing the growth of a tumour, in reducing its size or even in the complete elimination thereof,



compared with treatment with an individual active compound;

- the possibility of using the chemotherapeutics used in a lower dosage than in the case of monotherapy;

5

- the possibility of a more tolerable therapy with fewer side effects compared with individual administration;

10

- the possibility of treatment of a broader spectrum of tumour diseases;

- the achievement of a higher rate of response to the therapy;

15

- a longer survival time of the patient compared with present-day standard therapy.

In addition, the compounds according to the invention can also be used in combination with radiotherapy and/or surgical intervention.

20

The present invention further provides medicaments which comprise at least one compound according to the invention, typically together with one or more inert, nontoxic, pharmaceutically suitable excipients, and the use thereof for the aforementioned purposes.

25

The compounds according to the invention can act systemically and/or locally. For this purpose, they can be administered in a suitable manner, for example parenterally, possibly inhalatively or as implants or stents.

30

The compounds according to the invention can be administered in suitable administration forms for these administration routes.

35

Parenteral administration can bypass an absorption step (for example intravenously, intraarterially, intracardially, intraspinally or intralumbally) or include an absorption (for example intramuscularly, subcutaneously, intracutaneously,

percutaneously or intraperitoneally). Administration forms suitable for parenteral administration include preparations for injection and infusion in the form of solutions, suspensions, emulsions or lyophilizates. Preference is given to parenteral administration, especially intravenous administration.

In general, it has been found to be advantageous in the case of parenteral administration to administer amounts of from about 0.001 to 1 mg/kg, preferably about 0.01 to 0.5 mg/kg, of body weight to achieve effective results.

It may nevertheless be necessary where appropriate to deviate from the stated amounts, specifically as a function of body weight, route of administration, individual response to the active compound, nature of the preparation and time or interval over which administration takes place. Thus, in some cases less than the abovementioned minimum amount may be sufficient, while in other cases the upper limit mentioned must be exceeded. In the case of administration of greater amounts, it may be advisable to divide them into several individual doses over the day.

### ***Examples***

The examples which follow illustrate the invention. The invention is not restricted to the examples.

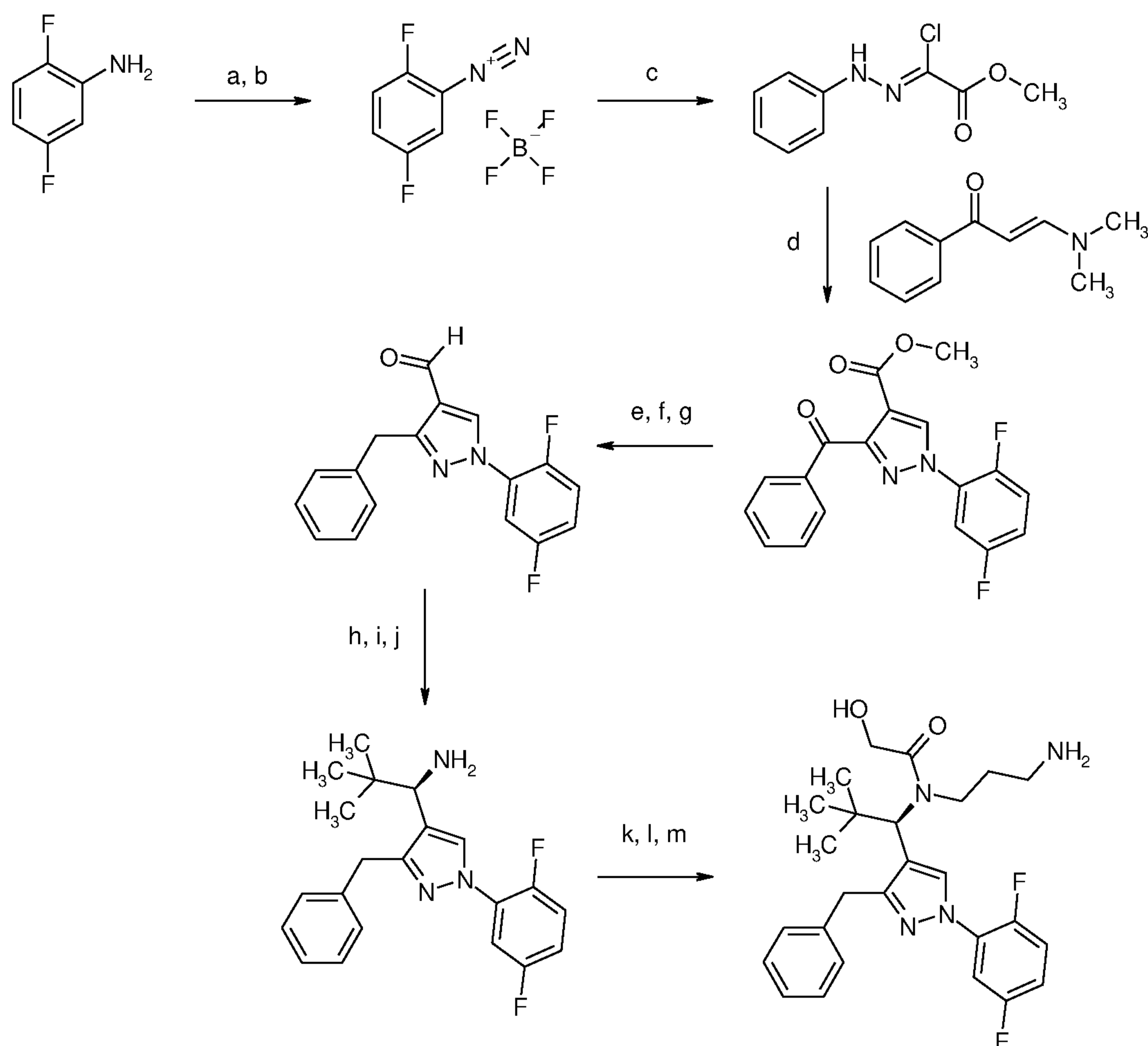
Unless stated otherwise, the percentages in the tests and examples which follow are percentages by weight; parts are parts by weight. Solvent ratios, dilution ratios and concentration data for the liquid/liquid solutions are in each case based on volume.

### Synthesis routes:

35

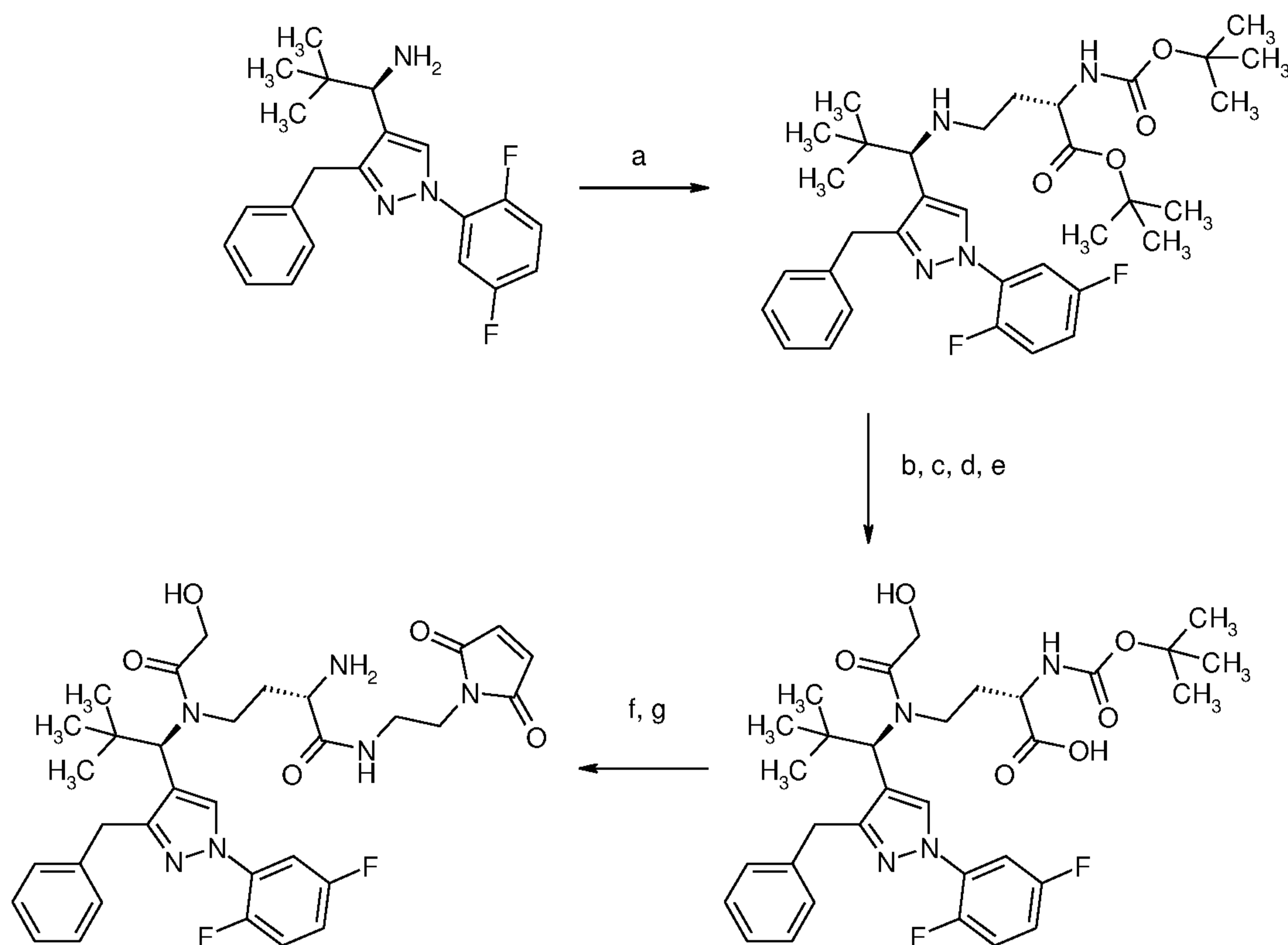
Exemplary for the working examples, the schemes below show exemplary synthesis routes leading to the working examples:

Scheme 1:



- 5 [a]: for example  $\text{BF}_3\text{OEt}_2$ , THF,  $0^\circ\text{C}$ ; b): for example isoamyl nitrite,  $-10^\circ\text{C}$ , 0.5 h; c): for example methyl 2-chloro-3-oxobutanoate, pyridine, water,  $-5^\circ\text{C}$ ; d): for example  $\text{NEt}_3$ , toluene, RT; e): for example  $\text{Et}_3\text{SiH}$ , TFA, RT; f): for example  $\text{LiBH}_4$ , THF,  $60^\circ\text{C}$ ; g): for example Dess-Martin periodinane, DCM, RT; h): for example (R)-(+)-methyl-2-propanesulphinamide, titanium(IV) isopropoxide, THF, RT; i): for example tert-BuLi, pentane, THF,  $-78^\circ\text{C}$ ; j): for example HCl in dioxane, THF, MeOH, RT; k): for example 3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propanal,  $\text{NaB}(\text{OAc})_3\text{H}$ , AcOH, DCM, RT; l): for example 2-chloro-2-oxoethyl acetate,  $\text{NEt}_3$ , DCM, RT; m): for example methylamine, water, EtOH,  $50^\circ\text{C}$ ]
- 10
- 15

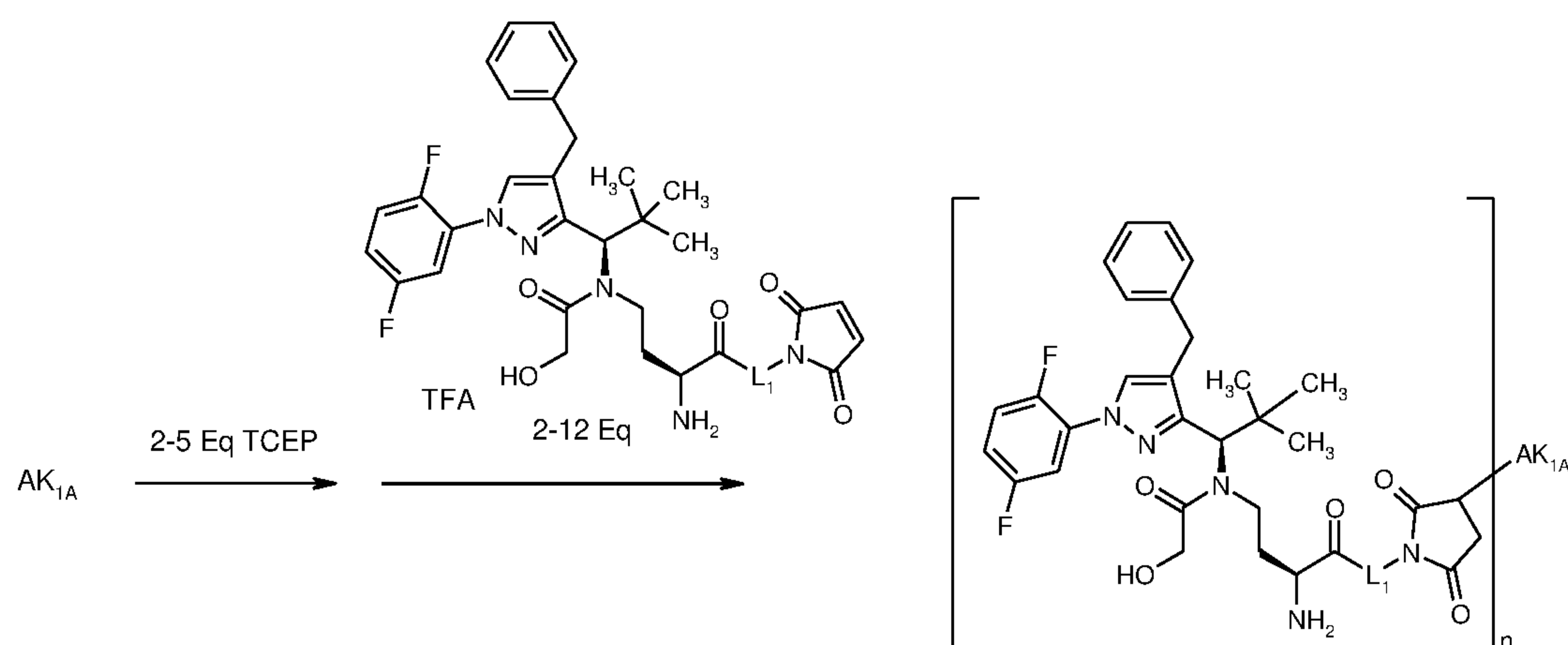
Scheme 2



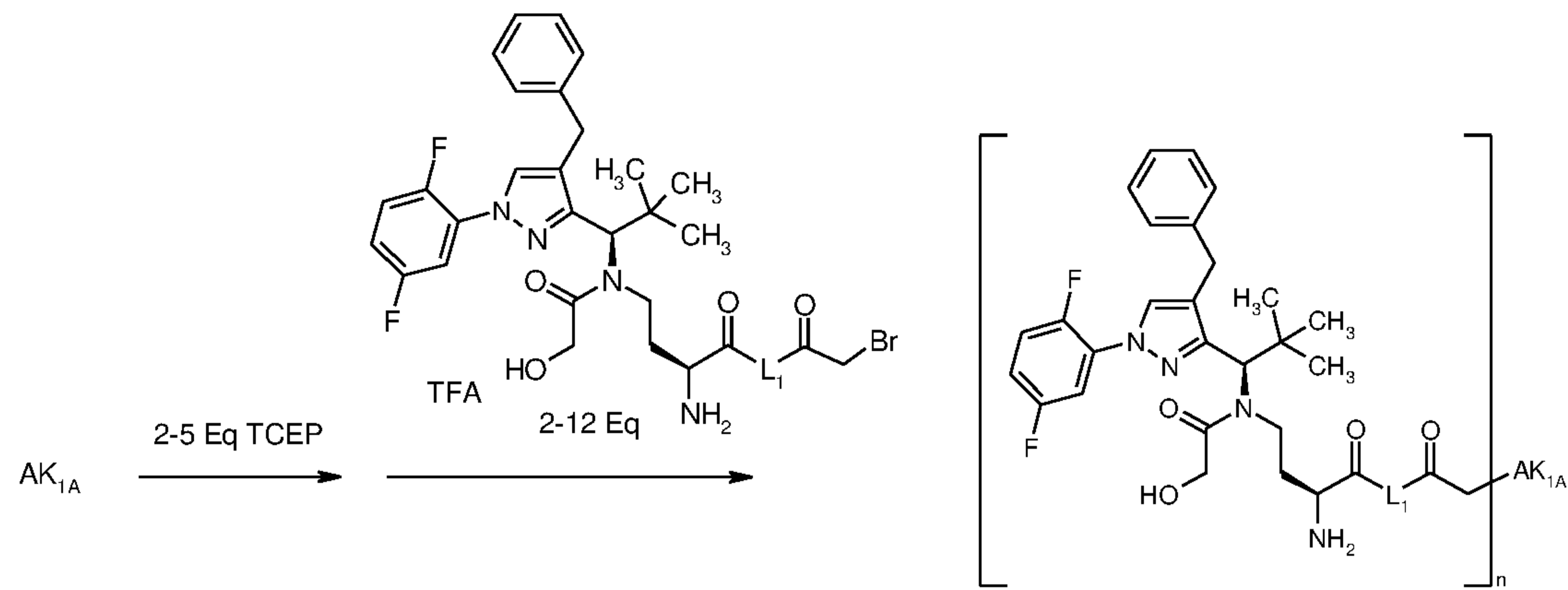
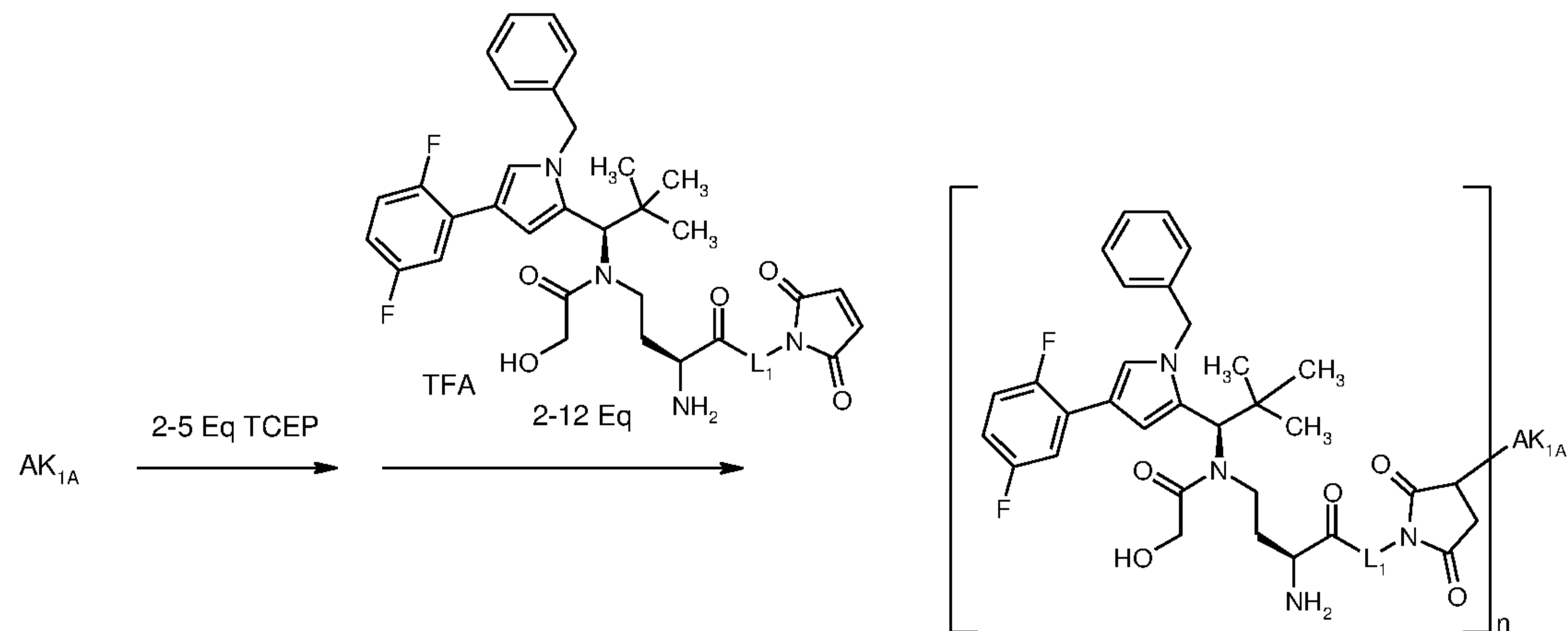
[a]: for example tert-butyl N-(tert-butoxycarbonyl)-5-oxo-L-norvalinate, NaB(OAc)<sub>3</sub>H, AcOH, DCM, RT; b): for example 2-chloro-2-oxoethyl acetate, NEt<sub>3</sub>, DCM, RT; c): for example methylamine, water, EtOH, 60°C; d): for example THF, DCM, 50°C; e): for example Boc<sub>2</sub>O, NEt<sub>3</sub>, DCM, RT; f): for example trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1), HATU, diisopropylethylamine, DMF, RT; g): for example TFA, DCM, RT]

10

Scheme 3: Synthesis of cysteine-linked ADCs

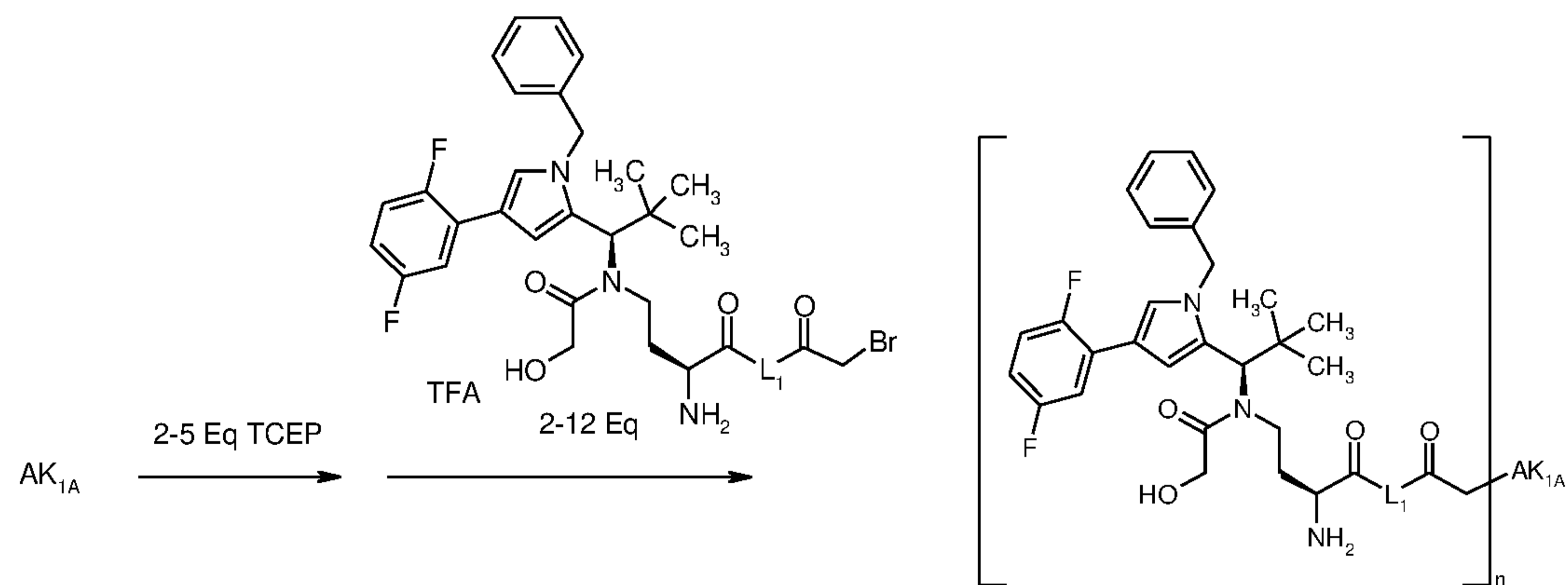


Scheme 4: Synthesis of cysteine-linked ADCs

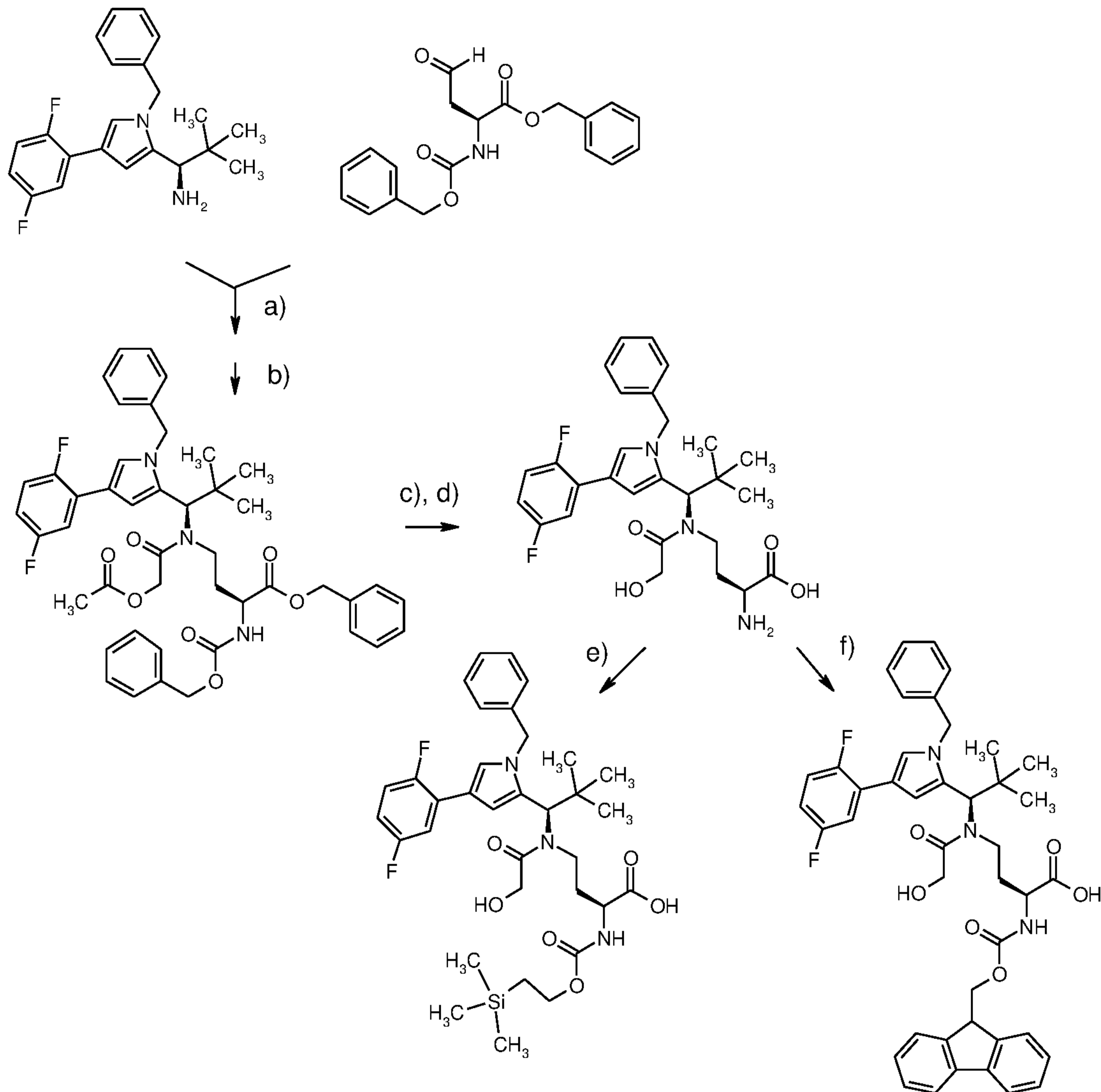
5 Scheme 5: Synthesis of cysteine-linked ADCs

Scheme 6: Synthesis of cysteine-linked ADCs

10

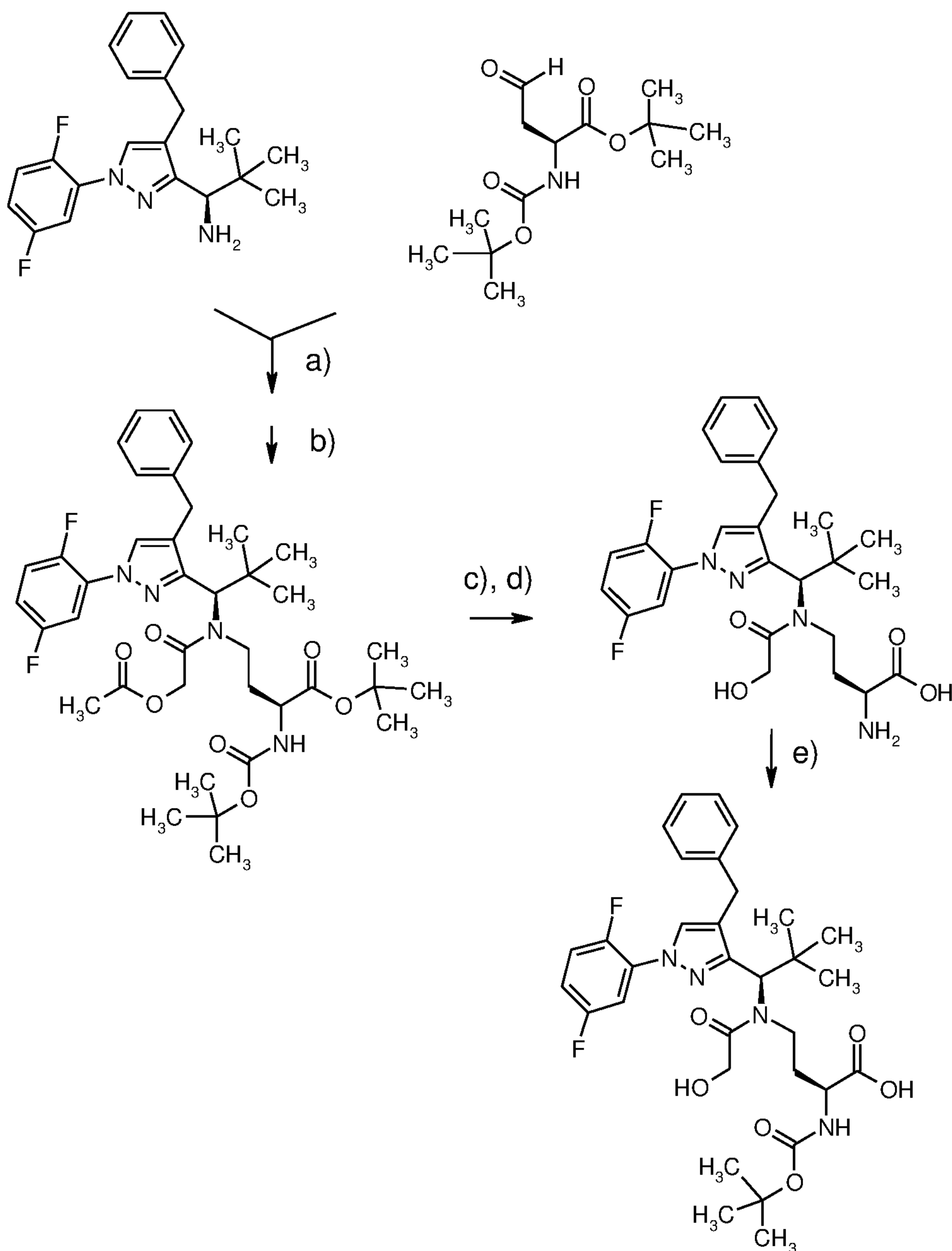


Scheme 7



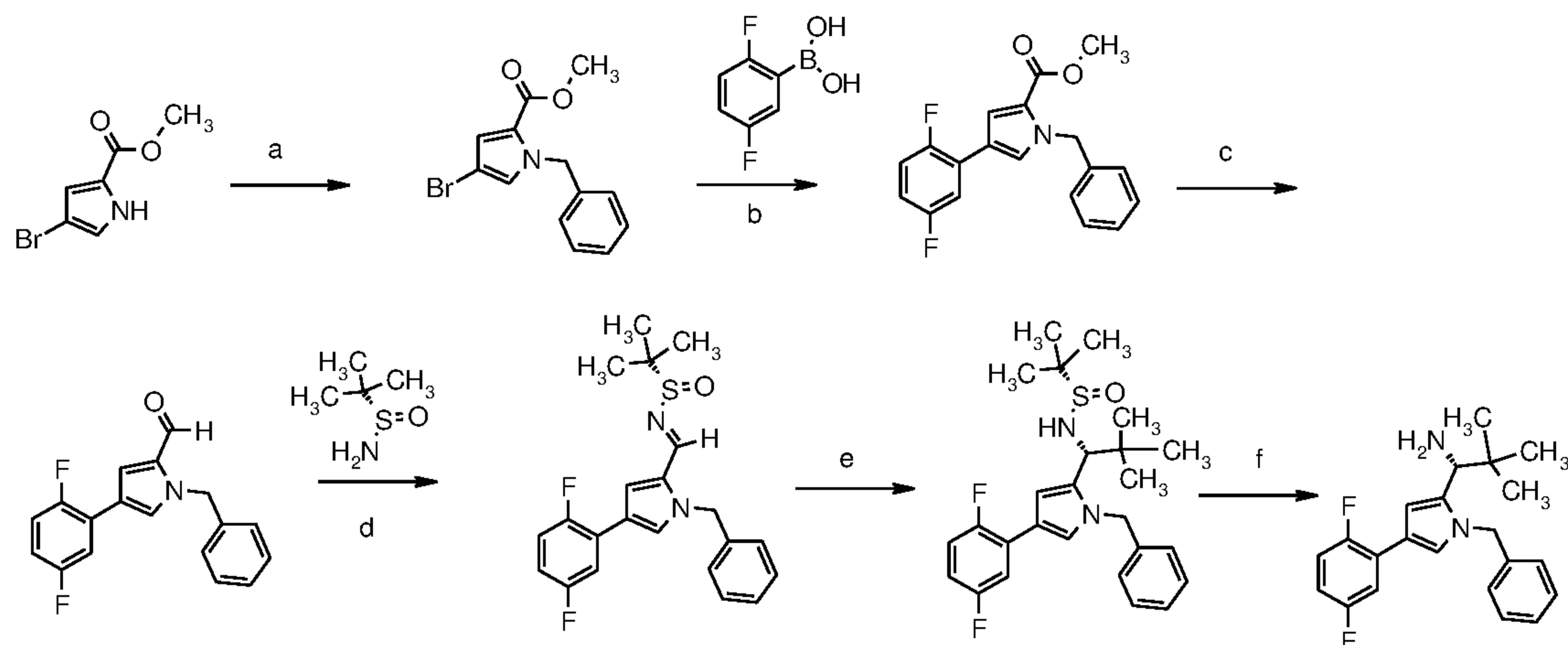
[a): for example sodium triacetoxyborohydride, acetic acid, DCM, RT; b) for example acetoxyacetyl chloride,  $\text{NEt}_3$ , DCM, RT; c) for example  $\text{LiOH}$ , THF/water, RT; d) for example  $\text{H}_2$ , Pd-C, EtOH, RT; e) for example Teoc-OSu,  $\text{NEt}_3$ , dioxane, RT; f) for example Fmoc-Cl, diisopropylethylamine, dioxane/water 2:1, RT]

Scheme 8



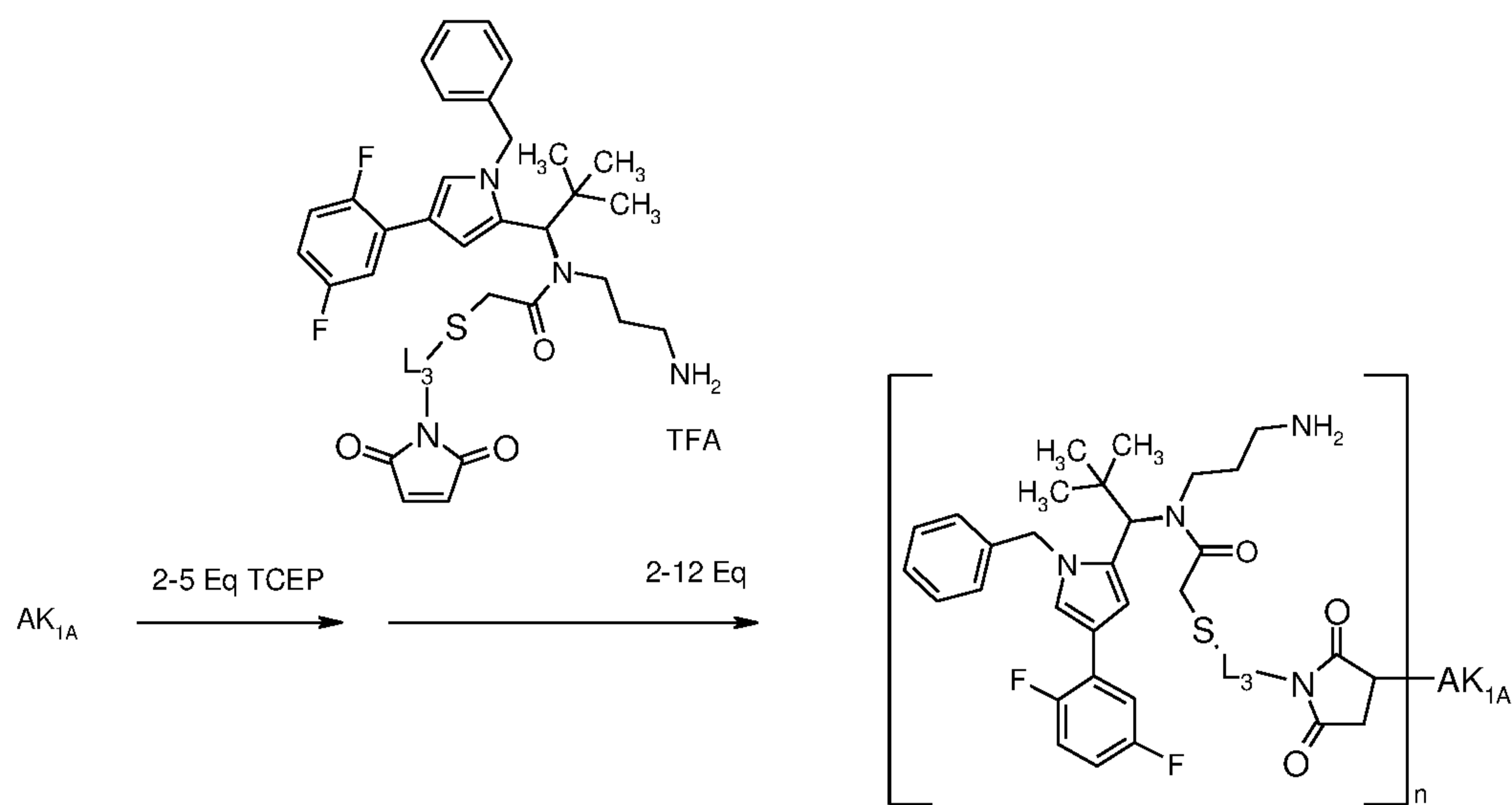
[a): for example sodium triacetoxyborohydride, acetic acid, DCM, RT; b) for example acetoxyacetyl chloride,  $\text{NEt}_3$ , DCM, RT; c) for example LiOH, methanol, RT; d) for example TFA, DCM, RT; e) for example  $\text{Boc}_2\text{O}$ , diisopropylethylamine, DCM, RT]

Scheme 9



[a): for example benzyl bromide,  $\text{Cs}_2\text{CO}_3$ , DMF, RT; b) for example  $\text{Pd}(\text{dppf})_2\text{Cl}_2$ , DMF,  $\text{Na}_2\text{CO}_3$ ,  $85^\circ\text{C}$ ; c) for example  $\text{LiAlH}_4$ , THF,  $0^\circ\text{C}$ ; MnO<sub>2</sub>, DCM, RT; d) for example  $\text{Ti}(\text{iOPr})_4$ , THF, RT; e) for example  $\text{tBuLi}$ , THF,  $-78^\circ\text{C}$ ; MeOH,  $\text{NH}_4\text{Cl}$ ; f) for example HCl/1,4-dioxane]

Scheme 10: Synthesis of cysteine-linked ADCs

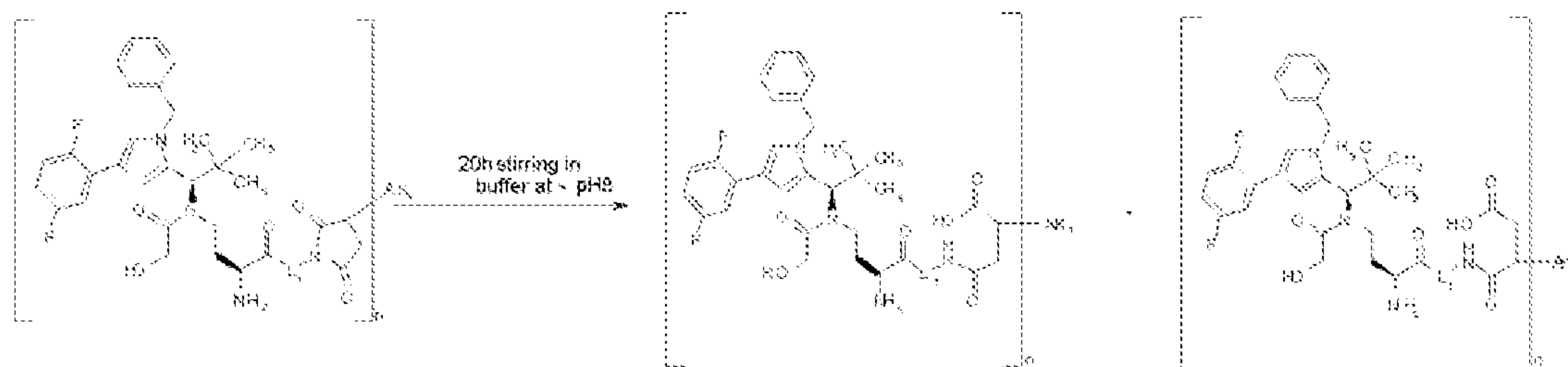


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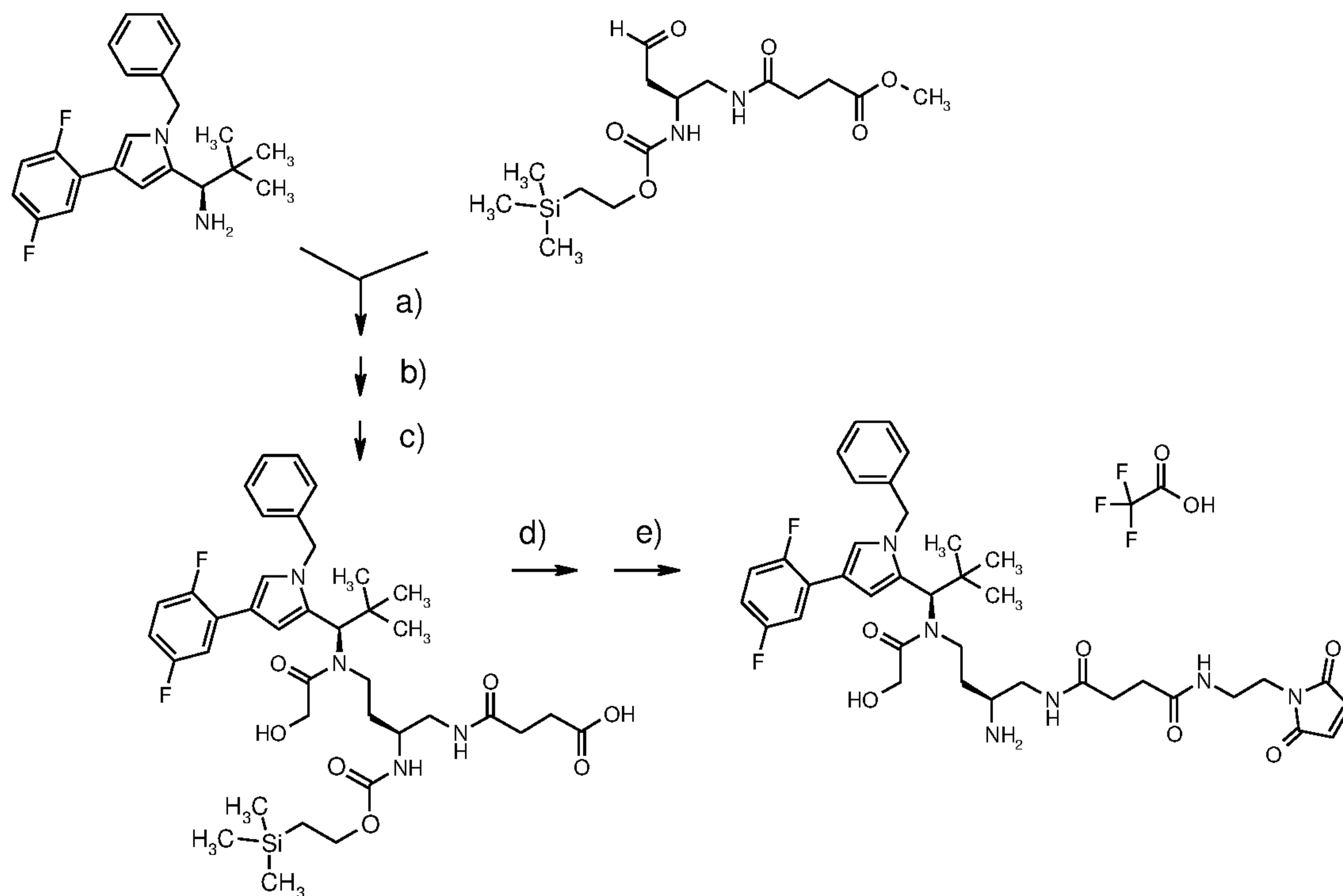
Scheme 11: Synthesis of cysteine-linked ADCs via hydrolysed succinamides

15 This process was used in particular for ADCs where  $\text{L}_1 = \text{CH}_2$  to convert these ADCs into the open-chain linking form.





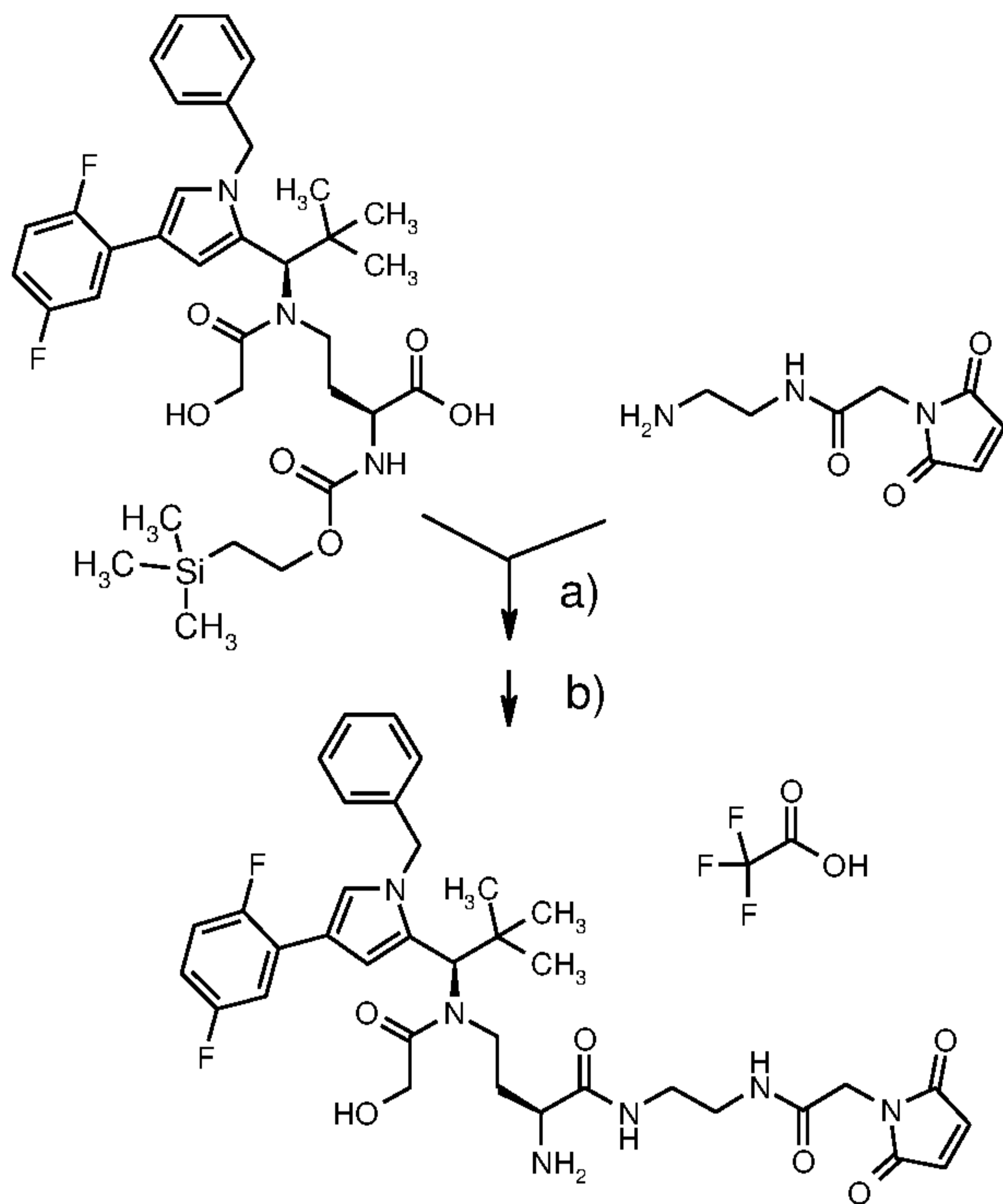
Scheme 12:



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[a): sodium triacetoxyborohydride, acetic acid, DCM, RT; b) acetoxyacetyl chloride, diisopropylethylamine, DCM, RT; c) LiOH, MeOH, RT; d) trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) HATU, DMF, diisopropylethylamine, RT; e) zinc chloride, trifluoroethanol, 50°C, EDTA.]

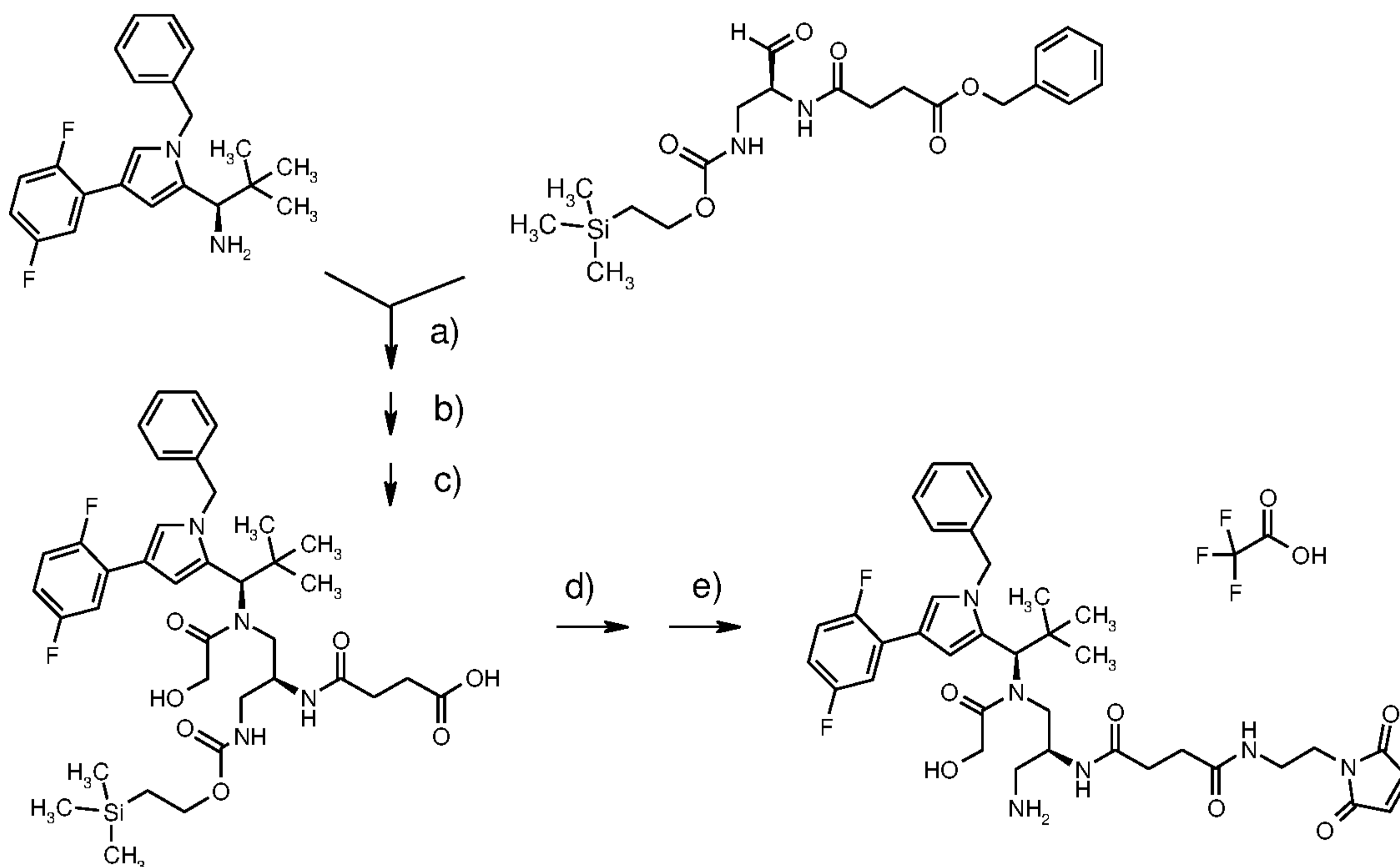
Scheme 13:



[a): HATU, DMF, diisopropylethylamine, RT; b) zinc chloride, trifluoroethanol, 50°C, EDTA.]

5

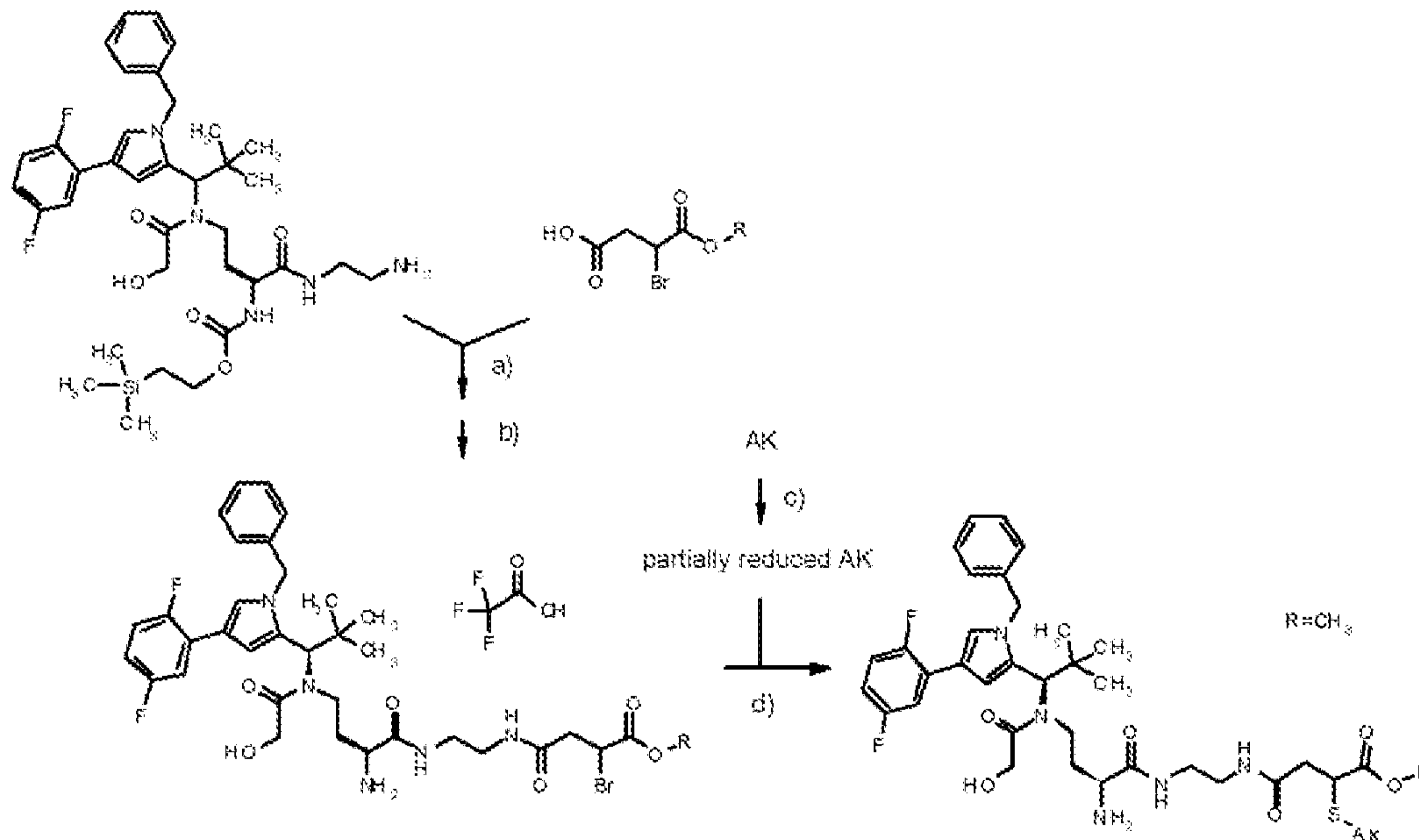
Scheme 14:



10 [a): sodium triacetoxyborohydride, acetic acid, DCM, RT; b) acetoxyacetyl chloride, triethylamine, DCM, RT; c) LiOH, MeOH,

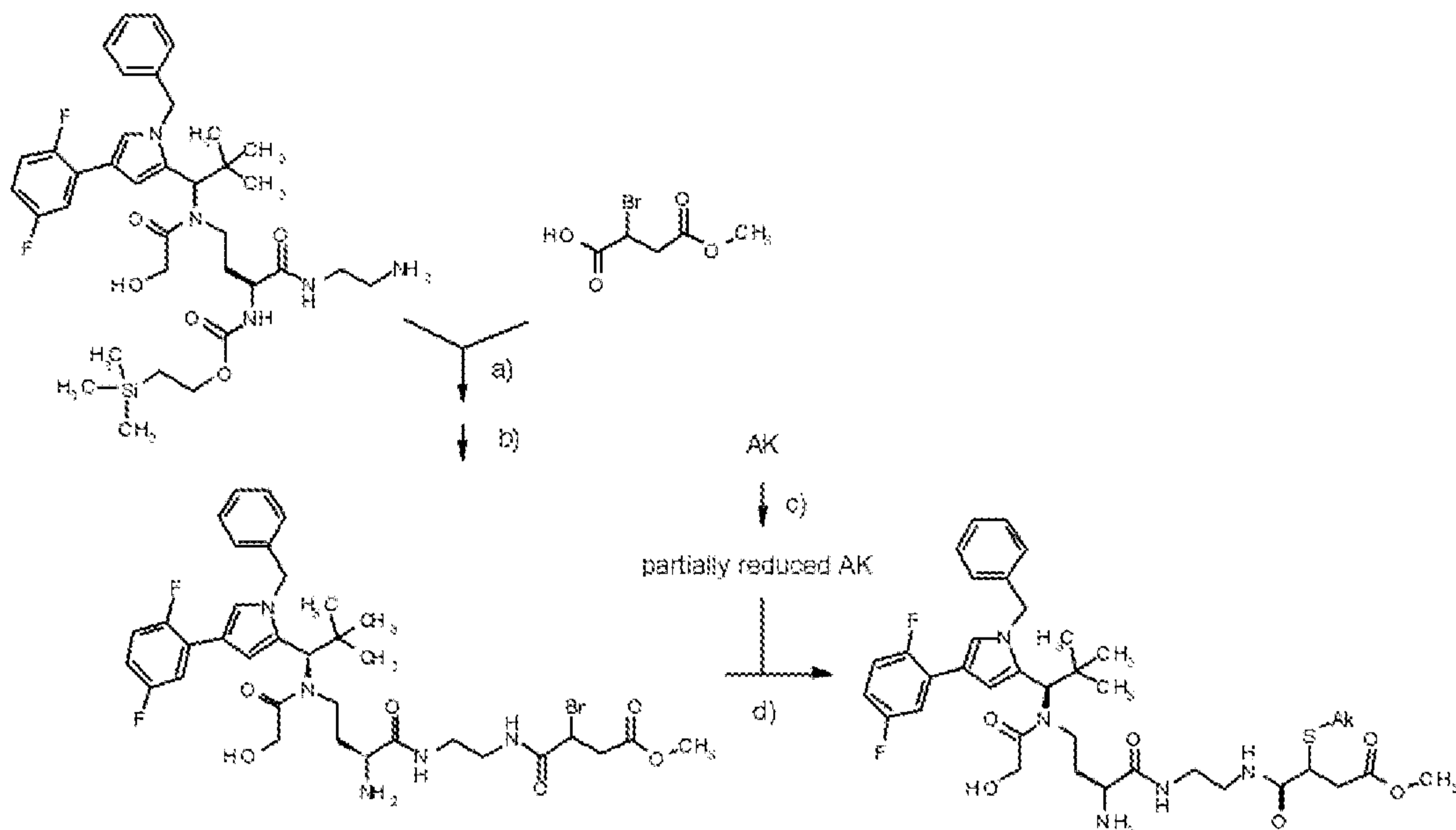
RT; d) trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) HATU, DMF, diisopropylethylamine, RT; e) zinc chloride, trifluoroethanol, 50°C, EDTA.]

5 Scheme 15:



[a): 2-bromo-1-ethylpyridinium tetrafluoroborate (BEP), DCM, pyridine, RT; b) zinc chloride, trifluoroethanol, 50°C, EDTA; c) 3-4 equivalents of TCEP, PBS buffer; d) PBS buffer, 20h RT.]

Scheme 16:



- [a): 2-bromo-1-ethylpyridinium tetrafluoroborate (BEP), DCM, pyridine, RT; b) zinc chloride, trifluoroethanol, 50°C, EDTA; 5 c) 3-4 equivalents of TCEP, PBS buffer; d) PBS buffer, 20h RT.]

## A. Examples

### Abbreviations and acronyms:

10

A431NS	human tumour cell line
A549	human tumour cell line
ABCB1	ATP-binding cassette sub-family B member 1 (synonym for P-gp and MDR1)
abs.	absolute
Ac	acetyl
ACN	acetonitrile
aq.	aqueous, aqueous solution
ATP	adenosine triphosphate
BCRP	breast cancer resistance protein, an efflux
BEP	transporter 2-bromo-1-ethylpyridinium tetrafluoroborate
Boc	<i>tert</i> -butoxycarbonyl
br.	broad (in NMR)
Ex.	Example

CI	chemical ionization (in MS)
d	doublet (in NMR)
d	day(s)
TLC	thin-layer chromatography
DCI	direct chemical ionization (in MS)
dd	doublet of doublets (in NMR)
DMAP	4- <i>N,N</i> -dimethylaminopyridine
DME	1,2-dimethoxyethane
DMEM	Dulbecco's Modified Eagle Medium (standardized nutrient medium for cell culture)
DMF	<i>N,N</i> -dimethylformamide
DMSO	dimethyl sulphoxide
DPBS, D-PBS, PBS	Dulbecco's phosphate-buffered salt solution PBS = DPBS = D-PBS, pH 7.4, from Sigma, No D8537  composition:  0.2 g KCl 0.2 g KH <sub>2</sub> PO <sub>4</sub> (anhyd) 8.0 g NaCl 1.15 g Na <sub>2</sub> HPO <sub>4</sub> (anhyd) made up ad 1 l with H <sub>2</sub> O
dt	doublet of triplets (in NMR)
DTT	DL-dithiothreitol
EDC	<i>N'</i> -(3-dimethylaminopropyl)- <i>N</i> -ethylcarbodiimide hydrochloride
EGFR	epidermal growth factor receptor
EI	electron impact ionization (in MS)
ELISA	enzyme-linked immunosorbent assay
eq.	equivalent(s)
ESI	electrospray ionization (in MS)
ESI-MicroTofq	<u>ESI</u> - MicroTofq (name of the mass spectrometer with Tof = time of flight and q = quadrupol)
FCS	foetal calf serum

Fmoc	(9 <i>H</i> -fluoren-9-ylmethoxy) carbonyl
GTP	guanosine-5'-triphosphate
h	hour(s)
HATU	<i>O</i> -(7-azabenzotriazol-1-yl)- <i>N,N,N',N'</i> -tetramethyluronium hexafluorophosphate
HCT-116	human tumour cell line
HEPES	4-(2-hydroxyethyl)piperazine-1-ethanesulphonic acid
HOAc	acetic acid
HOAt	1-hydroxy-7-azabenzotriazole
HOBt	1-hydroxy-1 <i>H</i> -benzotriazole hydrate
HOSu	<i>N</i> -hydroxysuccinimide
HPLC	high-pressure high-performance liquid chromatography
HT29	human tumour cell line
IC <sub>50</sub>	half-maximal inhibitory concentration
i.m.	intramuscularly, administration into the muscle
i.v.	intravenously, administration into the vein
conc.	concentrated
LC-MS	liquid chromatography-coupled mass spectroscopy
LLC-PK1 cells	Lewis lung carcinoma pork kidney cell line
L-MDR	human MDR1 transfected LLC-PK1 cells
m	multiplet (in NMR)
MDR1	multidrug resistance protein 1
MeCN	acetonitrile
min	minute(s)
MS	mass spectrometry
MTT	3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl-2 <i>H</i> -tetrazolium bromide
NCI-H292	human tumour cell line
NCI-H520	human tumour cell line
NMM	<i>N</i> -methyilmorpholine
NMP	<i>N</i> -methyl-2-pyrrolidinone
NMR	nuclear magnetic resonance spectrometry

NMRI	mouse strain, originates from the Naval Medical Research Institute (NMRI)
NSCLC	non small cell lung cancer
PBS	phosphate-buffered salt solution
Pd/C	palladium on activated carbon
P-gp	P-glycoprotein, a transporter protein
PNGaseF	enzyme for cleaving sugar
quant.	quantitative (in yield)
quart	quartet (in NMR)
quint	quintet (in NMR)
R <sub>f</sub>	retention index (in TLC)
RT	room temperature
R <sub>t</sub>	retention time (in HPLC)
s	singlet (in NMR)
s.c.	subcutaneously, administration under the skin
SCC-4	human tumour cell line
SCC-9	human tumour cell line
SCID mice	test mice with severe combined immunodeficiency
t	triplet (in NMR)
TBAF	tetra-n-butylammonium fluoride
TEMPO	(2,2,6,6-tetramethylpiperidin-1-yl)oxyl
tert.	tertiary
TFA	trifluoroacetic acid
THF	tetrahydrofuran
T3P <sup>®</sup>	2,4,6-tripropyl-1,3,5,2,4,6-trioxatriphosphinane 2,4,6-trioxide
UV	ultraviolet spectrometry
v/v	ratio by volume (of a solution)
Z	benzyloxycarbonyl

If, in the context of the present disclosure, no temperature is given in the description of a reaction, room temperature should always be assumed.

5

#### HPLC and LC-MS methods:

Method 1 (LC-MS):

Instrument: Waters ACQUITY SQD UPLC system; column: Waters Acquity UPLC HSS T3 1.8  $\mu$  50 x 1 mm; mobile phase A: 1 l of water + 0.25 ml of 99% strength formic acid; mobile phase B: 1 l of acetonitrile + 0.25 ml of 99% strength formic acid; gradient: 0.0 min 90% A  $\rightarrow$  1.2 min 5% A  $\rightarrow$  2.0 min 5% A oven: 50°C; flow rate: 0.40 ml/min; UV detection: 208 - 400 nm.

10 Method 2 (LC-MS):

MS instrument type: Waters Synapt G2S; UPLC instrument type: Waters Acquity I-CLASS; column: Waters, BEH300, 2.1 x 150 mm, C18 1.7  $\mu$ m; mobile phase A: 1 l of water + 0.01% formic acid; mobile phase B: 1 l of acetonitrile + 0.01% formic acid; gradient: 0.0 min 2% B  $\rightarrow$  1.5 min 2% B  $\rightarrow$  8.5 min 95% B  $\rightarrow$  10.0 min 95% B; oven: 50°C; flow rate: 0.50 ml/min; UV detection: 220 nm

20 Method 3 (LC-MS):

MS instrument: Waters (Micromass) QM; HPLC instrument: Agilent 1100 series; column: Agilent ZORBAX Extend-C18 3.0 x 50 mm 3.5 micron; mobile phase A: 1 l of water + 0.01 mol of ammonium carbonate, mobile phase B: 1 l of acetonitrile; gradient: 0.0 min 98% A  $\rightarrow$  0.2min 98% A  $\rightarrow$  3.0 min 5% A  $\rightarrow$  4.5 min 5% A ; oven: 40°C; flow rate: 1.75 ml/min; UV detection: 210 nm

30 Method 4 (LC-MS):

MS instrument type: Waters Synapt G2S; UPLC instrument type: Waters Acquity I-CLASS; column: Waters, HSST3, 2.1 x 50 mm, C18 1.8  $\mu$ m; mobile phase A: 1 l of water + 0.01% formic acid; mobile phase B: 1 l of acetonitrile + 0.01% formic acid; gradient: 0.0 min 10% B  $\rightarrow$  0.3 min 10% B  $\rightarrow$  1.7 min 95% B  $\rightarrow$  2.5 min 95% B; oven: 50°C; flow rate: 1.20 ml/min; UV detection: 210 nm

35 Method 5 (LC-MS):



Instrument: Waters ACQUITY SQD UPLC system; column: Waters Acquity UPLC HSS T3 1.8  $\mu$  50 x 1 mm; mobile phase A: 1 l of water + 0.25 ml of 99% strength formic acid; mobile phase B: 1 l of acetonitrile + 0.25 ml of 99% strength formic acid; gradient: 0.0 min 95% A  $\rightarrow$  6.0 min 5% A  $\rightarrow$  7.5 min 5% A oven: 50°C; flow rate: 0.35 ml/min; UV detection: 210 - 400 nm.

Method 6 (LC-MS):

Instrument: Micromass Quattro Premier with Waters UPLC Acquity; column: Thermo Hypersil GOLD 1.9  $\mu$  50 x 1 mm; mobile phase A: 1 l of water + 0.5 ml of 50% strength formic acid; mobile phase B: 1 l of acetonitrile + 0.5 ml of 50% strength formic acid; gradient: 0.0 min 97% A  $\rightarrow$  0.5 min 97% A  $\rightarrow$  3.2 min 5% A  $\rightarrow$  4.0 min 5% A; oven: 50°C; flow rate: 0.3 ml/min; UV detection: 210 nm.

Method 7 (LC-MS):

Instrument: Agilent MS Quad 6150; HPLC: Agilent 1290; column: Waters Acquity UPLC HSS T3 1.8  $\mu$  50 x 2.1 mm; mobile phase A: 1 l of water + 0.25 ml of 99% strength formic acid; mobile phase B: 1 l of acetonitrile + 0.25 ml of 99% strength formic acid; gradient: 0.0 min 90% A  $\rightarrow$  0.3 min 90% A  $\rightarrow$  1.7 min 5% A  $\rightarrow$  3.0 min 5% A; oven: 50°C; flow rate: 1.20 ml/min; UV detection: 205 - 305 nm.

Method 8 (LC-MS):

MS instrument type: Waters Synapt G2S; UPLC instrument type: Waters Acquity I-CLASS; column: Waters, HSST3, 2.1 x 50 mm, C18 1.8  $\mu$ m; mobile phase A: 1 l of water + 0.01% formic acid; mobile phase B: 1 l of acetonitrile + 0.01% formic acid; gradient: 0.0 min 2% B  $\rightarrow$  2.0 min 2% B  $\rightarrow$  13.0 min 90% B  $\rightarrow$  15.0 min 90% B; oven: 50°C; flow rate: 1.20 ml/min; UV detection: 210 nm

Method 9: LC-MS-Prep purification method for Examples 181-191 (Method LIND-LC-MS-Prep)

MS instrument: Waters; HPLC instrument: Waters (column Waters X-Bridge C18, 19 mm x 50 mm, 5 µm, mobile phase A: water + 0.05% ammonia, mobile phase B: acetonitrile (ULC) with gradient; flow rate: 40 ml/min; UV detection: DAD; 210 - 400 nm).

5

or:

MS instrument: Waters; HPLC instrument: Waters (column Phenomenex Luna 5µ C18(2) 100A, AXIA Tech. 50 x 21.2 mm, mobile phase A: water + 0.05% formic acid, mobile phase B: acetonitrile (ULC) with gradient; flow rate: 40 ml/min; UV detection: DAD; 210 - 400 nm).

Method 10: LC-MS analysis method for Examples 181-191 (LIND\_SQD\_SB\_AQ)

MS instrument: Waters SQD; HPLC instrument: Waters UPLC; column: Zorbax SB-Aq (Agilent), 50 mm x 2.1 mm, 1.8 µm; mobile phase A: water + 0.025% formic acid, mobile phase B: acetonitrile (ULC) + 0.025% formic acid; gradient: 0.0 min 98%A - 0.9 min 25%A - 1.0 min 5%A - 1.4 min 5%A - 1.41 min 98%A - 1.5 min 98%A; oven: 40°C; flow rate: 0.600 ml/min; UV detection: DAD; 210 nm.

Method 11 (HPLC):

25

Instrument: HP1100 Series

Column: Merck Chromolith SpeedROD RP-18e, 50-4.6 mm, Cat.

No.1.51450.0001, precolumn Chromolith Guard Cartridge Kit, RP-18e,

5-4.6mm, Cat. No. 1.51470.0001

35 Gradient: flow rate 5 ml/min

injection volume 5 µl

solvent A: HClO<sub>4</sub> (70% strength) in water (4 ml/l)

solvent B: acetonitrile

5 start 20% B

0.50 min 20% B

3.00 min 90% B

10

3.50 min 90% B

3.51 min 20% B

15 4.00 min 20% B

column temperature: 40°C

Wavelength: 210 nm

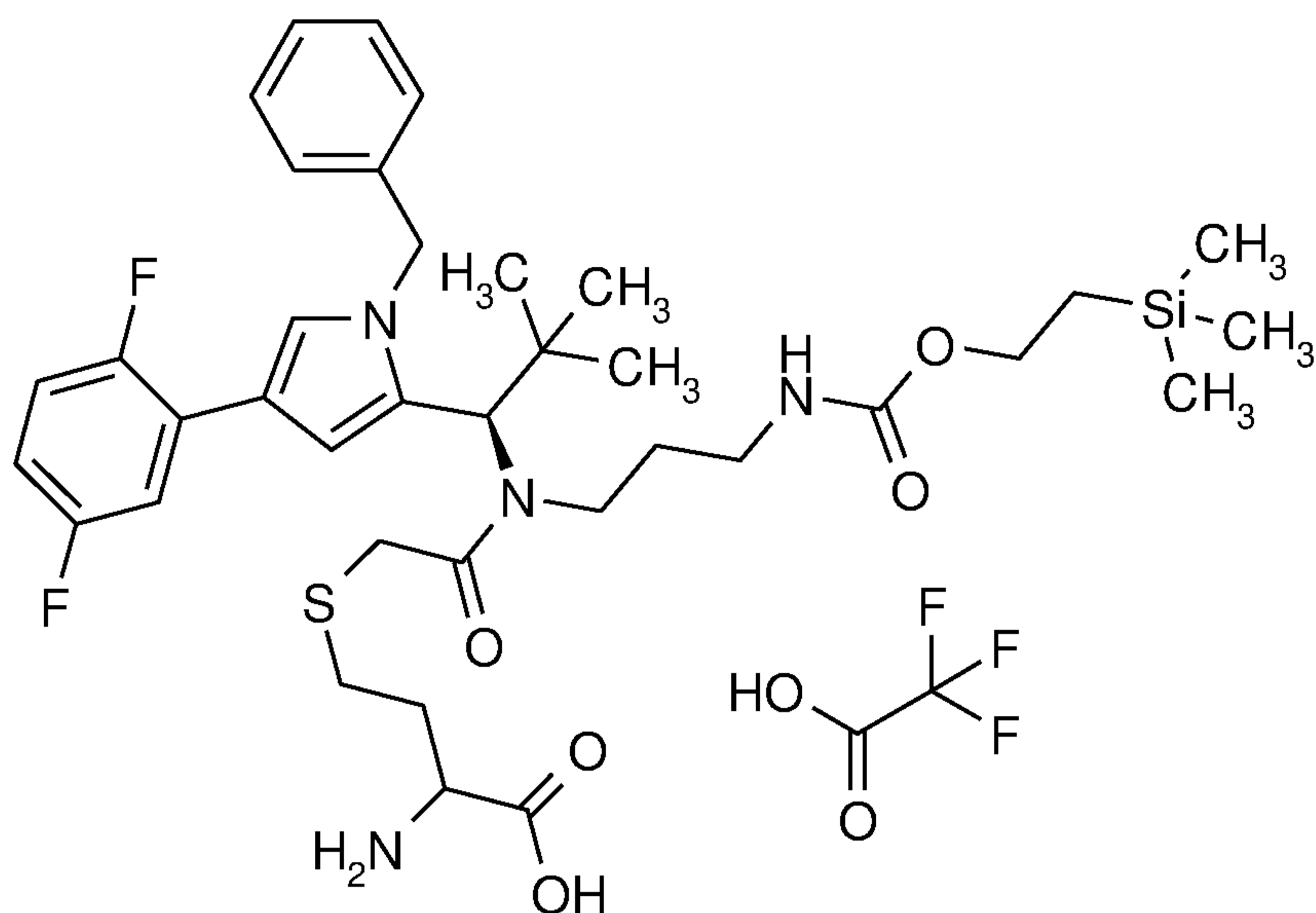
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All reactants or reagents whose preparation is not described explicitly hereinafter were purchased commercially from generally accessible sources. For all other reactants or reagents whose preparation likewise is not described hereinafter and which were not commercially obtainable or were obtained from sources which are not generally accessible, a reference is given to the published literature in which their preparation is described.

30 **Starting materials and intermediates:**

**Intermediate C11**

35 R/S-(11-{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-homocysteine / trifluoroacetate (1:1)



990.0 mg (2.79 mmol) of (1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropan-1-amine  
 5 were initially charged in 15.0 ml of dichloromethane, and 828.8 mg (3.91 mmol) of sodium triacetoxyborohydride and 129.9 mg (3.21 mmol) of acetic acid were added, and the mixture was stirred at RT for 5 min. 698.1 mg (3.21 mmol) of 2-(trimethylsilyl)ethyl (3-oxopropyl)carbamate (Intermediate L58)  
 10 dissolved in 15.0 ml of dichloromethane were added, and the reaction mixture was stirred at RT overnight. The reaction mixture was diluted with ethyl acetate and the organic phase was washed in each case twice with saturated sodium carbonate solution and saturated NaCl solution. The organic phase was  
 15 dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was purified using silica gel (mobile phase: dichloromethane/methanol 100:2). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 1.25 g (73% of theory) of the  
 20 compound 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino]propyl]carbamate.

LC-MS (Method 1):  $R_t = 1.09$  min; MS (ESIpos):  $m/z = 556$  (M+H)<sup>+</sup>.  
 25

151.4 mg (1.5 mmol) of triethylamine and 161.6 mg (1.43 mmol) of chloroacetyl chloride were added to 400.0 mg (0.65 mmol) of

2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-

5 dimethylpropyl}amino)propyl]carbamate. The reaction mixture was stirred at RT overnight. Ethyl acetate was added to the reaction mixture and the organic phase was washed three times with water and once with saturated NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was purified using silica gel (mobile phase: cyclohexane/ethyl acetate 3:1). The solvents  
10 were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 254.4 mg (57% of theory) of the compound 2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate.

15 LC-MS (Method 1):  $R_t = 1.49$  min; MS (ESI<sup>neg</sup>):  $m/z = 676$  ( $M+HCOO^-$ ).

117.4 mg (0.19 mmol) of 2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate were  
20 dissolved in 10.0 ml of isopropanol, and 928.4  $\mu$ l of 1M NaOH and 50.2 mg (0.37 mmol) of DL-homocysteine were added. The reaction mixture was stirred at 50°C for 4.5 h. Ethyl acetate was added to the reaction mixture and the organic phase was washed with  
25 saturated sodium bicarbonate solution and sat. NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was purified by preparative RP-HPLC (column: Reprosil 250x40; 10 $\mu$ ,  
30 flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 75.3 mg (48% of theory) of the title compound.

35 LC-MS (Method 1):  $R_t = 1.24$  min; MS (ESI<sup>pos</sup>):  $m/z = 731$  ( $M+H$ )<sup>+</sup>.

<sup>1</sup>H NMR (400 MHz, DMSO- $d_6$ ):  $\delta$  [ppm] = 0.03 (s, 9H), 0.40 (m, 1H), 0.75-0.91 (m, 11H), 1.30 (m, 1H), 1.99-2.23 (m, 2H), 2.63-2.88

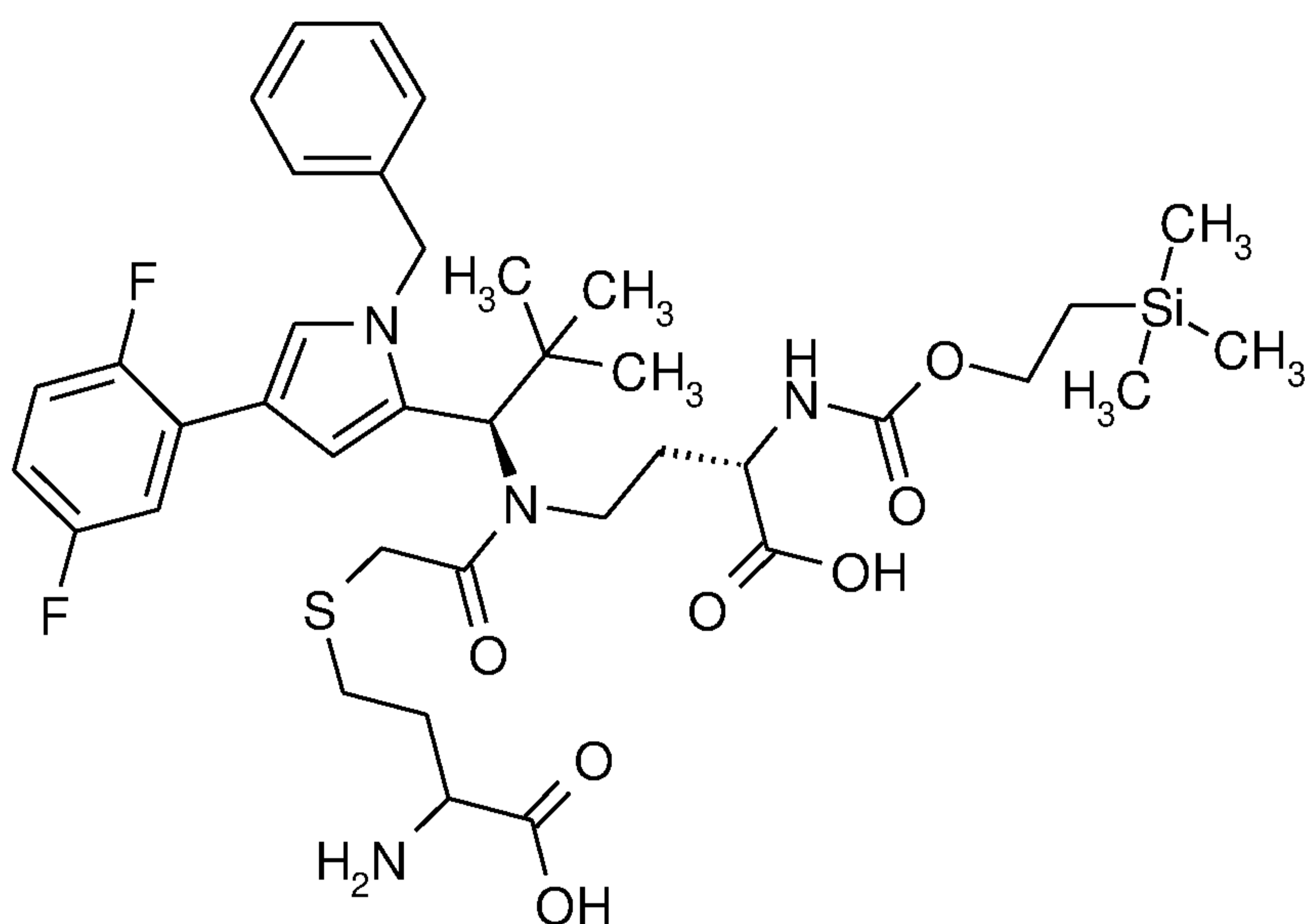
(m, 4H), 3.18-3.61 (m, 5H), 3.79-4.10 (m, 3H), 4.89 (d, 1H), 4.89 (d, 1H), 5.16 (d, 1H), 5.56 (s, 1H), 6.82 (m, 1H), 6.91 (s, 1H), 6.97 (m, 1H), 7.13-7.38 (m, 6H), 7.49 (s, 1H), 7.63 (m, 1H), 8.26 (s, 3H).

5

### Intermediate C12

R/S-[(8S)-11-{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-8-carboxy-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl]homocysteine

10



The synthesis was carried out analogously to the synthesis of Intermediate C11 using methyl (2S)-4-oxo-2-([2-(trimethylsilyl)ethoxy]carbonyl)amino)butanoate (Intermediate L57) and Intermediate C52 as starting materials.

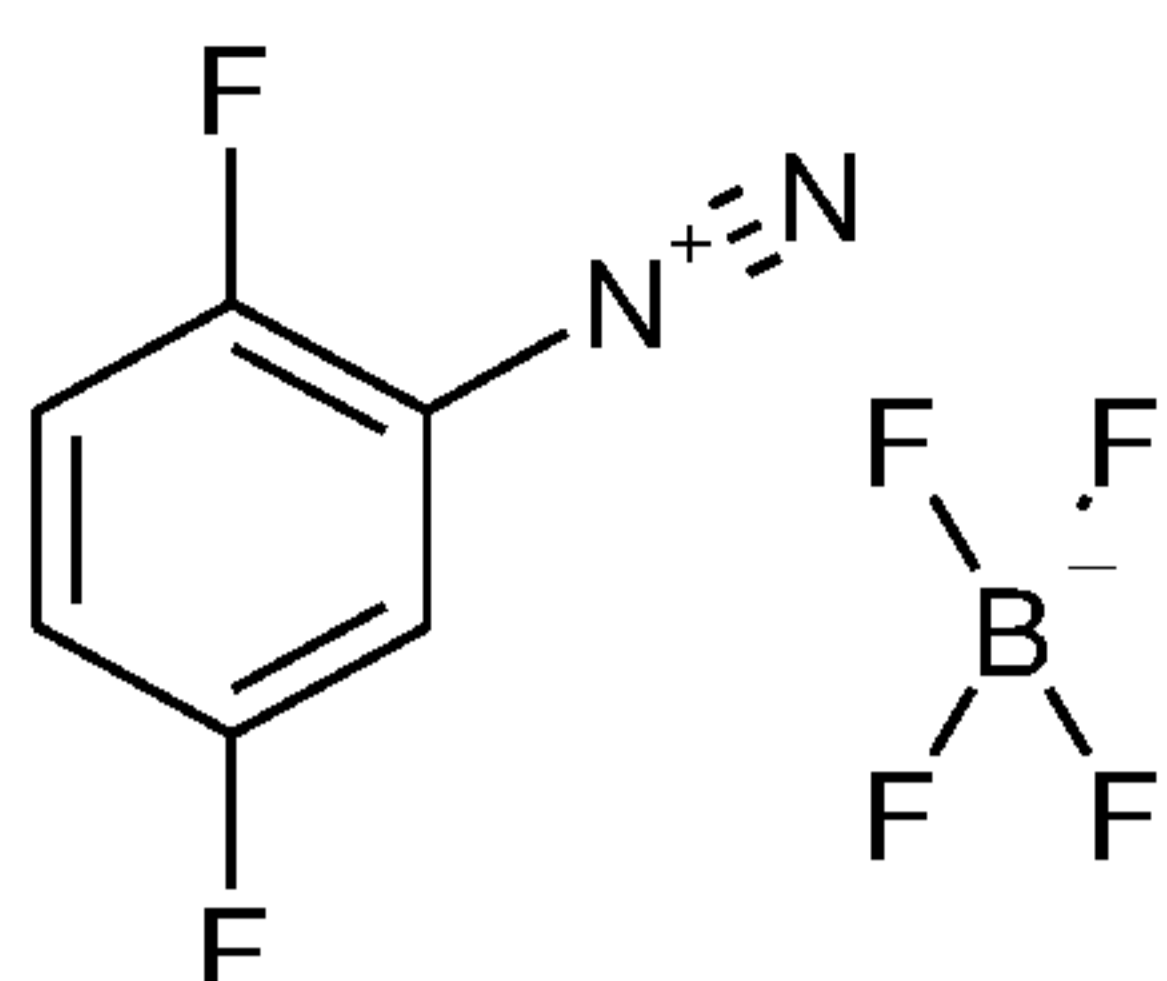
15

LC-MS (Method 1):  $R_t = 1.18$  min; MS (ESIpos):  $m/z = 775$  (M+H)<sup>+</sup>.

20

### Intermediate C42

2,5-Difluorobenzediazonium tetrafluoroborate



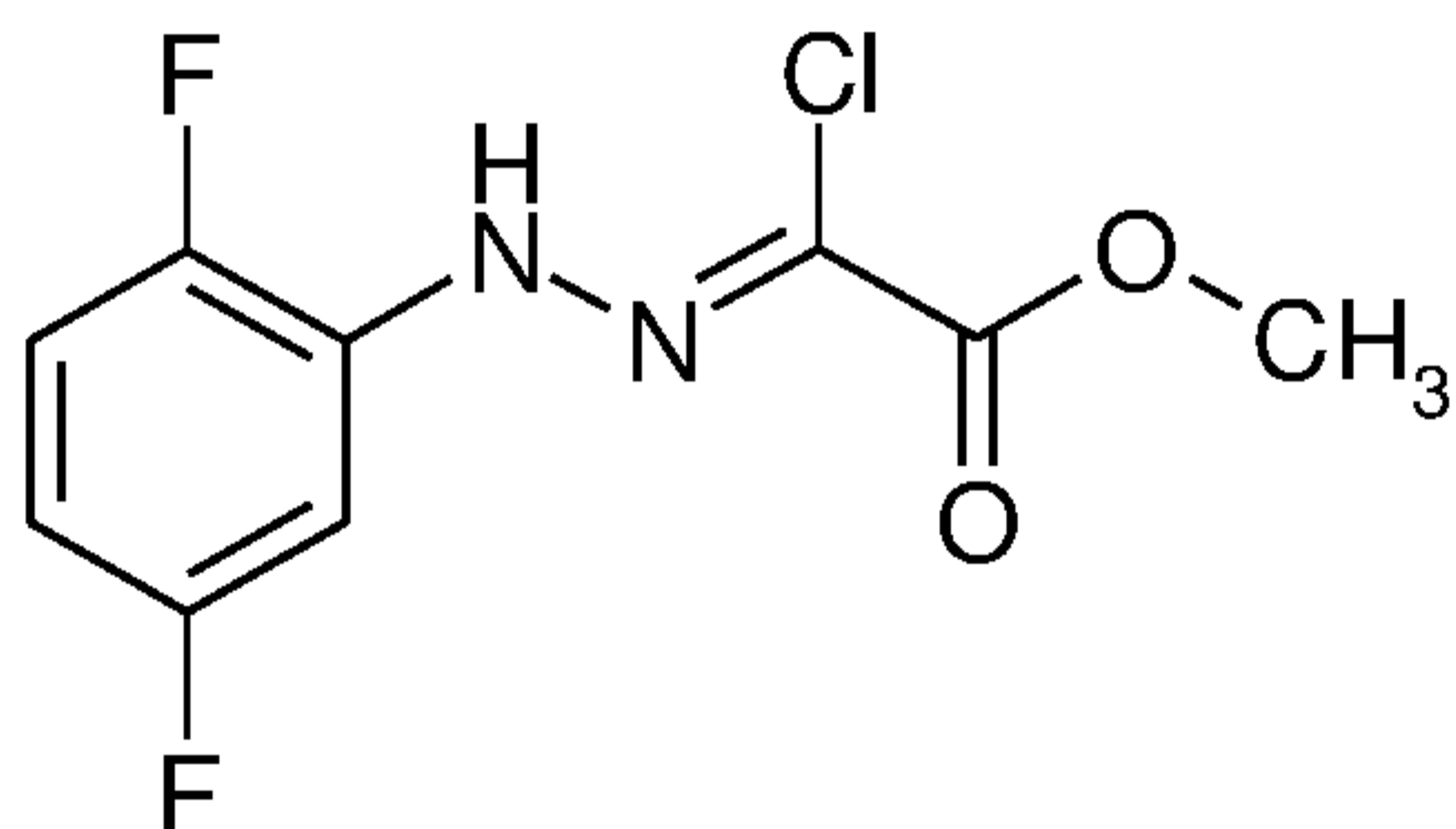
3.00 g (21.16 mmol, 2.68 ml) of boron trifluoride-diethyl ether complex were initially charged, and 1.37 g (10.58 mmol) of 2,5-difluoroaniline dissolved in 27 ml of absolute THF were slowly added dropwise at 0°C. At -10°C, a solution of 1.61 g (13.75 mmol, 1.85 ml) of isoamyl nitrite dissolved in 3 ml of absolute THF was added dropwise, and stirring was continued at the same temperature for 30 min. 15 ml of diethyl ether were added and the precipitated diazonium salt was filtered off, washed with a little diethyl ether and dried under high vacuum. This gave 2.27 g of the target compound (94% of theory).

LC-MS (Method 6):  $R_t = 0.24$  min; MS (ESIpos):  $m/z = 141$  [M]<sup>+</sup>.

<sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>):  $\delta$  [ppm] = 8.11-8.17 (m, 1H), 8.36-8.43 (m, 1H), 8.69-8.73 (m, 1H).

### Intermediate C43

Methyl chloro[2-(2,5-difluorophenyl)hydrazinyldene]acetate



Under an atmosphere of argon, 3.63 g (24.13 mmol) of methyl 2-chloro-3-oxobutanoate were initially charged in 100 ml of water, and 48.90 g (618.19 mmol, 50.00 ml) of pyridine were added at -5°C and the mixture was stirred at this temperature for 10 min. At -5°C, 5.00 g (21.94 mmol) of 2,5-difluorobenzenediazonium tetrafluoroborate were then added, resulting in the formation

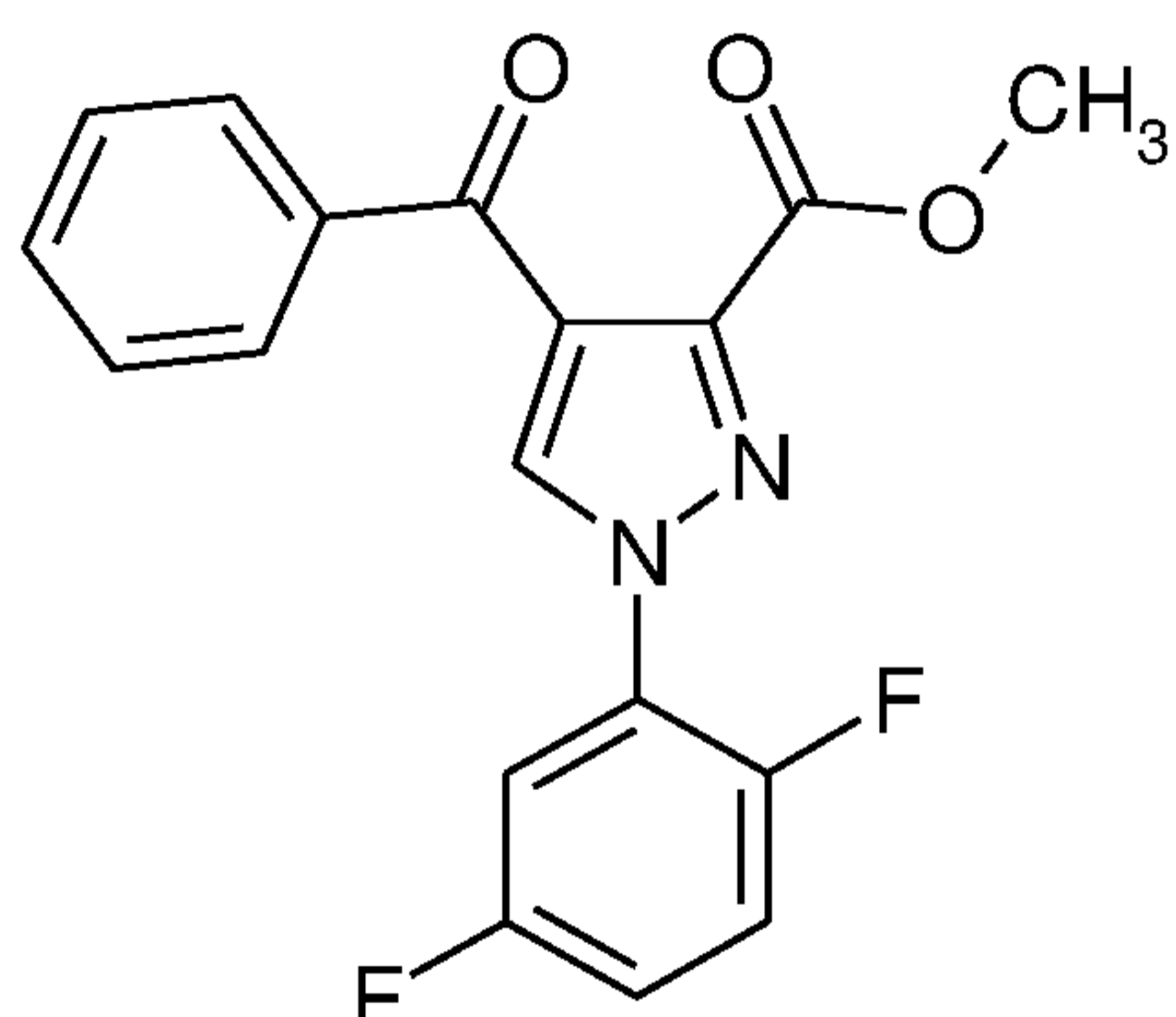
of an orange suspension. The mixture was stirred at this temperature for 30 min and the reaction was diluted with water and extracted three times with dichloromethane. The combined organic phases were washed with saturated sodium chloride solution, dried over sodium sulphate, concentrated on a rotary evaporator and dried under high vacuum. This gave 5.52 g of the target compound (97% of theory, purity according to LC/MS = 96%).

10 LC-MS (Method 1):  $R_t = 1.03$  min; MS (ESIpos):  $m/z = 249$   $[M+H]^+$ .

$^1\text{H}$  NMR (400 MHz,  $\text{DMSO-d}_6$ ):  $\delta$  [ppm] = 3.85 (s, 3H), 6.88-6.94 (m, 1H), 7.16-7.21 (m, 1H), 7.31-7.37 (m, 1H), 10.00 (s, 1H).

15 **Intermediate C44**

Methyl 4-benzoyl-1-(2,5-difluorophenyl)-1H-pyrazole-3-carboxylate



20

3.50 g (13.52 mmol) of methyl chloro[2-(2,5-difluorophenyl)hydrazinyliden]acetate (purity according to LC/MS 96%) were dissolved in 9 ml of absolute toluene, 2.61 g (14.87 mmol) of (2E)-3-(dimethylamino)-1-phenylprop-2-en-1-one and 3.01 g (29.73 mmol), 4.14 ml) of triethylamine were added and the mixture was stirred at room temperature for 16 h. The reaction mixture was concentrated on a rotary evaporator and the residue separated by preparative HPLC (mobile phase: ACN/water with 0.1% formic acid, gradient). This gave 1.79 g (39% of theory) of the target compound.

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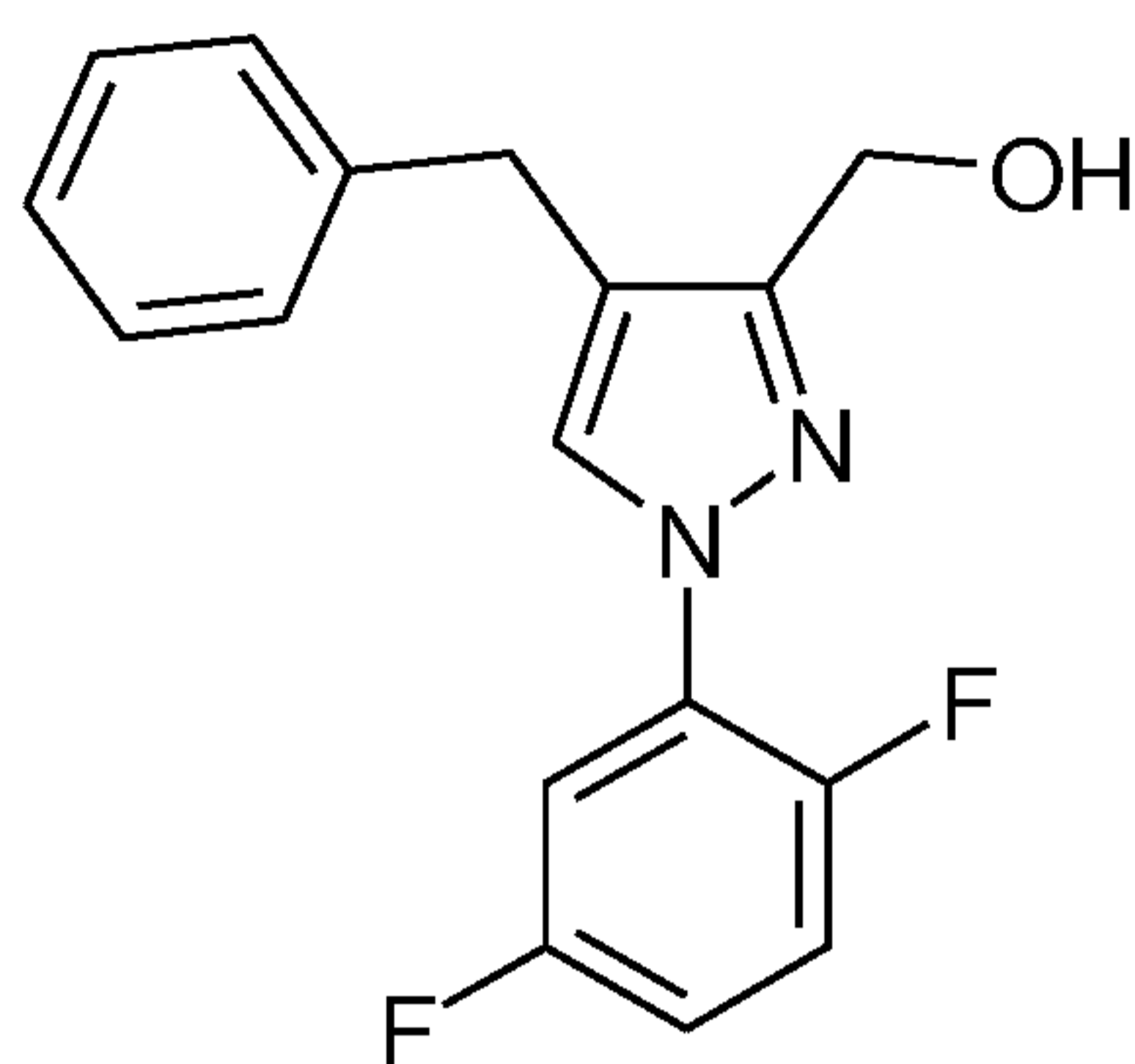
LC-MS (Method 1):  $R_t = 1.07$  min; MS (ESIpos):  $m/z = 343$   $[M+H]^+$ .

$^1\text{H}$  NMR (400 MHz,  $\text{DMSO-d}_6$ ):  $\delta$  [ppm] = 3.86 (s, 3H), 7.44-7.50 (m, 1H), 7.55-7.72 (m, 4H), 7.81-7.87 (m, 3H), 8.80 (d, 1H).

5

### Intermediate C45

[4-Benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]methanol



10

3.18 g (8.92 mmol) of methyl 4-benzoyl-1-(2,5-difluorophenyl)-1H-pyrazole-3-carboxylate (purity according to LC/MS = 96%) were initially charged in 50 ml of trifluoroacetic acid, 8.74 g (75.13 mmol, 12 ml) of triethylsilane were added dropwise and the mixture was stirred at room temperature for 1 h. The reaction mixture was concentrated on a rotary evaporator and dried under high vacuum. The residue obtained was taken up in 120 ml of absolute THF, and 2.89 g (33.63 mmol, 33.63 ml) of borane-tetrahydrofuran complex were added dropwise at 0°C. The mixture was stirred at RT overnight. Owing to the low conversion, another 12.33 ml (12.33 mmol) of a 1M lithium borohydride solution in THF were added. The mixture was stirred at room temperature for 1 h, at 60°C for 30 min and at 80°C for 2 h. At 0°C, the reaction was carefully quenched with 60 ml of saturated sodium bicarbonate solution. The mixture was extracted twice with in each case 100 ml of ethyl acetate, the combined organic phases were dried over sodium sulphate and concentrated on a rotary evaporator and the residue was dried under high vacuum. This gave 2.67 g (76% of theory, purity = 96%) of the target compound.

30

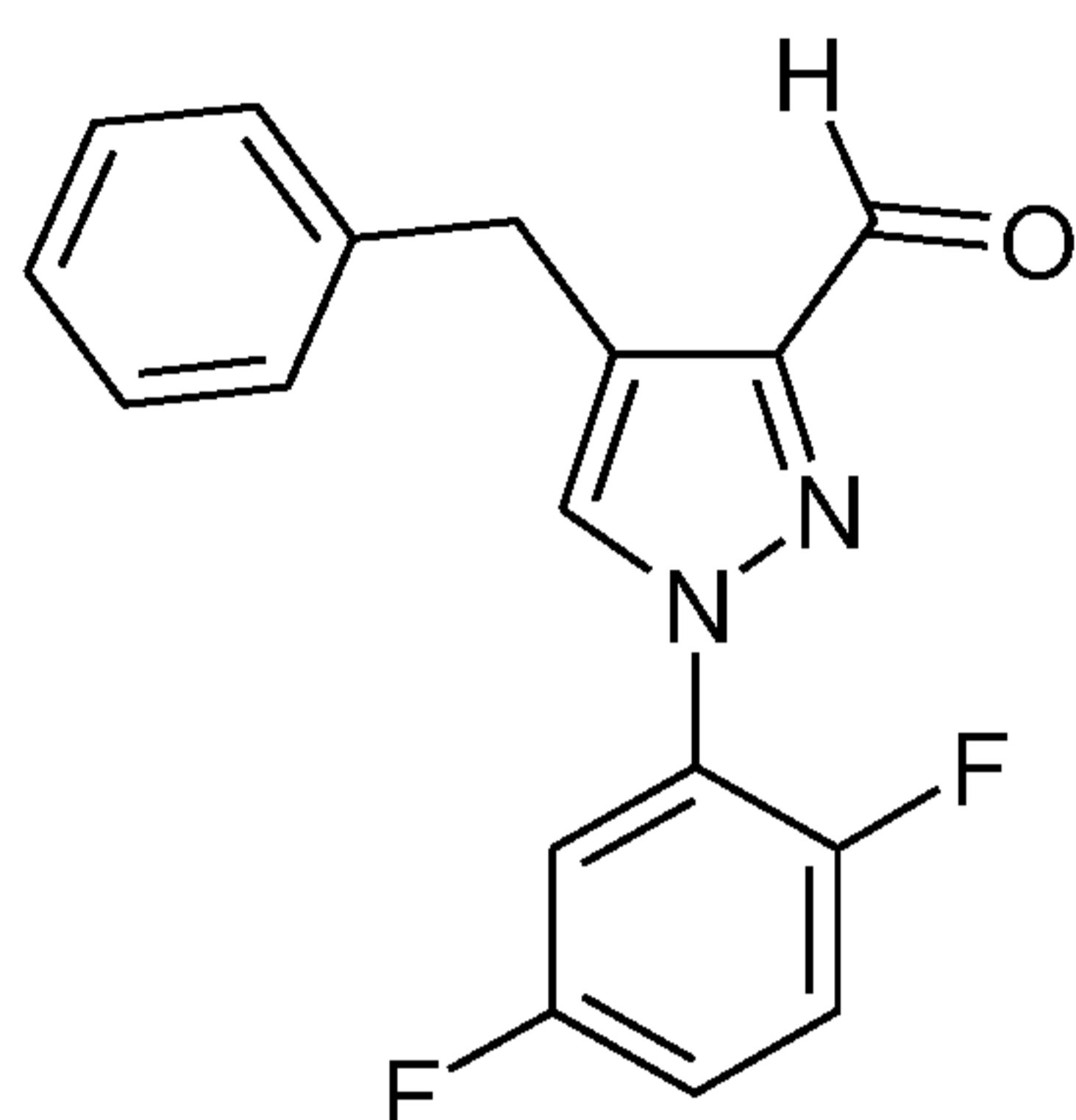
LC-MS (Method 3):  $R_t = 2.79$  min; MS (ESIpos):  $m/z = 329$   $[M+H]^+$ .

$^1\text{H}$  NMR (400 MHz, DMSO- $d_6$ ):  $\delta$  [ppm] = 3.91 (s, 2H), 4.45 (d, 2H), 6.51 (s, 1H), 7.18-7.23 (m, 2H), 7.27-7.32 (m, 4H), 7.46-7.53 (m, 1H), 7.60-7.65 (m, 1H), 7.95 (d, 1H).

5

### Intermediate C46

4-Benzyl-1-(2,5-difluorophenyl)-1H-pyrazole-3-carbaldehyde



10

2.66 g (8.50 mmol) of [4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]methanol (purity 96%) were dissolved in 150 ml of dichloromethane, and 4.33 g (10.20 mmol) of Dess-Martin periodinane were added a little at a time. The mixture was stirred at room temperature for 2 h, 100 ml of a semiconcentrated sodium bicarbonate solution and 100 ml of a 10% strength sodium thiosulphate solution were then added and the mixture was stirred for 20 min. The organic phase was separated off, dried over sodium sulphate and concentrated under high vacuum. This gave 2.35 g (88% of theory, purity = 95%) of the target compound.

15

20

LC-MS (Method 7):  $R_t$  = 1.49 min; MS (ESIpos):  $m/z$  = 299  $[\text{M}+\text{H}]^+$ .

$^1\text{H}$  NMR (400 MHz, DMSO- $d_6$ ):  $\delta$  [ppm] = 4.12 (s, 2H), 7.17-7.21 (m, 1H), 7.27-7.31 (m, 4H), 7.37-7.42 (m, 1H), 7.57-7.62 (m, 1H), 7.75-7.78 (m, 1H), 8.22 (d, 1H), 10.06 (s, 1H).

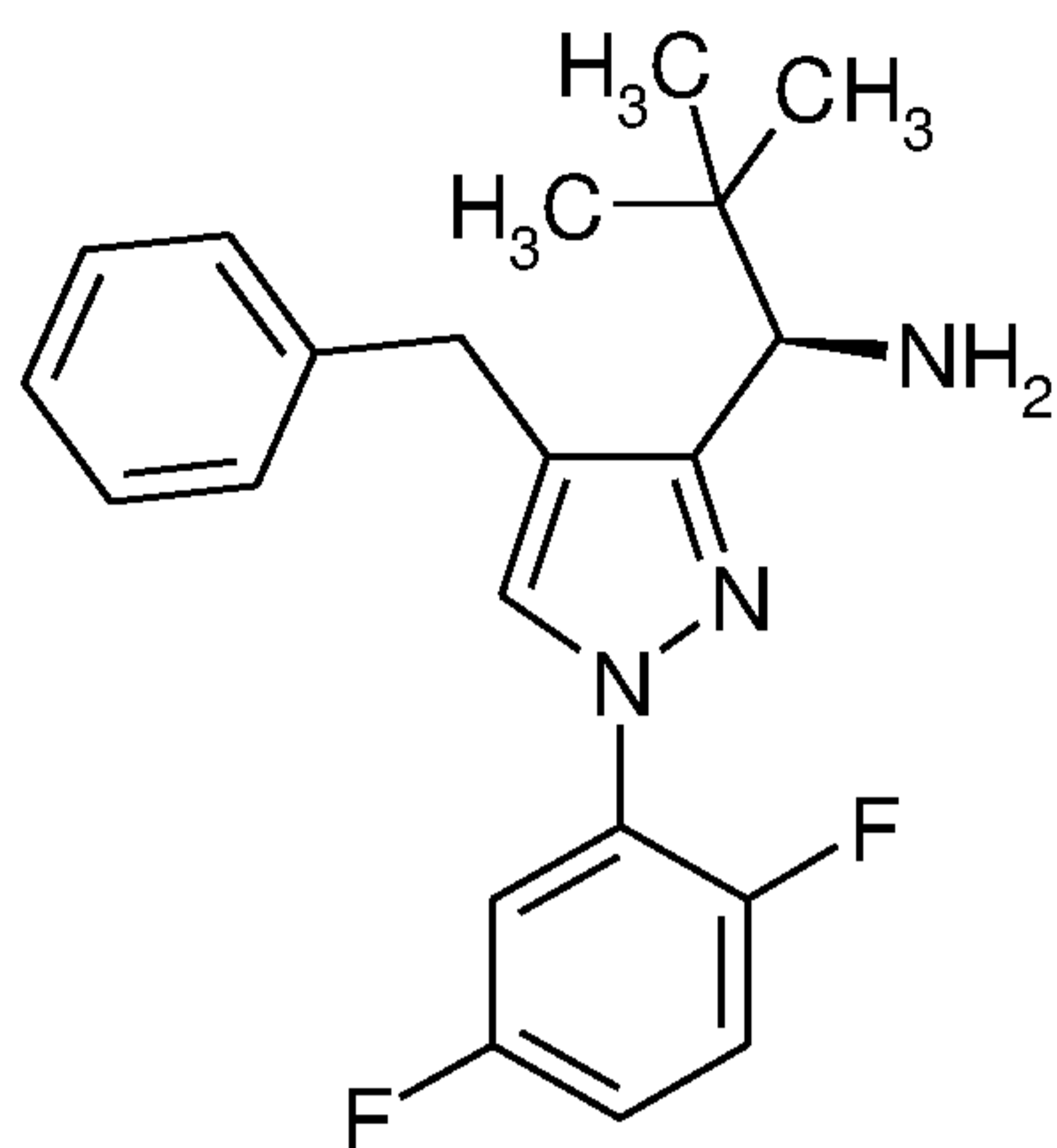
25

### Intermediate C47

30

(1R)-1-[4-Benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-

dimethylpropan-1-amine



5 2.35 g (7.56 mmol) of 4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazole-3-carbaldehyde were dissolved in 25 ml of absolute THF, and 1.10 g (9.08 mmol) of (R)-(+)-2-methyl-2-propanesulphinamide and 4.73 g (16.64 mmol) of titanium(IV) isopropoxide were added. The reaction mixture was stirred at room temperature for 16 h,  
10 and 20 ml of a saturated sodium chloride solution and 30 ml of ethyl acetate were added. About 3 g of kieselguhr were then added, and the mixture was boiled under reflux for 1 h. The mixture was filtered and the organic phase was separated from the filtrate. The aqueous phase was extracted with ethyl acetate  
15 and the combined organic phases were washed with saturated sodium chloride solution, dried over sodium sulphate, concentrated on a rotary evaporator and dried under high vacuum.

The residue was used further without further purification.

20 Under an atmosphere of argon, the residue was dissolved in 60 ml of absolute THF and cooled to  $-78^{\circ}\text{C}$ , and 14.5 ml (23.24 mmol) of a solution of tert-butyllithium in pentane ( $c = 1.6 \text{ mol/l}$ ) were added dropwise. The reaction was stirred at  $-78^{\circ}\text{C}$  for 3 h  
25 and then quenched with 5 ml of methanol and 15 ml of a saturated ammonium chloride solution. With stirring, the reaction mixture was allowed to warm to room temperature (about 30 min.). The mixture was extracted with ethyl acetate and the organic phase was extracted with saturated sodium chloride solution,  
30 concentrated on a rotary evaporator and dried under high vacuum.

The residue was used further without further purification.

The residue was taken up in 30 ml of THF and 6 ml of methanol, 6 ml (24.00 mmol) of a 4N hydrogen chloride solution in dioxane were added and the mixture was stirred at room temperature for 1 h. 15 ml of saturated sodium carbonate solution were then added, and the mixture was extracted with ethyl acetate. The organic phase was separated off, concentrated on a rotary evaporator and dried under high vacuum. The residue was separated by preparative HPLC (mobile phase: ACN/water, gradient). This gave two fractions of the target compound. The first fraction yielded 1.31 g (72% of theory, LC/MS purity = 97%) and the second 0.37 g (17% of theory, LC/MS purity = 83%) of product.

15

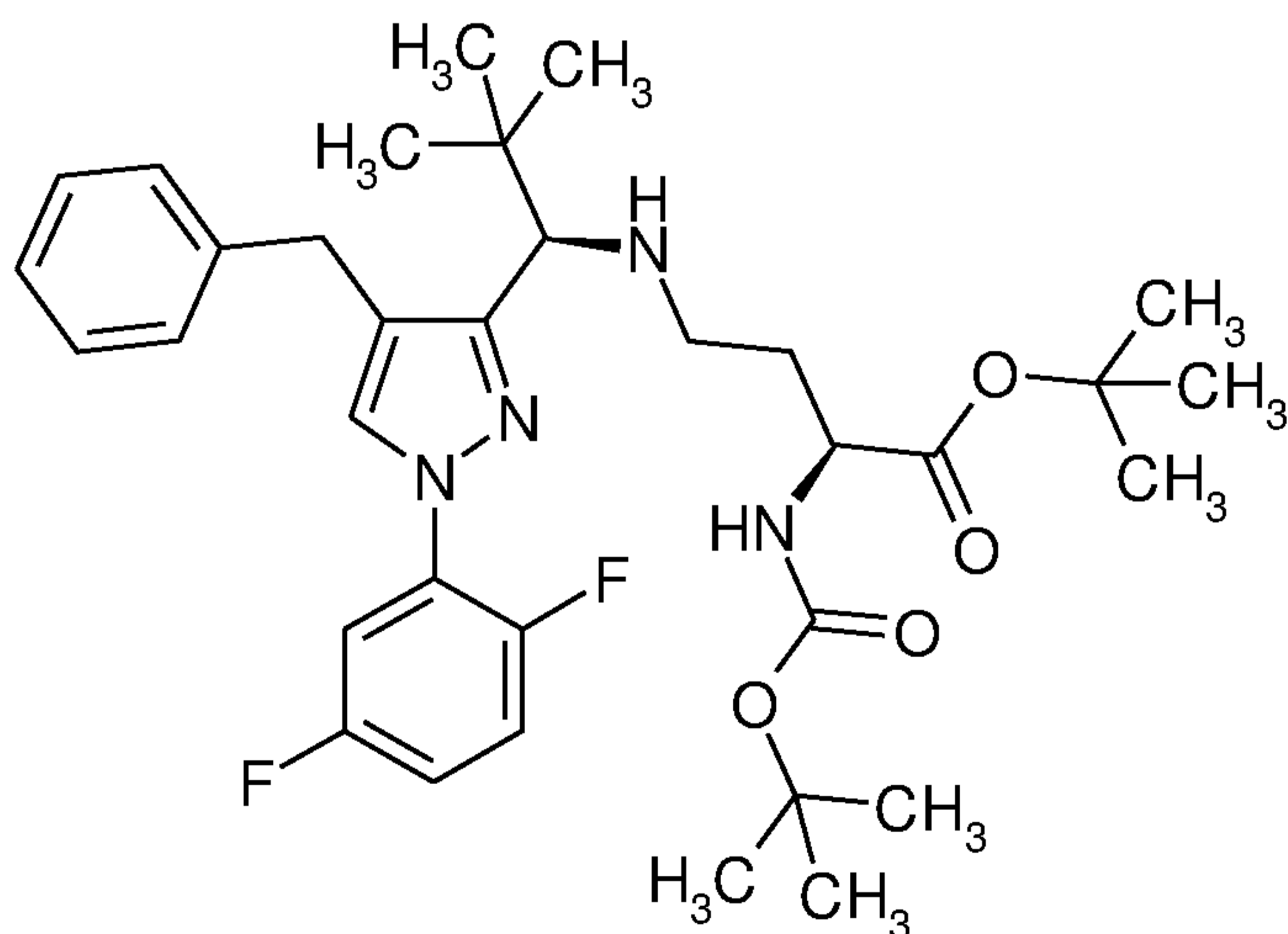
LC-MS (Method 1):  $R_t = 0.88$  min; MS (ESIpos):  $m/z = 356$   $[M+H]^+$ .

$^1H$  NMR (400 MHz, DMSO- $d_6$ ):  $\delta$  [ppm] = 0.91 (s, 9H), 1.71 (s, 2H), 3.59 (s, 1H), 3.87 (s, 2H), 7.17-7.32 (m, 6H), 7.45-7.51 (m, 1H), 7.61-7.65 (m, 1H), 7.84 (s br, 1H).

20

### Intermediate C48

tert-Butyl (2S)-4-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)amino)-2-[(tert-butoxycarbonyl)amino]butanoate



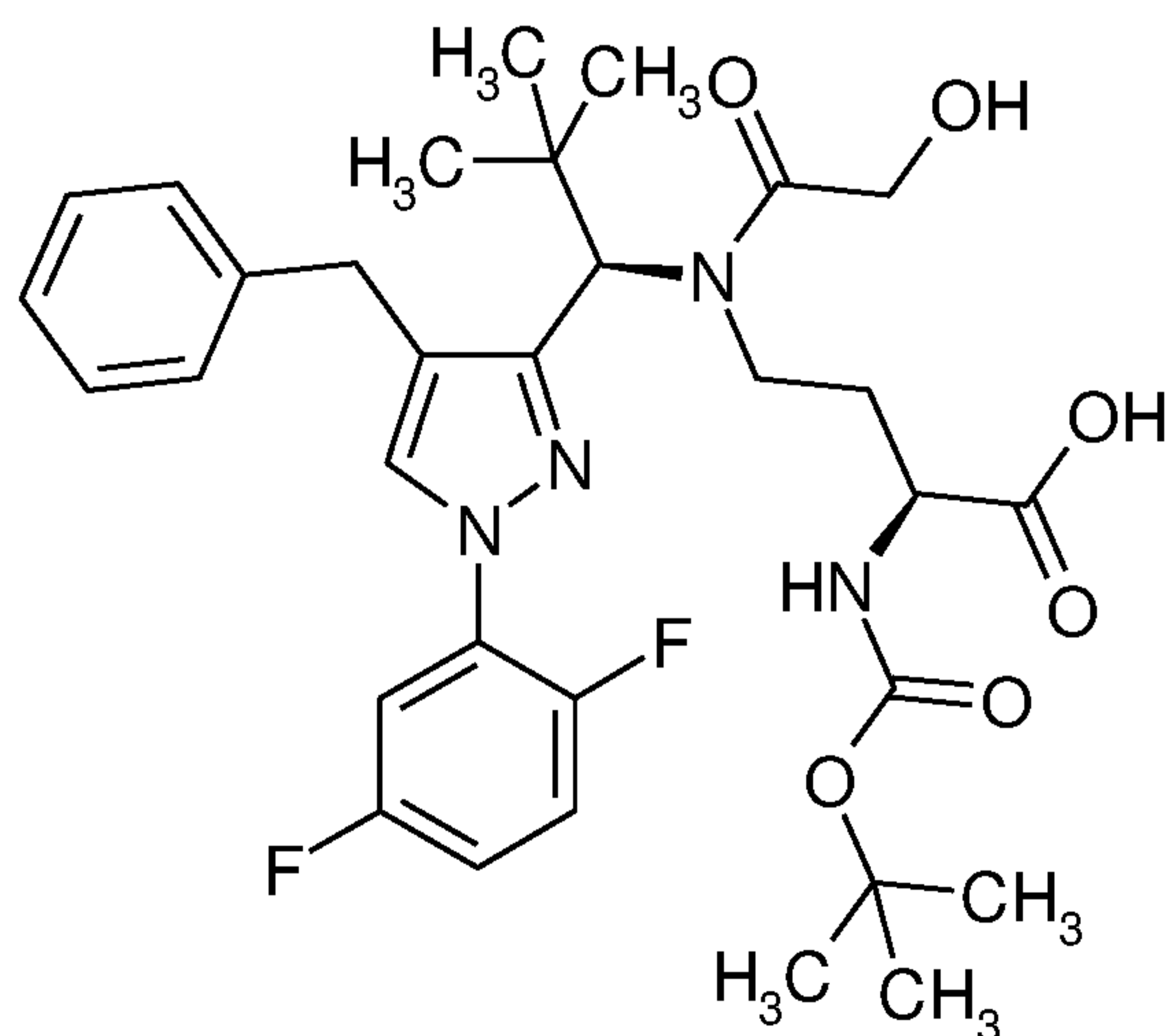
1.28 g (3.35 mmol, LC/MS purity 93%) of (1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropan-1-amine were dissolved in 100 ml of absolute dichloromethane, and 261 mg (4.35 mmol, 250  $\mu$ l) of acetic acid and 1.14 g (4.34 mmol) of sodium triacetoxyborohydride were added at room temperature followed after 5 min of stirring by 1.19 g (4.35 mmol) of tert-butyl (2S)-2-[(tert-butoxycarbonyl)amino]-4-oxobutanoate. The mixture was stirred at room temperature for 15 min, concentrated on a rotary evaporator, taken up in acetonitrile and water and purified by preparative HPLC (mobile phase: ACN/water + 0.1% TFA, gradient). This gave 1.64 g (80% of theory) of the target compound.

LC-MS (Method 1):  $R_t$  = 1.10 min; MS (ESIpos):  $m/z$  = 613  $[M+H]^+$ .

$^1H$  NMR (400 MHz, DMSO- $d_6$ ):  $\delta$  [ppm] = 1.01 (s, 9H), 1.32 (s, 9H), 1.35 (s, 9H), 1.80-1.89 (m, 1H), 2.01-2.11 (m, 1H), 2.54-2.71 (m, 2H), 3.75-3.81 (m, 1H), 3.90 (s, 2H), 4.18 (d, 1H), 7.13 (d, 1H), 7.20-7.24 (m, 1H), 7.28-7.34 (m, 5H), 7.52-7.58 (m, 1H), 7.76-7.80 (m, 1H), 8.10 (s br, 1H), 8.23 (s br, 1H).

### Intermediate C49

(2S)-4-[(1R)-1-[4-Benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]-2-[(tert-butoxycarbonyl)amino]butanoic acid



225 mg (0.37 mmol) of tert-butyl (2S)-4-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)amino)-2-[(tert-butoxycarbonyl)amino]butanoate were dissolved in 10 ml of absolute dichloromethane, and 156 mg (1.54 mmol) of triethylamine were added. At 0°C, 125 mg (0.92 mmol) of acetoxyacetyl chloride were added, and the mixture was stirred at RT for 16 h. Another 251 mg (1.84 mmol) of acetoxyacetyl chloride and 186 mg (1.84 mmol) of triethylamine were added, and the mixture was stirred at RT for 3 h. A little dichloromethane was added and the mixture was washed with saturated sodium bicarbonate solution and saturated sodium chloride solution. The organic phase was dried over sodium sulphate, concentrated on a rotary evaporator and dried under high vacuum. The residue was taken up in 10 ml of ethanol, 0.91 ml (12.67 mmol) of a 40% strength aqueous methylamine solution was added and the mixture was stirred at 50°C for 3 h. The mixture was concentrated on a rotary evaporator, the residue was taken up in dichloromethane and the organic phase was washed twice with water. The organic phase was dried over sodium sulphate, concentrated on a rotary evaporator and dried under high vacuum. The residue was taken up in 2 ml of dichloromethane, 2 ml (25.96 mmol) of trifluoroacetic acid were added and the mixture was stirred at 50°C for 4 h. The mixture was concentrated on a rotary evaporator and the residue was dried under high vacuum. The residue was taken up in 10 ml of absolute dichloromethane, 298 mg (2.95 mmol) of triethylamine and 429 mg (1.97 mmol) of di-tert-butyl dicarbonate were added and the mixture was stirred at RT for 1 h. The mixture was concentrated on a rotary evaporator and the residue was purified by preparative HPLC (mobile phase: ACN/water, gradient). This gave 62 mg (27% of theory) of the target compound.

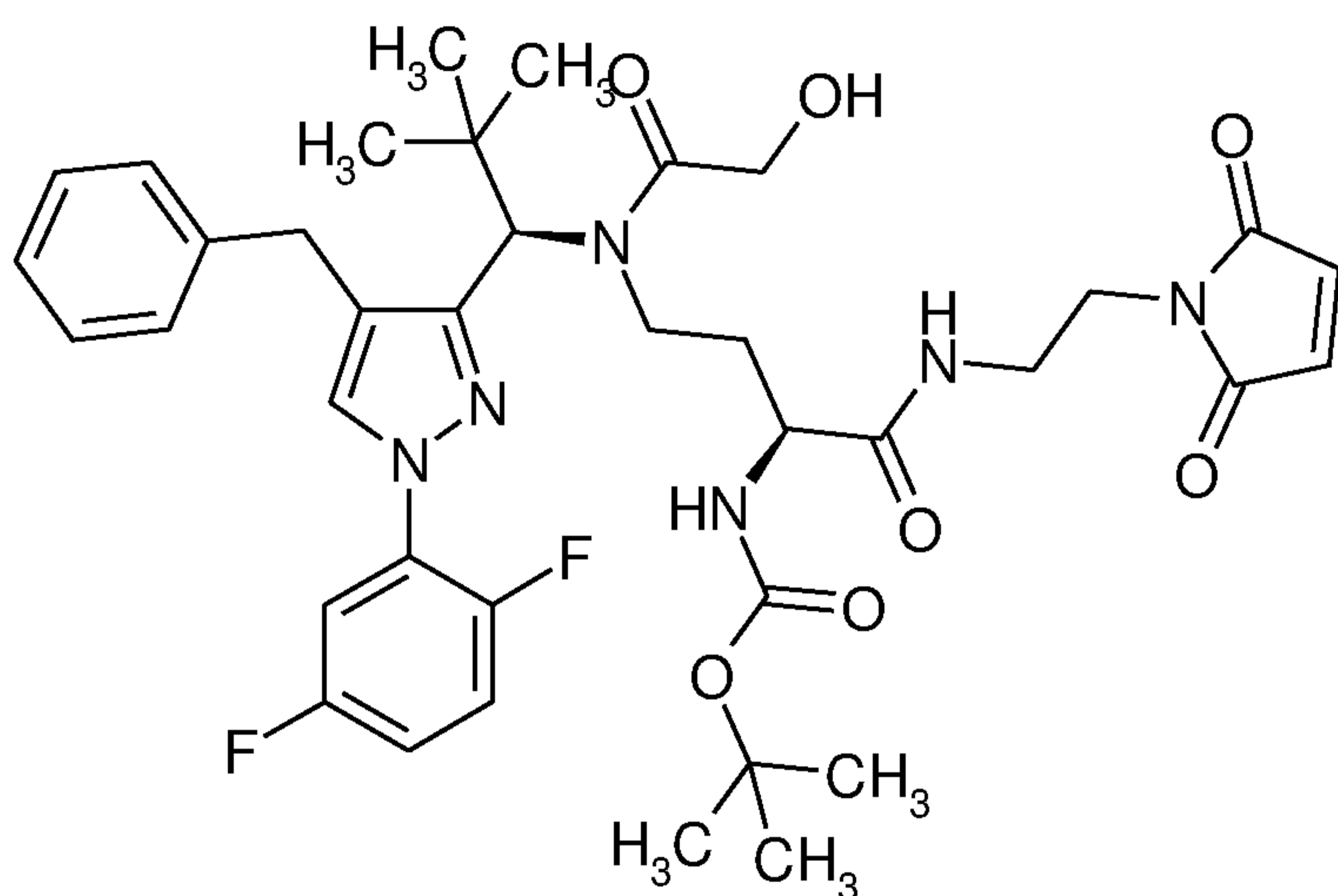
LC-MS (Method 1):  $R_t = 1.32$  min; MS (ESIpos):  $m/z = 615$  [M+H]<sup>+</sup>.

<sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>):  $\delta$  [ppm] = 0.91 (s, 9H), 1.32 (s, 9H), 2.64-2.72 (m, 4H), 3.50-3.58 (m, 1H), 3.72 (dd, 2H), 4.07-4.22 (m, 2H), 4.47-4.54 (m, 1H), 5.75 (s, 1H), 6.84-6.89 (m, 1H), 7.15-7.30 (m, 6H), 7.47-7.53 (m, 1H), 7.70-7.75 (m, 1H), 8.09-

8.13 (m, 1H), 11.66 (s br, 1H).

### Intermediate C50

5 tert-Butyl [(2S)-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]-1-[[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]amino]-1-oxobutan-2-yl]carbamate



10

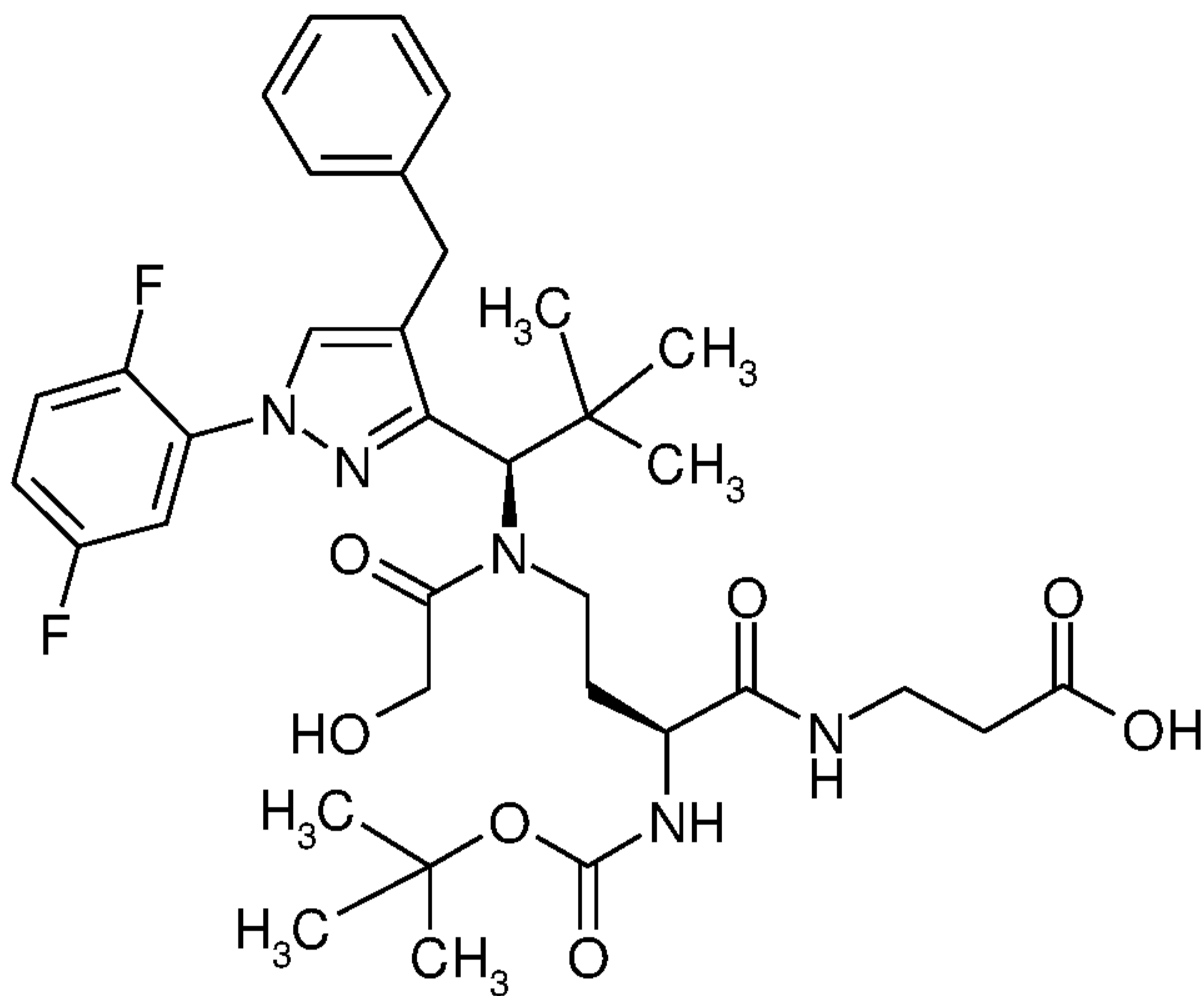
60 mg (0.1 mmol) of (2S)-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]-2-[(tert-butoxycarbonyl)amino]butanoic acid were dissolved in 10 ml of absolute DMF, and 74 mg (0.20 mmol) of HATU were added. 74 mg (0.29 mmol) of trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) were dissolved separately in 2 ml of absolute DMF, 38 mg (0.29 mmol) of *N,N*-diisopropylethylamine were added and the mixture was added dropwise to the reaction mixture. The reaction was stirred at RT for 3 d. The mixture was purified directly by preparative HPLC mobile phase: ACN/water + 0.1% TFA, gradient). This gave 9.3 mg (13% of theory) of the target compound.

25

LC-MS (Method 1):  $R_t = 1.34$  min; MS (ESIpos):  $m/z = 737$  [M+H]<sup>+</sup>.

### Intermediate C51

N-{(2S)-4-[[{(1R)-1-[4-Benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-2-[(tert-butoxycarbonyl)amino]butanoyl]-beta-alanine



5

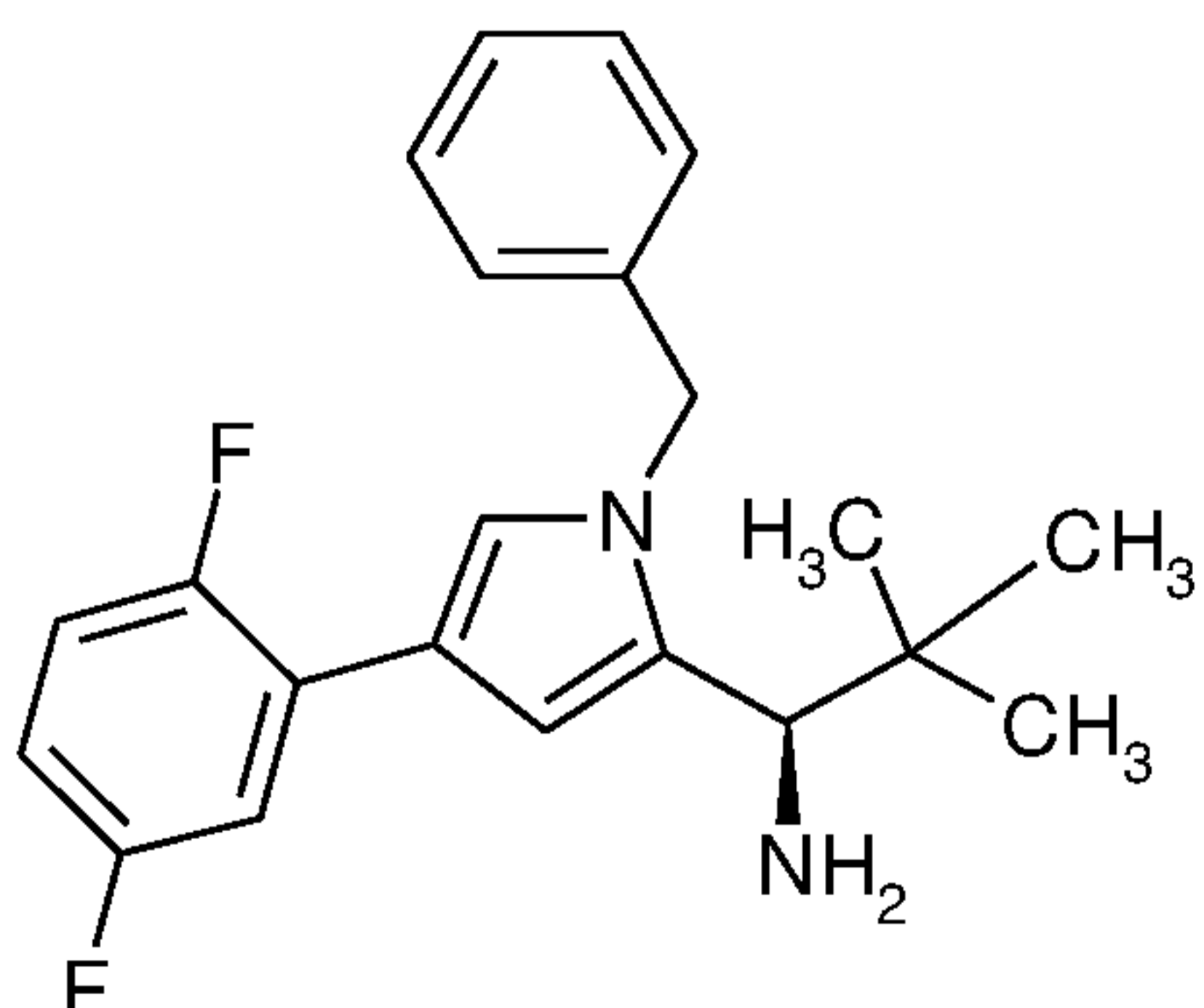
First, Intermediate C47 was reductively alkylated with benzyl N-{(2S)-2-[(tert-butoxycarbonyl)amino]-4-oxobutanoyl}-beta-alaninate analogously to Intermediate C2. The secondary amino group was then acylated with 2-chloro-2-oxoethyl acetate as described for Intermediate C27, and the two ester groups were then hydrolysed with 2M lithium hydroxide solution in methanol. This gave 23 mg of the title compound.

15 LC-MS (Method 1):  $R_t = 1.24$  min; MS (ESIpos):  $m/z = 686$  (M+H)<sup>+</sup>.

### Intermediate C52

(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropan-1-amine

20





10.00 g (49.01 mmol) of methyl 4-bromo-1H-pyrrole-2-carboxylate were initially charged in 100.0 ml of DMF, and 20.76 g (63.72 mmol) of caesium carbonate and 9.22 g (53.91 mmol) of benzyl bromide were added. The reaction mixture was stirred at RT overnight. The reaction mixture was partitioned between water and ethyl acetate and the aqueous phase was extracted with ethyl acetate. The combined organic phases were dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The reaction was repeated with 90.0 g of methyl 4-bromo-1H-pyrrole-2-carboxylate.

The two combined reactions were purified by preparative RP-HPLC (column: Daiso 300x100; 10 $\mu$ , flow rate: 250 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 125.15 g (87% of theory) of the compound methyl 1-benzyl-4-bromo-1H-pyrrole-2-carboxylate.

LC-MS (Method 1):  $R_t$  = 1.18 min; MS (ESIpos):  $m/z$  = 295 [M+H]<sup>+</sup>.

Under argon, 4.80 g (16.32 mmol) of methyl 1-benzyl-4-bromo-1H-pyrrole-2-carboxylate were initially charged in DMF, and 3.61 g (22.85 mmol) of (2,5-difluorophenyl)boronic acid, 19.20 ml of saturated sodium carbonate solution and 1.33 g (1.63 mmol) of [1,1'-bis(diphenylphosphino)ferrocene]-dichloropalladium(II):dichloromethane were added. The reaction mixture was stirred at 85°C overnight. The reaction mixture was filtered through Celite and the filter cake was washed with ethyl acetate. The organic phase was extracted with water and then washed with saturated NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was purified on silica gel (mobile phase: cyclohexane/ethyl acetate 100:3). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 3.60 g (67% of theory) of the compound methyl 1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrole-2-carboxylate.

LC-MS (Method 7):  $R_t = 1.59$  min; MS (ESIpos):  $m/z = 328$  [M+H]<sup>+</sup>.

3.60 g (11.00 mmol) of methyl 1-benzyl-4-(2,5-difluorophenyl)-  
5 1H-pyrrole-2-carboxylate were initially charged in 90.0 ml of  
THF, and 1.04 g (27.50 mmol) of lithium aluminium hydride (2.4  
M in THF) were added at 0°C. The reaction mixture was stirred  
at 0°C for 30 minutes. At 0°C, saturated potassium sodium  
tartrate solution was added, and ethyl acetate was added to the  
10 reaction mixture. The organic phase was extracted three times  
with saturated potassium sodium tartrate solution. The organic  
phase was washed once with saturated NaCl solution and dried  
over magnesium sulphate. The solvent was evaporated under  
reduced pressure and the residue was dissolved in 30.0 ml of  
15 dichloromethane. 3.38 g (32.99 mmol) of manganese(IV) oxide were  
added and the mixture was stirred at RT for 48 h. Another 2.20  
g (21.47 mmol) of manganese(IV) oxide were added and the mixture  
was stirred at RT overnight. The reaction mixture was filtered  
through Celite and the filter cake was washed with  
20 dichloromethane. The solvent was evaporated under reduced  
pressure and the residue 2.80 g of (1-benzyl-4-(2,5-  
difluorophenyl)-1H-pyrrole-2-carbaldehyde) was used without  
further purification in the next step of the synthesis.

25 LC-MS (Method 7):  $R_t = 1.48$  min; MS (ESIpos):  $m/z = 298$  [M+H]<sup>+</sup>.

28.21 g (94.88 mmol) of 1-benzyl-4-(2,5-difluorophenyl)-1H-  
pyrrole-2-carbaldehyde together with 23.00 g (189.77 mmol) of  
(R)-2-methylpropane-2-sulphinamide were initially charged in  
30 403.0 ml of absolute THF, and 7.42 g (237.21 mmol) of  
titanium(IV) isopropoxide were added and the mixture was stirred  
at RT overnight. 500.0 ml of saturated NaCl solution and 1000.0  
ml of ethyl acetate were added, and the mixture was stirred at  
RT for 1 h. The mixture was filtered through kieselguhr and the  
35 filtrate was washed twice with saturated NaCl solution. The  
organic phase was dried over magnesium sulphate, the solvent was  
evaporated under reduced pressure and the residue was purified  
using Biotage Isolera (silica gel, column 1500+340 g SNAP, flow

rate 200 ml/min, ethyl acetate/cyclohexane 1:10).

LC-MS (Method 7):  $R_t = 1.63$  min; MS (ESIpos):  $m/z = 401$   $[M+H]^+$ .

5 25.00 g (62.42 mmol) of (R)-N-{(E/Z)-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]methylene}-2-methylpropane-2-sulphinamide were initially charged in absolute THF under argon and cooled to  $-78^\circ\text{C}$ . 12.00 g (187.27 mmol) of tert-butyllithium (1.7 M solution in pentane) were then added at  $-78^\circ\text{C}$  and the  
10 mixture was stirred at this temperature for 3 h. At  $-78^\circ\text{C}$ , 71.4 ml of methanol and 214.3 ml of saturated ammonium chloride solution were then added in succession, and the reaction mixture was allowed to warm to RT and stirred at RT for 1 h. The mixture was diluted with ethyl acetate and washed with water. The organic  
15 phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue (R)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-methylpropane-2-sulphinamide was used without further purification in the next step of the synthesis.

20

LC-MS (Method 6):  $R_t = 2.97$  min; MS (ESIpos):  $m/z = 459$   $[M+H]^+$ .

28.00 g (61.05 mmol) of (R)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-  
25 methylpropane-2-sulphinamide were initially charged in 186.7 ml of 1,4-dioxane, and 45.8 ml of HCl in 1,4-dioxane solution (4.0 M) were then added. The reaction mixture was stirred at RT for 2 h and the solvent was evaporated under reduced pressure. The residue was purified by preparative RP-HPLC (column: (column:  
30 Kinetix 100x30; flow rate: 60 ml/min, MeCN/water). The acetonitrile was evaporated under reduced pressure and dichloromethane was added to the aqueous residue. The organic phase was washed with sodium bicarbonate solution and dried over magnesium sulphate. The solvent was evaporated under reduced  
35 pressure and the residue was dried under high vacuum. This gave 16.2 g (75% of theory) of the title compound.

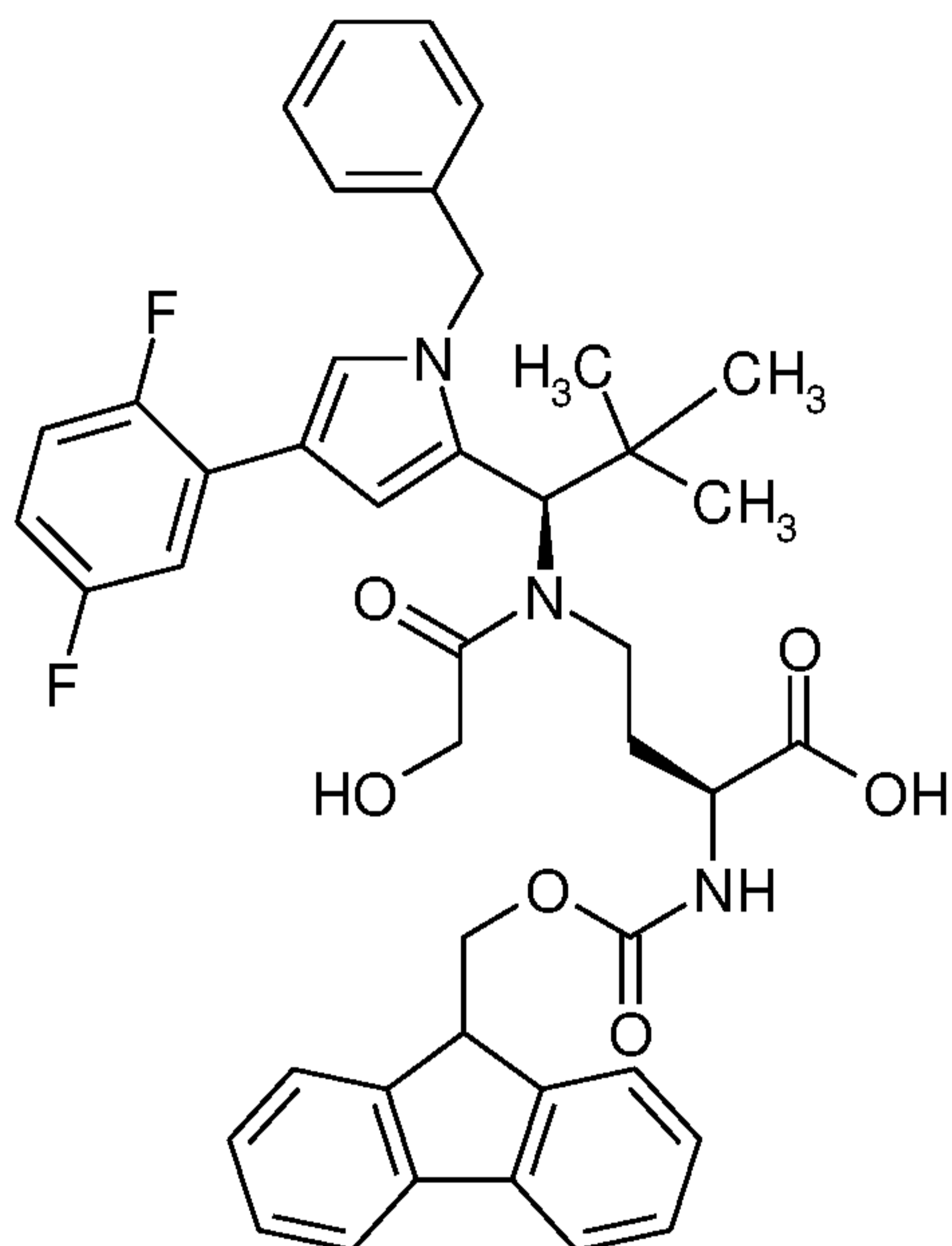
LC-MS (Method 6):  $R_t = 2.10$  min; MS (ESIpos):  $m/z = 338$   $[M-NH_2]^+$ ,

709 [2M+H]<sup>+</sup>.

<sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>): δ [ppm] = 0.87 (s, 9H), 1.53 (s, 2H), 3.59 (s, 1H), 5.24 (d, 2H), 6.56 (s, 1H), 6.94 (m, 1H), 7.10 (d, 2H), 7.20 (m, 1H), 7.26 (m, 2H), 7.34 (m, 2H), 7.46 (m, 1H).

### Intermediate C53

(2S)-4-[[{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-2-[[{(9H-fluoren-9-ylmethoxy)carbonyl]amino}butanoic acid



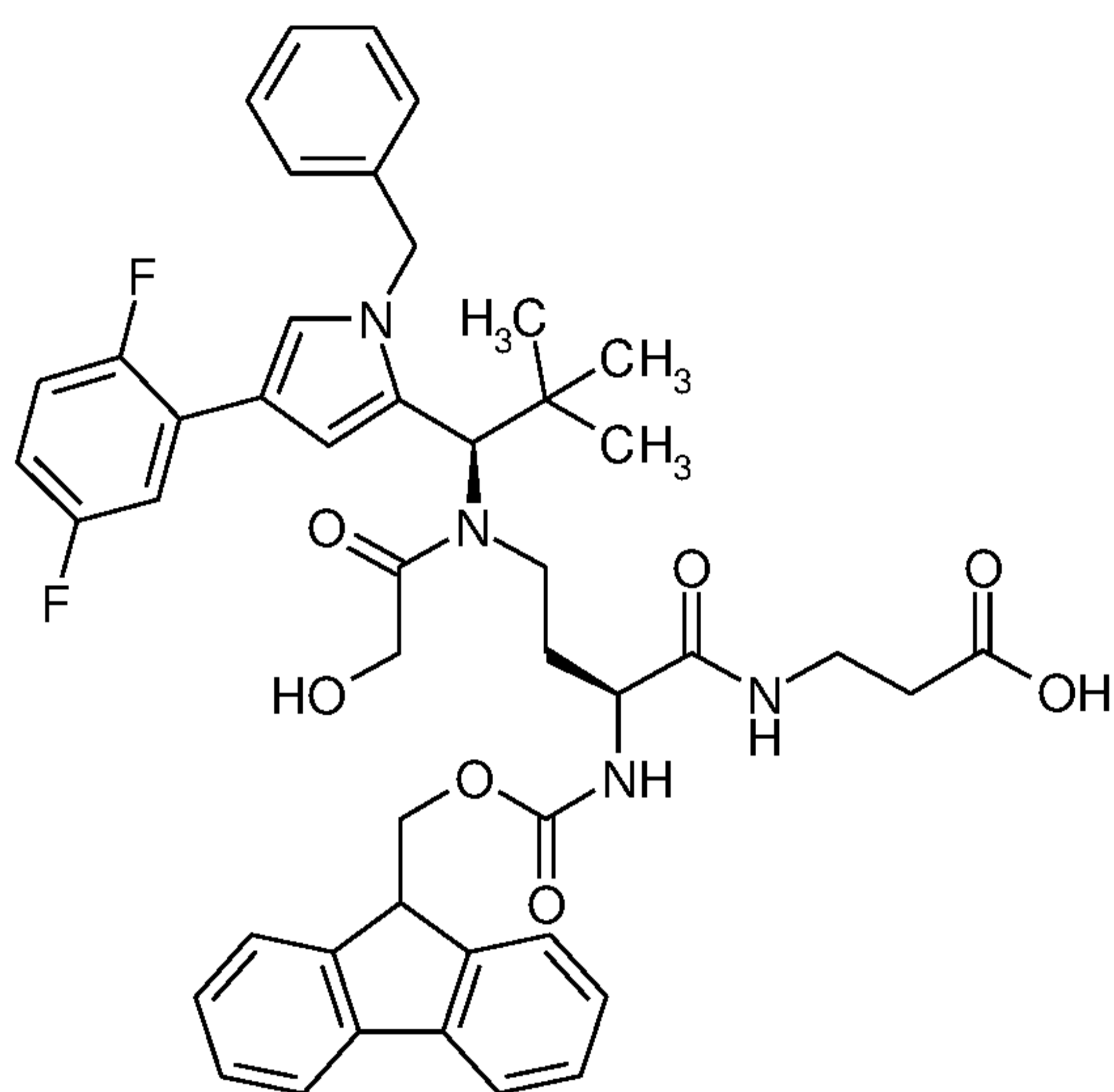
15 First, Intermediate C52 was reductively alkylated with benzyl  
 (2S)-2-[[ (benzyloxy) carbonyl] amino]-4-oxobutanoate analogously  
 to C2. The secondary amino group was then acylated with 2-chloro-  
 2-oxoethyl acetate as described for Intermediate C27, and the  
 two ester groups were then hydrolysed with 2M lithium hydroxide  
 20 solution in methanol. The intermediate obtained in this manner  
 was dissolved in ethanol, palladium on carbon (10%) was added  
 and the mixture was hydrogenated at RT with hydrogen under  
 standard pressure for 1 h. The deprotected compound was taken  
 up in dioxane/water 2:1 and in the last step the Fmoc protective  
 25 group was introduced using 9H-fluoren-9-ylmethyl  
 chlorocarbonate in the presence of *N,N*-diisopropylethylamine.

LC-MS (Method 1):  $R_t = 1.37$  min; MS (ESIpos):  $m/z = 734$  (M-H)<sup>-</sup>.

### Intermediate C54

5

N-[(2S)-4-[[{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-2-[[{(9H-fluoren-9-ylmethoxy)carbonyl]amino}butanoyl]-beta-alanine



10

First, Intermediate C52 was reductively alkylated with benzyl N-[(2S)-2-[[ (benzyloxy) carbonyl] amino]-4-oxobutanoyl]-beta-alaninate analogously to Intermediate C2. The secondary amino group was then acylated with 2-chloro-2-oxoethyl acetate as described for Intermediate C27. The intermediate obtained in this manner was dissolved in ethanol, palladium on carbon (10%) was added and the mixture was hydrogenated at RT with hydrogen under standard pressure for 1 h. The two ester groups were then hydrolysed with 2M lithium hydroxide solution in methanol. The deprotected compound was taken up in dioxane/water 2:1 and in the last step the Fmoc protective group was introduced using 9H-fluoren-9-ylmethyl chlorocarbonate in the presence of *N,N*-diisopropylethylamine. This gave 48 mg of the title compound.

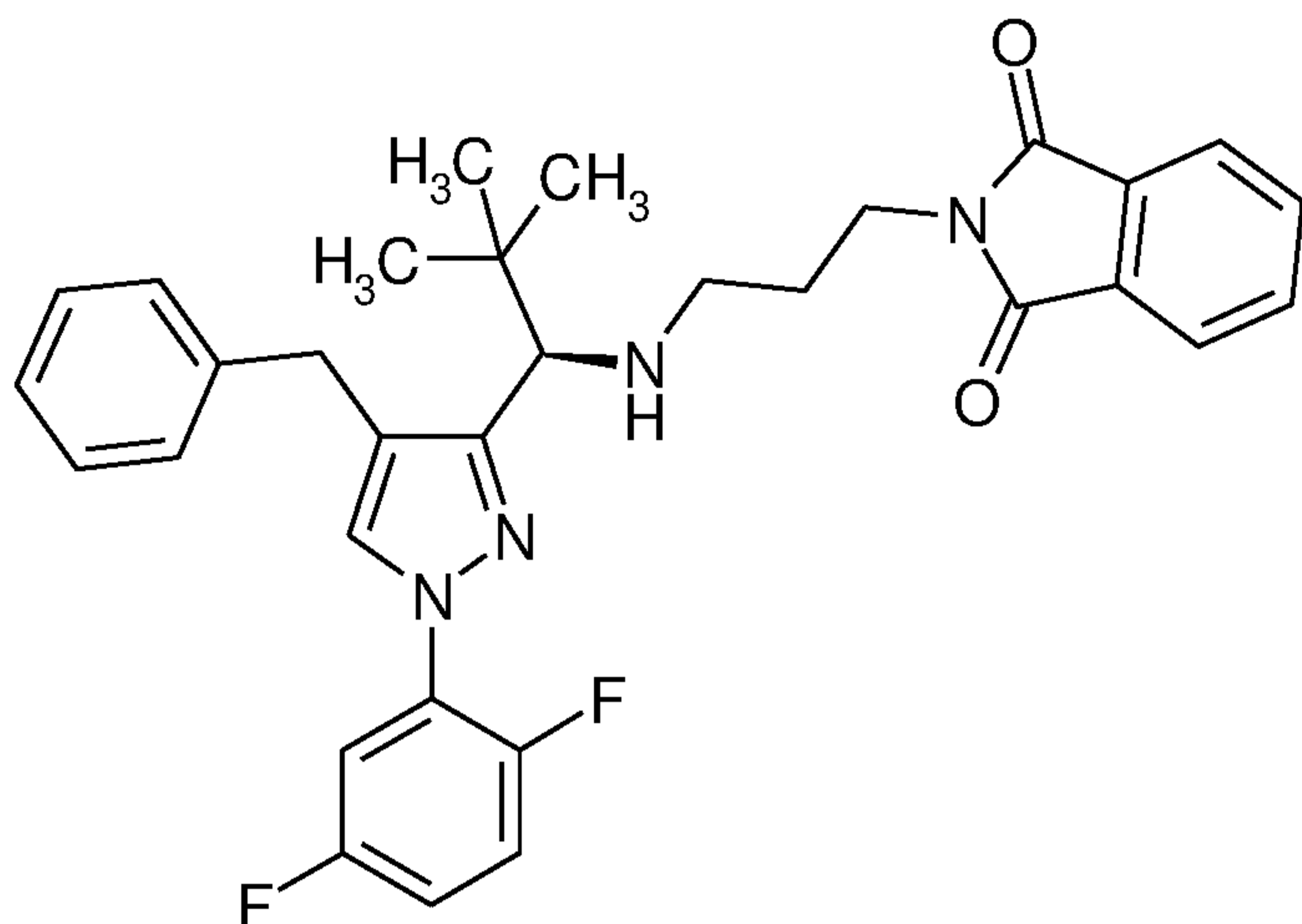
25

LC-MS (Method 1):  $R_t = 1.38$  min; MS (ESIpos):  $m/z = 807$  (M+H)<sup>+</sup>.

### Intermediate C55

2-[3-((1R)-1-[4-Benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)amino)propyl]-1H-isoindole-1,3(2H)-dione

5



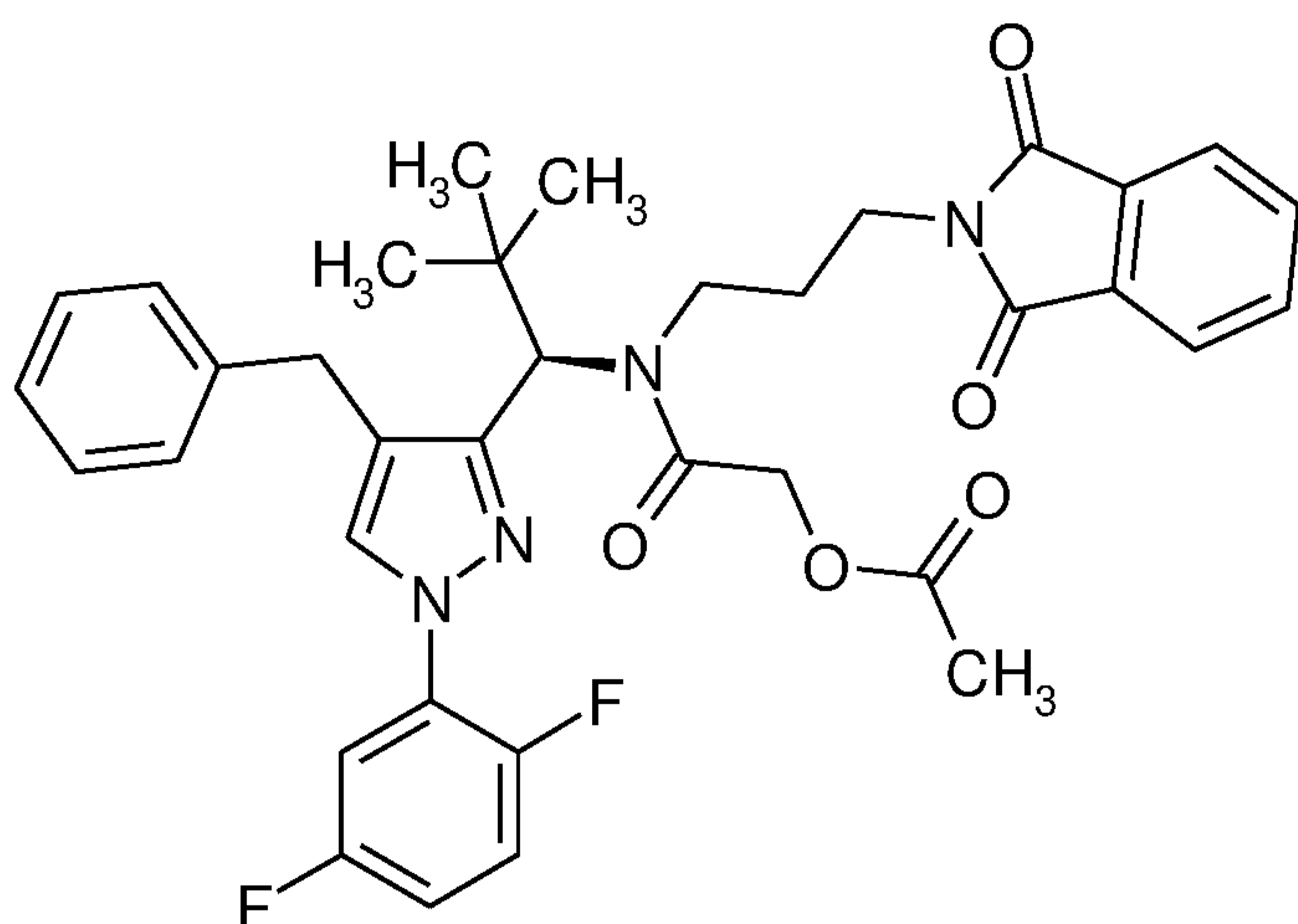
340 mg (0.96 mmol) of (1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropan-1-amine were dissolved in 7 ml of absolute DCM, and 69 mg (1.15 mmol, 60  $\mu$ l) acetic acid and 284 mg (1.34 mmol) of sodium triacetoxyborohydride were added at RT. The mixture was stirred for 15 min, and 233 mg (1.15 mmol) of 3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propanal were then added. The mixture was stirred at RT for 4.5 h. Another 233 mg (1.15 mmol) of 3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propanal, 69 mg (1.15 mmol, 60  $\mu$ l) acetic acid and 284 mg (1.34 mmol) of sodium triacetoxyborohydride were added, and the mixture was stirred at RT for 7 h. Ethyl acetate was added and the reaction mixture was washed with saturated sodium carbonate solution. The organic phase was concentrated and the residue was purified twice by preparative HPLC [1.) mobile phase: ACN/water + 0.1% TFA, gradient; 2.) mobile phase: ACN/water + 1% TFA+1.0% NEt<sub>3</sub>]. This gave 108 mg (21% of theory) of the target compound.

LC-MS (Method 1): R<sub>t</sub> = 0.96 min; MS (ESIpos): m/z = 543 [M+H]<sup>+</sup>.

#### Intermediate C56

2-((1R)-1-[4-Benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-

2,2-dimethylpropyl}[3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propyl]amino)-2-oxoethyl acetate



5

102 mg (0.19 mmol) of 2-[3-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)amino)propyl]-1H-isoindole-1,3(2H)-dione were initially charged in 2 ml of absolute DCM, and 44 mg (0.43 mmol) of triethylamine were added at RT. At 0°C, 31 mg (0.23 mmol) of 2-chloro-2-oxoethyl acetate dissolved in 1 ml of absolute DCM were added. The mixture was stirred at RT for 40 min. Another 26 mg of 2-chloro-2-oxoethyl acetate dissolved in 0.5 ml of absolute DCM and 19 mg (0.19 mmol) of triethylamine were added, and the mixture was stirred at RT for 60 min.

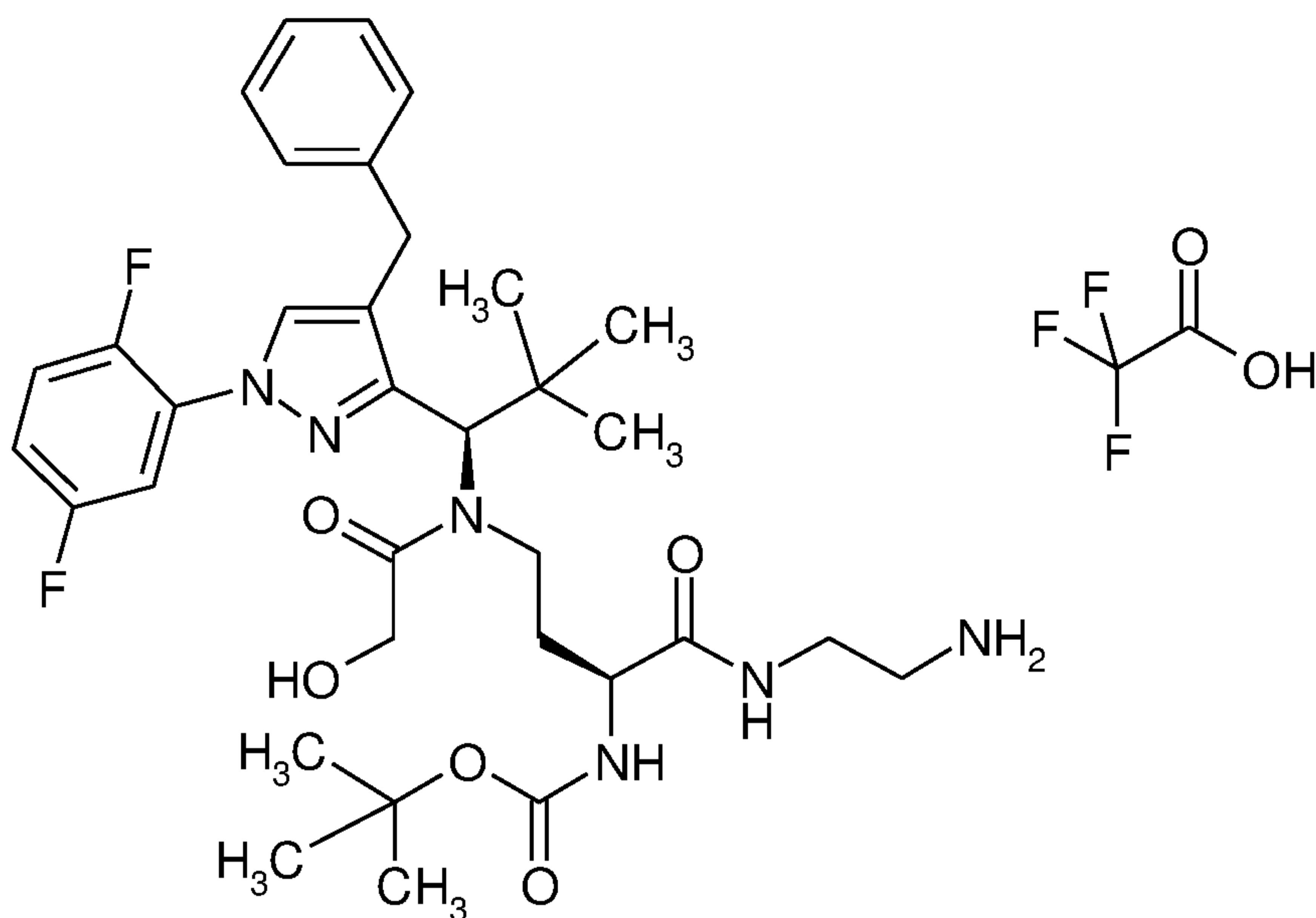
Water was added, the mixture was concentrated on a rotary evaporator and the residue was purified by preparative HPLC (mobile phase: ACN/water + 0.1% TFA, gradient). This gave 106 mg (88% of theory) of the target compound.

LC-MS (Method 1):  $R_t = 1.37$  min; MS (ESIpos):  $m/z = 643$   $[M+H]^+$ .

#### Intermediate C57

25

Trifluoroacetic acid / tert-butyl {(2S)-1-[(2-aminoethyl)amino]-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]-1-oxobutan-2-yl}carbamate (1:1)



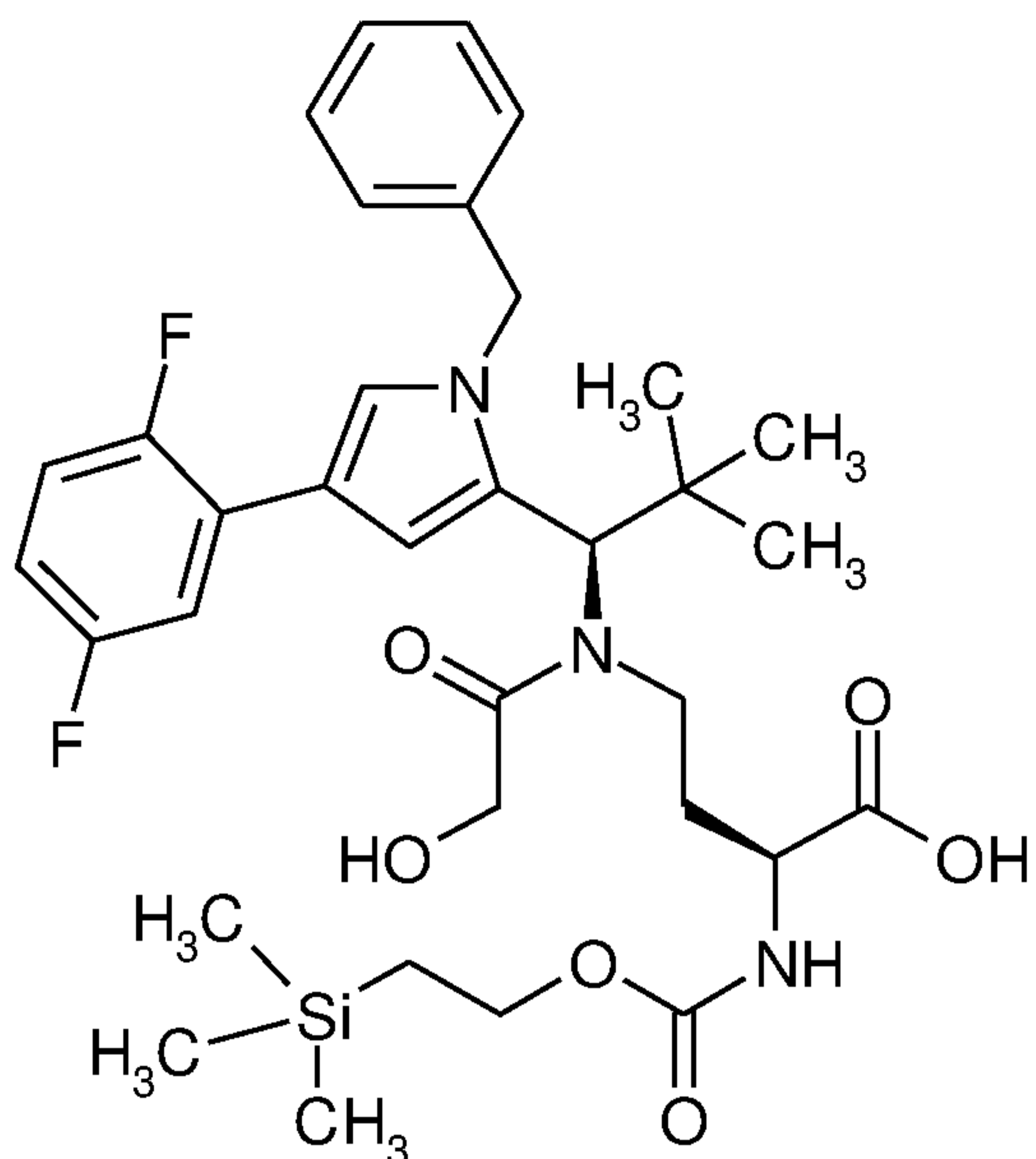
The title compound was prepared according to standard methods  
 5 by coupling Intermediate C49 with 9H-fluoren-9-ylmethyl (2-  
 aminoethyl)carbamate in the presence of HATU and subsequent  
 removal of the Fmoc protective group with piperidine. This gave  
 14 mg of the title compound (40% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t = 0.98$  min; MS (ESIpos):  $m/z = 657$  (M+H)<sup>+</sup>.

#### Intermediate C58

(2S)-4-[[{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-  
 15 yl]-2,2-dimethylpropyl}(glycoloyl)amino]-2-({[2-  
 (trimethylsilyl)ethoxy]carbonyl}amino)butanoic acid





First, Intermediate C52 was reductively alkylated with benzyl  
 (2S)-2-{[(benzyloxy)carbonyl]amino}-4-oxobutanoate analogously  
 5 to Intermediate C2. First, Intermediate C52 was reductively  
 alkylated with benzyl (2S)-2-{[(benzyloxy)carbonyl]amino}-4-  
 oxobutanoate analogously to C2. The secondary amino group was  
 then acylated with 2-chloro-2-oxoethyl acetate as described for  
 Intermediate C27, and the two ester groups were then hydrolysed  
 10 with 2M lithium hydroxide solution in methanol. The intermediate  
 obtained in this manner was dissolved in ethanol, palladium on  
 carbon (10%) was added and the mixture was hydrogenated at RT  
 with hydrogen under standard pressure for 1 h.

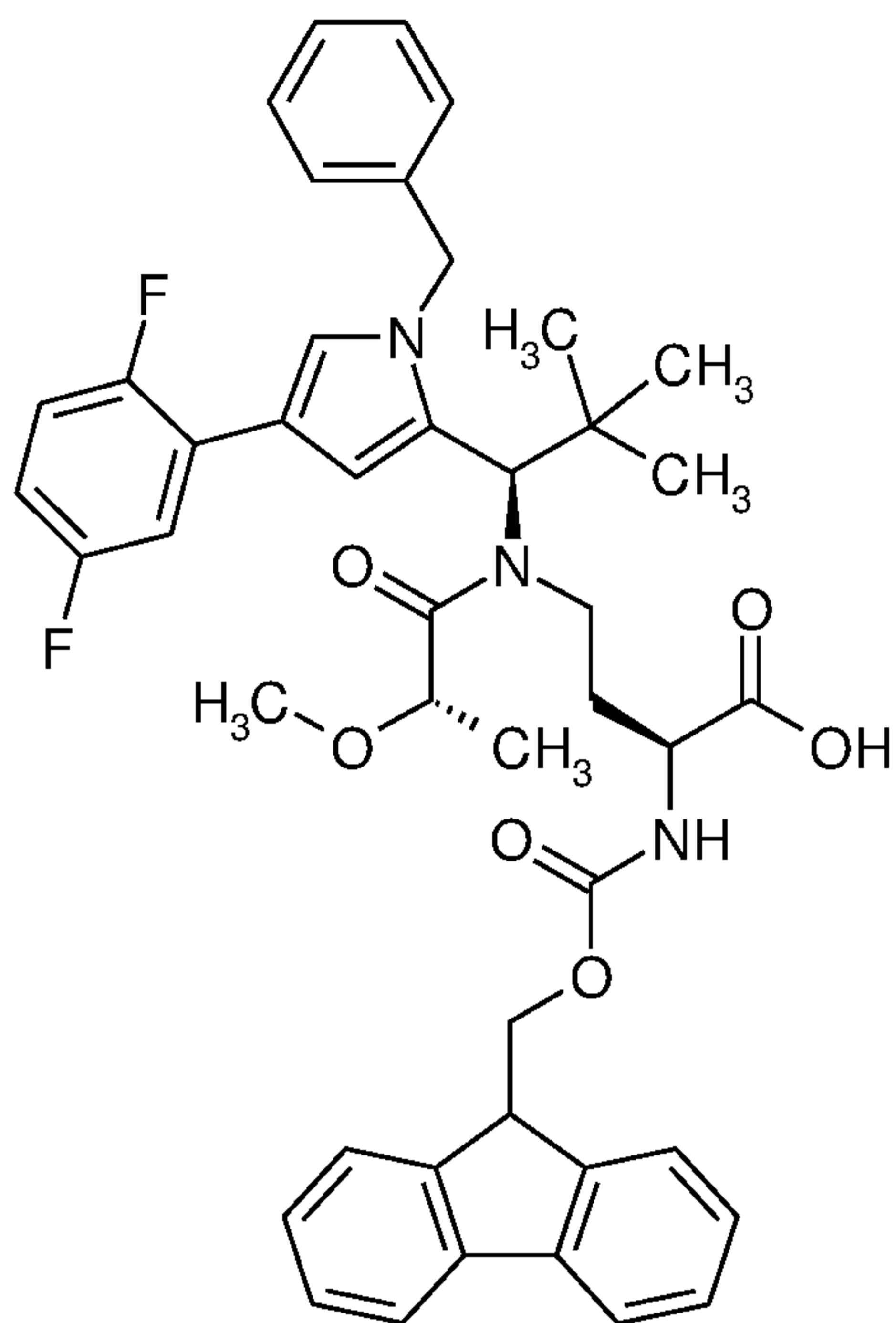
15 500 mg (0.886 mmol) of this fully deprotected intermediate were  
 taken up in 60 ml of dioxane, and 253 mg (0.975 mmol) of 1-({[2-  
 (trimethylsilyl)ethoxy]carbonyl}oxy)pyrrolidine-2,5-dione and  
 198  $\mu$ l of triethylamine were added. After 24 h of stirring at  
 RT, the reaction was concentrated and the residue was purified  
 20 by preparative HPLC. Combination of the appropriate fractions,  
 concentration under reduced pressure and drying under high  
 vacuum gave 312 mg (50% of theory) of the title compound.

LC-MS (Method 5):  $R_t$  = 4.61 min; MS (ESIpos):  $m/z$  = 658 (M+H)<sup>-</sup>.

25

**Intermediate C59**

(2S)-4-({(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[(2S)-2-methoxypropanoyl]amino)-2-{{[(9H-fluoren-9-ylmethoxy)carbonyl]amino}butanoic acid



5

Initially, the secondary amino group of benzyl (2S)-4-({(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)-2-{{[(benzyloxy)carbonyl]amino}butanoate  
 10 was acylated with (2S)-2-methoxypropanoyl chloride (intermediate of Intermediate C53) in the presence of triethylamine as described for Intermediate C53. The intermediate obtained was taken up in ethanol, palladium on carbon (10%) was added and the mixture was hydrogenated at RT  
 15 with hydrogen under standard pressure for 1 h. The deprotected compound was taken up in dioxane/water 2:1 and in the last step the Fmoc protective group was introduced using 9H-fluoren-9-ylmethyl chlorocarbonate in the presence of *N,N*-diisopropylethylamine.

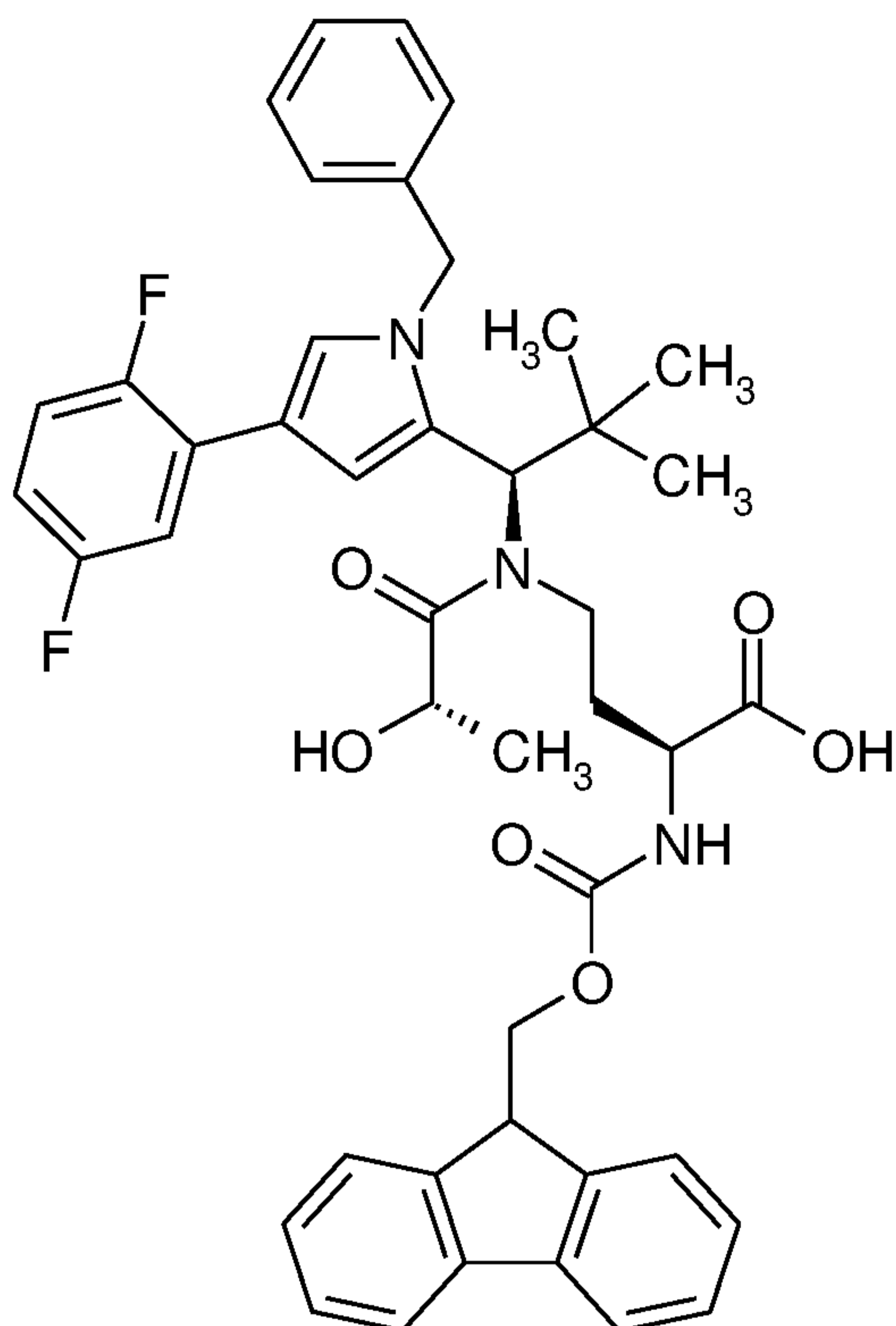
20

LC-MS (Method 1):  $R_t = 1.39$  min; MS (ESIpos):  $m/z = 764$  (M-H)<sup>-</sup>.

#### Intermediate C60

25 (2S)-4-({(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-

yl]-2,2-dimethylpropyl} [(2S)-2-methoxypropanoyl] amino)-2-  
 {[ (9H-fluoren-9-ylmethoxy) carbonyl] amino}butanoic acid



5

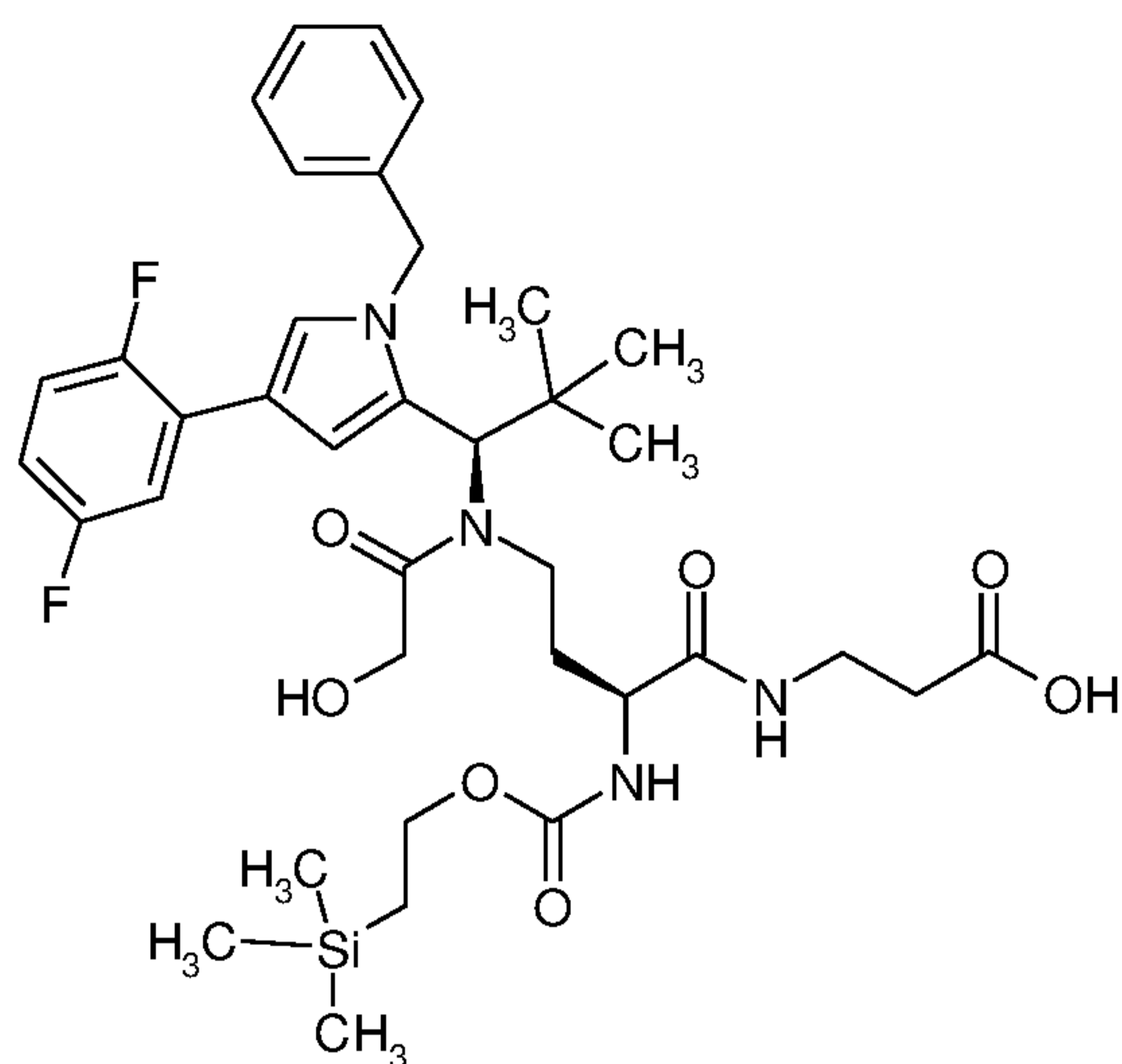
The synthesis was carried out analogously to Intermediate C53.

LC-MS (Method 1):  $R_t = 1.41$  min; MS (ESIpos):  $m/z = 750$  (M+H)<sup>+</sup>.

#### 10 Intermediate C61

N-[(2S)-4-[(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl) amino]-2-([2-(trimethylsilyl)ethoxy] carbonyl) amino]butanoyl]-beta-alanine

15



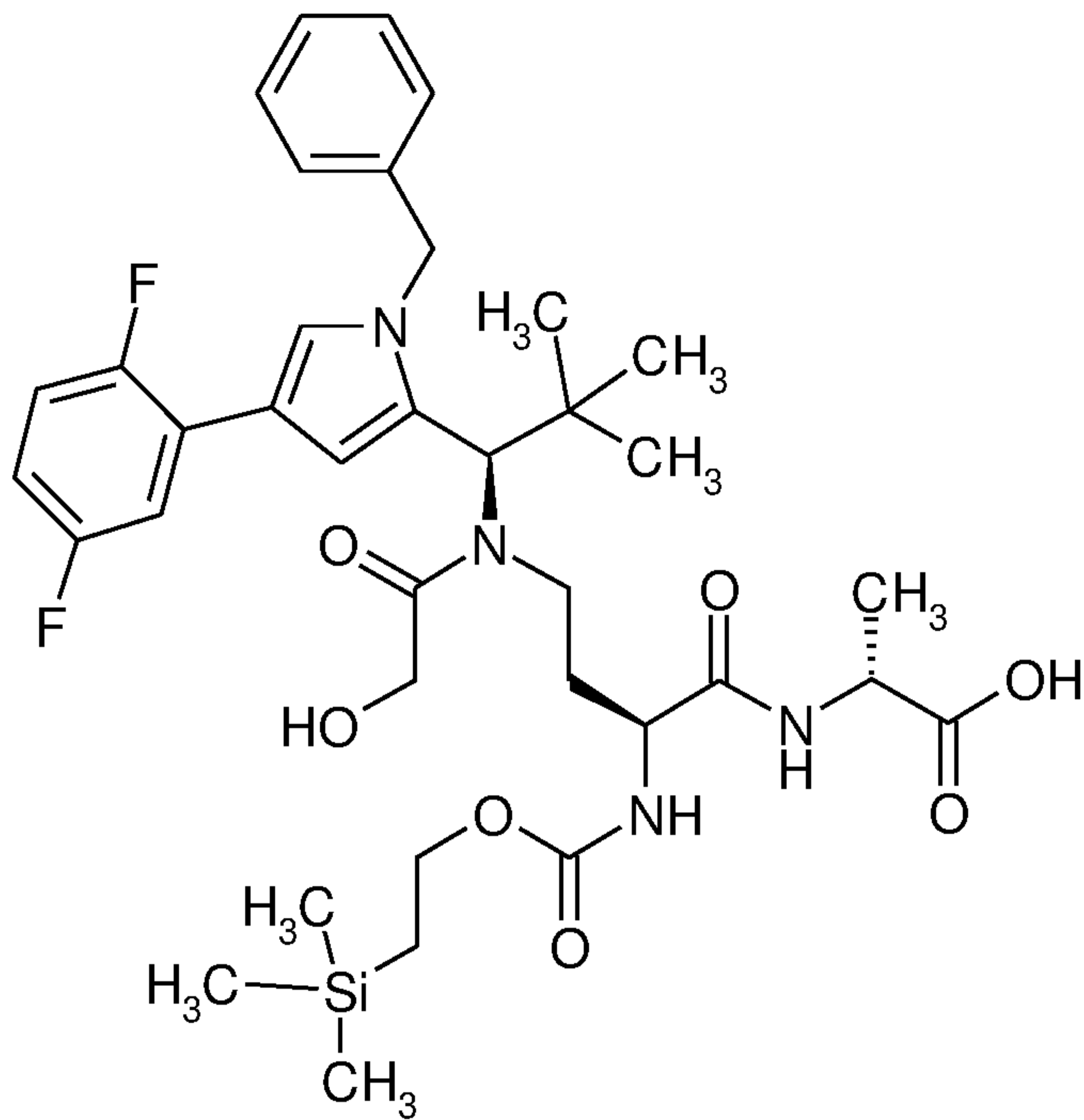
The title compound was prepared by coupling 60 mg (0.091 mmol) of Intermediate C58 with methyl  $\beta$ -alaninate, followed by ester cleavage with 2M lithium hydroxide solution. This gave 67mg (61% of theory) of the title compound over 2 steps.

LC-MS (Method 1):  $R_t = 1.29$  min; MS (ESIpos):  $m/z = 729$  (M+H)<sup>+</sup>.

#### 10 Intermediate C62

N-[(2S)-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]-2-([2-(trimethylsilyl)ethoxy]carbonyl)amino)butanoyl]-D-alanine

15



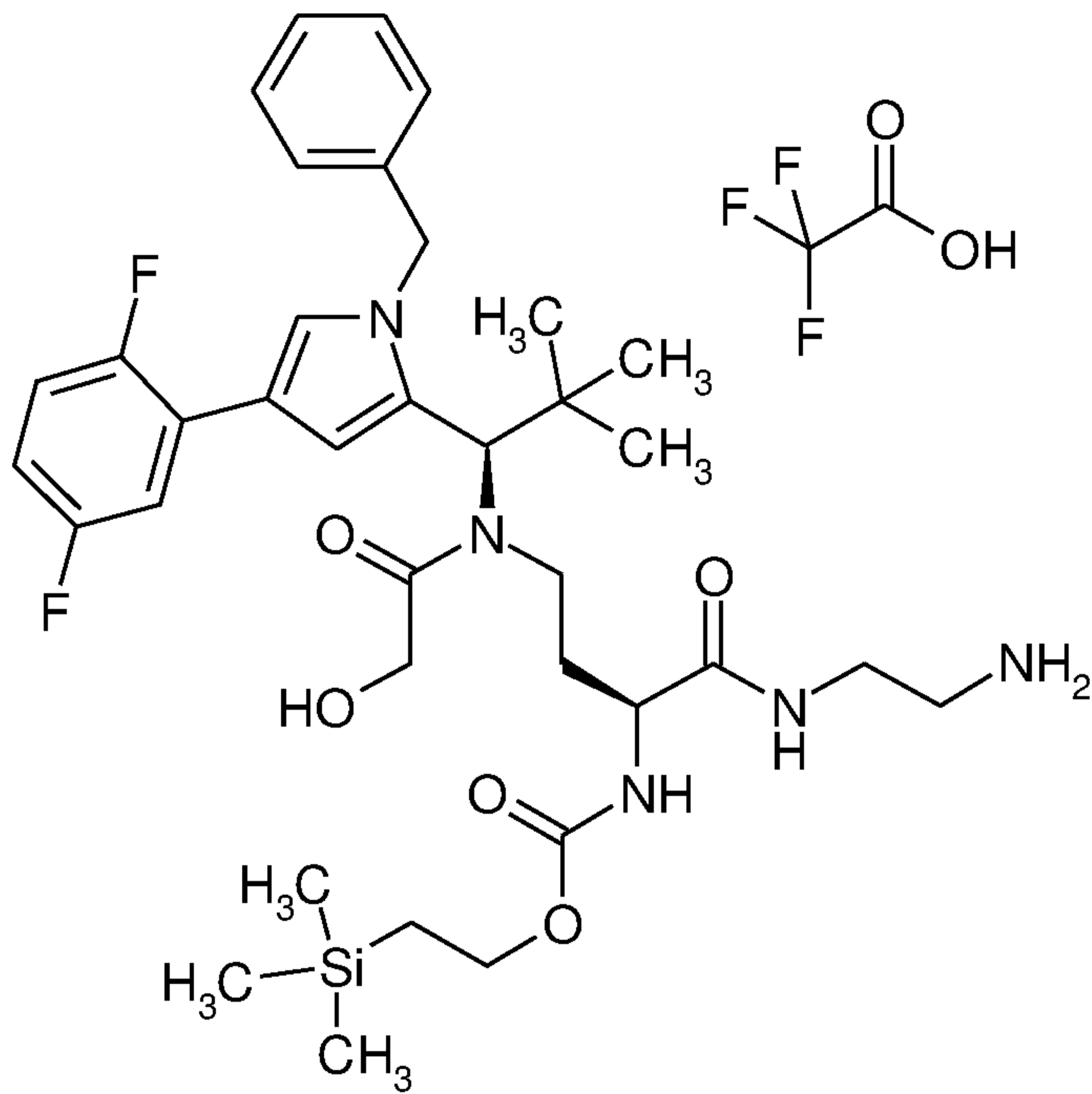
The title compound was prepared analogously to Intermediate C61 from Intermediate C58 and methyl D-alaninate.

5

LC-MS (Method 1):  $R_t = 1.32$  min; MS (ESIpos):  $m/z = 729$  (M+H)<sup>+</sup>.

#### Intermediate C64

10 Trifluoroacetic acid / 2-(trimethylsilyl)ethyl {(2S)-1-[(2-aminoethyl)amino]-4-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl)amino]-1-oxobutan-2-yl}carbamate (1:1)



The title compound was prepared from Intermediate C58 analogously to Intermediate C63.

5

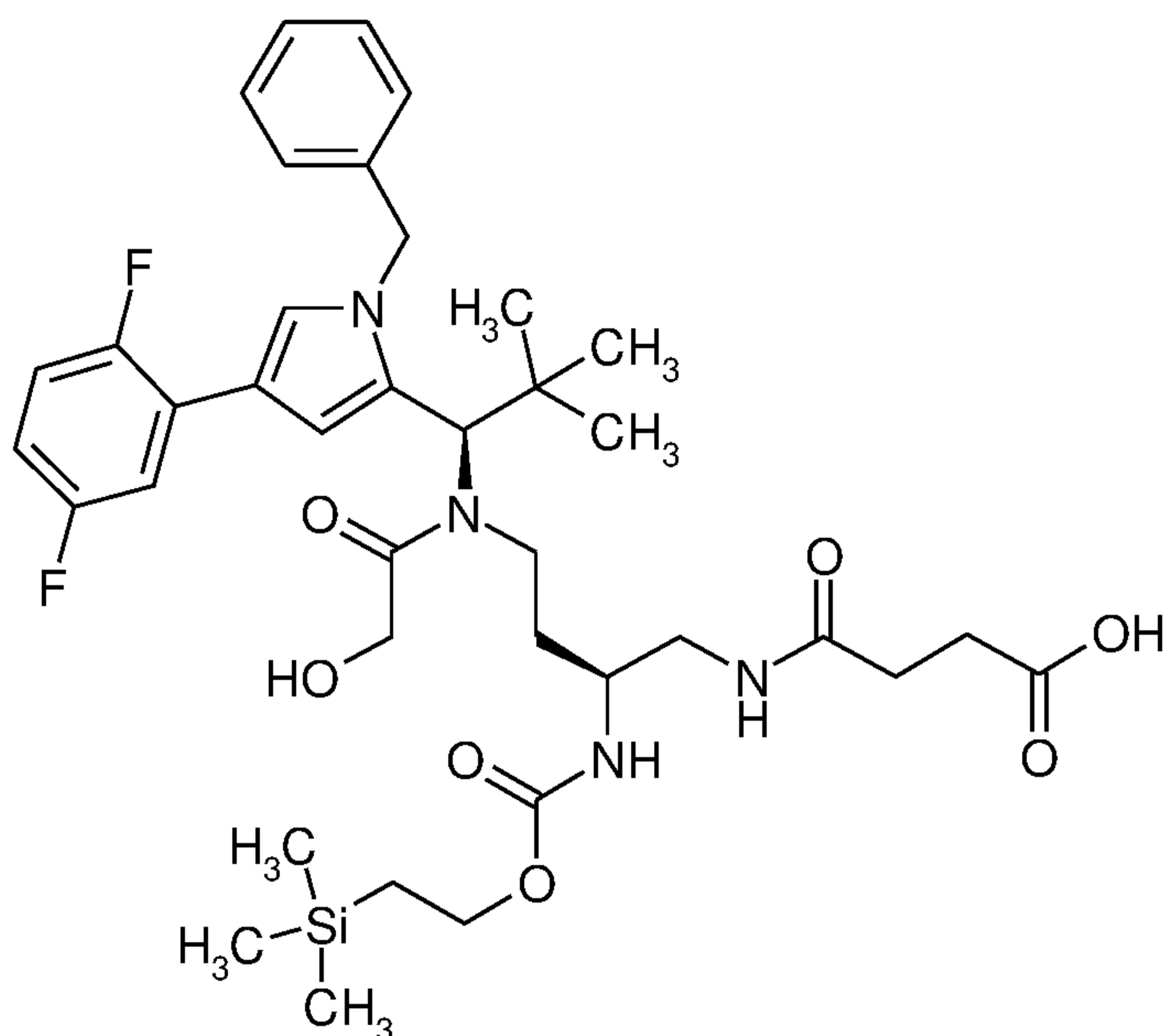
HPLC (Method 11):  $R_t = 2.4$  min;

LC-MS (Method 1):  $R_t = 1.01$  min; MS (ESIpos):  $m/z = 700$  (M+H)<sup>+</sup>.

10 **Intermediate C65**

(8S)-8-{2-[[{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-(glycoloyl)amino]ethyl]-2,2-dimethyl-6,11-dioxo-5-oxa-7,10-diaza-2-silatetradecan-14-oic acid

15



215 mg (0.59 mmol) of Intermediate L66 were initially charged in 25 ml of dichloromethane, and 377 mg (0.89 mmol) of Dess-  
 5 Martin periodinane and 144  $\mu$ l (1.78 mmol) of pyridine were added. The mixture was stirred at RT for 30 min. The reaction was then diluted with 300 ml of dichloromethane and the organic phase was washed in each case twice with 10% strength  $\text{Na}_2\text{S}_2\text{O}_3$  solution, 10% strength citric acid solution and saturated sodium bicarbonate  
 10 solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. This gave 305 mg of the aldehyde which was reacted without further purification.

15 175 mg (0.49 mmol) of Intermediate C52 were dissolved in 50 ml of dichloromethane, and 147mg (0.69 mmol) of sodium triacetoxyborohydride and 32.5  $\mu$ l of acetic acid were added. After 5 min of stirring at RT, 214 mg (0.593 mmol) of the aldehyde described above were added, and the reaction was  
 20 stirred at RT overnight. Here, instead of the expected product, 2-(trimethylsilyl)ethyl [(2S)-4-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino)-1-(2,5-dioxopyrrolidin-1-yl)butan-2-yl]carbamate was formed. Since this imide can also be converted into the title compound,  
 25 the reaction was concentrated and the residue was purified by preparative HPLC. After combination of the appropriate imide-

containing fractions, the solvent was evaporated under reduced pressure and the residue was dried under high vacuum. This gave 195 mg (58%) of the imide named above.

5 LC-MS (Method 5):  $R_t = 3.32$  min; MS (ESIpos):  $m/z = 667$  (M+H)<sup>+</sup>.

65 mg (97.5  $\mu$ mol) of this imide were taken up in 15 ml of dichloromethane, and 367  $\mu$ l (3.4 mmol) of acetoxyacetyl chloride and 595  $\mu$ l of *N,N*-diisopropylethylamine were added. After 30 min  
 10 of stirring at RT, the reaction was concentrated without heating under reduced pressure and the residue was purified by preparative HPLC. The appropriate fractions were combined giving, after evaporation of the solvents and drying under high vacuum, 28 mg (37% of theory) of (8S)-11-{(1R)-1-[1-benzyl-4-  
 15 (2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-8-[(2,5-dioxopyrrolidin-1-yl)methyl]-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl acetate.

LC-MS (Method 1):  $R_t = 1.44$  min; MS (ESIpos):  $m/z = 767$  (M+H)<sup>+</sup>.

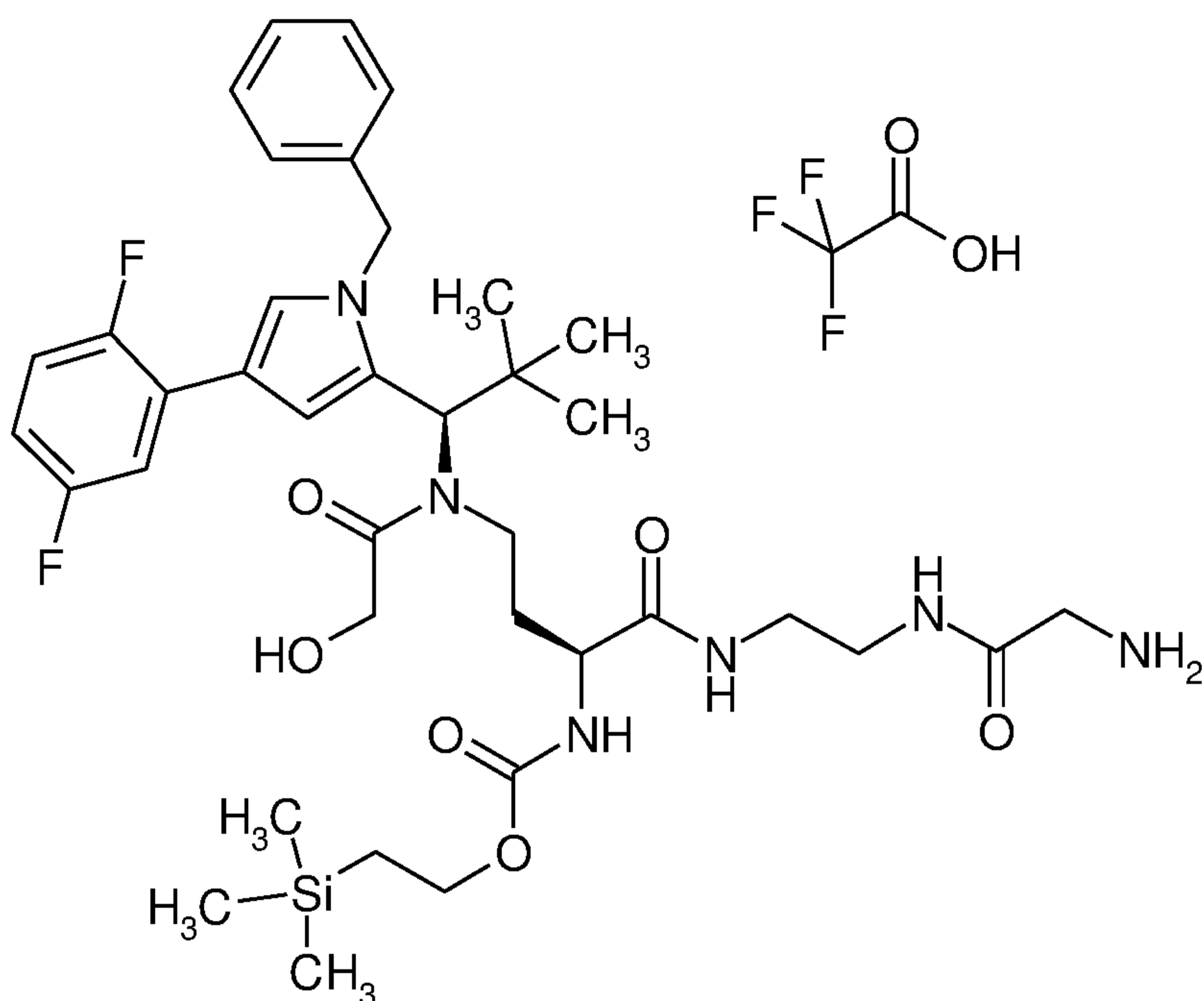
20 28 mg (37  $\mu$ mol) of this intermediate were dissolved in 3 ml of methanol, and 548  $\mu$ l of a 2M lithium hydroxide solution were added. After 10 min of stirring at RT, the reaction was adjusted to pH 4 with trifluoroacetic acid and then concentrated. The  
 25 residue was purified by preparative HPLC. The appropriate fractions were combined, the solvent was evaporated and the residue was dried under high vacuum, giving 26 mg (96% of theory) of the title compound as a white solid.

30 LC-MS (Method 1):  $R_t = 1.33$  min; MS (ESIpos):  $m/z = 743$  (M+H)<sup>+</sup>.

### Intermediate C66

2-(Trimethylsilyl)ethyl [(2S)-4-[(1R)-1-[1-benzyl-4-(2,5-  
 35 difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-1-[[2-(glycylamino)ethyl]amino]-1-oxobutan-2-yl]carbamate





First, trifluoroacetic acid / benzyl {2-[(2-aminoethyl)amino]-  
 2-oxoethyl}carbamate (1:1) was prepared from N-  
 5 [(benzyloxy)carbonyl]glycine and tert-butyl (2-  
 aminoethyl)carbamate according to classical methods of peptide  
 chemistry (HATU coupling and Boc removal).

13 mg (0.036 mmol) of this Intermediate and 25 mg (0.033 mmol)  
 10 of Intermediate C58 were taken up in 3 ml of DMF, and 19 mg  
 (0.05 mmol) of HATU and 17  $\mu$ l of *N,N*-diisopropylethylamine were  
 added.

After 10 min of stirring at RT, the mixture was concentrated and  
 15 the residue was purified by preparative HPLC. This gave 17.8 mg  
 (60% of theory) of the intermediate.

LC-MS (Method 1):  $R_t$  = 1.36 min; MS (ESIpos):  $m/z$  = 891 (M+H)<sup>+</sup>.

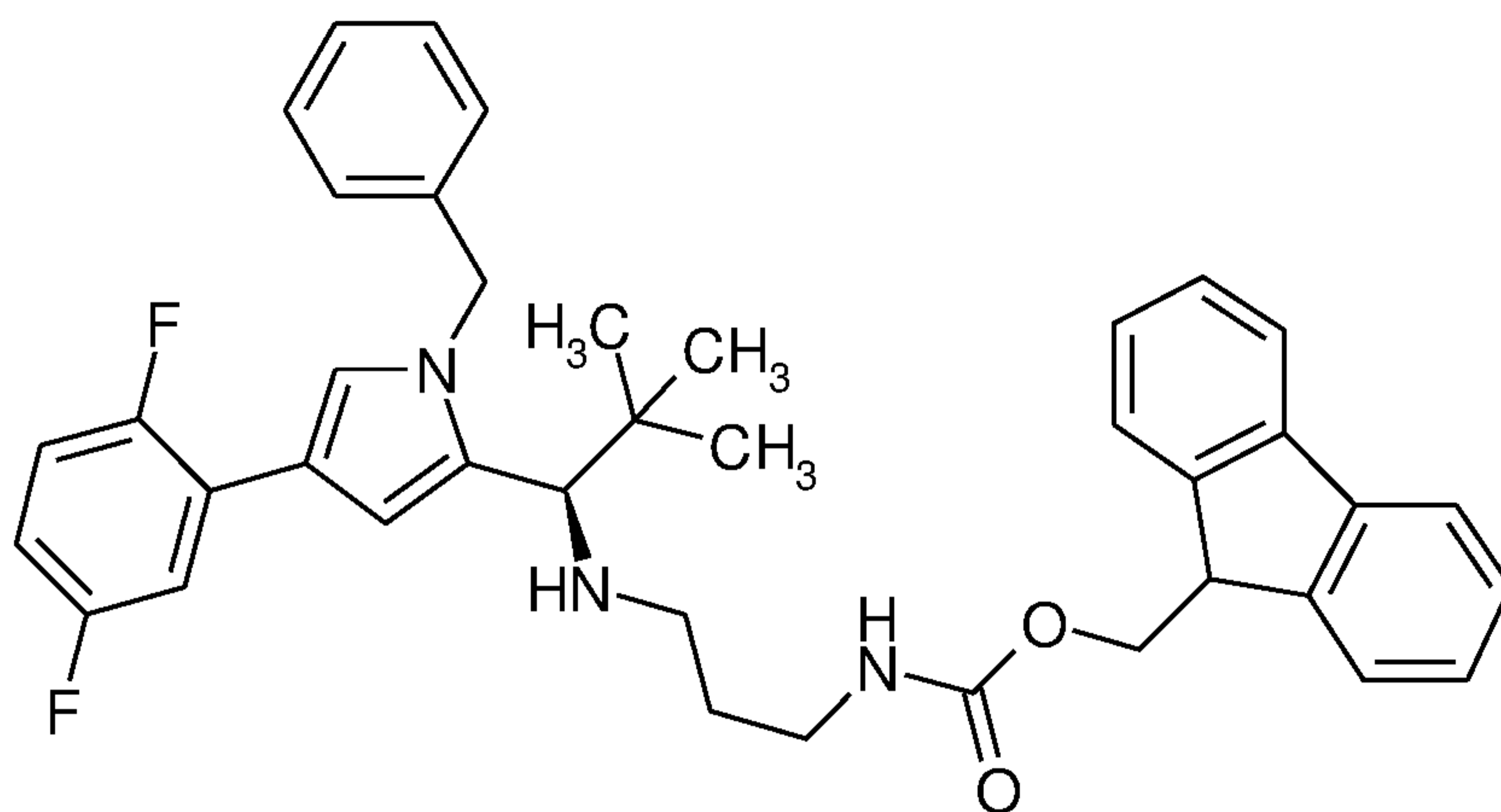
20 17 mg (0.019 mmol) of this intermediate were dissolved in 10 ml  
 of ethanol, palladium on carbon (10%) was added and the mixture  
 was hydrogenated at RT with hydrogen at standard pressure for 2  
 h. The catalyst was filtered off, the solvents were evaporated  
 under reduced pressure and the residue was dried under high  
 25 vacuum. This gave 9 mg (62% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 1.03$  min; MS (ESIpos):  $m/z = 757$  (M+H)<sup>+</sup>.

### Intermediate C67

5

9H-Fluoren-9-ylmethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino)propyl]carbamate



10

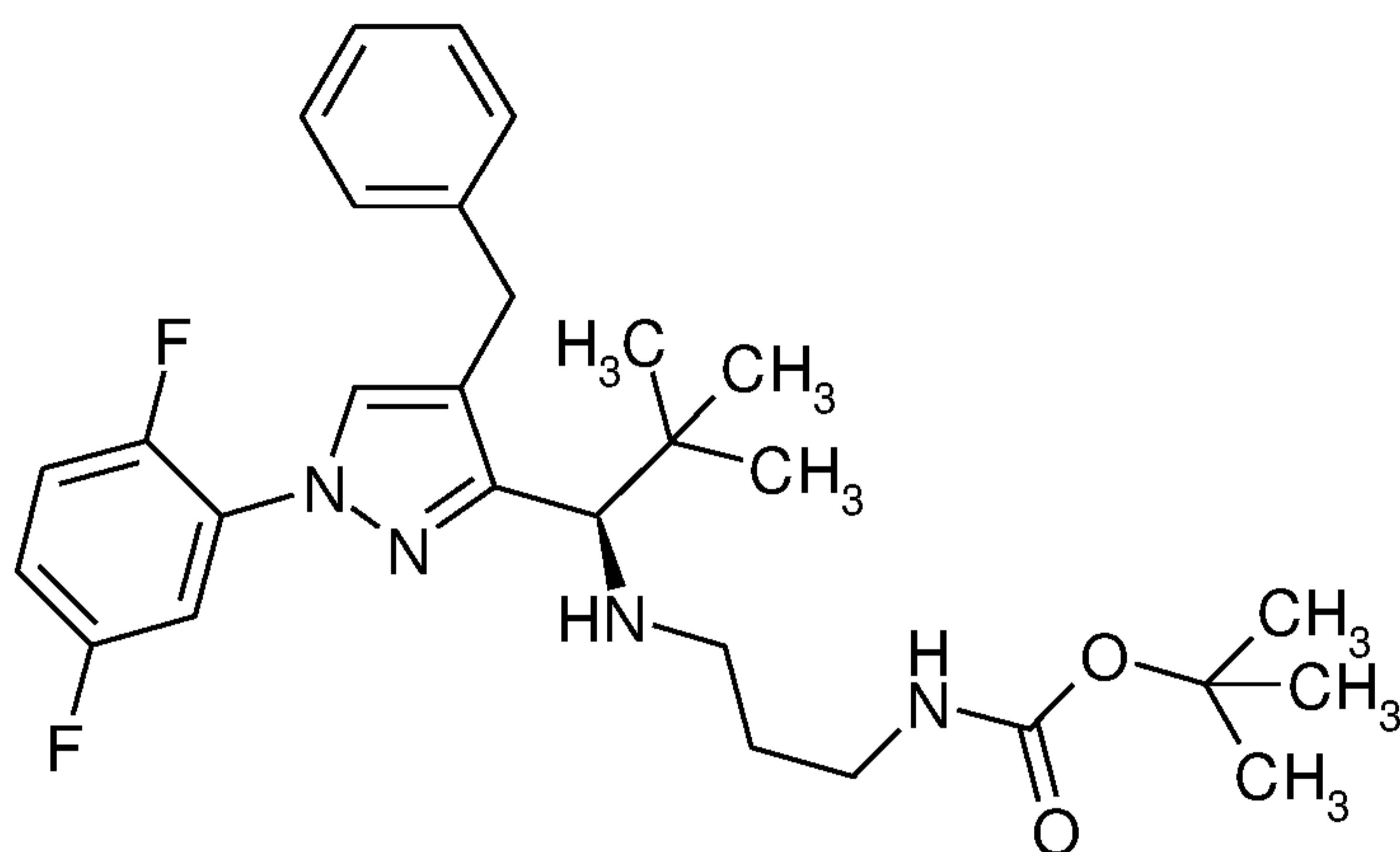
605.3 mg (1.71 mmol) of (1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropan-1-amine (Intermediate C52) were initially charged in 10.0 ml of dichloromethane, and 506.7 mg (2.39 mmol) of sodium triacetoxyborohydride and 117.9 mg (1.96 mmol) of acetic acid were added and the mixture was stirred at RT for 5 min. 580.0 mg (1.96 mmol) of 9H-fluoren-9-ylmethyl (3-oxopropyl)carbamate (Intermediate L70) dissolved in 10.0 ml of dichloromethane were added and the reaction mixture stirred at RT overnight. The reaction mixture was diluted with ethyl acetate and the organic phase was washed in each case twice with saturated sodium carbonate solution and saturated NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was purified on silica gel (mobile phase: cyclohexane/ethyl acetate 3:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 514.7 mg (46% of theory) of the title compound.

30

LC-MS (Method 1):  $R_t = 1.10$  min; MS (ESIpos):  $m/z = 634$  (M+H)<sup>+</sup>.

**Intermediate C68**

5 tert-Butyl [3-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)amino)propyl]carbamate



10 The synthesis was carried out analogously to the synthesis of the compound Intermediate C67.

1000.0 mg (2.81 mmol) of (1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropan-1-amine  
 15 (Intermediate C47)

835.0 mg (3.94 mmol) of sodium triacetoxyborohydride

194.0 mg (3.24 mmol) of acetic acid

20

560.0 mg (3.24 mmol) of tert-butyl (3-oxopropyl)carbamate

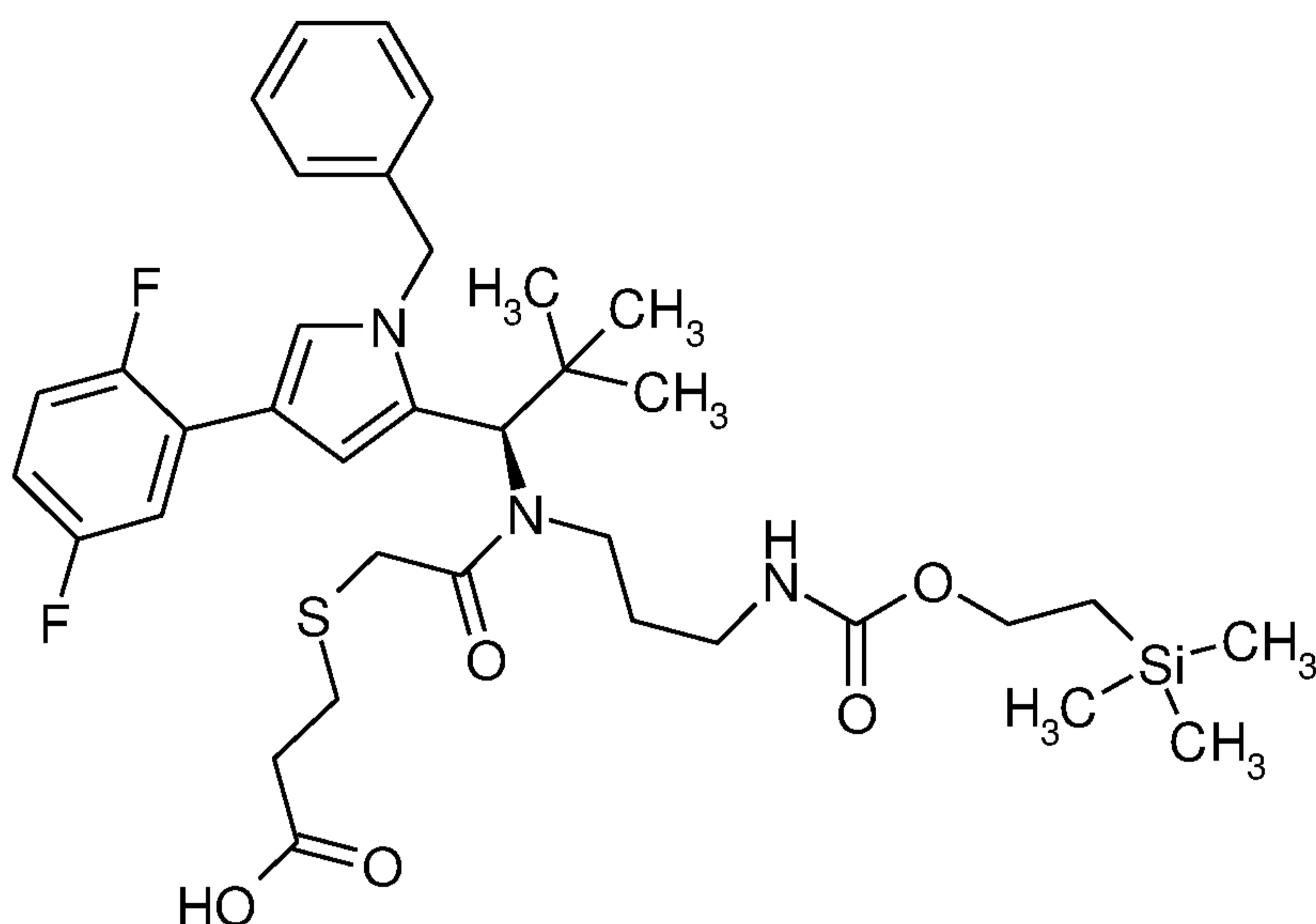
This gave 695.8 mg (48% of theory) of the title compound.

25 LC-MS (Method 1):  $R_t = 1.02$  min; MS (ESIpos):  $m/z = 513$  (M+H)<sup>+</sup>.

**Intermediate C69**

11-((1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-  
 30 2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-

7,11-diaza-2-silaheptadecan-17-oic acid

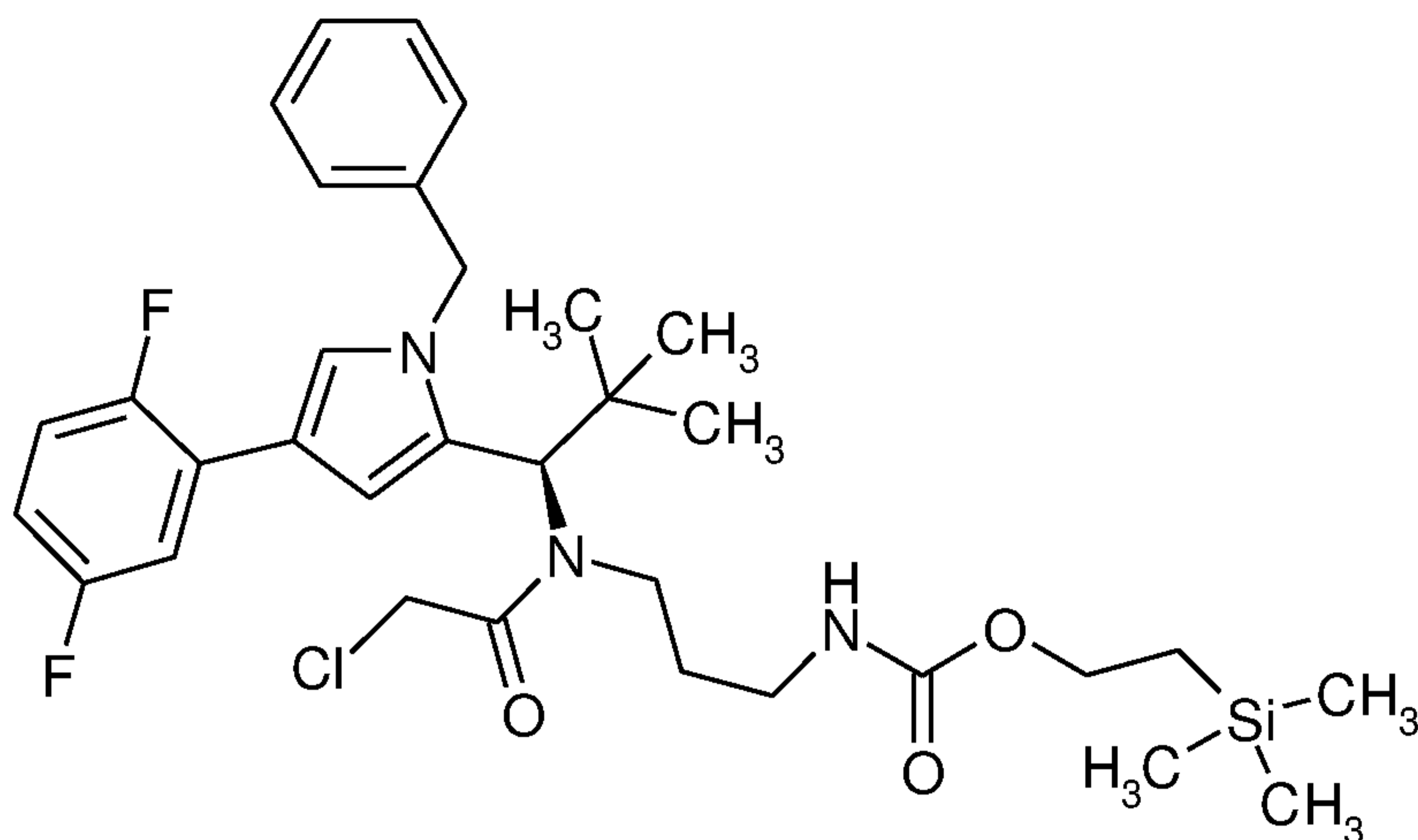


5 117.0 mg (0.19 mmol) of (2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate (Intermediate C70) and 21.6 mg (0.20 mmol) of 3-sulphanylpropanoic acid were initially charged in 3.0 ml of  
 10 methanol, 89.5 mg (0.65 mmol) of potassium carbonate were added and the mixture was stirred at 50°C for 4 h. The reaction mixture was diluted with ethyl acetate and the organic phase was washed with water and saturated NaCl solution. The organic phase was dried over magnesium sulphate, the solvent was evaporated under  
 15 reduced pressure and the residue was dried under high vacuum. The residue was used without further purification in the next step of the synthesis. This gave 106.1 mg (73% of theory) of the title compound.

20 LC-MS (Method 1):  $R_t = 1.42$  min; MS (ESI<sup>neg</sup>):  $m/z = 700$  (M-H)<sup>-</sup>.

### Intermediate C70

(2-(Trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate  
 25

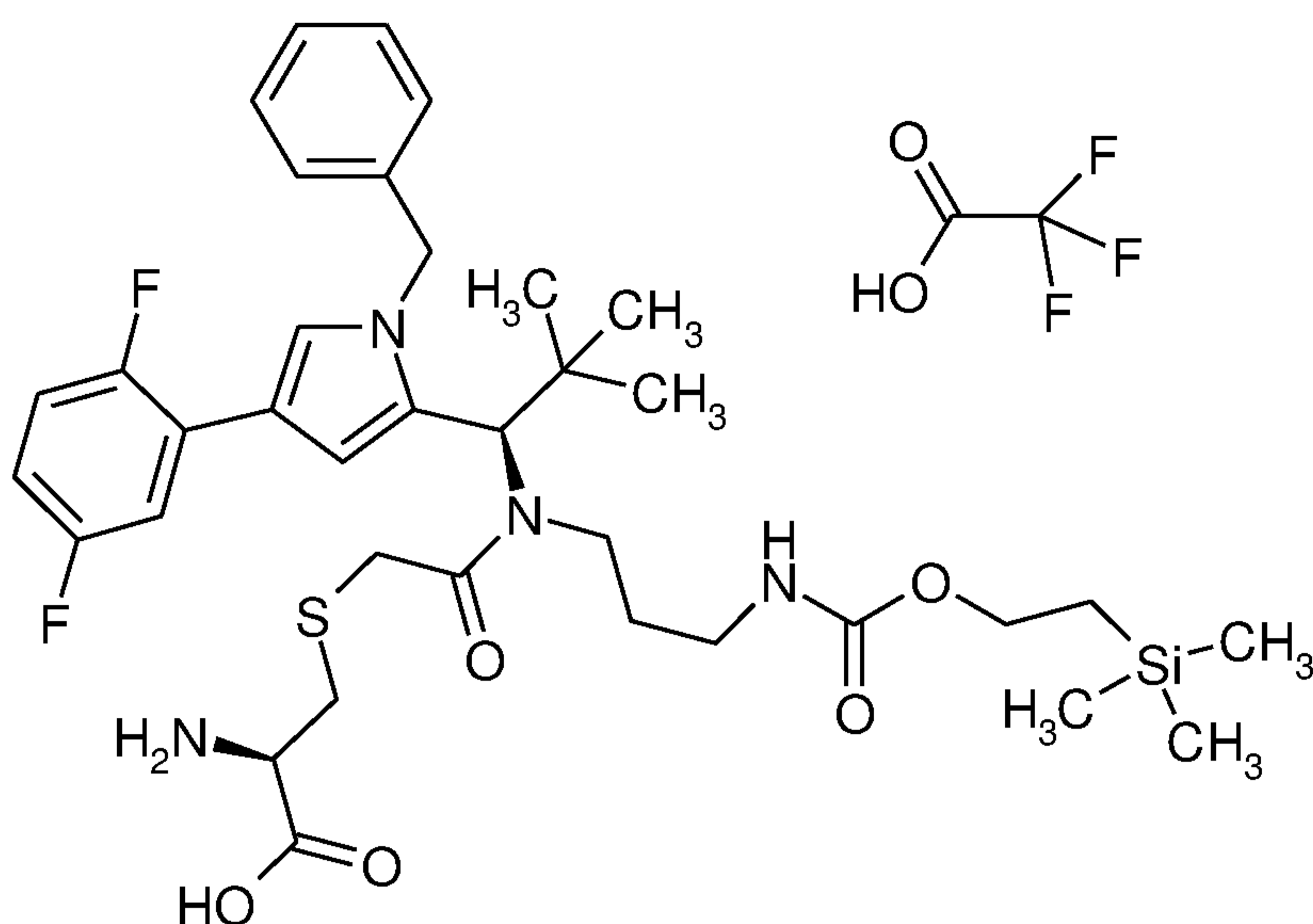


908.1 mg (1.63 mmol) of 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino]propyl carbamate (see synthesis of Intermediate C11) and 545.6 mg (5.39 mmol) of triethylamine were initially charged in 10.0 ml of dichloromethane, and the mixture was cooled to 0°C. At this temperature, 590.5 mg (5.23 mmol) of chloroacetyl chloride were added and the mixture was stirred at RT overnight. The reaction mixture was diluted with ethyl acetate and the organic phase was washed in each case three times with saturated sodium bicarbonate solution and saturated ammonium chloride solution. The organic phase was washed with saturated NaCl solution and dried over magnesium sulphate. The residue was purified by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 673.8 mg (65% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 1.53 min; MS (ESI<sup>neg</sup>):  $m/z$  = 676 ( $M+HCOO^-$ ).

### Intermediate C71

S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1)



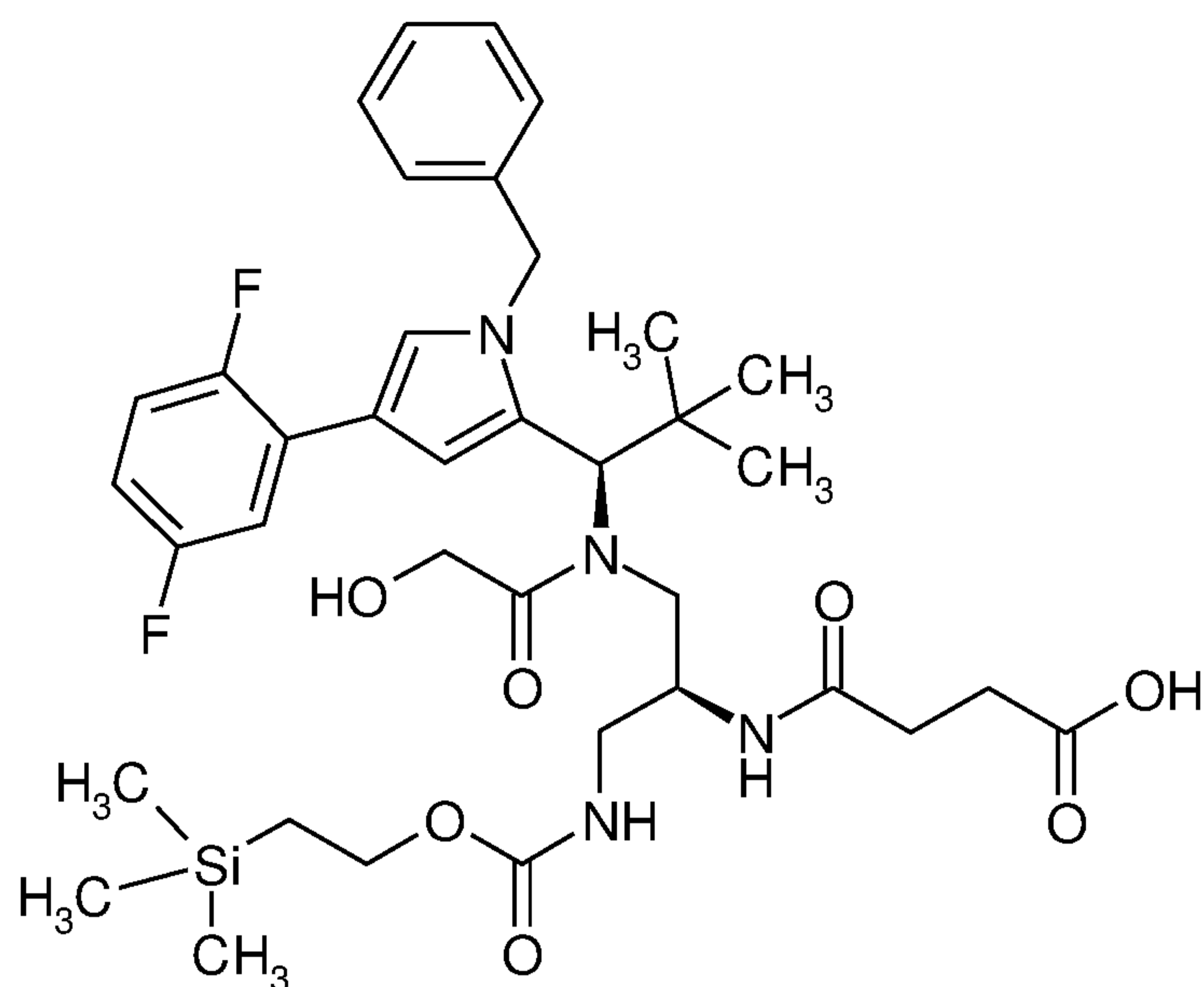
536.6 mg (4.43 mmol) of L-cysteine were suspended in 2.5 ml of  
 5 water together with 531.5 mg (6.33 mmol) of sodium bicarbonate.  
 400.0 mg (0.63 mmol) of 2-(trimethylsilyl)ethyl {3-[[{(1R)-1-[1-  
 benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl}(chloroacetyl)amino]propyl}carbamate  
 (Intermediate C70) dissolved in 25.0 ml of isopropanol and 1.16  
 10 g (7.59 mmol) of 1,8-diazabicyclo[5.4.0]undec-7-ene were added.  
 The reaction mixture was stirred at 50°C for 1.5 h. Ethyl acetate  
 was added to the reaction mixture and the organic phase was  
 washed repeatedly with saturated sodium bicarbonate solution and  
 once with sat. NaCl solution. The organic phase was dried over  
 15 magnesium sulphate, the solvent was evaporated under reduced  
 pressure and the residue was dried under high vacuum. The residue  
 was purified by preparative RP-HPLC (column: Reprosil 250x30;  
 10µ, flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents  
 were evaporated under reduced pressure and the residue was dried  
 20 under high vacuum. This gave 449.5 mg (86% of theory) of the  
 title compound.

LC-MS (Method 1):  $R_t = 1.20$  min; MS (ESIpos):  $m/z = 717$  (M+H)<sup>+</sup>.

## 25 Intermediate C72

(9S)-9-[[{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-

yl]-2,2-dimethylpropyl}(glycoloyl) amino]methyl}-2,2-dimethyl-6,11-dioxo-5-oxa-7,10-diaza-2-silatetradecan-14-oic acid



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90 mg (0.212 mmol) of Intermediate L72 were initially charged in 6 ml of dichloromethane, and 86  $\mu$ l (1.06 mmol) of pyridine and 135 mg (0.318 mmol) of Dess-Martin periodinane were added. The mixture was stirred at RT for 30 min. The reaction was then diluted with 30 ml of dichloromethane and the organic phase was washed twice with 10% strength  $\text{Na}_2\text{S}_2\text{O}_3$  solution and once with 5% strength citric acid solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The aldehyde obtained in this manner was reacted without further purification.

63 mg (0.177 mmol) of Intermediate C52 were dissolved in 15 ml of dichloromethane, and 52.4 mg (0.247 mmol) of sodium triacetoxyborohydride and 20.2  $\mu$ l of acetic acid were added. After 5 min of stirring at RT, 89.6 mg (0.212 mmol) of the aldehyde described above were added, and the reaction was stirred at RT for 20 min. The reaction was concentrated under reduced pressure and the residue was purified by preparative HPLC. After combination of the appropriate fractions, the solvent was evaporated under reduced pressure and the residue was lyophilized from acetonitrile/water. This gave 71 mg (53% of theory over 2 steps) of benzyl (9R)-9-[(1R)-1-[1-benzyl-4-

(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)methyl]-2,2-dimethyl-6,11-dioxo-5-oxa-7,10-diaza-2-silatetradecan-14-oate.

5 LC-MS (Method 1):  $R_t = 1.21$  min; MS (ESIpos):  $m/z = 761$  (M+H)<sup>+</sup>.

70 mg (92  $\mu$ mol) of this intermediate were taken up in 15 ml of dichloromethane, the mixture was cooled to 10°C and 54  $\mu$ l of triethylamine and 25.5  $\mu$ l (0.23 mmol) of acetoxyacetyl chloride  
10 were added. After 1 h of stirring at RT, the same amounts of acid chloride and triethylamine were added, and once more after a further hour of stirring at RT. The reaction was then stirred at RT for a further 30 min and then concentrated under reduced pressure, and the residue was purified by preparative HPLC. The  
15 appropriate fractions were combined giving, after evaporation of the solvents and lyophilization of the residue from acetonitrile/water, 46.5 mg (59% of theory) of the acylated intermediate.

20 LC-MS (Method 1):  $R_t = 1.53$  min; MS (ESIpos):  $m/z = 861$  (M+H)<sup>+</sup>.

46 mg (53  $\mu$ mol) of this intermediate were dissolved in 5 ml of methanol, and 2.7 ml of a 2M lithium hydroxide solution were added. After 10 min of stirring at RT, the reaction was adjusted  
25 to pH 3-4 with acetic acid and then diluted with 15 ml of water. The aqueous phase was extracted with ethyl acetate and the organic phase was dried over magnesium sulphate and concentrated. The residue was lyophilized from acetonitrile/water giving, after drying of the residue under  
30 high vacuum, 37 mg (90% of theory) of the title compound as a white solid.

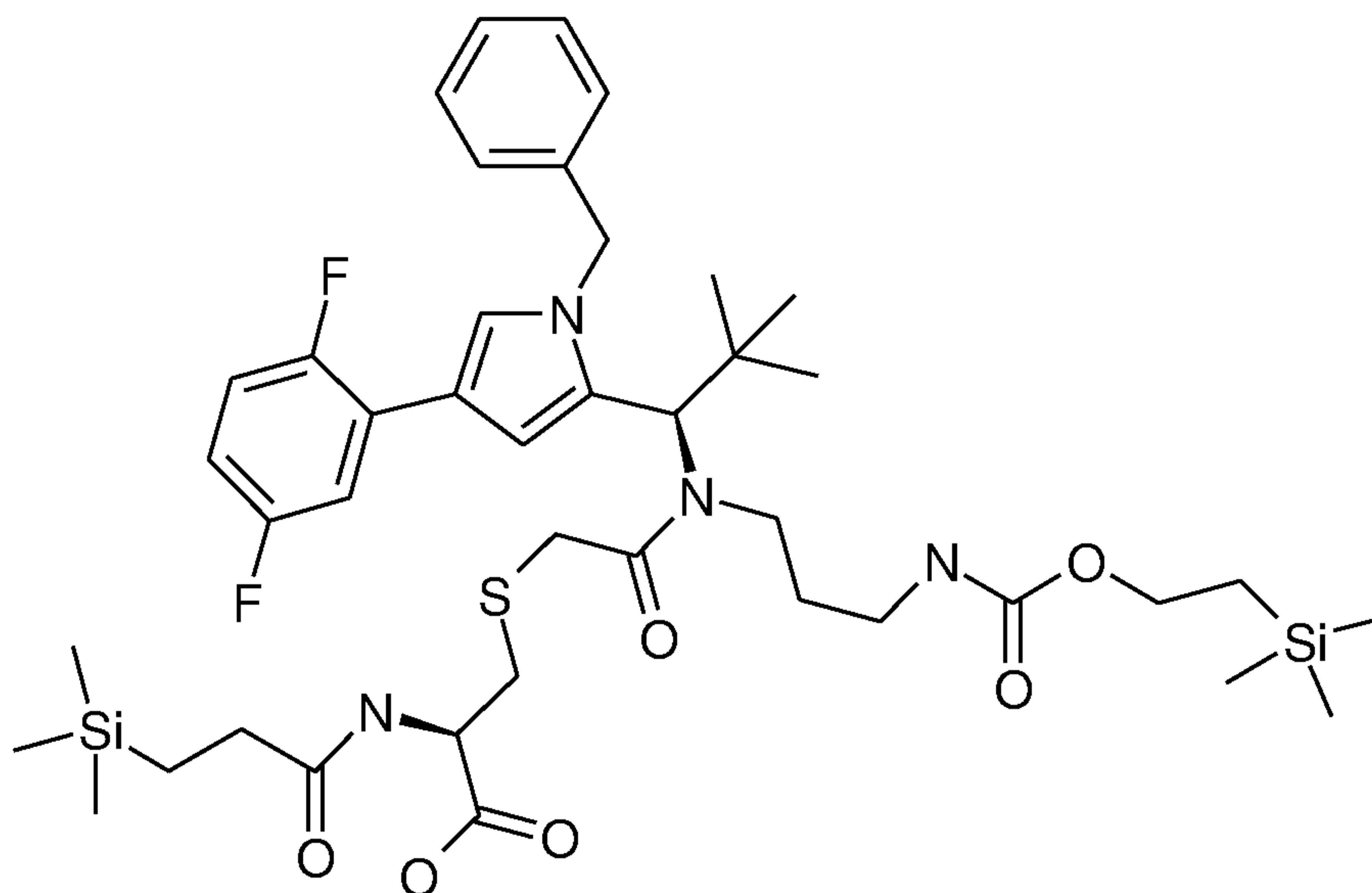
LC-MS (Method 1):  $R_t = 1.32$  min; MS (ESIpos):  $m/z = 729$  (M+H)<sup>+</sup>.

35 **Intermediate C73**

S-(11-{(1R)-1-[1-Benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-



diaza-2-silatridecan-13-yl)-N-[3-(trimethylsilyl)propanoyl]-L-cysteine



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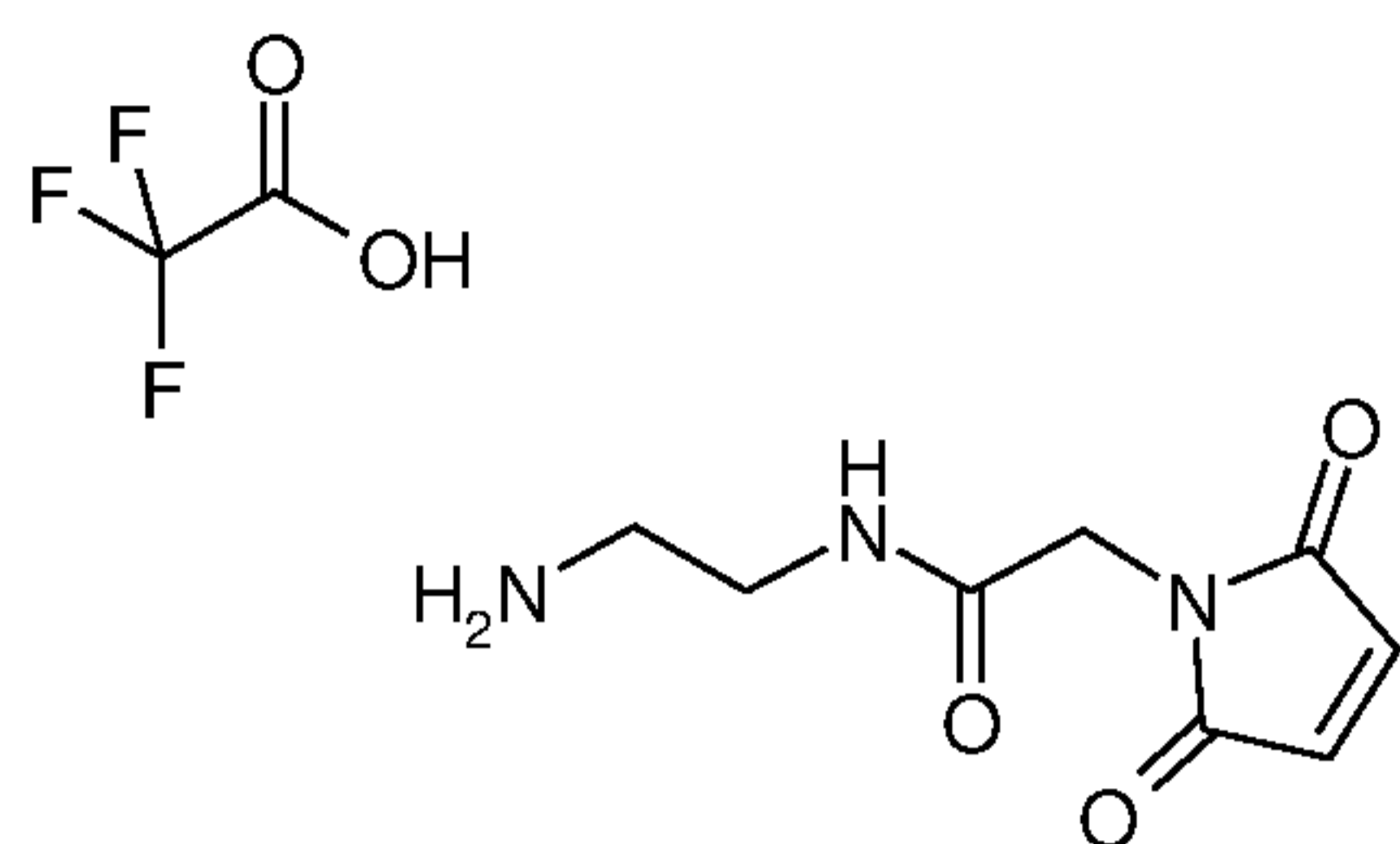
619 mg (0.86 mmol) of S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1) (Intermediate C71) were initially charged in 8.8 ml of dichloromethane, and 87 mg (0.86 mmol) of triethylamine and 224 mg (0.86 mmol) of N-[2-(trimethylsilyl)ethoxycarbonyloxy]pyrrolidine-2,5-dione were added. After 1 h, 45 mg (0.17 mmol) of N-[2-(trimethylsilyl)ethoxycarbonyloxy]pyrrolidine-2,5-dione were added. The reaction mixture was stirred at RT for 1 h. The mixture was concentrated under reduced pressure, the residue was taken up in dichloromethane and the organic phase was then washed twice with water and a saturated sodium bicarbonate solution. The organic phase was dried over magnesium sulphate, concentrated on a rotary evaporator and dried under high vacuum. The residue was used further without further purification. This gave 602 mg (71%, purity 87%) of the title compound.

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LC-MS (Method 1):  $R_t = 1.58$  min; MS (ESIpos):  $m/z = 861$  (M+H)<sup>+</sup>.

**Intermediate L1**

Trifluoroacetic acid / N-(2-aminoethyl)-2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetamide (1:1)



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The title compound was prepared by classical methods of peptide chemistry from commercially available (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetic acid and tert-butyl (2-aminoethyl)carbamate.

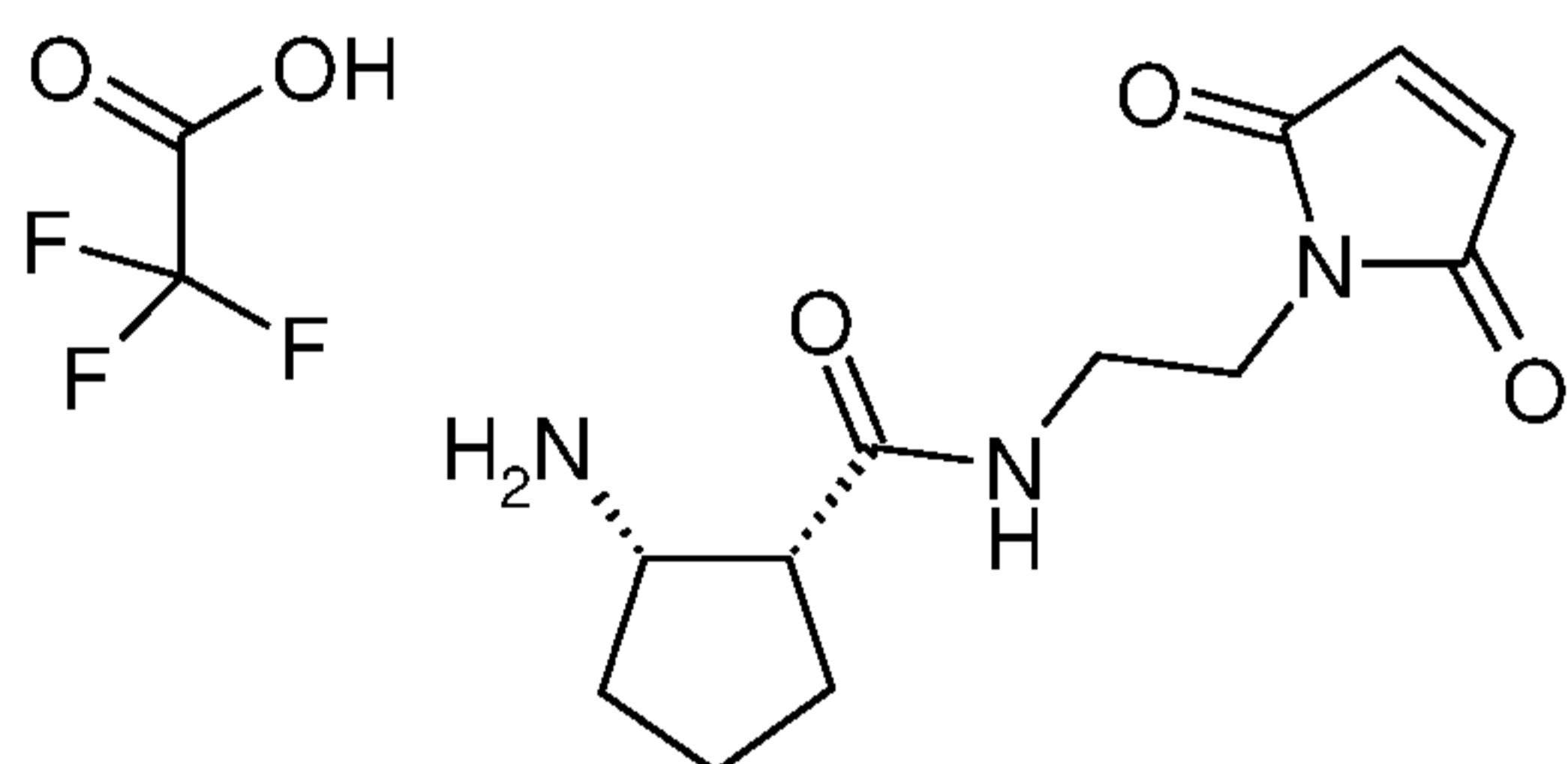
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HPLC (Method 11):  $R_t = 0.19$  min;

LC-MS (Method 1):  $R_t = 0.17$  min; MS (ESIpos):  $m/z = 198$  (M+H)<sup>+</sup>.

### 15 Intermediate L2

Trifluoroacetic acid / rel-(1R,2S)-2-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)



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The title compound was prepared from 50 mg (0.214 mmol) of commercially available cis-2-[(tert-butoxycarbonyl)amino]-1-cyclopentanecarboxylic acid and 60 mg (0.235 mmol) of likewise commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) by coupling with EDC/HOBT and subsequent deprotection with TFA. This gave 36 mg (38% of theory over 2 steps) of the title compound.

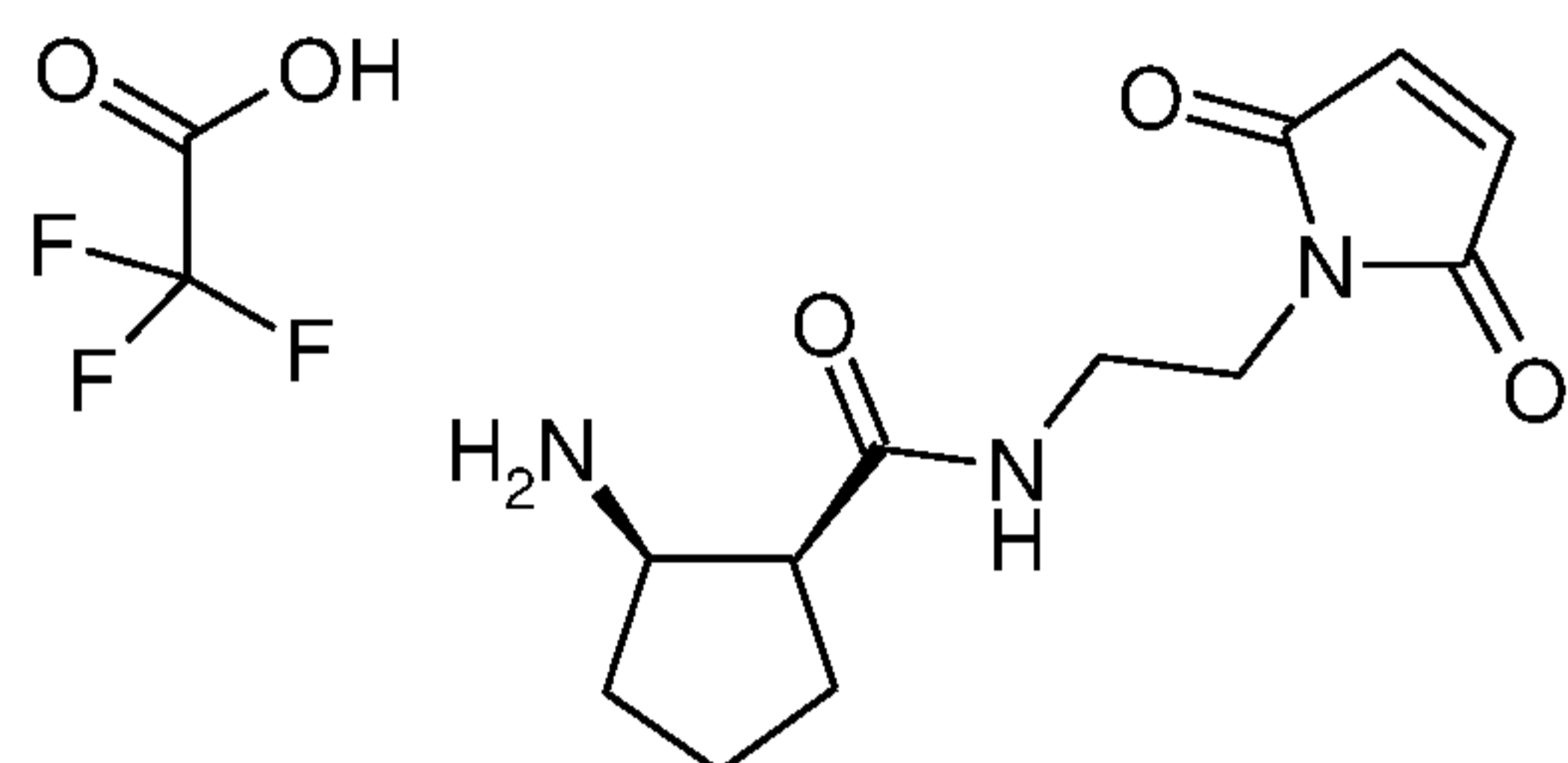
30 HPLC (Method 11):  $R_t = 0.2$  min;

LC-MS (Method 1):  $R_t = 0.17$  min; MS (ESIpos):  $m/z = 252$  (M+H)<sup>+</sup>.

### Intermediate L3

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Trifluoroacetic acid / (1S,2R)-2-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)



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The title compound was prepared from 50 mg (0.214 mmol) of commercially available (1S,2R)-2-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid with 72 mg (0.283 mmol) of likewise commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) by coupling with EDC/HOBT and subsequent deprotection with TFA. This gave 13 mg (16% of theory over 2 steps) of the title compound.

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HPLC (Method 11):  $R_t = 0.2$  min;

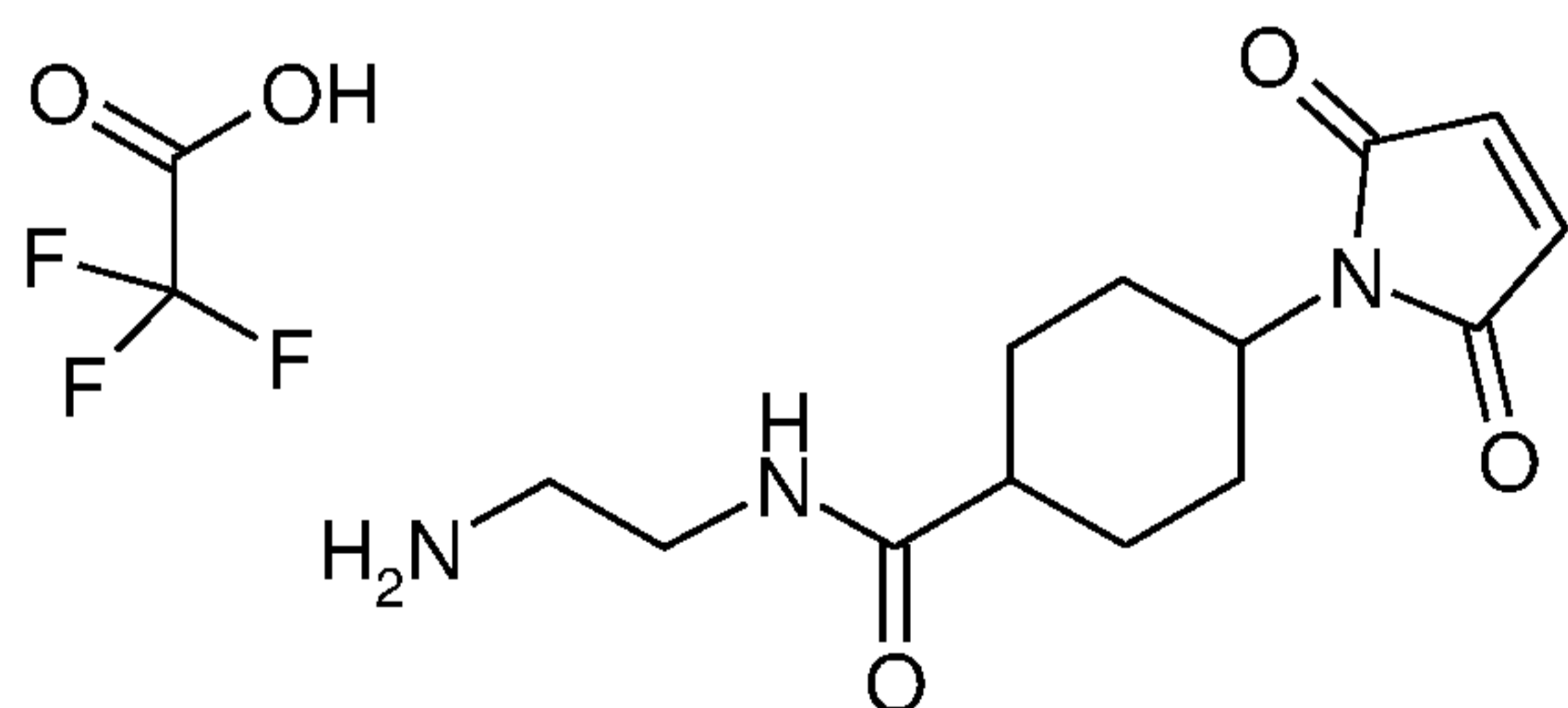
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LC-MS (Method 1):  $R_t = 0.2$  min; MS (ESIpos):  $m/z = 252$  (M+H)<sup>+</sup>.

### Intermediate L4

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Trifluoroacetic acid / N-(2-aminoethyl)-4-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)cyclohexanecarboxamide (1:1)



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The title compound was prepared by classical methods of peptide

chemistry from commercially available 1-[(4-[(2,5-dioxopyrrolidin-1-yl)oxy]carbonyl)cyclohexyl)methyl]-1H-pyrrole-2,5-dione and tert-butyl (2-aminoethyl) carbamate.

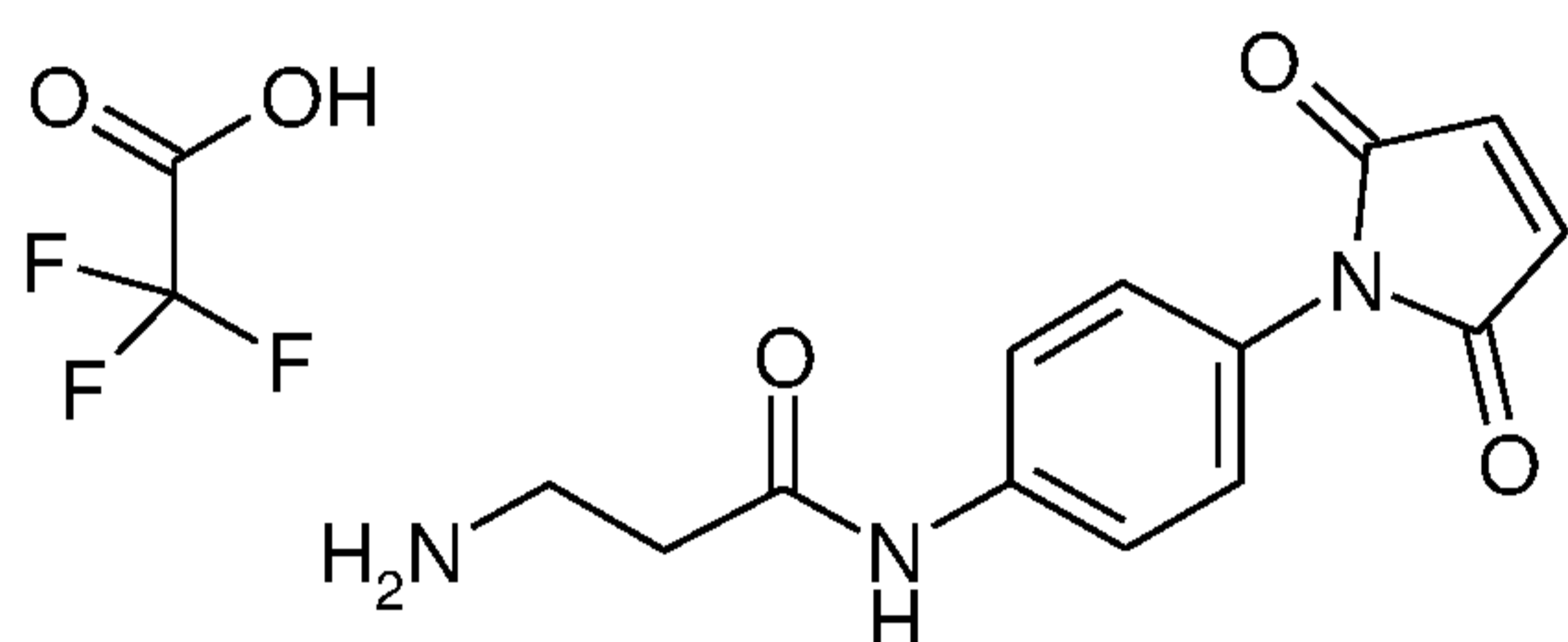
5 HPLC (Method 11):  $R_t = 0.26$  min;

LC-MS (Method 1):  $R_t = 0.25$  min; MS (ESIpos):  $m/z = 280$  (M+H)<sup>+</sup>.

### Intermediate L5

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Trifluoroacetic acid / N-[4-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)phenyl]-beta-alaninamide (1:1)



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The title compound was prepared by classical methods of peptide chemistry from commercially available 1-(4-aminophenyl)-1H-pyrrole-2,5-dione and N-(tert-butoxycarbonyl)-beta-alanine.

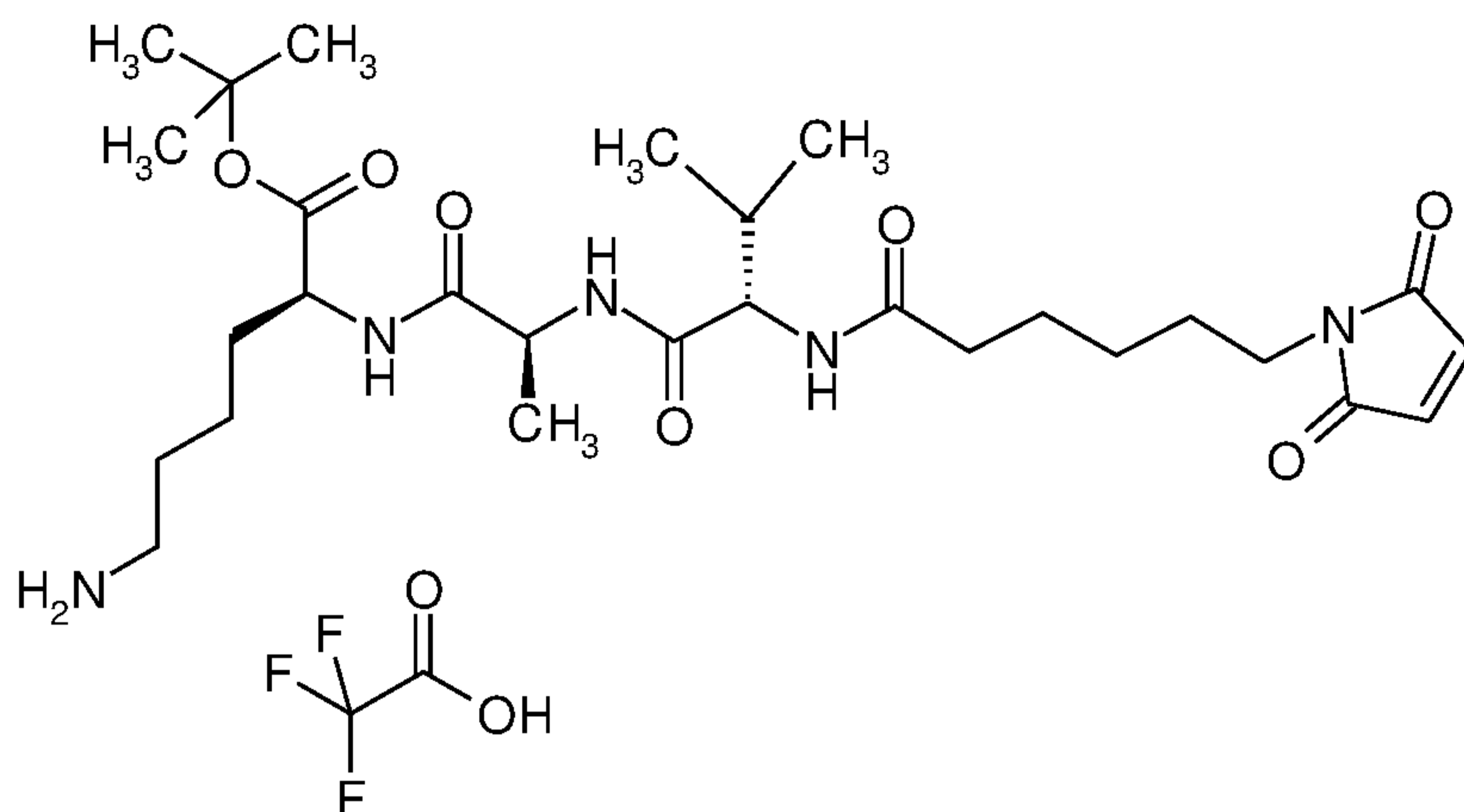
20 HPLC (Method 11):  $R_t = 0.22$  min;

LC-MS (Method 1):  $R_t = 0.22$  min; MS (ESIpos):  $m/z = 260$  (M+H)<sup>+</sup>.

### Intermediate L6

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Trifluoroacetic acid / tert-butyl-N-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-L-alanyl-L-lysinate (1:1)



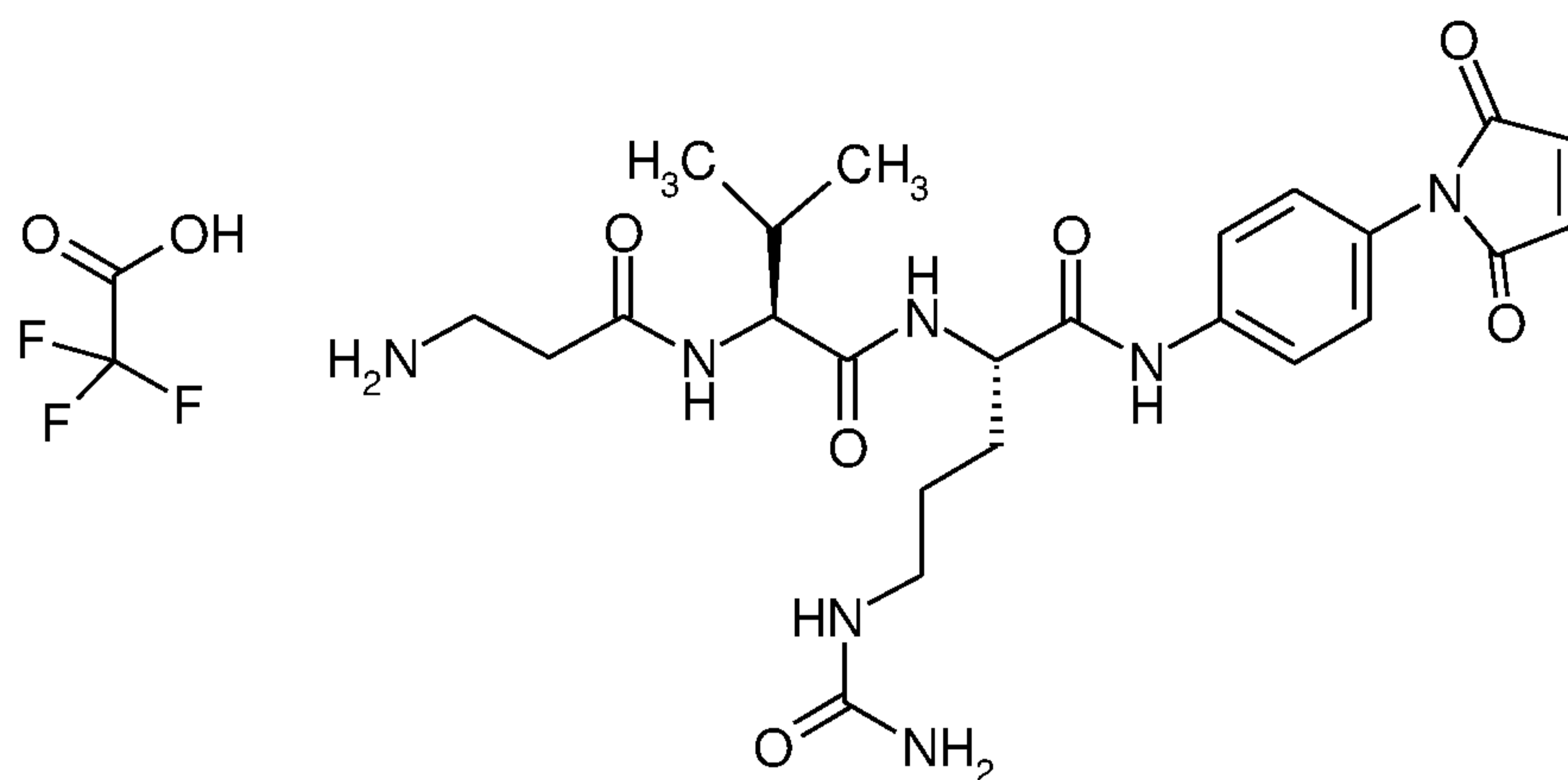
The title compound was prepared by initially coupling, in the presence of EDC/HOBT, commercially available 6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoic acid with the partially protected peptide tert-butyl L-valyl-L-alanyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysinate, prepared by classical methods of peptide chemistry. This was followed by deprotection at the amino group under gentle conditions by stirring in 5% strength trifluoroacetic acid in DCM at RT, which gave the title compound in a yield of 37%.

HPLC (Method 11):  $R_t = 1.29$  min;

LC-MS (Method 1):  $R_t = 0.62$  min; MS (ESIpos):  $m/z = 566$  (M+H)<sup>+</sup>.

### Intermediate L7

Trifluoroacetic acid / beta-alanyl-L-valyl-N<sup>5</sup>-carbamoyl-N-[4-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)phenyl]-L-ornithinamide (1:1)



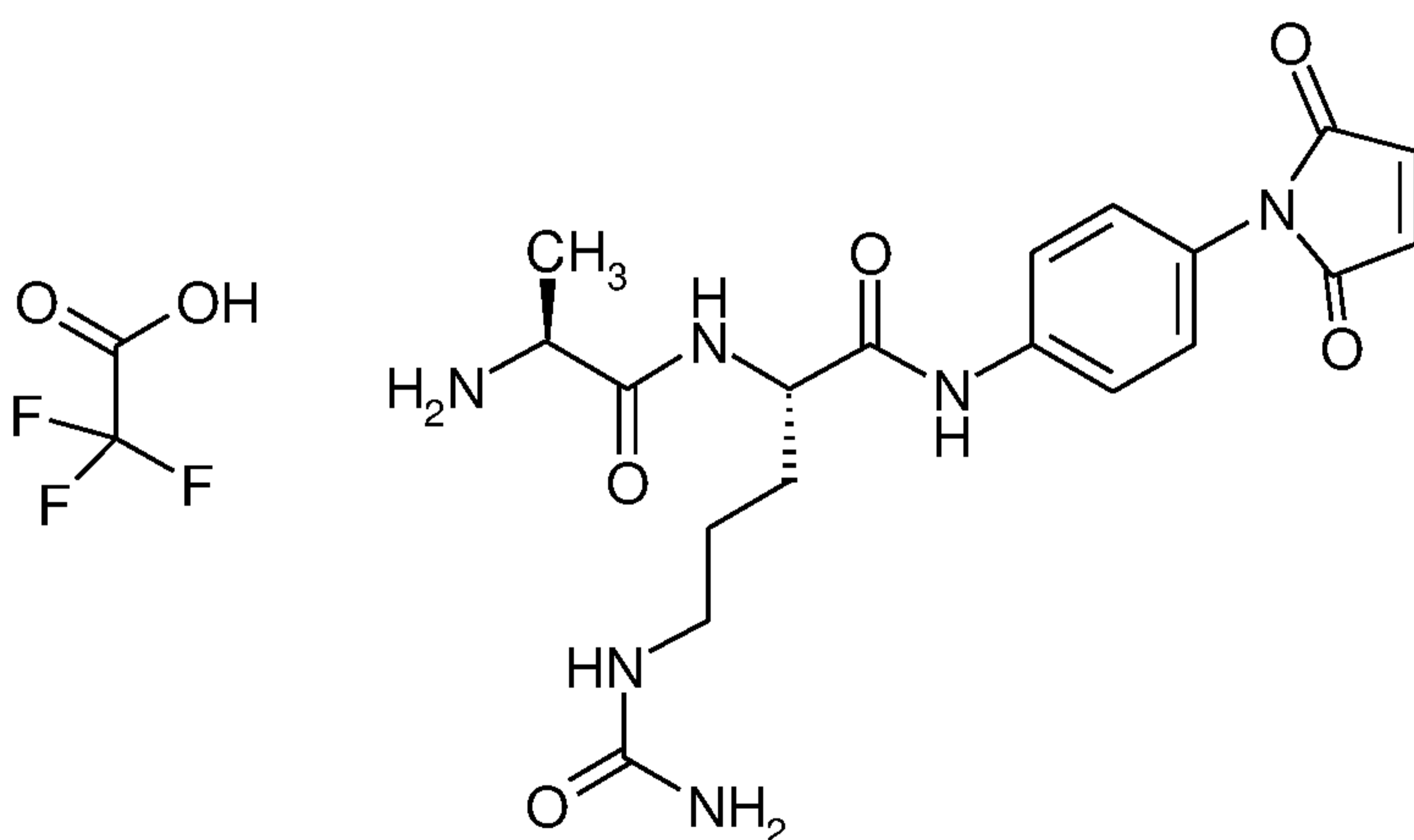
The title compound was prepared according to classical methods of peptide chemistry from commercially available 1-(4-aminophenyl)-1H-pyrrole-2,5-dione by sequential coupling with N2-(tert-butoxycarbonyl)-N5-carbamoyl-L-ornithine in the presence of HATU, deprotection with TFA, coupling with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-L-valinate, deprotection with TFA, coupling with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-beta-alaninate and another deprotection with TFA. This gave 32 mg of the title compound.

HPLC (Method 11):  $R_t = 0.31$  min;

15 LC-MS (Method 1):  $R_t = 0.47$  min; MS (ESIpos):  $m/z = 516$  (M+H)<sup>+</sup>.

### Intermediate I8

20 Trifluoroacetic acid / L-alanyl-N<sup>5</sup>-carbamoyl-N-[4-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)phenyl]-L-ornithinamide (1:1)



The title compound was prepared according to classical methods of peptide chemistry from commercially available 1-(4-aminophenyl)-1H-pyrrole-2,5-dione by sequential coupling with  
 5 N<sup>2</sup>-(tert-butoxycarbonyl)-N<sup>5</sup>-carbamoyl-L-ornithine in the presence of HATU, deprotection with TFA, coupling with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-L-alaninate and another deprotection with TFA. This gave 171 mg of the title compound.

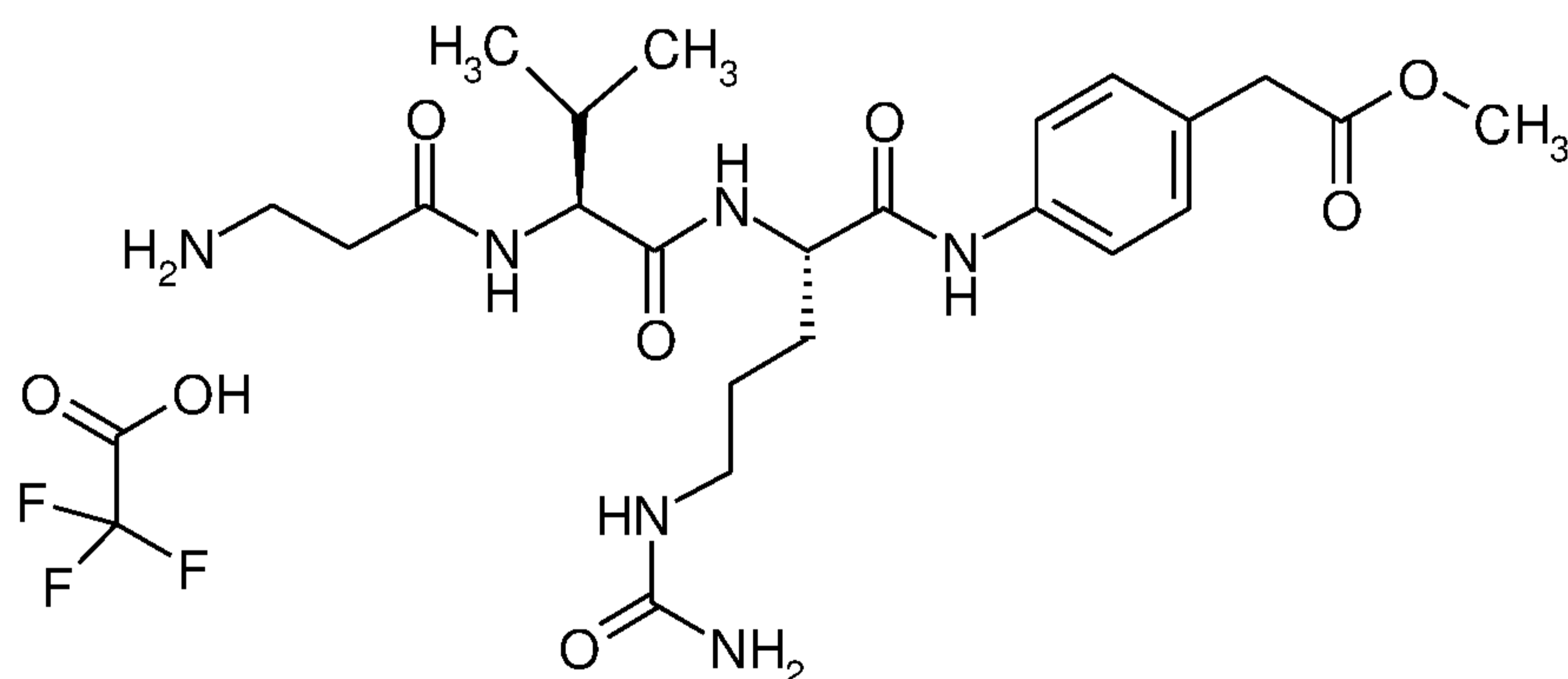
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HPLC (Method 11): R<sub>t</sub> = 0.23 min;

LC-MS (Method 7): R<sub>t</sub> = 0.3 min; MS (ESIpos): m/z = 417 (M+H)<sup>+</sup>.

### 15 Intermediate L9

Trifluoroacetic acid / beta-alanyl-L-valyl-N<sup>5</sup>-carbamoyl-N-[4-(2-methoxy-2-oxoethyl)phenyl]-L-ornithinamide (1:1)



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The title compound was prepared analogously to Intermediate L7 from commercially available methyl (4-aminophenyl)acetate. This gave 320 mg of the title compound.

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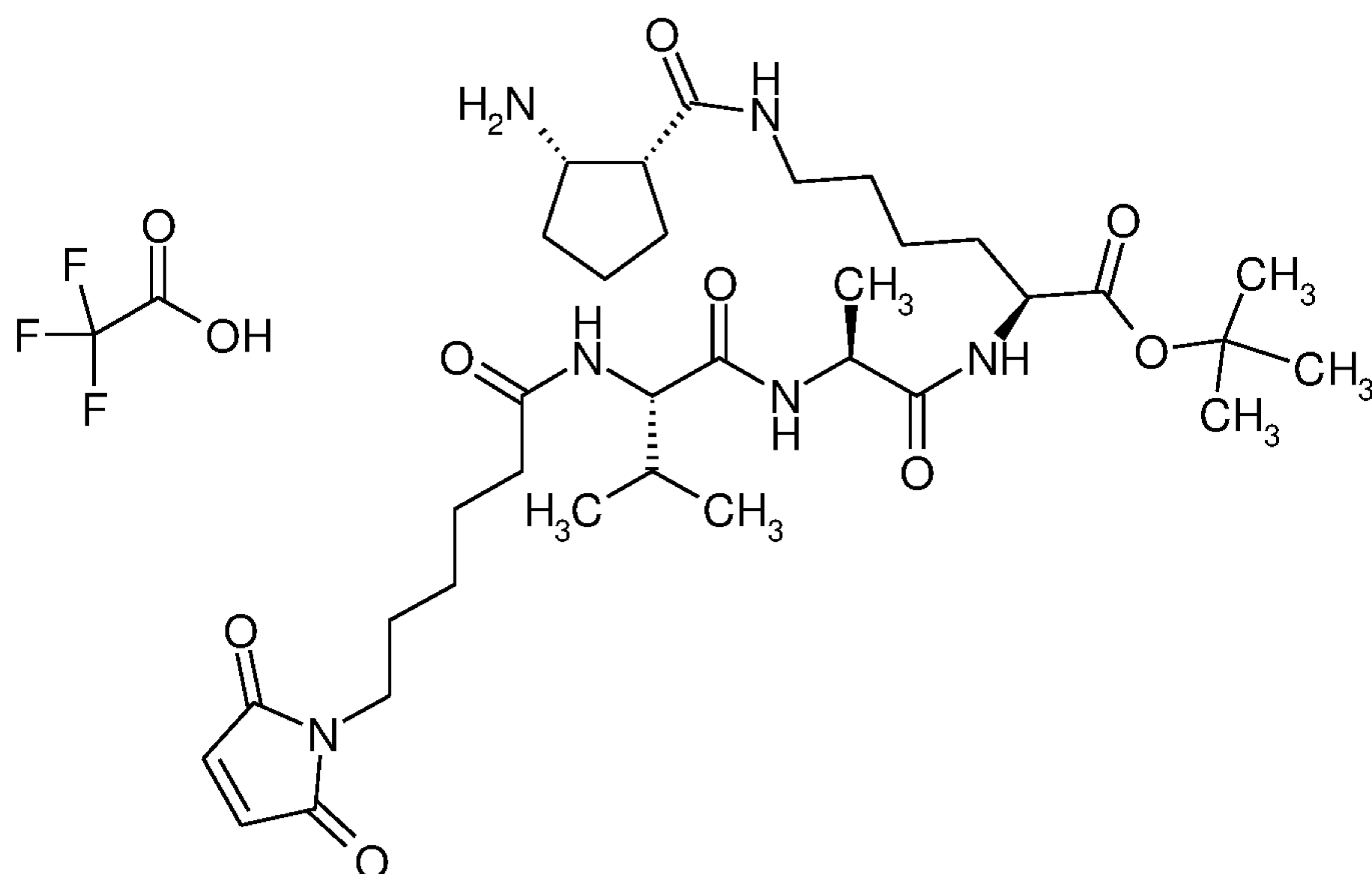
HPLC (Method 11): R<sub>t</sub> = 0.45 min;

LC-MS (Method 1): R<sub>t</sub> = 0.48 min; MS (ESIpos): m/z = 493 (M+H)<sup>+</sup>.

### 30 Intermediate L10

N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-

L-alanyl-rel-N<sup>6</sup>-{[(1R,2S)-2-aminocyclopentyl]carbonyl}-L-lysine  
/ trifluoroacetic acid (1:2)



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The title compound was prepared from Intermediate L6 by coupling with *cis*-2-[(*tert*-butoxycarbonyl)amino]-1-cyclopentanecarboxylic acid with EDC/HOBT and subsequent deprotection with TFA. This gave 12 mg (52% of theory over 2 steps) of the title compound.

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HPLC (Method 11):  $R_t = 1.45$  min;

LC-MS (Method 1):  $R_t = 0.73$  min; MS (ESIpos):  $m/z = 677$  (M+H)<sup>+</sup>.

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### Intermediate L11

N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-L-alanyl-N<sup>6</sup>-{[(1S,2R)-2-aminocyclopentyl]carbonyl}-L-lysine /  
trifluoroacetic acid (1:2)

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dioxane/water 1:1.

1.2 ml of a saturated sodium bicarbonate solution were then added and the reaction was stirred at RT. After a total of 5 days of stirring and 2 further additions of the same amounts of the sodium bicarbonate solution, the reaction was worked up by acidification with trifluoroacetic acid, concentration on a rotary evaporator and purification by preparative HPLC. The appropriate fractions were combined, the solvent was removed under reduced pressure and the residue was lyophilized from acetonitrile/water 1:1.

The residue was taken up in 3 ml of dichloromethane, and 1 ml of trifluoroacetic acid was added. After 15 min of stirring at RT, the solvent was removed under reduced pressure and the residue was lyophilized from acetonitrile/water 1:1. This gave 70 mg (67% of theory over 2 steps) of the title compound as a resinous residue.

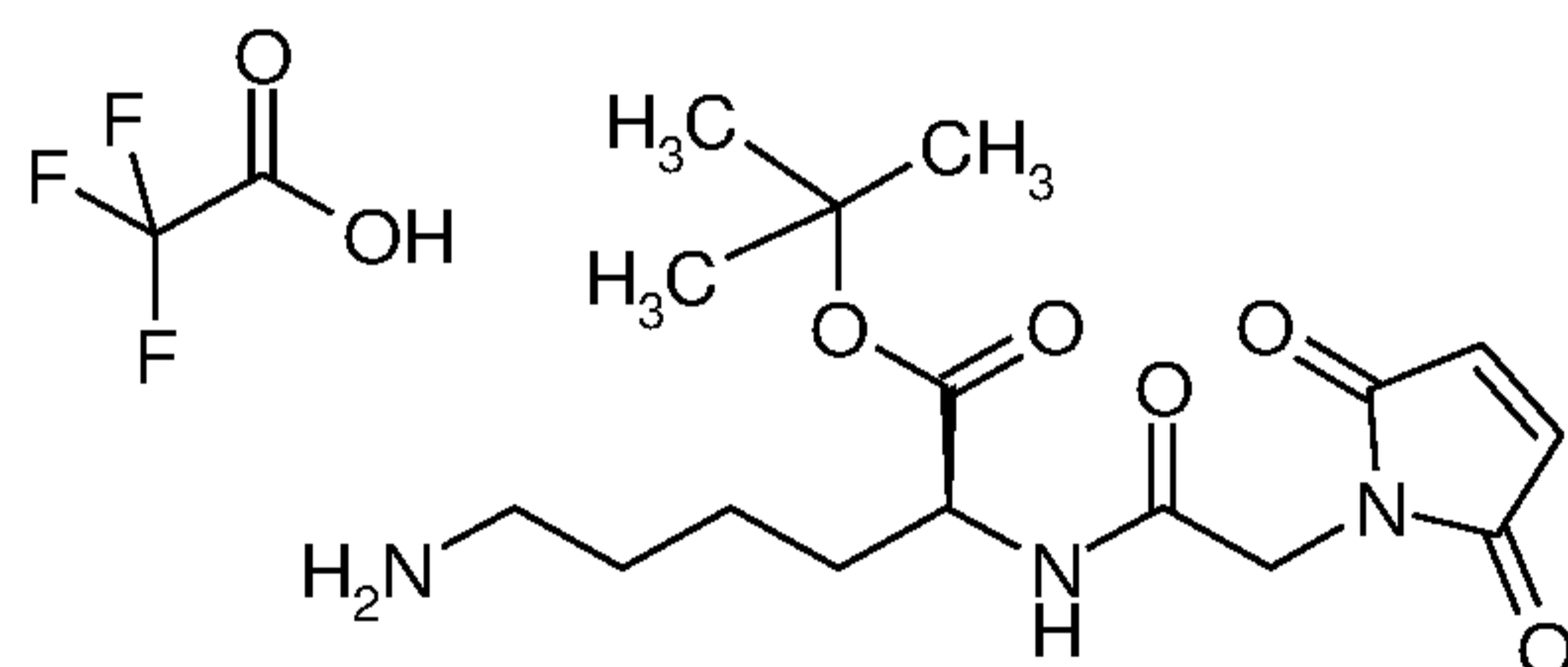
HPLC (Method 11):  $R_t = 0.2$  min;

LC-MS (Method 1):  $R_t = 0.18$  min; MS (ESIpos):  $m/z = 185$  (M+H)<sup>+</sup>.

### Intermediate L13

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Trifluoroacetic acid / tert-butyl N2-[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]-L-lysinate (1:1)



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The title compound was prepared by coupling of (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetic acid with tert-butyl N6-(tert-butoxycarbonyl)-L-lysinate hydrochloride (1:1) in the presence of EDC/HOBT and subsequent gentle removal of the tert-butoxycarbonyl protective group analogously to Intermediate L6.

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HPLC (Method 11):  $R_t = 0.42$  min;

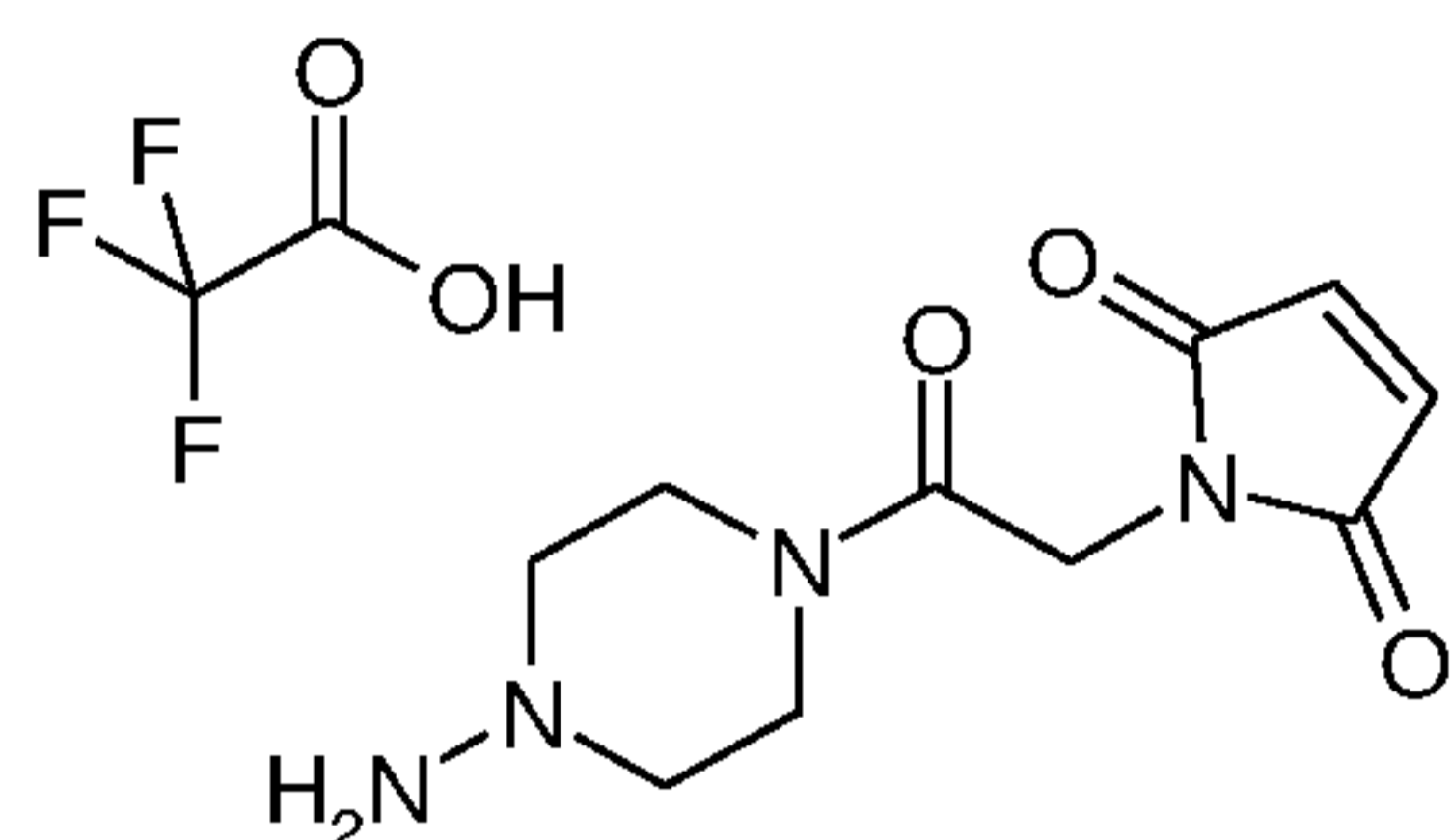
LC-MS (Method 1):  $R_t = 0.43$  min; MS (ESIpos):  $m/z = 340$  (M+H)<sup>+</sup>.

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### Intermediate L14

Trifluoroacetic acid / 1-[2-(4-aminopiperazin-1-yl)-2-oxoethyl]-1H-pyrrole-2,5-dione (1:1)

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The title compound was prepared analogously to Intermediate L2 over 2 steps from tert-butyl piperazin-1-ylcarbamate and (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetic acid.

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HPLC (Method 11):  $R_t = 0.2$  min;

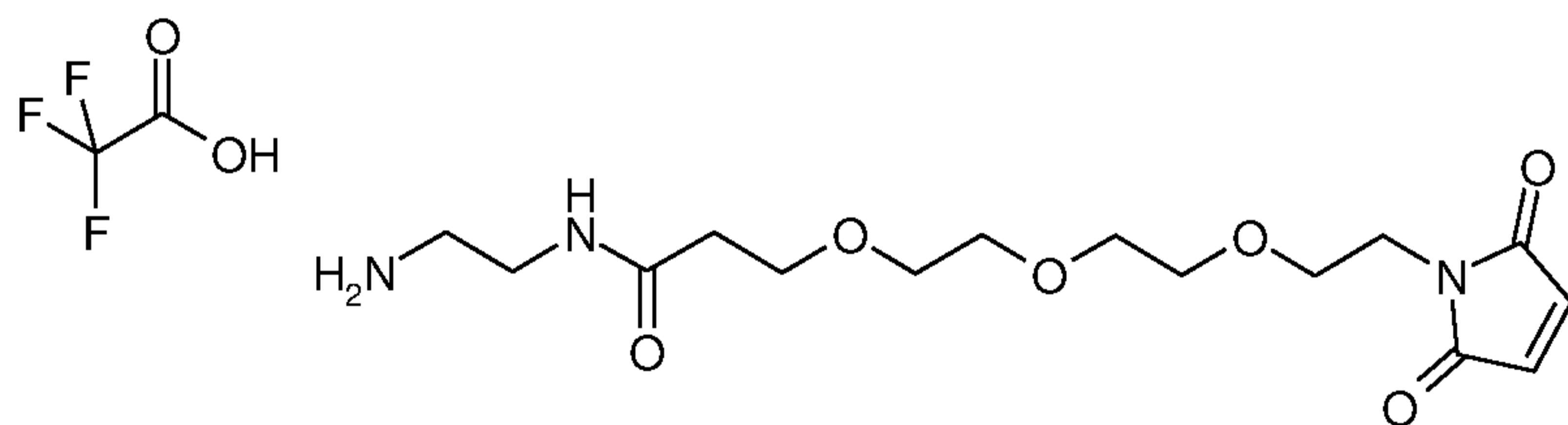
LC-MS (Method 3):  $R_t = 0.25$  min; MS (ESIpos):  $m/z = 239$  (M+H)<sup>+</sup>.

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### Intermediate L15

Trifluoroacetic acid / N-(2-aminoethyl)-3-(2-{2-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethoxy]ethoxy}ethoxy)propanamide (1:1)

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2.93 g (10.58 mmol) of tert-butyl 3-{2-[2-(2-aminoethoxy)ethoxy]ethoxy}propanoate were dissolved in 100 ml of dioxane/water 1:1, and 3.28 g (21.15 mmol) of methyl 2,5-

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dioxo-2,5-dihydro-1H-pyrrole-1-carboxylate and a saturated sodium bicarbonate solution were added until a pH of 6-7 had been reached. The solution was stirred at RT for 30 min and the 1,4-dioxane was then evaporated under reduced pressure. 200 ml  
 5 of water were then added, and the mixture was extracted three times with in each case 300 ml of ethyl acetate. The organic extracts were combined, dried over magnesium sulphate and filtered. Concentration gave tert-butyl 3-(2-{2-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethoxy]ethoxy}ethoxy)propanoate as a  
 10 brown oil which was then dried under high vacuum.

HPLC (Method 11):  $R_t = 1.5$  min;

LC-MS (Method 3):  $R_t = 0.88$  min; MS (ESIpos):  $m/z = 375$  ( $M+NH_4$ )<sup>+</sup>.  
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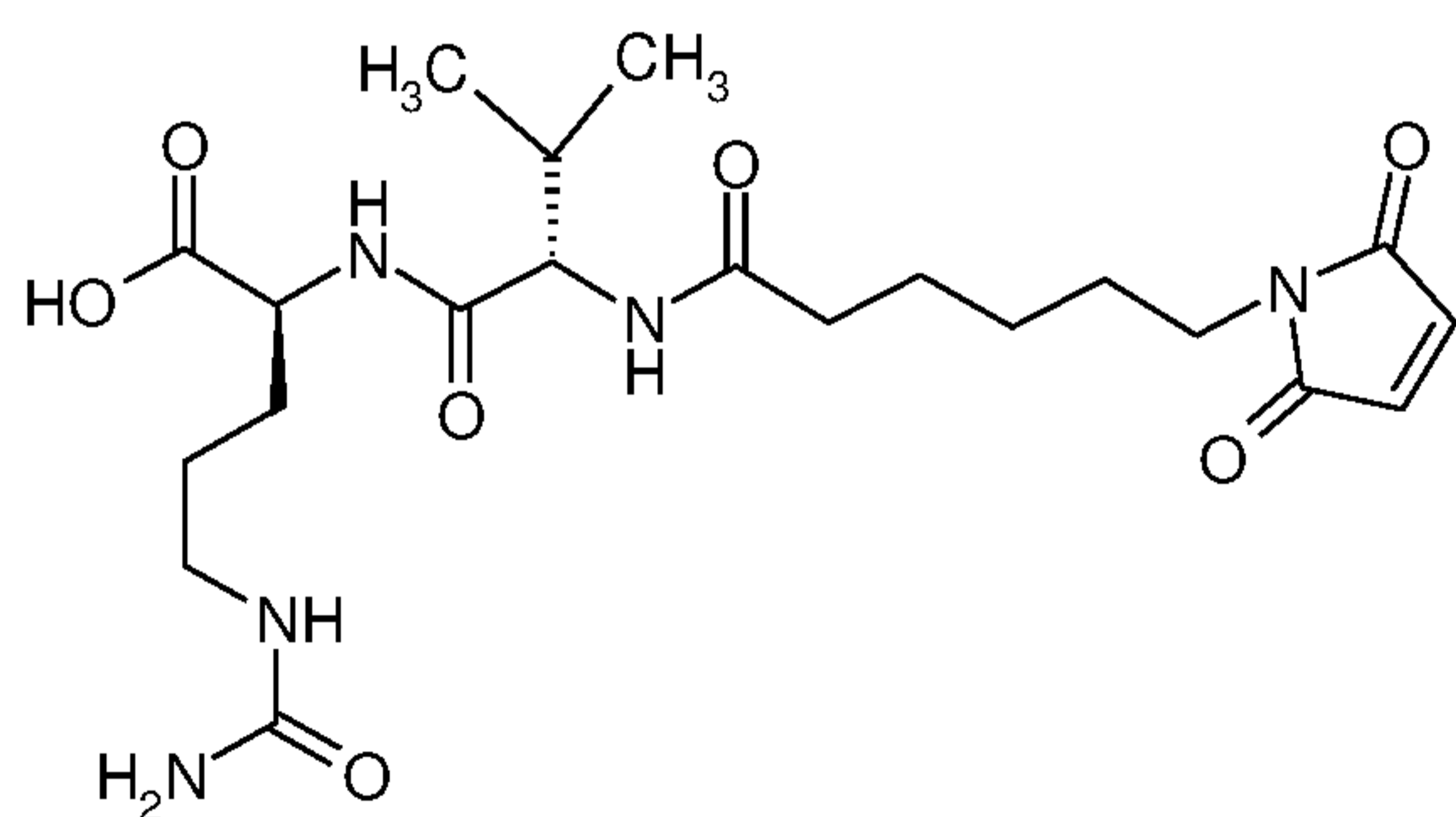
This intermediate was converted by standard methods (deprotection with TFA, coupling with tert-butyl (2-aminoethyl)carbamate and another deprotection with TFA) into the title compound.

20 HPLC (Method 11):  $R_t = 0.2$  min;

LC-MS (Method 3):  $R_t = 0.25$  min; MS (ESIpos):  $m/z = 344$  ( $M+H$ )<sup>+</sup>.

### 25 Intermediate L16

N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithine



30 535 mg (1.73 mmol) of commercially available 1-{6-[(2,5-dioxopyrrolidin-1-yl)oxy]-6-oxohexyl}-1H-pyrrole-2,5-dione and

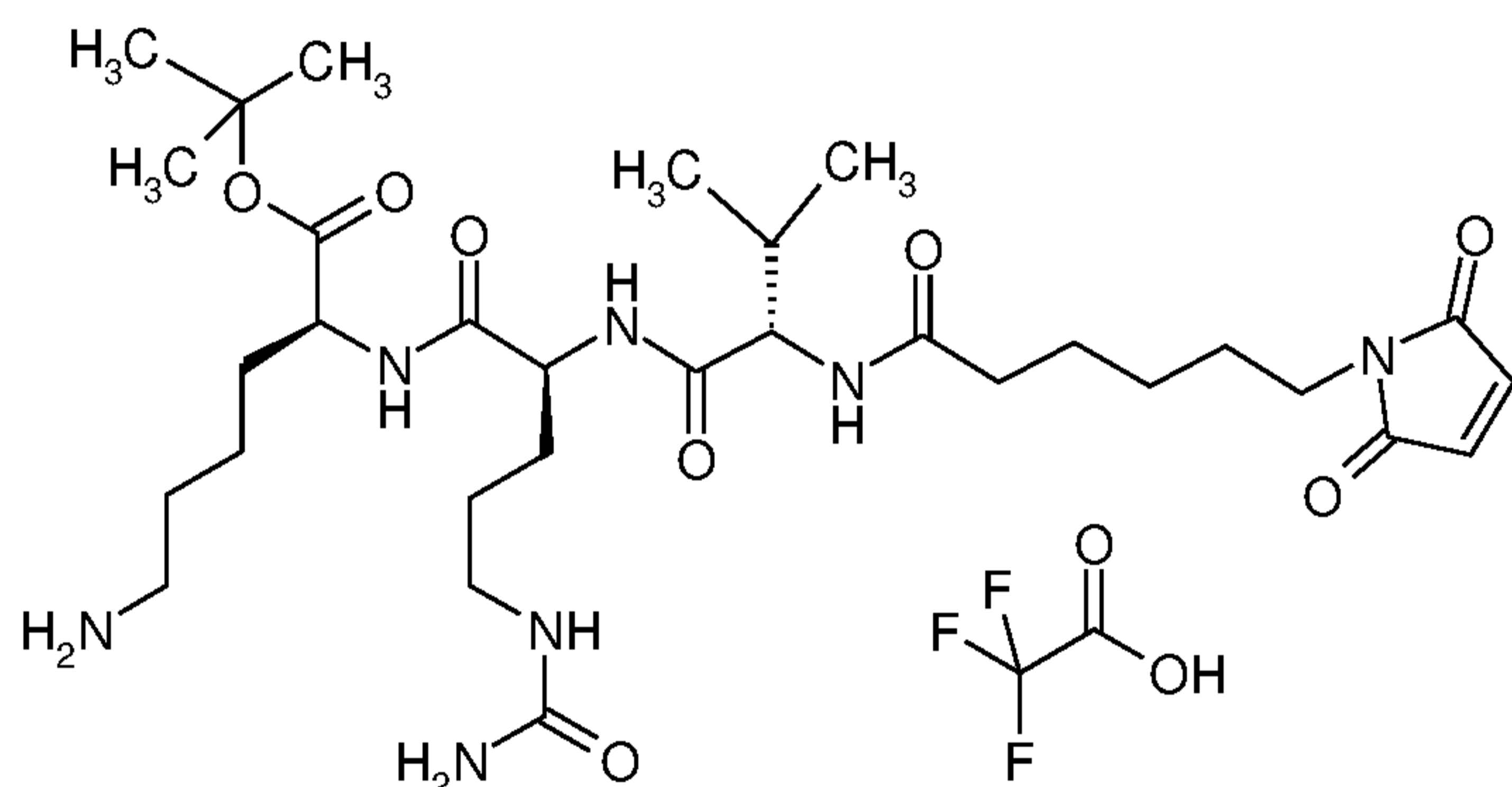
930 ml of *N,N*-diisopropylethylamine were added to a solution of 266 mg (1.33 mmol) of L-valyl-*N*<sup>5</sup>-carbamoyl-L-ornithine in 24 ml of DMF. The reaction was treated in an ultrasonic bath for 24 h and then concentrated to dryness under reduced pressure. The residue that remained was purified by preparative HPLC and gave, after concentration of the appropriate fractions and drying of the residue under high vacuum, 337 mg (50% of theory) of the title compound.

HPLC (Method 11):  $R_t = 0.4$  min;

LC-MS (Method 3):  $R_t = 0.58$  min; MS (ESIpos):  $m/z = 468$  (M+H)<sup>+</sup>.

### Intermediate L17

Trifluoroacetic acid / tert-butyl *N*-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-*N*<sup>5</sup>-carbamoyl-L-ornithyl-L-lysinate (1:1)



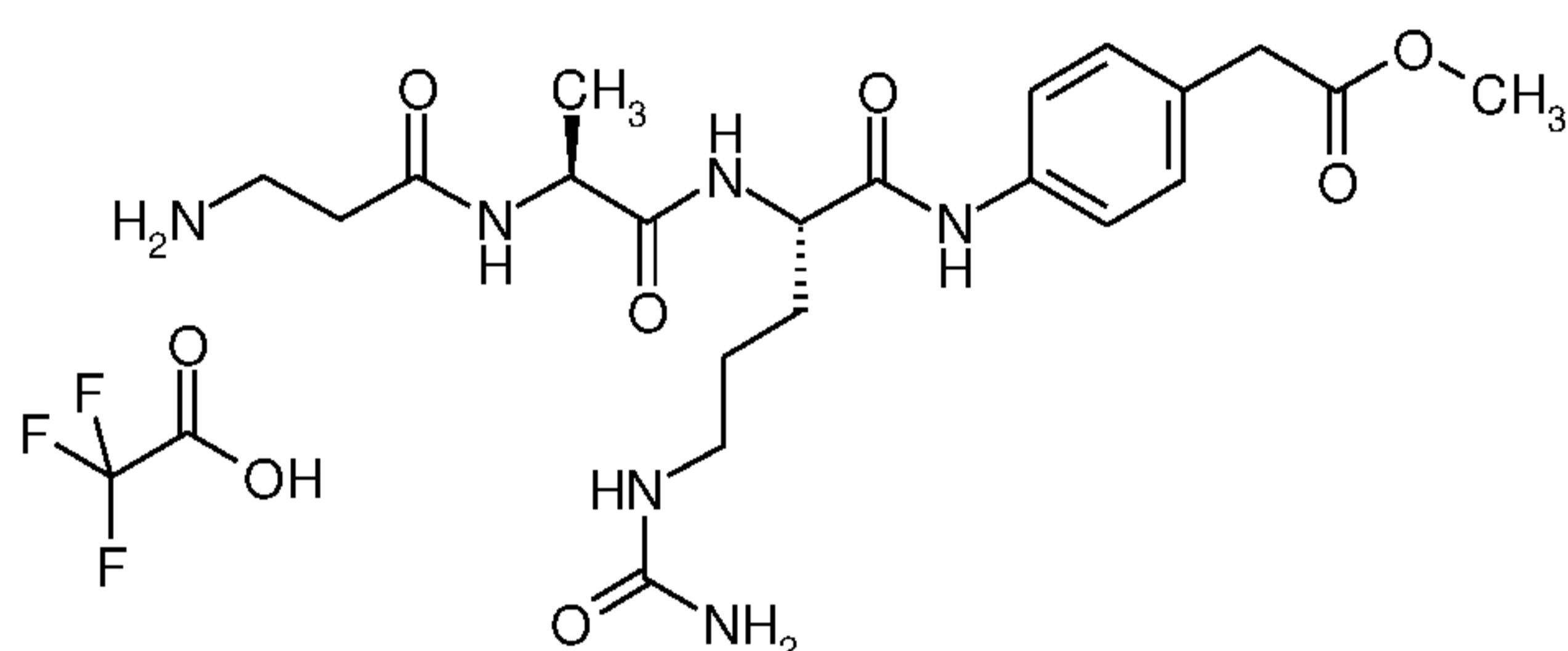
The title compound was prepared by initially coupling 172 mg (0.37 mmol) of Intermediate L16 and 125 mg (0.37 mmol) of tert-butyl *N*<sup>6</sup>-(tert-butoxycarbonyl)-L-lysinate hydrochloride (1:1) in the presence of EDC/HOBT and *N,N*-diisopropylethylamine and then deprotecting the amino group under gentle conditions by stirring for 2 h in 10% strength trifluoroacetic acid in DCM at RT. Freeze-drying from acetonitrile/water gave 194 mg (49% of theory) of the title compound over 2 steps.

HPLC (Method 11):  $R_t = 1.1$  min;

LC-MS (Method 1):  $R_t = 0.58$  min; MS (ESIpos):  $m/z = 652$  (M+H)<sup>+</sup>.

### Intermediate L18

- 5 Trifluoroacetic acid / beta-alanyl-L-alanyl-N<sup>5</sup>-carbamoyl-N-[4-(2-methoxy-2-oxoethyl)phenyl]-L-ornithinamide (1:1)



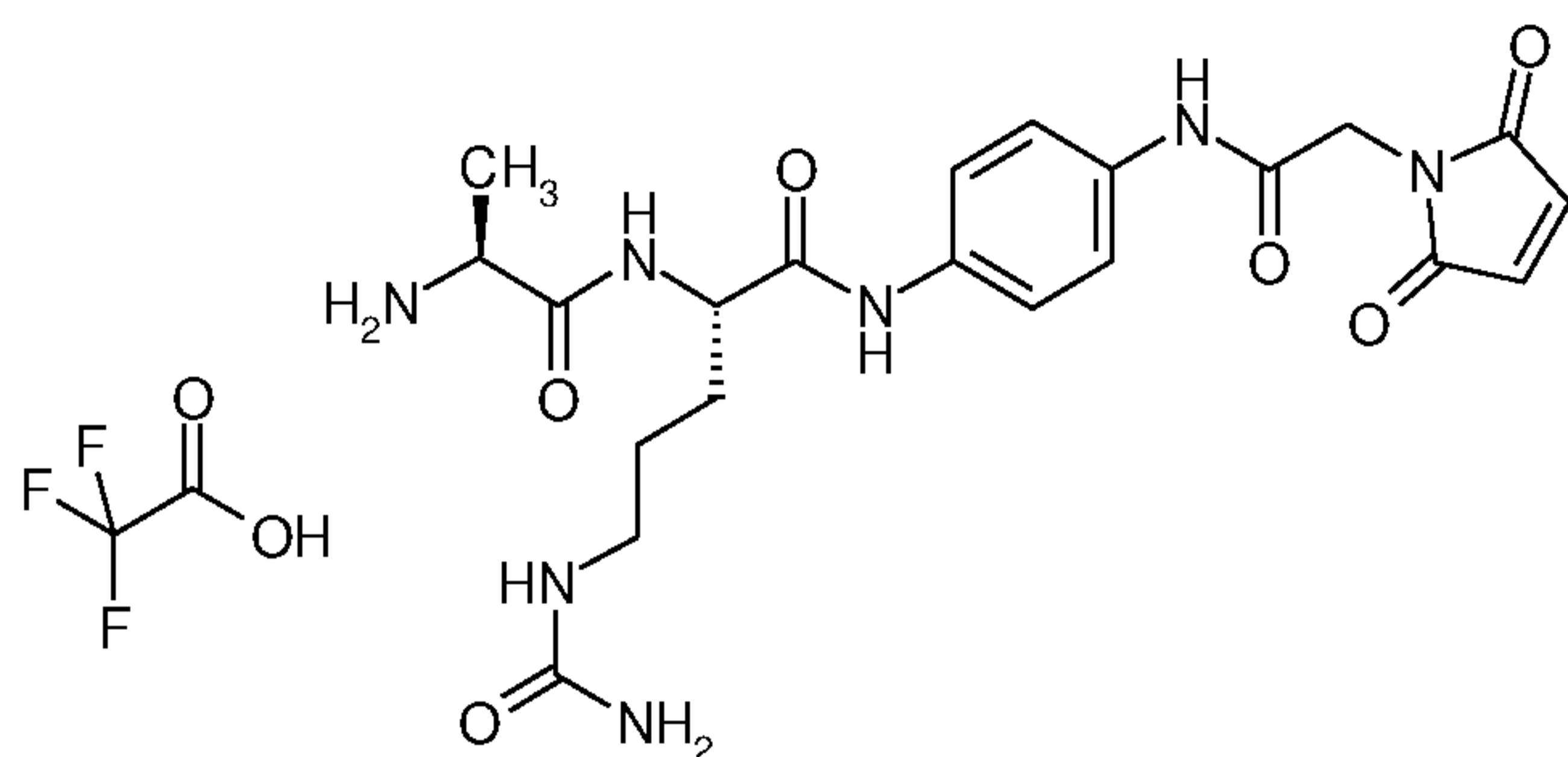
- 10 The title compound was prepared from methyl (4-aminophenyl)acetate analogously to Intermediate L7 sequentially according to classical methods of peptide chemistry by linking N<sup>2</sup>-(tert-butoxycarbonyl)-N<sup>5</sup>-carbamoyl-L-ornithine in the presence of HATU, deprotection with TFA, coupling with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-L-alaninate, deprotection with TFA, coupling with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-beta-alaninate and another deprotection with TFA. This gave 330 mg of the title compound.

- 20 HPLC (Method 11):  $R_t = 0.29$  min;

LC-MS (Method 1):  $R_t = 0.41$  min; MS (ESIpos):  $m/z = 465$  (M+H)<sup>+</sup>.

### Intermediate L19

- 25 Trifluoroacetic acid / L-alanyl-N<sup>5</sup>-carbamoyl-N-(4-{{(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl}amino}phenyl)-L-ornithinamide (1:1)



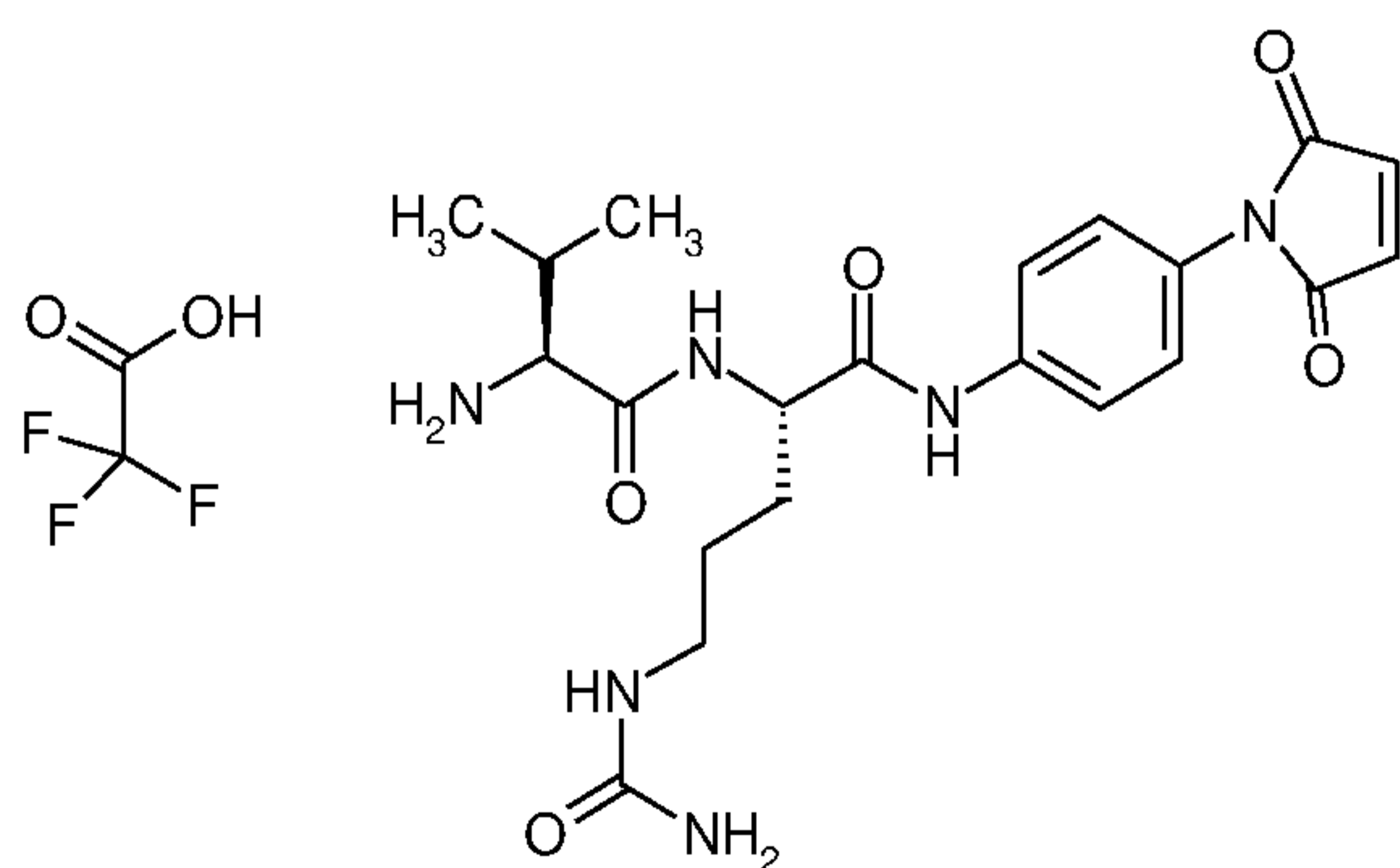
The title compound was prepared from 1,4-phenylenediamine sequentially according to classical methods of peptide chemistry. In the first step, 942 mg (8.72 mmol) of 1,4-phenylenediamine were monoacylated with 0.8 g (2.9 mmol) of N<sup>2</sup>-(tert-butoxycarbonyl)-N<sup>5</sup>-carbamoyl-L-ornithine in the presence of HATU and *N,N*-diisopropylethylamine. In the second step, in an analogous manner, the second anilinic amino group was acylated with (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetic acid in the presence of HATU and *N,N*-diisopropylethylamine. Deprotection with TFA, coupling with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-L-alaninate and another deprotection with TFA then gave, in 3 further synthesis steps, the title compound, 148 mg of which were obtained by this route.

LC-MS (Method 1):  $R_t = 0.21$  min; MS (ESIpos):  $m/z = 474$  (M+H)<sup>+</sup>.

LC-MS (Method 4):  $R_t = 0.2$  min; MS (ESIpos):  $m/z = 474$  (M+H)<sup>+</sup>.

### Intermediate L20

Trifluoroacetic acid / L-valyl-N<sup>5</sup>-carbamoyl-N-[4-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)phenyl]-L-ornithinamide (1:1)



The title compound was prepared according to classical methods of peptide chemistry analogously to Intermediate L8 from commercially available 1-(4-aminophenyl)-1H-pyrrole-2,5-dione by sequential coupling with N<sup>2</sup>-(tert-butoxycarbonyl)-N<sup>5</sup>-carbamoyl-L-ornithine in the presence of HATU, deprotection with TFA, coupling with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-L-valinate and another deprotection with TFA. This gave 171 mg of the title compound.

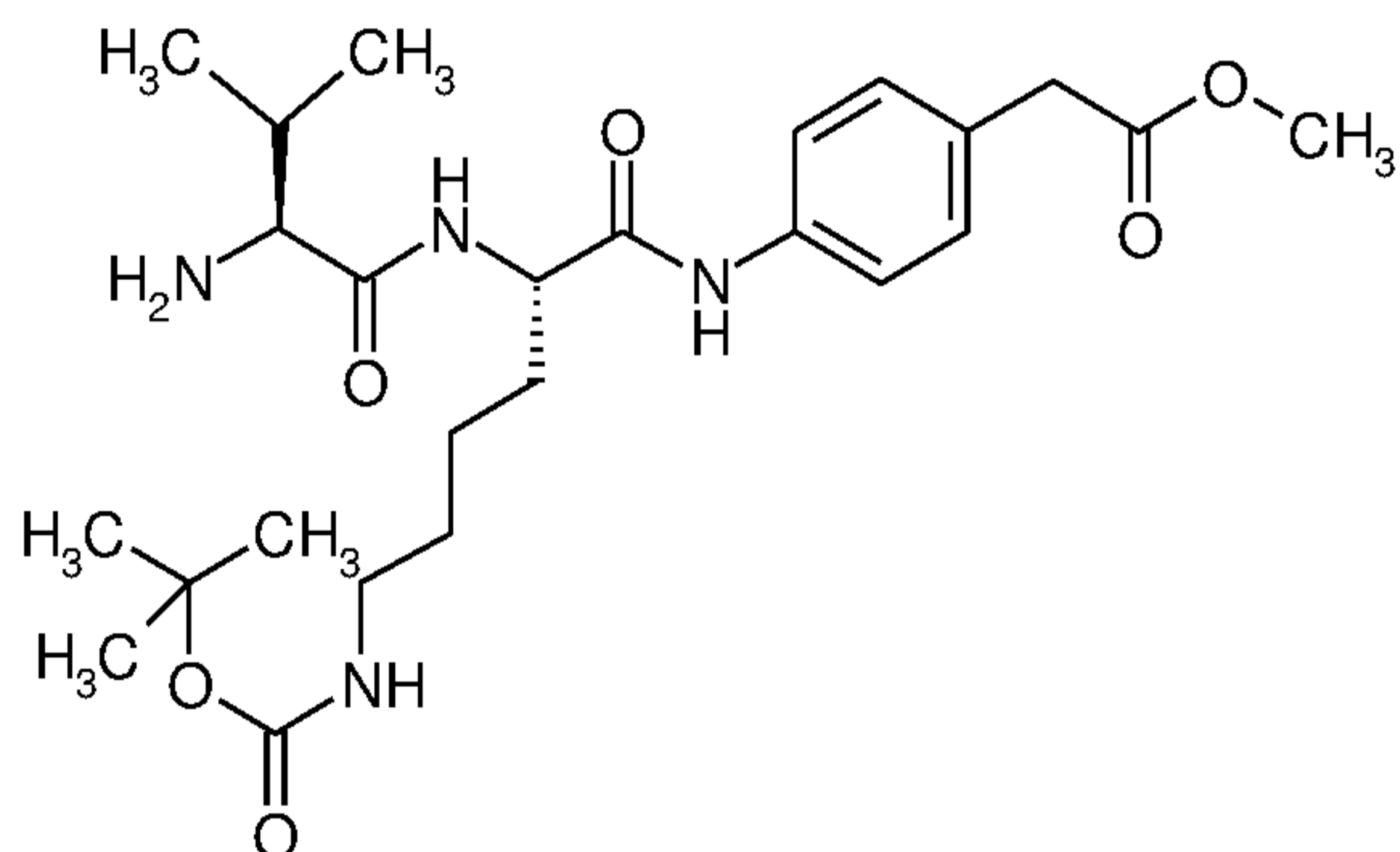
10 HPLC (Method 11): R<sub>t</sub> = 0.28 min;

LC-MS (Method 1): R<sub>t</sub> = 0.39 min; MS (ESIpos): m/z = 445 (M+H)<sup>+</sup>.

### Intermediate L21

15

L-Valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-N-[4-(2-methoxy-2-oxoethyl)phenyl]-L-lysineamide



20

The title compound was prepared according to classical methods of peptide chemistry from commercially available 0.42 g (2.56 mmol) of methyl (4-aminophenyl)acetate by sequential coupling with N<sup>6</sup>-(tert-butoxycarbonyl)-N<sup>2</sup>-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-lysine in the presence of HATU and *N,N*-diisopropylethylamine, deprotection with piperidine, coupling with 2,5-dioxopyrrolidin-1-yl N-[(benzyloxy)carbonyl]-L-valinate in the presence of *N,N*-diisopropylethylamine and subsequent hydrogenolytic removal of the benzyloxycarbonyl protective group over 10% palladium on activated carbon. This gave 360 mg (32% of theory over 4 steps) of the title compound.

30



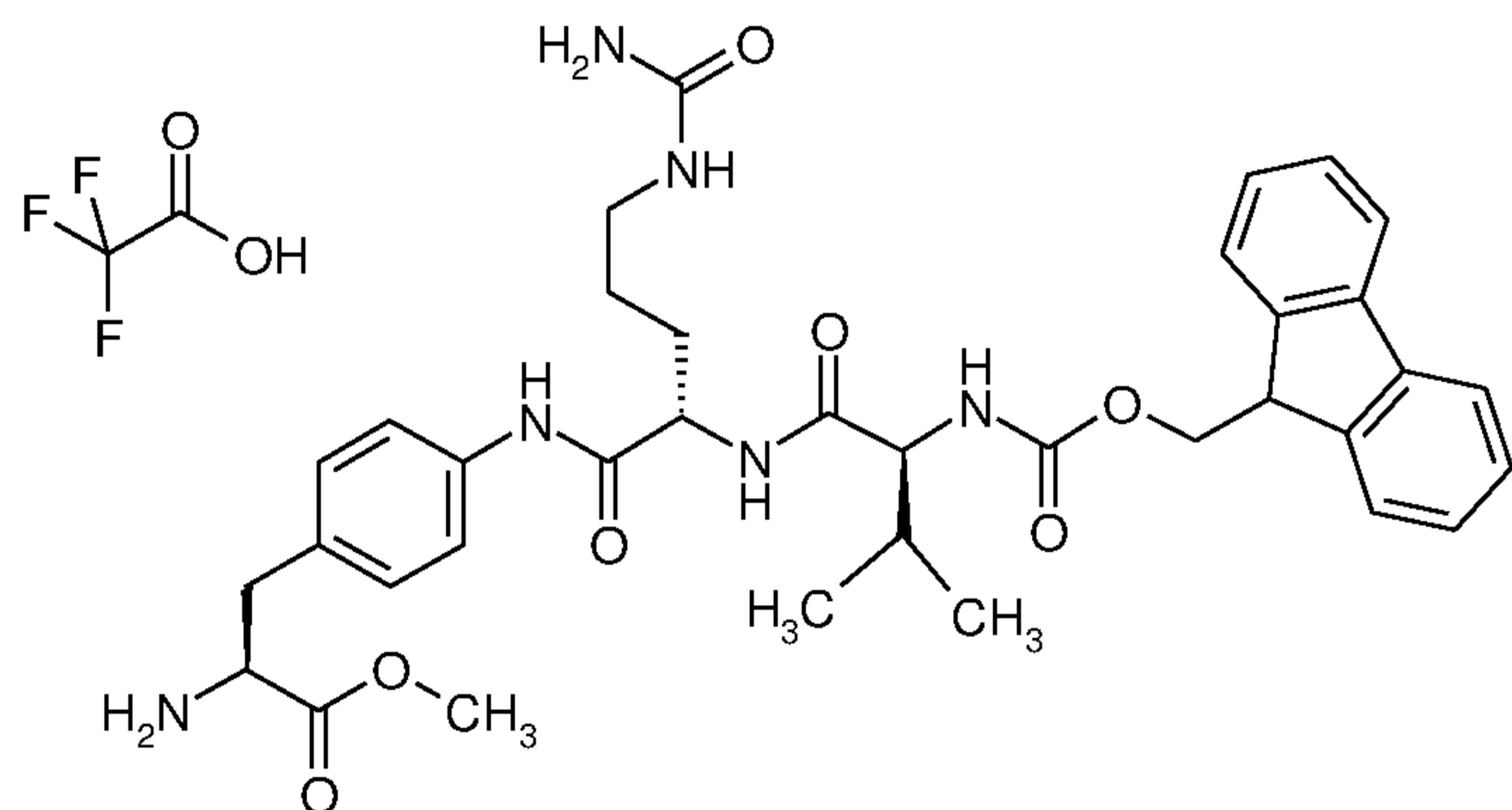
HPLC (Method 11):  $R_t = 1.5$  min;

LC-MS (Method 1):  $R_t = 0.73$  min; MS (ESIpos):  $m/z = 493$  (M+H)<sup>+</sup>.

## 5 Intermediate L22

Trifluoroacetic acid / N-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-valyl-N-{4-[(2S)-2-amino-3-methoxy-3-oxopropyl]phenyl}-N<sup>5</sup>-carbamoyl-L-ornithinamide (1:1)

10



The title compound was prepared from N-(tert-butoxycarbonyl)-4-nitro-L-phenylalanine sequentially according to classical methods of peptide chemistry. 2.5 g (8.06 mmol) of this starting material were in the first step initially converted into the caesium salt and then with iodomethane in DMF into the methyl ester.

20 Hydrogenolytically in methanol over 10% palladium on activated carbon, the nitro group was then converted into an amino group.

The amino group generated in this manner was then acylated with N<sup>5</sup>-carbamoyl-N<sup>2</sup>-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-ornithine in DMF in the presence of HATU and *N,N*-diisopropylethylamine. In the next step, the Fmoc group was removed with piperidine in DMF.

30 Coupling was then carried out in DMF with N-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-valine in the presence of 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide hydrochloride, 1-hydroxy-1*H*-benzotriazole hydrate and *N,N*-diisopropylethylamine

and finally removal of the tert-butoxycarbonyl group with trifluoroacetic acid.

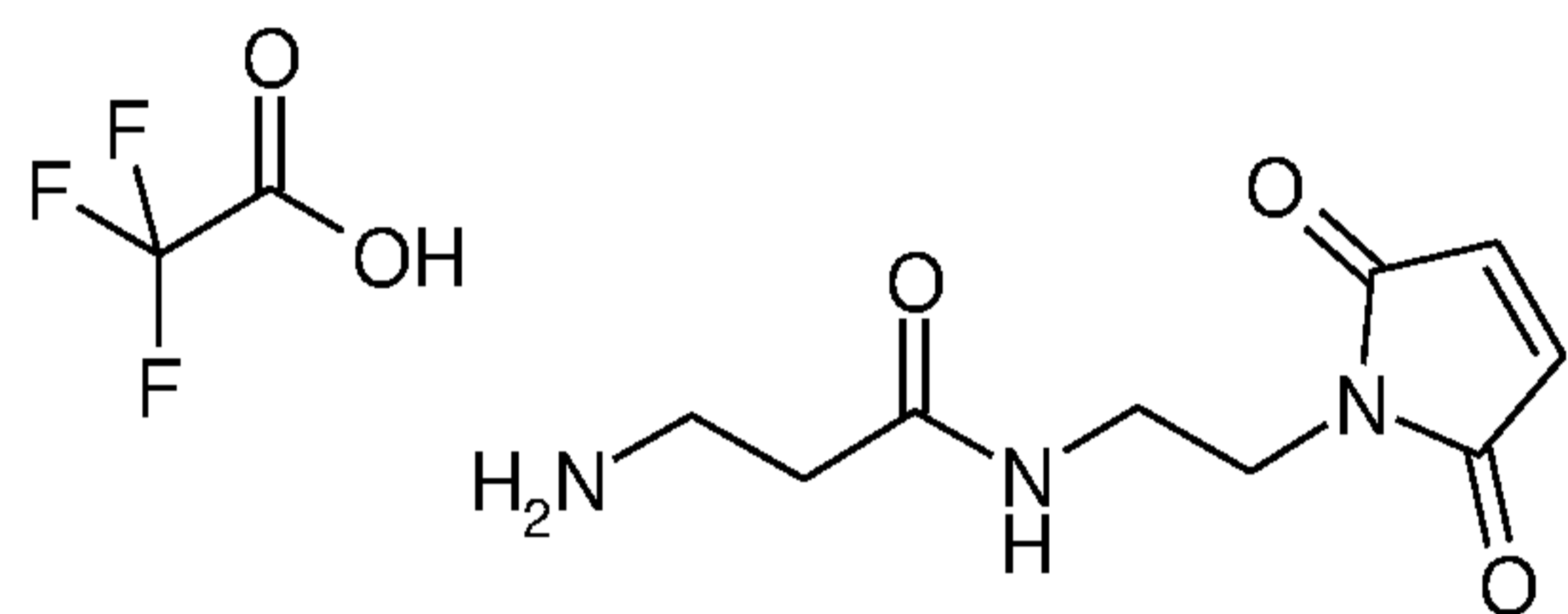
HPLC (Method 11):  $R_t = 1.6$  min;

5

LC-MS (Method 1):  $R_t = 0.77$  min; MS (ESIpos):  $m/z = 673$  (M+H)<sup>+</sup>.

### Intermediate L23

10 Trifluoroacetic acid / N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]-beta-alaninamide (1:1)



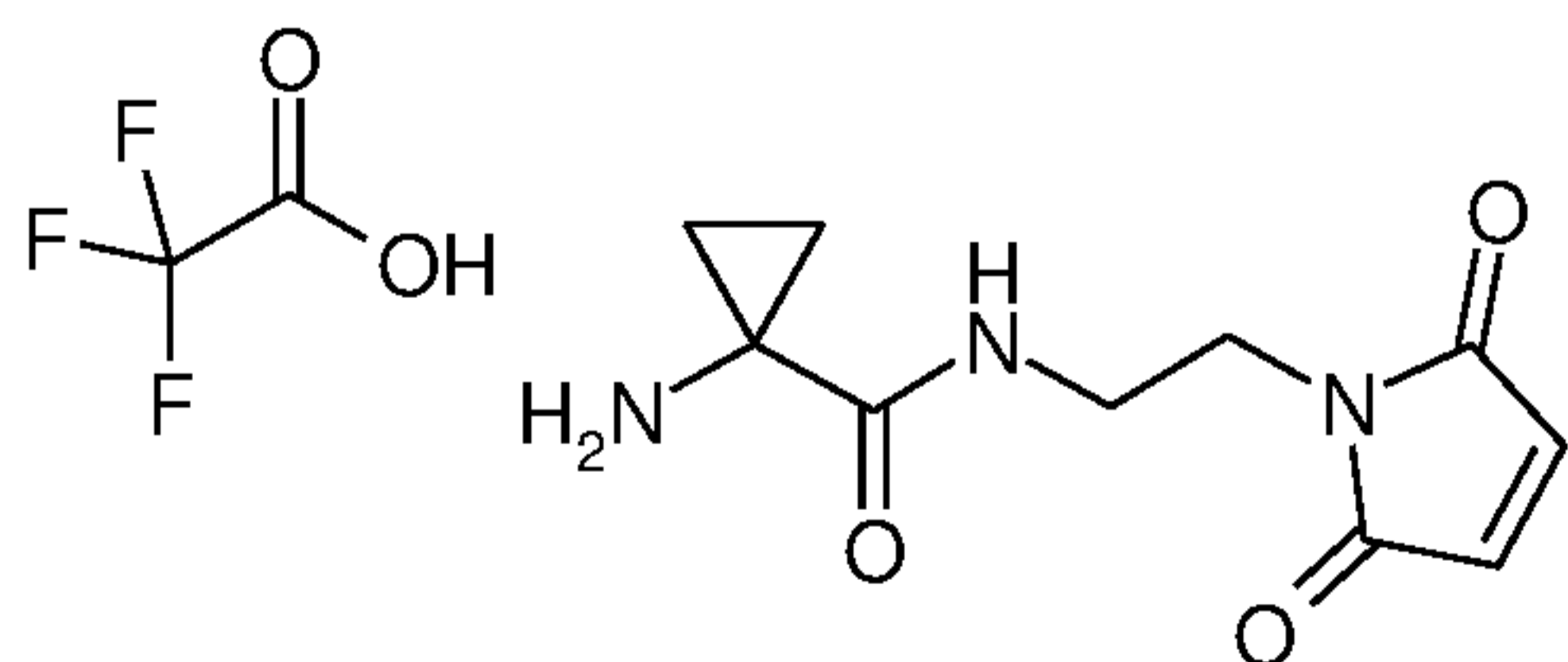
15 The title compound was prepared from commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) by coupling with N-(tert-butoxycarbonyl)-beta-alanine in the presence of EDCI/HOBT and *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid.

20

HPLC (Method 11):  $R_t = 0.19$  min.

### Intermediate L24

25 Trifluoroacetic acid / 1-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl] cyclopropanecarboxamide (1:1)



30 114 mg (0.67 mmol) of commercially available 1-[(tert-butoxycarbonyl)amino]cyclopropane-carboxylic acid were

dissolved in 25 ml of DCM, 110 mg (0.623 mmol) of commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) and 395  $\mu$ l of *N,N*-diisopropylethylamine were added and the mixture was cooled to  $-10^{\circ}\text{C}$ . 217 mg (0.793 mmol) of 2-bromo-1-ethylpyridinium tetrafluoroborate were then added, and the mixture was stirred at RT for 2 h. The mixture was then diluted with ethyl acetate and extracted successively with 10% strength citric acid, saturated sodium bicarbonate solution and saturated sodium chloride solution, then dried over magnesium sulphate and concentrated. Drying under high vacuum gave 152 mg of the protected intermediate.

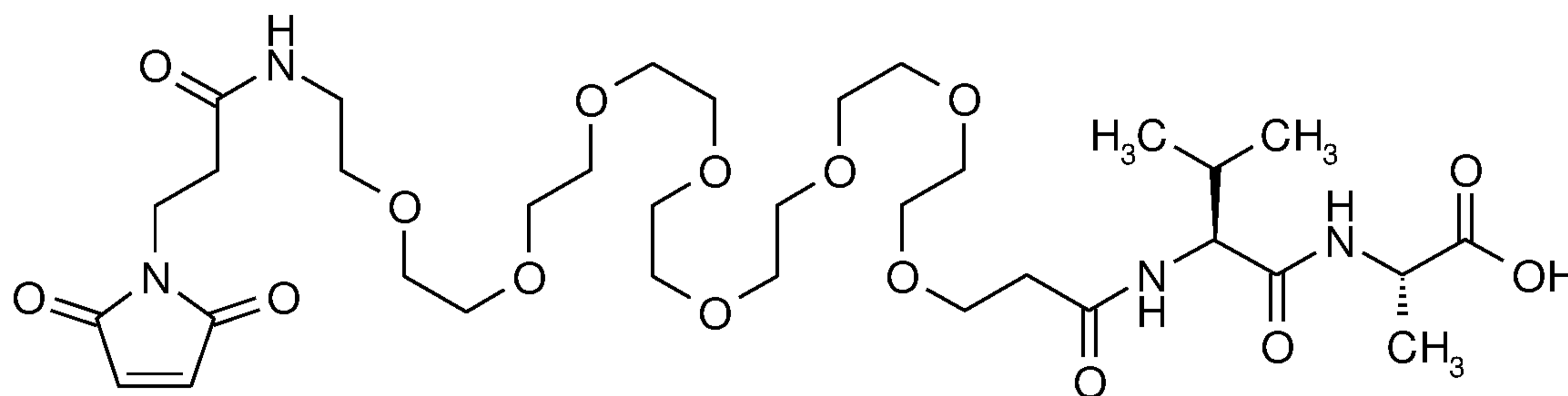
These were then taken up in 10 ml of DCM and deprotected with 1 ml of trifluoroacetic acid. Lyophilization from acetonitrile/water gave 158 mg (71% of theory over 2 steps) of the title compound.

HPLC (Method 11):  $R_t = 0.19$  min.

LC-MS (Method 3):  $R_t = 0.98$  min; MS (ESIpos):  $m/z = 224$  (M+H)<sup>+</sup>.

### Intermediate L25

*N*-[31-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-29-oxo-4,7,10,13,16,19,22,25-octaoxa-28-azahentriacontan-1-oyl]-L-valyl-L-alanine



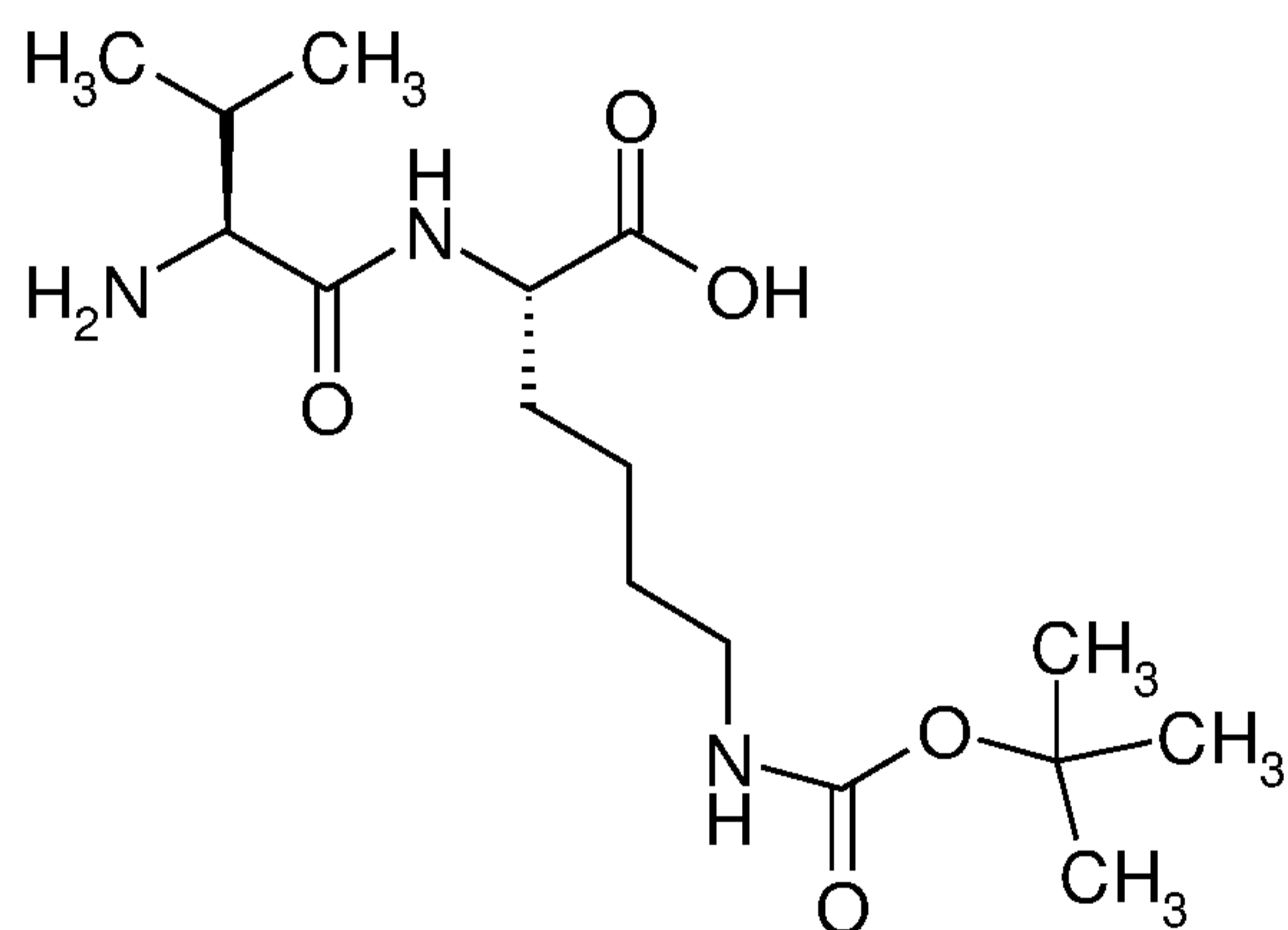
31.4 mg (0.17 mmol) of valyl-L-alanine were dissolved in 3.0 ml of DMF, and 115.0 mg (0.17 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-*N*-{27-[(2,5-dioxopyrrolidin-1-yl)oxy]-27-oxo-3,6,9,12,15,18,21,24-octaoxaheptacos-1-yl}propanamide and 33.7

mg (0.0.33 mmol) of triethylamine were added. The mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated  
5 under reduced pressure and the residue was dried under high vacuum. This gave 74.1 mg (58% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.61 min; MS (ESIpos):  $m/z$  = 763 [M+H]<sup>+</sup>.

## 10 Intermediate L26

L-Valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine



15

600.0 mg (1.58 mmol) of N<sup>2</sup>-[(benzyloxy)carbonyl]-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were suspended in 25.0 ml of water/ethanol/THF (1:1:0.5), palladium on carbon (10%) was added and the mixture was hydrogenated at RT with hydrogen under  
20 standard pressure for 5 h. The catalyst was filtered off and the solvents were evaporated under reduced pressure. The compound obtained was used in the next step without further purification.

LC-MS (Method 1):  $R_t$  = 0.42 min; MS (ESIpos):  $m/z$  = 247 [M+H]<sup>+</sup>.

25

180 mg (0.73 mmol) of N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were dissolved in 5.0 ml of DMF, and 74.0 mg (0.73 mmol) of triethylamine were added. 254.6 mg (0.73 mmol) of 2,5-dioxopyrrolidin-1-yl N-[(benzyloxy)carbonyl]-L-valinate and  
30 74.0 mg (0.73 mmol) of triethylamine were then added. The

reaction mixture was stirred at RT for 3.5 h. The reaction solution was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 294.1 mg (76% of theory) of N-[(benzyloxy)carbonyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine.

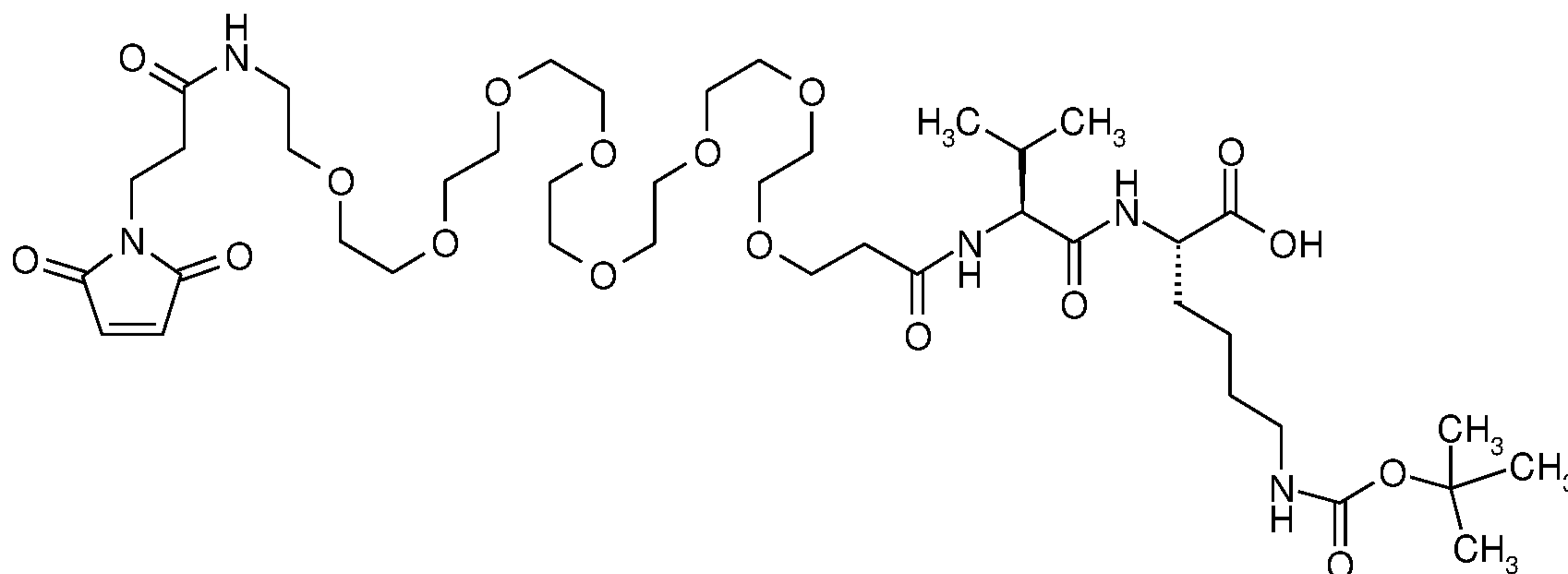
LC-MS (Method 1): R<sub>t</sub> = 0.97 min; MS (ESIpos): m/z = 480 [M+H]<sup>+</sup>.

272.2 mg (0.57 mmol) of N-[(benzyloxy)carbonyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were initially charged in 20.0 ml of ethyl acetate/ethanol/THF (1:1:1), and 27.2 mg of palladium on activated carbon were added. The mixture was hydrogenated with hydrogen at RT under standard pressure for 5 h. The mixture was filtered off with the aid of Celite<sup>(R)</sup> and the filter cake was washed with ethyl acetate/ethanol/THF (1:1:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. The title compound (182 mg, 72% of theory) was used in the next reaction step without further purification.

LC-MS (Method 1): R<sub>t</sub> = 0.53 min; MS (ESIpos): m/z = 346 [M+H]<sup>+</sup>.

### 25 Intermediate L27

N-[31-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-29-oxo-4,7,10,13,16,19,22,25-octaoxa-28-azahentriacontan-1-oyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine



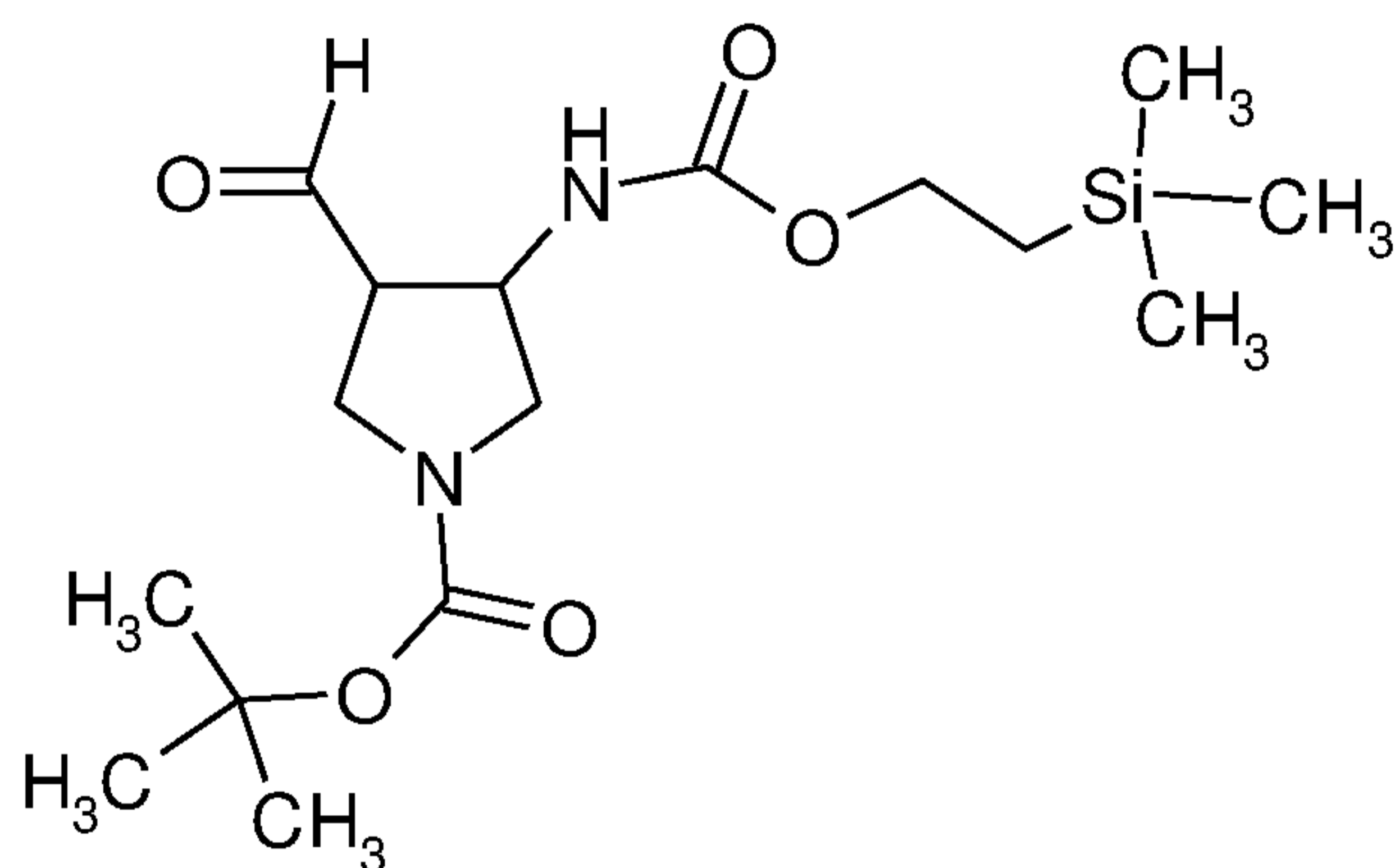
30 mg (0.07 mmol) of L-valyl-N6-(tert-butoxycarbonyl)-L-lysine (Intermediate L26) and 46.1 mg (0.07 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{27-[(2,5-dioxopyrrolidin-1-yl)oxy]-27-oxo-3,6,9,12,15,18,21,24-octaoxaheptacos-1-yl}propanamide were initially charged in 1.5 ml of DMF, and 6.8 mg (0.07 mmol) of 4-methylmorpholine were added. The reaction solution was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 55.6 mg (90% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.77 min; MS (ESIpos):  $m/z$  = 920 [M+H]<sup>+</sup>.

15

### Intermediate L28

tert-Butyl 3-formyl-4-({[2-(trimethylsilyl)ethoxy]carbonyl}amino)pyrrolidine-1-carboxylate



461.7 mg (1.15 mmol) of 1-tert-butyl 3-ethyl-4-({[2-(trimethylsilyl)ethoxy]carbonyl}amino)pyrrolidine-1,3-dicarboxylate (this compound was prepared according to the literature procedure of WO 2006/066896) were initially charged in 5.0 ml of absolute dichloromethane and the mixture was cooled to -78°C. 326.2 mg (2.29 mmol) of diisobutylaluminum hydride solution (1 M in THF) were then slowly added dropwise and the mixture was stirred at -78°C for 2 h (monitored by thin-layer

30

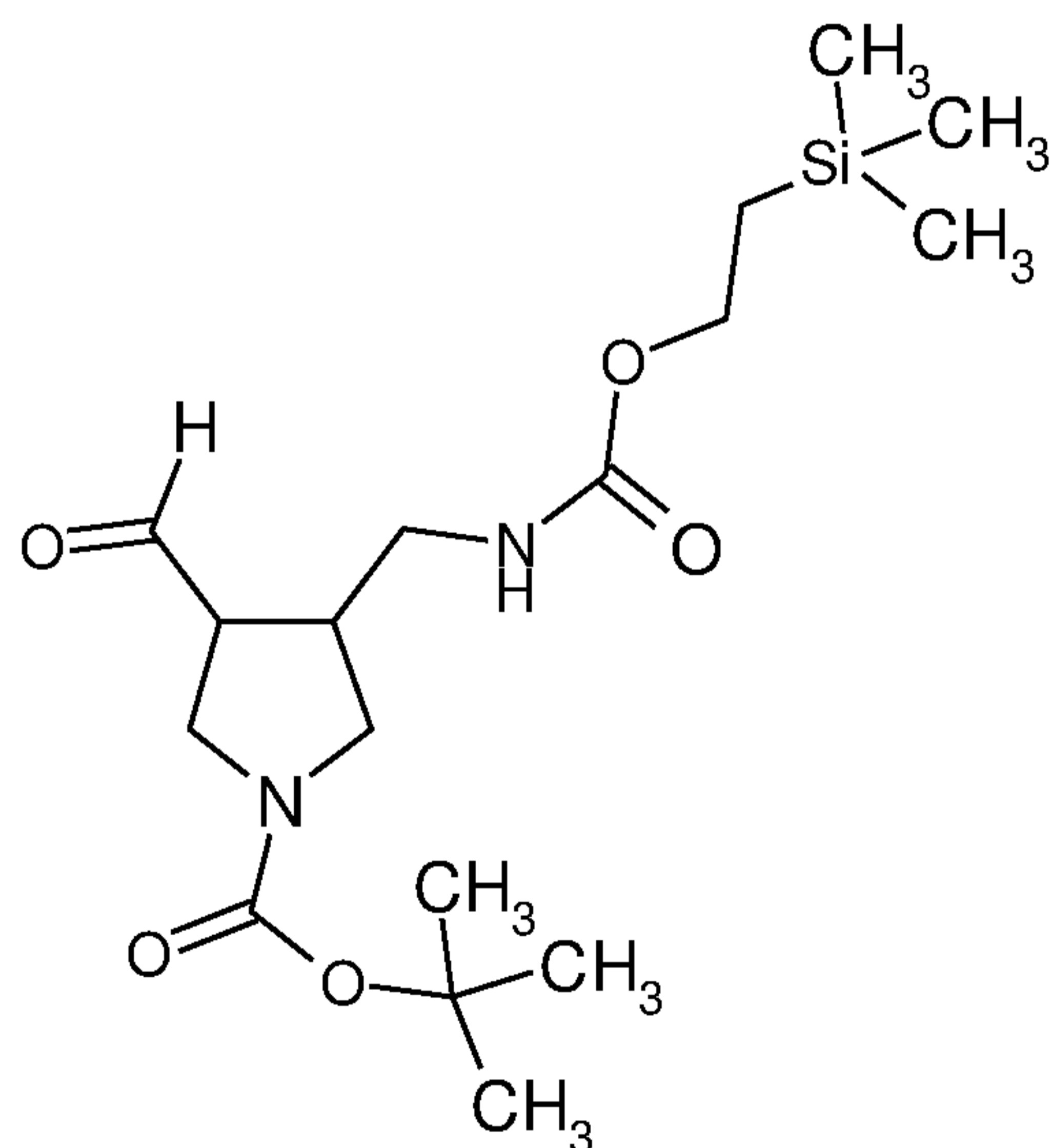
chromatography (petroleum ether/ethyl acetate = 3:1). 1.3 g (4.59 mmol) of potassium sodium tartrate dissolved in 60 ml of water were added dropwise and the reaction mixture was allowed to warm to RT. Ethyl acetate was added to the reaction mixture and the aqueous phase was extracted three times with ethyl acetate. The combined organic phases were washed once with sat. NaCl solution and dried over magnesium sulphate. The solvent was evaporated under reduced pressure and the residue was dried under high vacuum. This gave 629.0 mg of the title compound as a crude product which was used immediately without further purification in the next reaction step.

### Intermediate L29

15 tert-Butyl 3-formyl-4-[(2-(trimethylsilyl)ethoxy)carbonyl]amino)methylpyrrolidine-1-carboxylate

Mixture of diastereomers.

20



807.1 mg (2.34 mmol) of tert-butyl 3-([tert-butyl(dimethyl)silyl]oxy)methyl)-4-(hydroxymethyl)pyrrolidine-1-carboxylate (prepared according to the literature procedure of WO 2006/100036) were initially charged in 8.0 ml of dichloromethane, and 236.4 mg (2.34 mmol) of triethylamine were added. At 0°C, 267.6 mg (2.34 mmol) of methanesulphonyl chloride

were added dropwise, and the reaction mixture stirred at RT overnight. A further 133.8 mg (1.17 mmol) of methanesulphonyl chloride and 118.2 mg (1.17 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight. The mixture  
5 was diluted with dichloromethane and the organic phase was washed in each case once with saturated sodium bicarbonate solution, 5% strength potassium hydrogen sulphate solution and saturated NaCl solution. After drying over magnesium sulphate, the solvent was evaporated under reduced pressure and the  
10 residue was purified on Biotage Isolera (silica gel, column 50 g SNAP, flow rate 66 ml/min, cyclohexane/ethyl acetate). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 402.0 mg (41% of theory) of the compound tert-butyl 3-({[tert-  
15 butyl(dimethyl)silyloxy}methyl)-4-  
{[(methanesulphonyl)oxy]methyl}pyrrolidine-1-carboxylate.

LC-MS (Method 1):  $R_t = 1.38$  min; MS (ESIpos):  $m/z = 424$  [M+H]<sup>+</sup>.

20 400.0 mg (0.94 mmol) of tert-butyl 3-({[tert-  
butyl(dimethyl)silyloxy}methyl)-4-  
{[(methanesulphonyl)oxy]methyl}pyrrolidine-1-carboxylate were initially charged in 5.0 ml of DMF, and 98.2 mg (1.51 mmol) of sodium azide were added. The reaction mixture was stirred at  
25 40°C for 10 h. Another 30.7 mg (0.47 mmol) of sodium azide were then added, and the mixture was stirred at 40°C for a further 10 h. Ethyl acetate was added and the organic phase was washed repeatedly with water. After drying of the organic phase over magnesium sulphate, the solvent was evaporated under reduced  
30 pressure and the residue was dried under high vacuum. This gave 309.5 mg (89% of theory) of the compound tert-butyl 3-  
(azidomethyl)-4-({[tert-  
butyl(dimethyl)silyloxy}methyl)pyrrolidine-1-carboxylate. The compound was used without further purification in the next step  
35 of the synthesis.

LC-MS (Method 1):  $R_t = 1.50$  min; MS (ESIpos):  $m/z = 371$  [M+H]<sup>+</sup>.



250 mg (0.68 mmol) of tert-butyl 3-(azidomethyl)-4-([tert-butyl(dimethyl)silyloxy]methyl)pyrrolidine-1-carboxylate were dissolved in 10.0 ml of ethyl acetate/ethanol (1:1), and 25.0 mg of palladium on activated carbon (10%) were added. The mixture was hydrogenated with hydrogen at RT under standard pressure for 8 h. The reaction was filtered through Celite<sup>(R)</sup> and the filter cake was washed thoroughly with ethyl acetate. The solvent was evaporated under reduced pressure and the residue was dried under high vacuum. This gave 226.2 mg (82% of theory) of the compound tert-butyl 3-(aminomethyl)-4-([tert-butyl(dimethyl)silyloxy]methyl)pyrrolidine-1-carboxylate. The compound was used without further purification in the next step of the synthesis.

LC-MS (Method 1):  $R_t = 0.89$  min; MS (ESIpos):  $m/z = 345$  [M+H]<sup>+</sup>.

715.0 mg (2.08 mmol) of tert-butyl 3-(aminomethyl)-4-([tert-butyl(dimethyl)silyloxy]methyl)pyrrolidine-1-carboxylate were dissolved in 15.0 ml of THF, and 2.28 ml (2.28 mmol) of TBAF solution (1M in THF) were added. The reaction mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and the residue (1.54 g) used without further purification in the next step of the synthesis.

LC-MS (Method 1):  $R_t = 0.41$  min; MS (ESIpos):  $m/z = 231$  [M+H]<sup>+</sup>.

1.54 g (4.88 mmol) of tert-butyl 3-(aminomethyl)-4-(hydroxymethyl)pyrrolidine-1-carboxylate were initially charged in 1,4-dioxane, and 541.8 mg (4.88 mmol) of calcium chloride (anhydrous) and 488.6 mg (4.88 mmol) of calcium carbonate were added and the mixture was stirred vigorously. 592.8 mg (5.86 mmol) of triethylamine and 1.52 g (5.86 mmol) of 1-([2-(trimethylsilyl)ethoxy]carbonyl)oxy)pyrrolidine-2,5-dione were then added and the reaction mixture stirred at RT overnight. 644.9 mg (10.7 mmol) of HOAc and ethyl acetate were added. The organic phase was washed twice with water and once with saturated NaCl solution. After drying over magnesium sulphate, the solvent was evaporated under reduced pressure and the residue was

purified on silica gel (mobile phase: dichloromethane/methanol = 100:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 346.9 mg (19% of theory) of the compound tert-butyl 3-(hydroxymethyl)-4-  
5 [({[2-(trimethylsilyl)ethoxy]carbonyl}amino)methyl]pyrrolidine-1-carboxylate.

LC-MS (Method 1):  $R_t = 1.08$  min; MS (ESIpos):  $m/z = 375$  [M+H]<sup>+</sup>.

10

804.0 mg (2.15 mmol) of tert-butyl 3-(hydroxymethyl)-4-[(2-(trimethylsilyl)ethoxy)carbonyl]amino)methyl]pyrrolidine-1-carboxylate were initially charged in 20.0 ml of chloroform and 20.0 ml of 0.05 N potassium carbonate/0.05 N sodium bicarbonate  
15 solution (1:1). 59.7 mg (0.22 mmol) of tetra-n-butylammonium chloride, 429.9 mg (3.22 mmol) of N-chlorosuccinimide and 33.5 mg (0.22 mmol) of TEMPO were then added and the reaction mixture was stirred vigorously at RT overnight. The organic phase was separated off and freed from the solvent under reduced pressure.  
20 The residue was purified on silica gel (mobile phase: cyclohexane/ethyl acetate = 3:1). This gave 517.0 mg (46% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 1.13$  min; MS (ESIpos):  $m/z = 373$  [M+H]<sup>+</sup>.

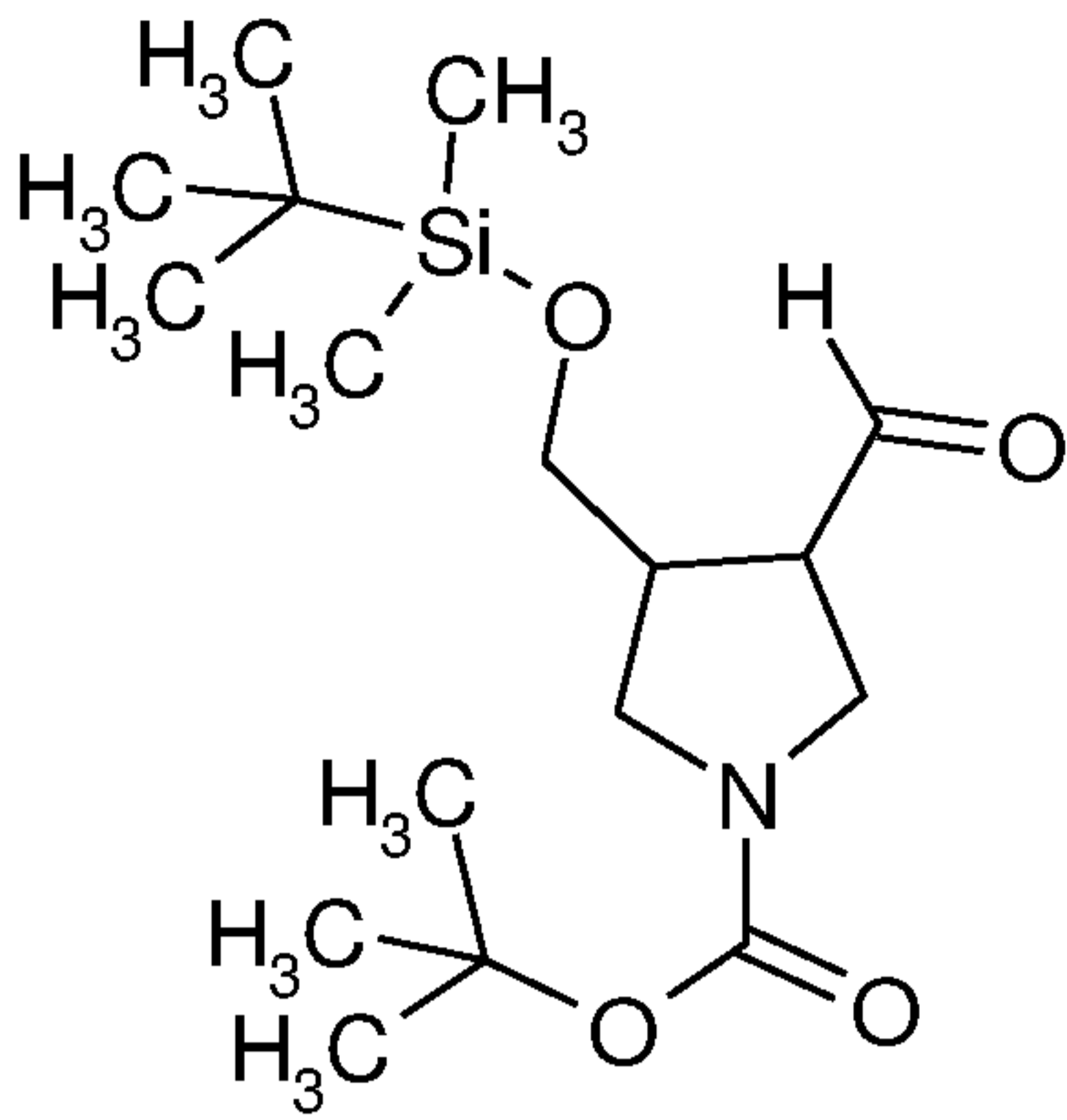
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### Intermediate L30

tert-Butyl 3-({[tert-butyl(dimethyl)silyl]oxy}methyl)-4-formylpyrrolidine-1-carboxylate

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Mixture of stereoisomers

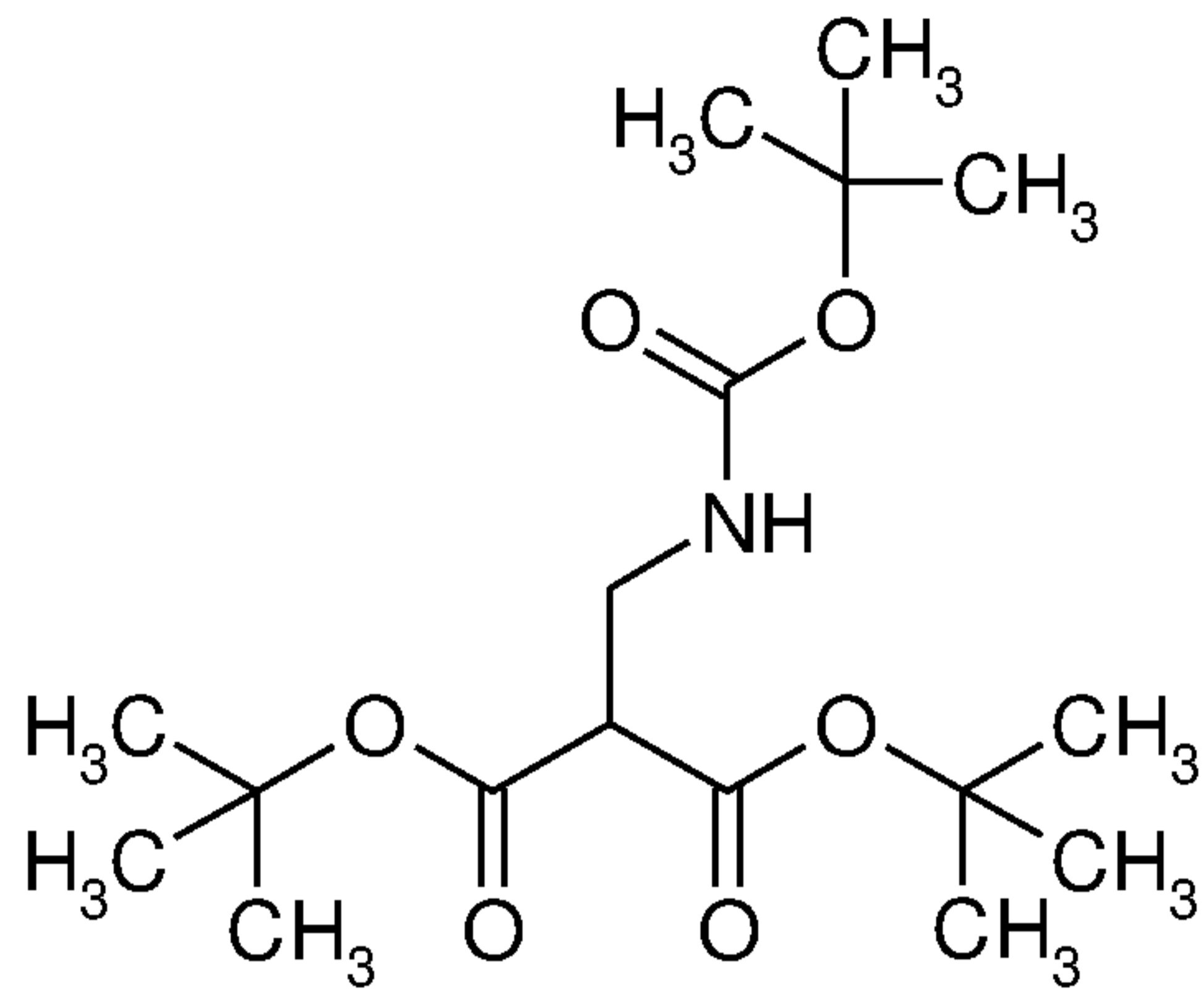


250.0 mg (0.72 mmol) of tert-butyl 3-((tert-butyl(dimethyl)silyloxy)methyl)-4-(hydroxymethyl)pyrrolidine-1-carboxylate (the compound was prepared according to the literature procedure of WO2006/100036) were initially charged in 12.5 ml of dichloromethane/DMSO (4:1), and 219.6 mg (2.17 mmol) of triethylamine were added. At 2°C, 345.5 mg (2.17 mmol) of sulphur trioxide-pyridine complex were added a little at a time and the mixture was stirred at 2°C for 3 h. Another 345.5 mg (2.17 mmol) of sulphur trioxide-pyridine complex were added a little at a time and the mixture was stirred at RT for 17 h. The reaction mixture was partitioned between dichloromethane and water. The aqueous phase was extracted three times with dichloromethane and the combined organic phases were washed once with water and dried over magnesium sulphate. The solvent was evaporated under reduced pressure and the residue was dried under high vacuum. The residue was used without further purification in the next step of the synthesis (thin-layer chromatography: petroleum ether/ethyl acetate 7:3).

### Intermediate L31

Di-tert-butyl {[(tert-butoxycarbonyl)amino]methyl}malonate

25



57.2 g (488.27 mmol) of tert-butyl carbamate, 51.2 ml (683.57 mmol) of a 37% strength solution of formaldehyde in water and  
5 25.9 g (244.13 mmol) of sodium carbonate were added to 600 ml of water. The mixture was warmed until a solution was formed and then stirred at RT for 16 h. The suspension formed was extracted with 500 ml of dichloromethane and the organic phase was separated off, washed with saturated sodium chloride solution and dried over sodium sulphate. The mixture was concentrated on  
10 a rotary evaporator and the residue was dried under high vacuum, giving a crystalline solid. The residue was taken up in 1000 ml of absolute THF, and a mixture of 322 ml (3.414 mol) of acetic anhydride and 138 ml (1.707 mol) of pyridine was added dropwise  
15 at RT. The reaction mixture was stirred at RT for 16 h and then concentrated on a rotary evaporator, with the water bath at room temperature. The residue was taken up in diethyl ether and washed three times with a saturated sodium bicarbonate solution and once with a saturated sodium chloride solution. The organic  
20 phase was dried over sodium sulphate and concentrated on a rotary evaporator and the residue was dried under high vacuum for 2 d. The residue was taken up in 2000 ml of absolute THF, and 456 ml (456.52 mmol) of a 1 M solution of potassium tert-butoxide in THF were added with ice cooling. The mixture was stirred at 0°C  
25 for 20 min, and 100.8 g (456.52 mmol) of di-tert-butyl malonate dissolved in 200 ml of absolute THF were then added dropwise. The mixture was stirred at RT for 48 h, and water was then added. The reaction mixture was concentrated on a rotary evaporator and taken up in 500 ml of ethyl acetate. The mixture was washed with  
30 500 ml of water and 100 ml of a saturated sodium chloride solution and the organic phase was dried over sodium sulphate.

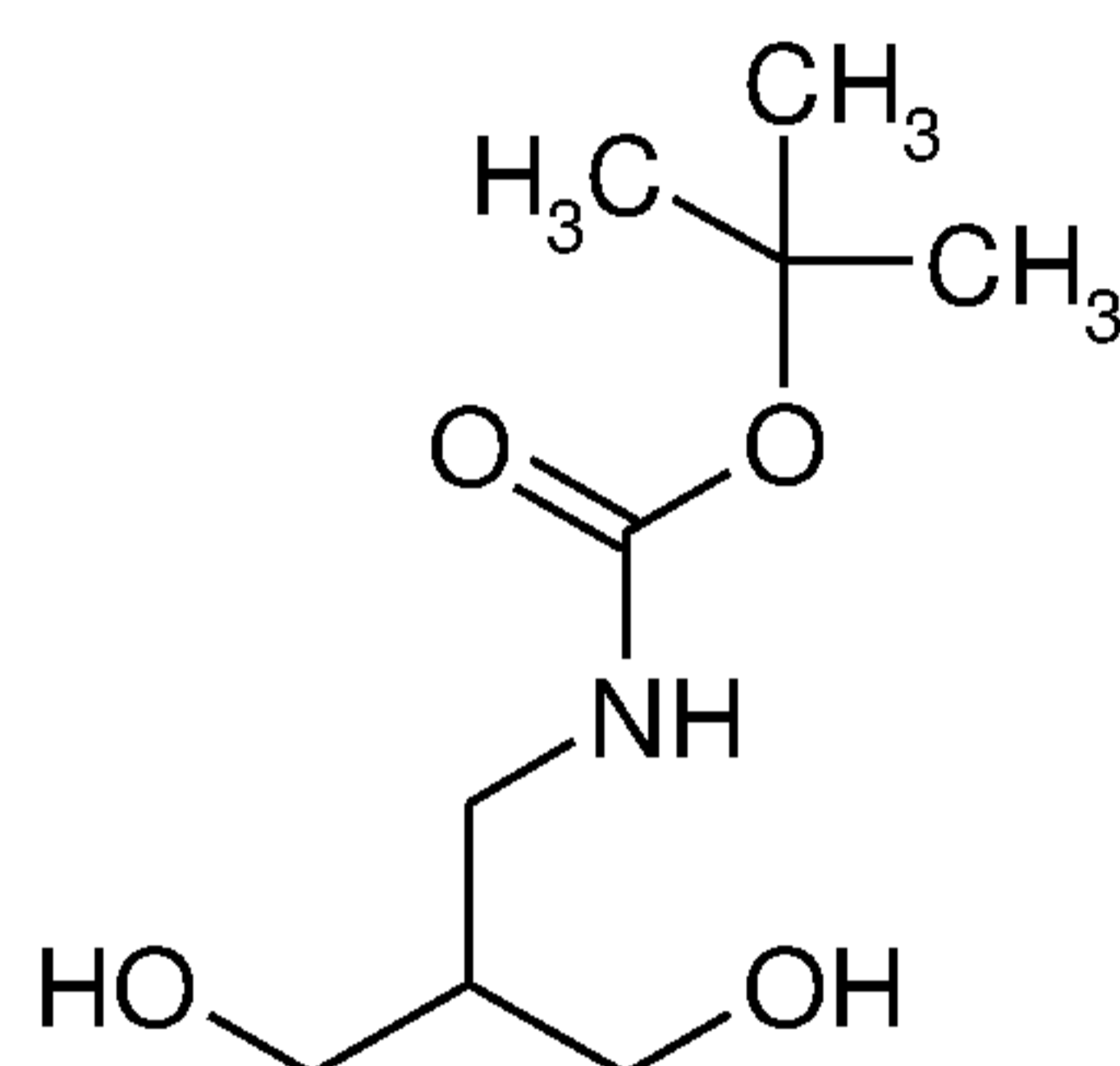
The organic phase was concentrated on a rotary evaporator and the residue was dried under high vacuum. The residue was purified by filtration through silica gel (mobile phase: cyclohexane/ethyl acetate, gradient = 30:1 → 5:1). This gave  
5 37.07 g (22% of theory) of the target compound.

LC-MS (Method 6):  $R_t = 2.87$  min; MS (ESIpos):  $m/z = 346$  [M+H]<sup>+</sup>.

### Intermediate L32

10

tert-Butyl [3-hydroxy-2-(hydroxymethyl)propyl]carbamate

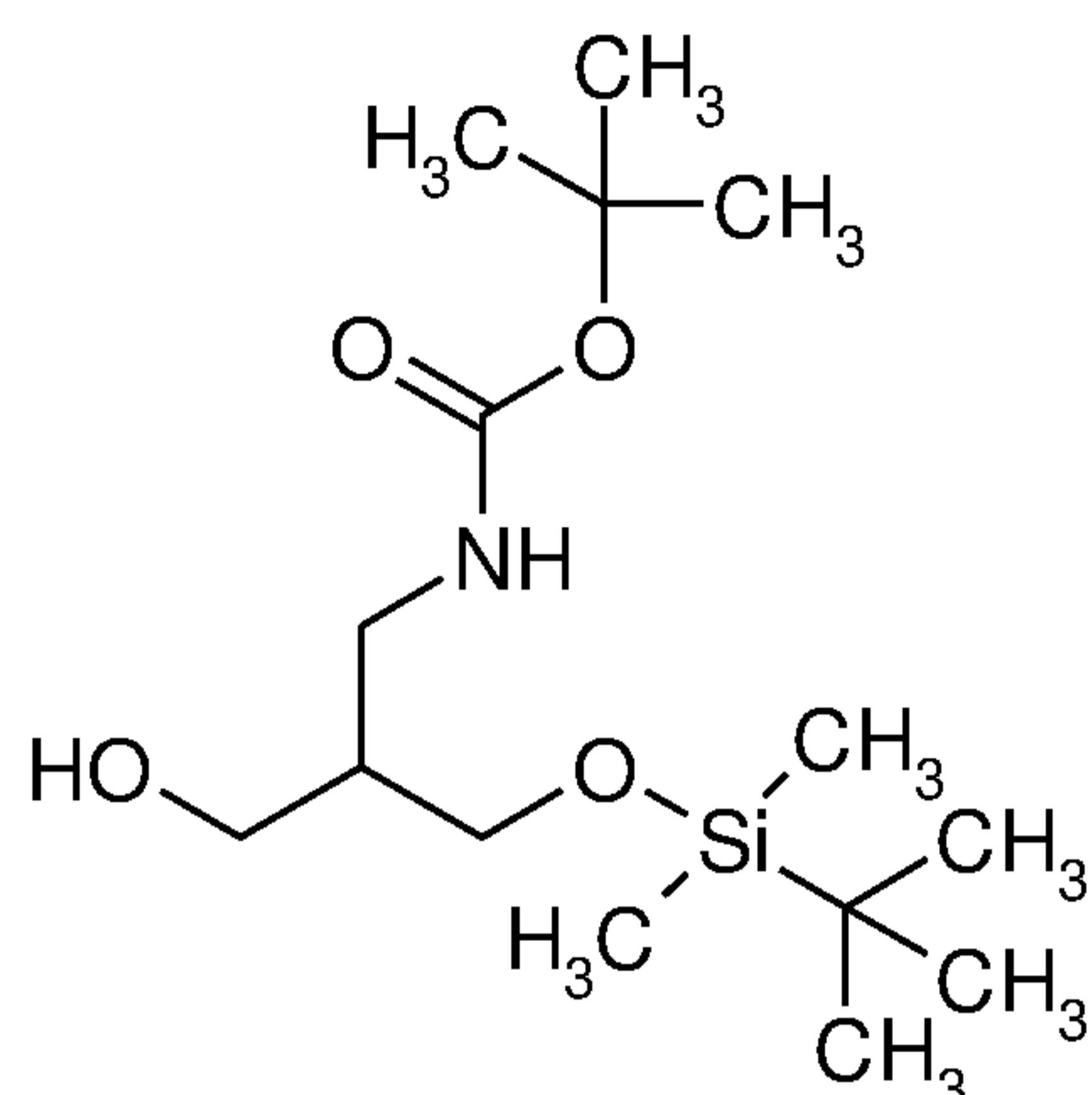


15 37.0 g (107.11 mmol) of di-tert-butyl (acetoxymethyl)malonate were dissolved in 1000 ml of absolute THF, and 535.5 ml (1071.10 mmol) of a 2 M solution of lithium borohydride in THF were added dropwise with ice cooling. 19.3 ml (1071.10 mmol) of water were added dropwise and the mixture was stirred at RT for 4.5 h. The  
20 reaction mixture was concentrated on a rotary evaporator and dried under high vacuum. The residue was taken up in 1500 ml of ethyl acetate, 100 ml of water were added and the mixture was stirred with water cooling (slightly exothermic) for 30 min. The organic phase was separated off and the aqueous phase was  
25 extracted twice with 500 ml of ethyl acetate. The organic phase was concentrated on a rotary evaporator and the residue was dried under high vacuum. This gave 20.7 g (94% of theory) of the target compound.

30 LC-MS (Method 6):  $R_t = 1.49$  min; MS (EIpos):  $m/z = 106$  [M-C<sub>5</sub>H<sub>8</sub>O<sub>2</sub>]<sup>+</sup>.

### Intermediate L33

tert-Butyl [3-{{tert-butyl(dimethyl)silyl}oxy}-2-(hydroxymethyl)propyl]carbamate



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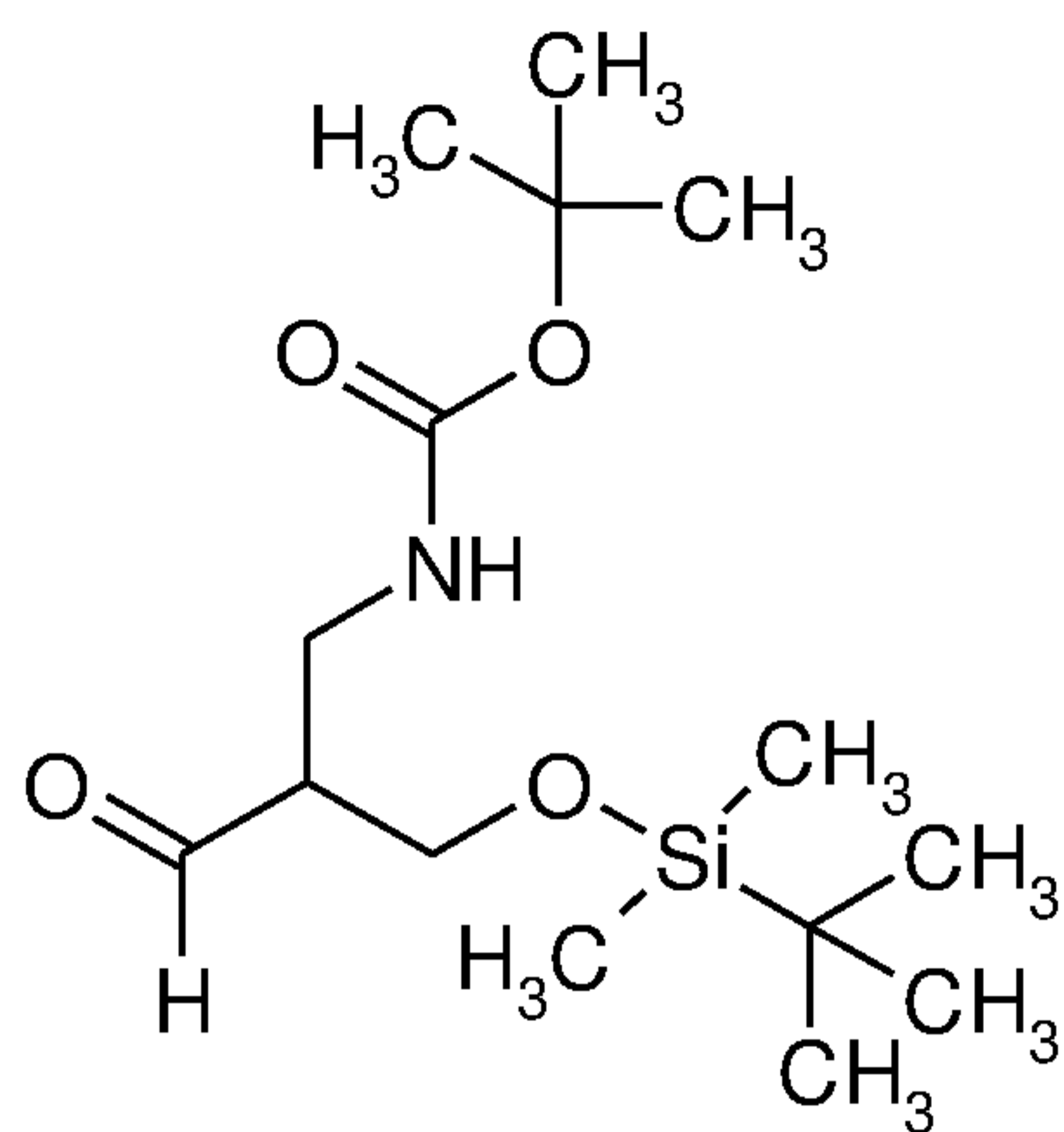
20.00 g (97.44 mmol) of tert-butyl [3-hydroxy-2-(hydroxymethyl)propyl]carbamate were dissolved in 1000 ml of absolute dichloromethane, and 6.63 g (97.44 mmol) of imidazole and 16.16 g (107.18 mmol) of tert-butyl(chloro)dimethylsilane were added at RT. The reaction mixture was stirred at RT for 16 h and washed with semiconcentrated sodium chloride solution. The aqueous phase was extracted with ethyl acetate and the combined organic phases were dried over sodium sulphate, concentrated on a rotary evaporator and dried under high vacuum. This gave 28.50 g (92% of theory) of the target compound.

<sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>): δ [ppm] = 0.02 (s, 6H), 0.86 (s, 9H), 1.37 (s, 9H), 1.58-1.73 (m, 1H), 2.91 (q, 2H), 3.33-3.36 [m, (2H, obscured)], 3.53-3.58 (m, 2H), 6.65-6.72 (m, 1H).

#### Intermediate L34

tert-Butyl (3-{{tert-butyl(dimethyl)silyl}oxy}-2-formylpropyl)carbamate

25

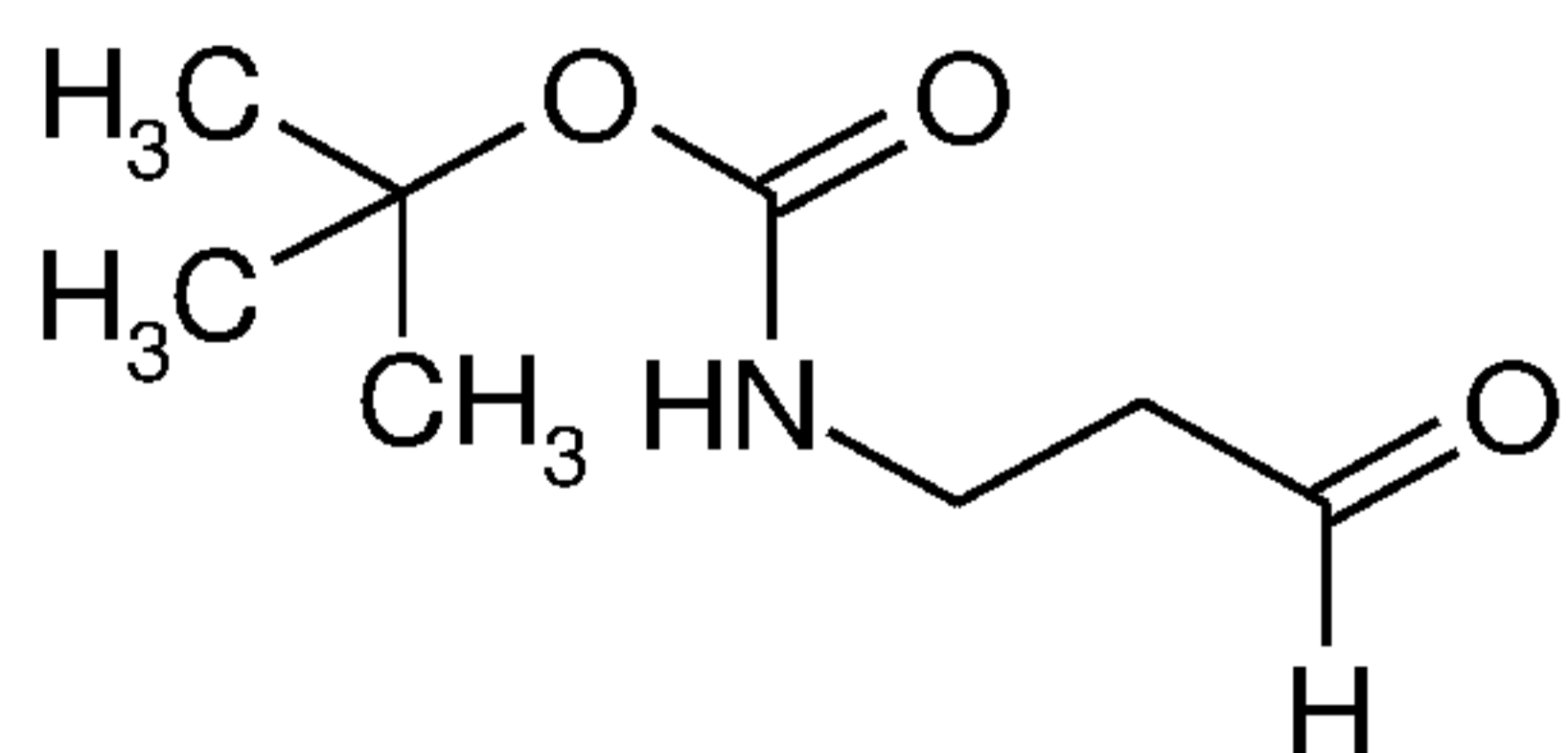


12.65 g (39.591 mmol) of tert-butyl [3-[[tert-butyl(dimethyl)silyloxy]-2-(hydroxy-methyl)propyl]carbamate  
 5 were dissolved in 200 ml of dichloromethane, and 19.31 g (45.53 mmol) of Dess-Martin periodinane dissolved in 150 ml of dichloromethane were added dropwise at RT. The mixture was stirred at room temperature for 2 h, 250 ml of a semiconcentrated sodium bicarbonate solution and 250 ml of a 10% strength sodium  
 10 thiosulphate solution were then added and the mixture was stirred for 20 min. The organic phase was separated off and the aqueous phase was extracted with ethyl acetate. The combined organic phases were washed with 300 ml of water, dried over sodium sulphate, concentrated on a rotary evaporator and dried  
 15 under high vacuum. This gave 11.35 g (90% of theory) of the target compound.

$^1\text{H}$  NMR (400 MHz, DMSO- $d_6$ ):  $\delta$  [ppm] = 0.02 (s, 6H), 0.84 (s, 9H), 1.36 (s, 9H), 1.48-1.51 (m, 1H), 3.08-3.32 [m, (1H, obscured)],  
 20 3.50-3.58 (m, 2H), 3.81-3.91 (m, 1H), 6.71 (t, 1H), 9.60 (d, 1H).

### Intermediate L35

25 tert-Butyl (3-oxopropyl)carbamate

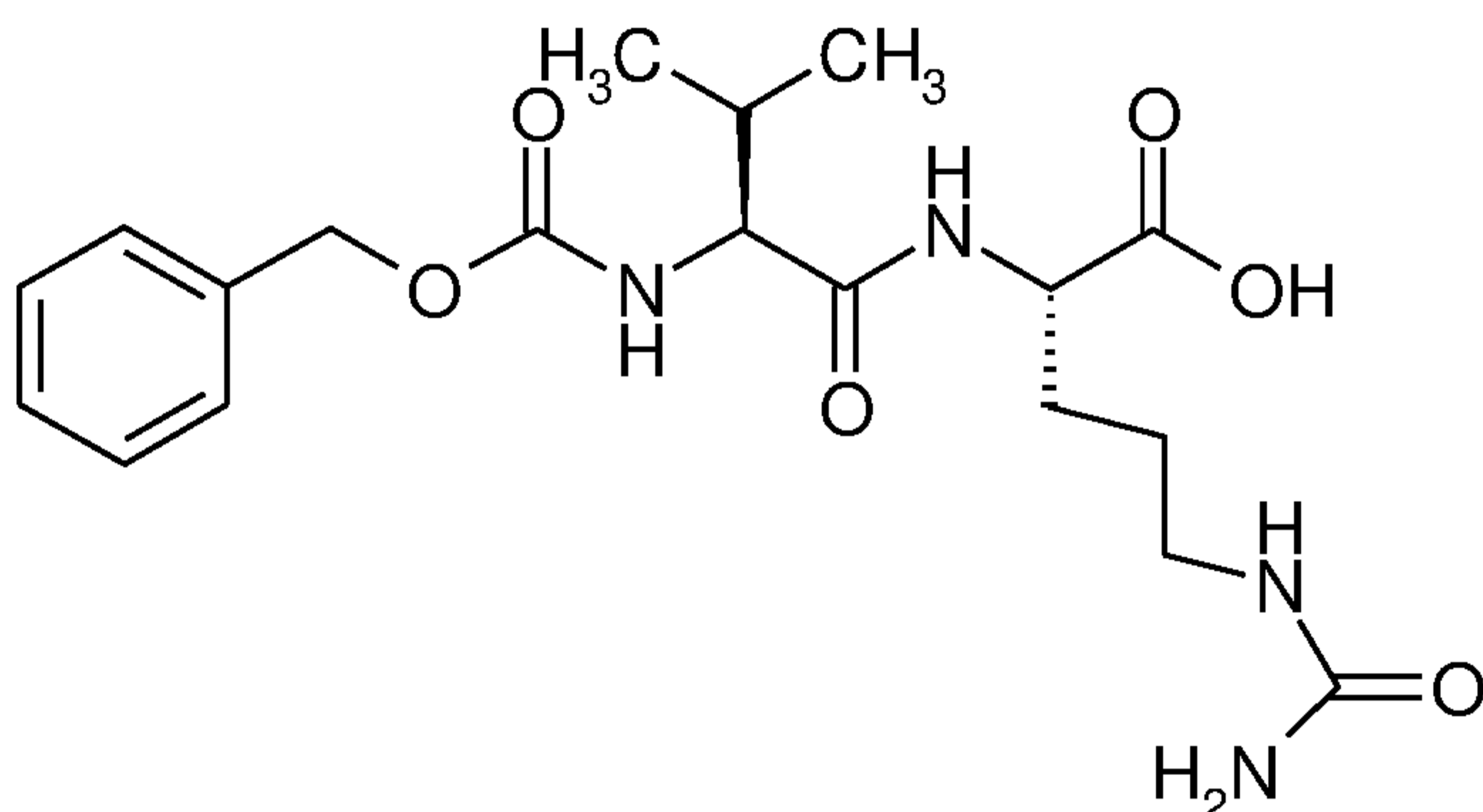


The title compound was prepared according to a method known from the literature (e.g. Jean Bastide et al. *J. Med. Chem.* **2003**, *46*(16), 3536-3545).

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### Intermediate L36

N-[(Benzyloxy)carbonyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithine



10

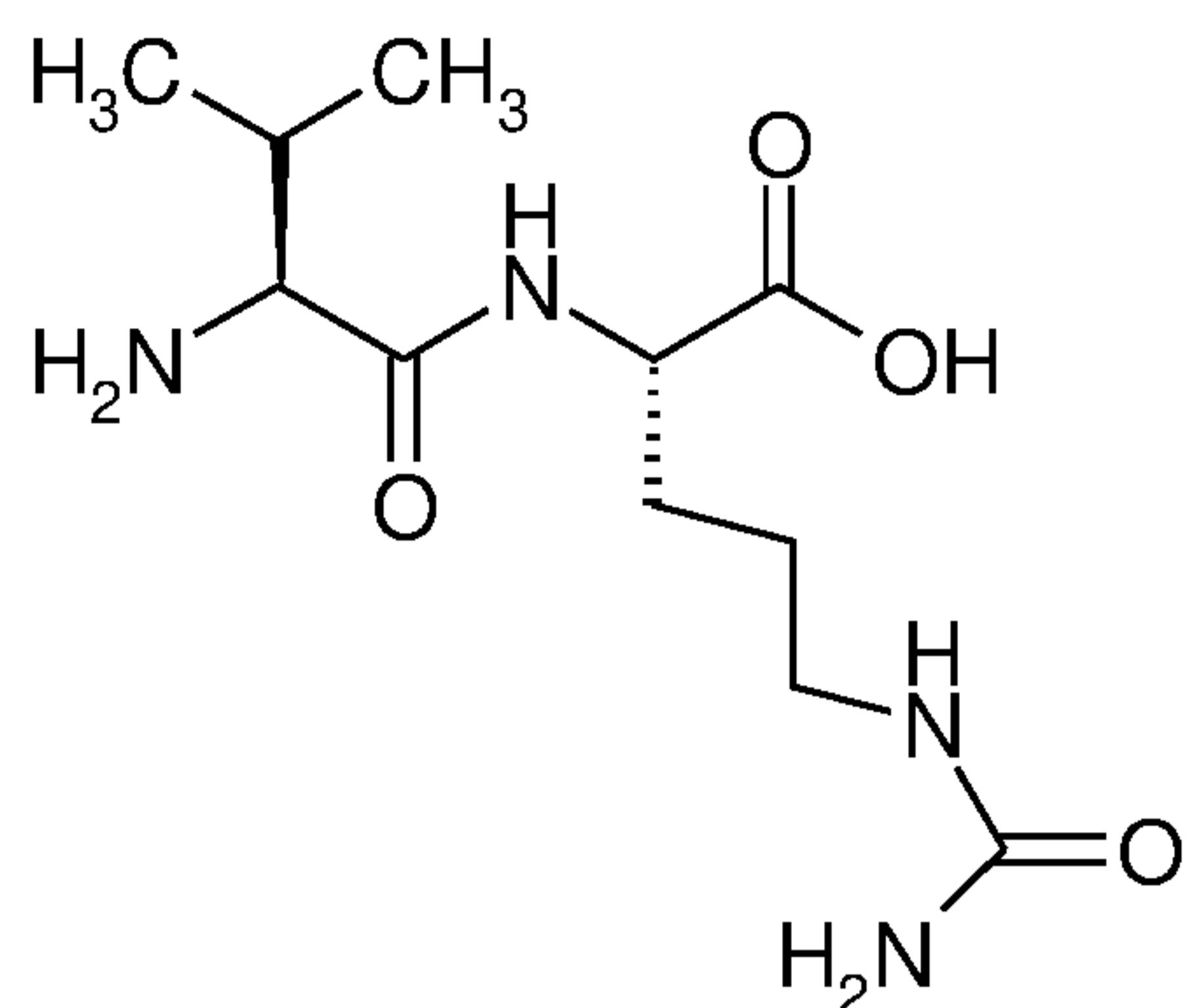
100 mg (0.57 mmol) of N<sup>5</sup>-carbamoyl-L-ornithine were taken up in 4.0 ml of DMF, and 0.08 ml (0.57 mmol) of triethylamine was added. 199.0 mg (0.57 mmol) of 2,5-dioxopyrrolidin-1-yl N-  
15 [(benzyloxy)carbonyl]-L-valine and 0.08 ml (0.57 mmol) of triethylamine were then added. The mixture was stirred at RT for 48 h. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water with 0.1% TFA). The solvents were evaporated under  
20 reduced pressure and the residue was dried under high vacuum. This gave 75.7 mg (33% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.69$  min; MS (ESIpos):  $m/z = 409$  [M+H]<sup>+</sup>.

### 25 Intermediate L37

L-Valyl-N<sup>5</sup>-carbamoyl-L-ornithine





75.7 mg (0.19 mmol) of Intermediate L36 were suspended in 25 ml  
 of water/ethanol/THF, and 7.5 mg of palladium on activated  
 5 carbon (10%) were added and the mixture was hydrogenated at RT  
 with hydrogen under standard pressure for 4.5 h. The catalyst  
 was filtered off and the reaction mixture was freed from the  
 solvent under reduced pressure and dried under high vacuum. The  
 residue was used for the next step without further purification.  
 10 This gave 64.9 mg (93% of theory) of the title compound.

LC-MS (Method 6):  $R_t = 0.25$  min; MS (ESIpos):  $m/z = 275$   $[M+H]^+$ .

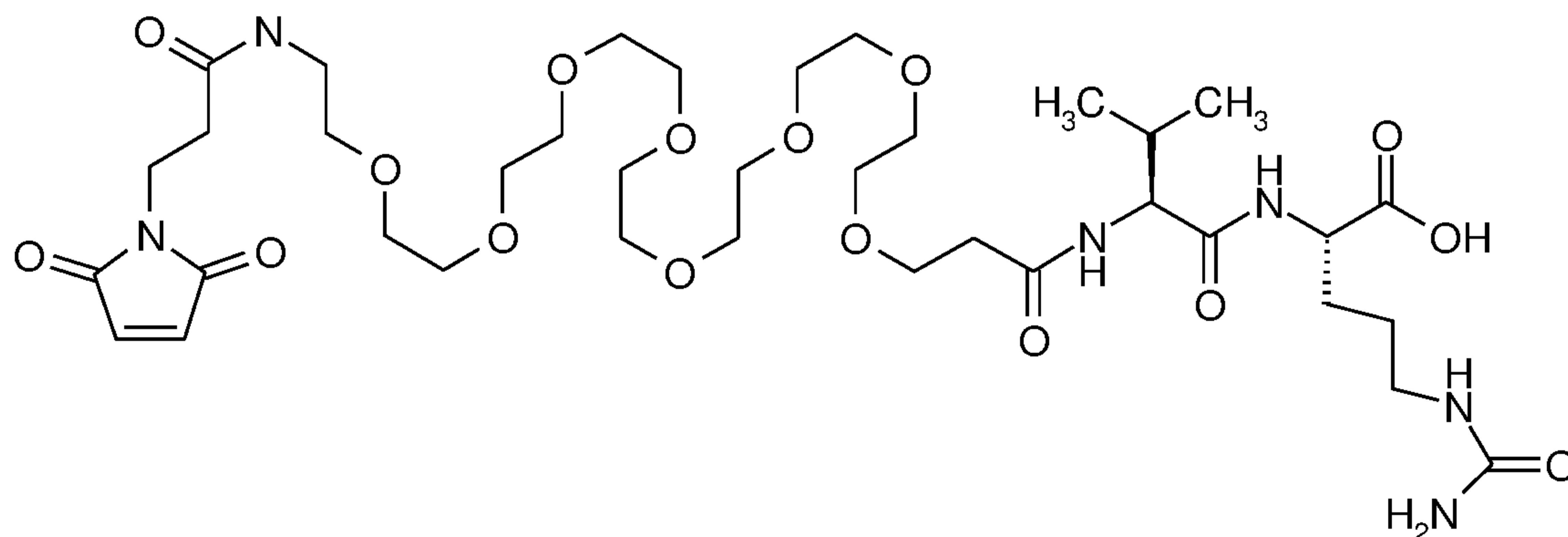
### Intermediate L38

15

N-[31-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-29-oxo-  
 4,7,10,13,16,19,22,25-octaoxa-28-

azahentriacontan-1-oyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithine

20



38.3 mg (0.14 mmol) of Intermediate L37 were initially charged  
 in 3.0 ml of DMF, and 96.4 mg (0.14 mmol) of 3-(2,5-dioxo-2,5-

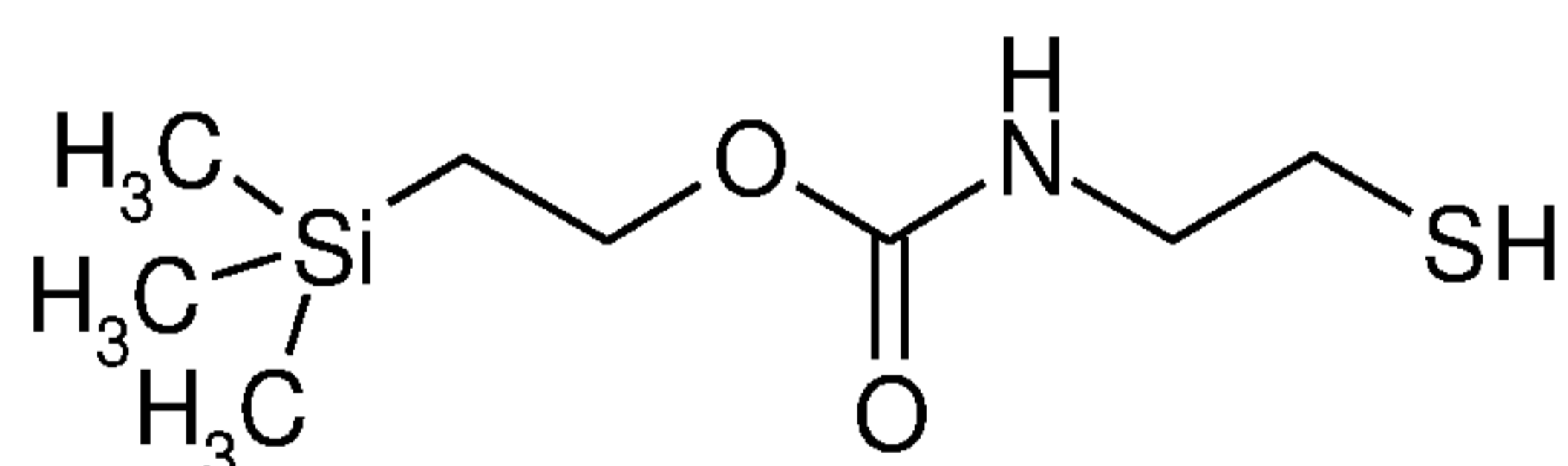
dihydro-1H-pyrrol-1-yl)-N-{27-[(2,5-dioxopyrrolidin-1-yl)oxy]-  
27-oxo-3,6,9,12,15,18,21,24-octaoxaheptacos-1-yl}propanamide  
and 39.0  $\mu$ l (0.28 mmol) of triethylamine were added. The mixture  
was stirred at RT overnight. 16.0  $\mu$ l (0.28 mmol) of HOAc were  
5 then added, and the reaction mixture was purified directly by  
preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate:  
50 ml/min, MeCN/water). The solvents were evaporated under  
reduced pressure and the residue was dried under high vacuum.  
This gave 58.9 mg (45% of theory) of the title compound.

10

LC-MS (Method 1):  $R_t$  = 0.61 min; MS (ESIpos):  $m/z$  = 849 [M+H]<sup>+</sup>.

### Intermediate L39

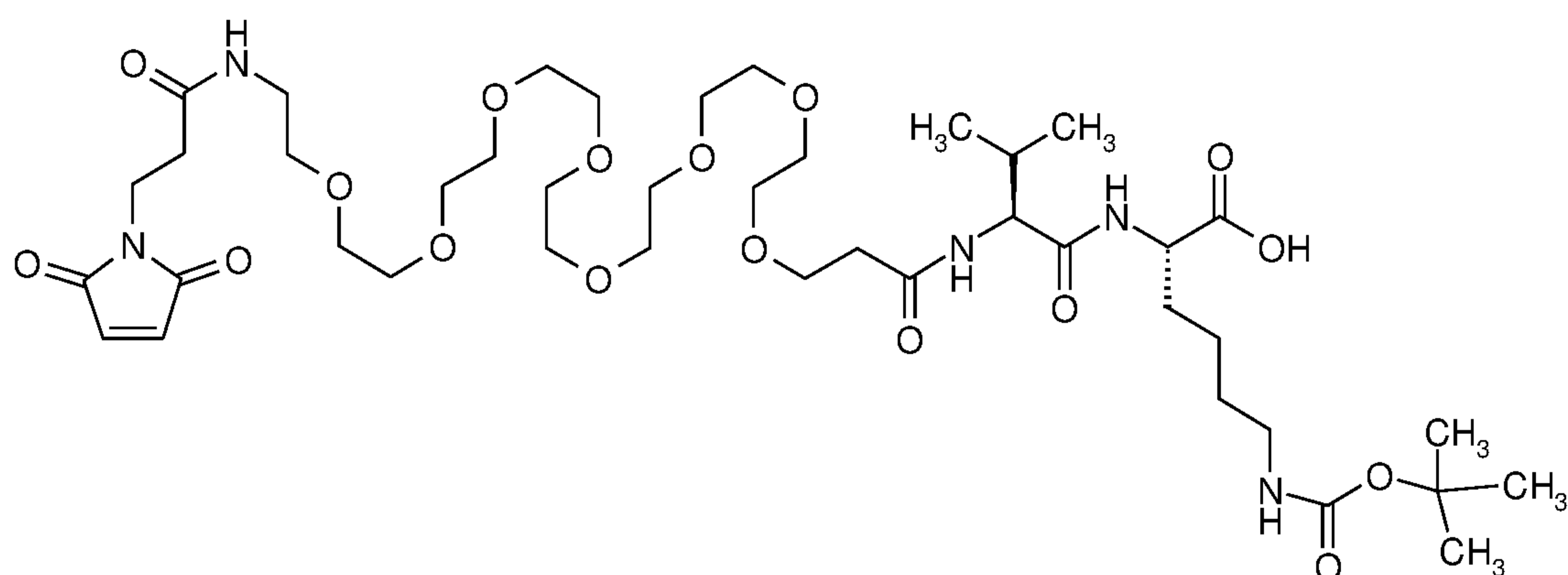
15 2-(Trimethylsilyl)ethyl (2-sulphanylethyl) carbamate



300 mg (2.64 mmol) of 2-aminoethanethiol hydrochloride (1:1)  
20 were initially charged in 3.0 ml of dichloromethane, and 668.0  
mg (6.60 mmol) of triethylamine and 719.1 mg (2.77 mmol) of 1-  
({[2-(trimethylsilyl)ethoxy]carbonyl}oxy)pyrrolidine-2,5-dione  
were added. The mixture was stirred at RT for 2 days (monitored  
by thin-layer chromatography: dichloromethane/methanol =  
25 100:1.5). Ethyl acetate was added and the reaction mixture was  
washed three times with water. The organic phase was washed  
twice with saturated NaCl solution and dried over magnesium  
sulphate. The solvent was evaporated under reduced pressure and  
the residue was dried under high vacuum. The compound was used  
30 without further purification in the next step of the synthesis.

### Intermediate L40

N-[31-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-29-oxo-  
35 4,7,10,13,16,19,22,25-octaoxa-28-azahentriacontan-1-oyl]-L-  
valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine



600 mg (1.58 mmol) of N<sup>2</sup>-[(benzyloxy)carbonyl]-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were hydrogenated in 25.0 ml of water/ethanol/THF (1:1:0.5) using palladium on carbon (10%) at RT under standard pressure with hydrogen. The compound N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine is used without further purification in the next step of the synthesis.

10

LC-MS (Method 1): R<sub>t</sub> = 0.99 min; MS (ESIpos): m/z = 247 [M+H]<sup>+</sup>.

180.0 g (0.73 mmol) of N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were dissolved in 5.0 ml of DMF, and 74.0 mg (0.73 mmol) of triethylamine were added. 254.6 mg (0.73 mmol) of 2,5-dioxopyrrolidin-1-yl N-[(benzyloxy)carbonyl]-L-valinate and 74.0 mg (0.73 mmol) of triethylamine were then added. The reaction mixture was stirred at RT for 3.5 h. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10μ, flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 294.1 mg (76% of theory) of the compound N-[(benzyloxy)carbonyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine.

25

LC-MS (Method 1): R<sub>t</sub> = 0.97 min; MS (ESIpos): m/z = 480 [M+H]<sup>+</sup>.

272.2 mg (0.57 mmol) of N-[(benzyloxy)carbonyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were dissolved in 20 ml of ethyl acetate/ethanol/THF (1:1:1), 27.2 mg of palladium on activated carbon were added and the mixture was hydrogenated under

30

standard pressure and at RT with hydrogen. The mixture was filtered through Celite<sup>(R)</sup> and the filter cake was washed thoroughly with ethyl acetate/ethanol/THF (1:1:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 182.0 mg (72% of theory) of the compound L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine.

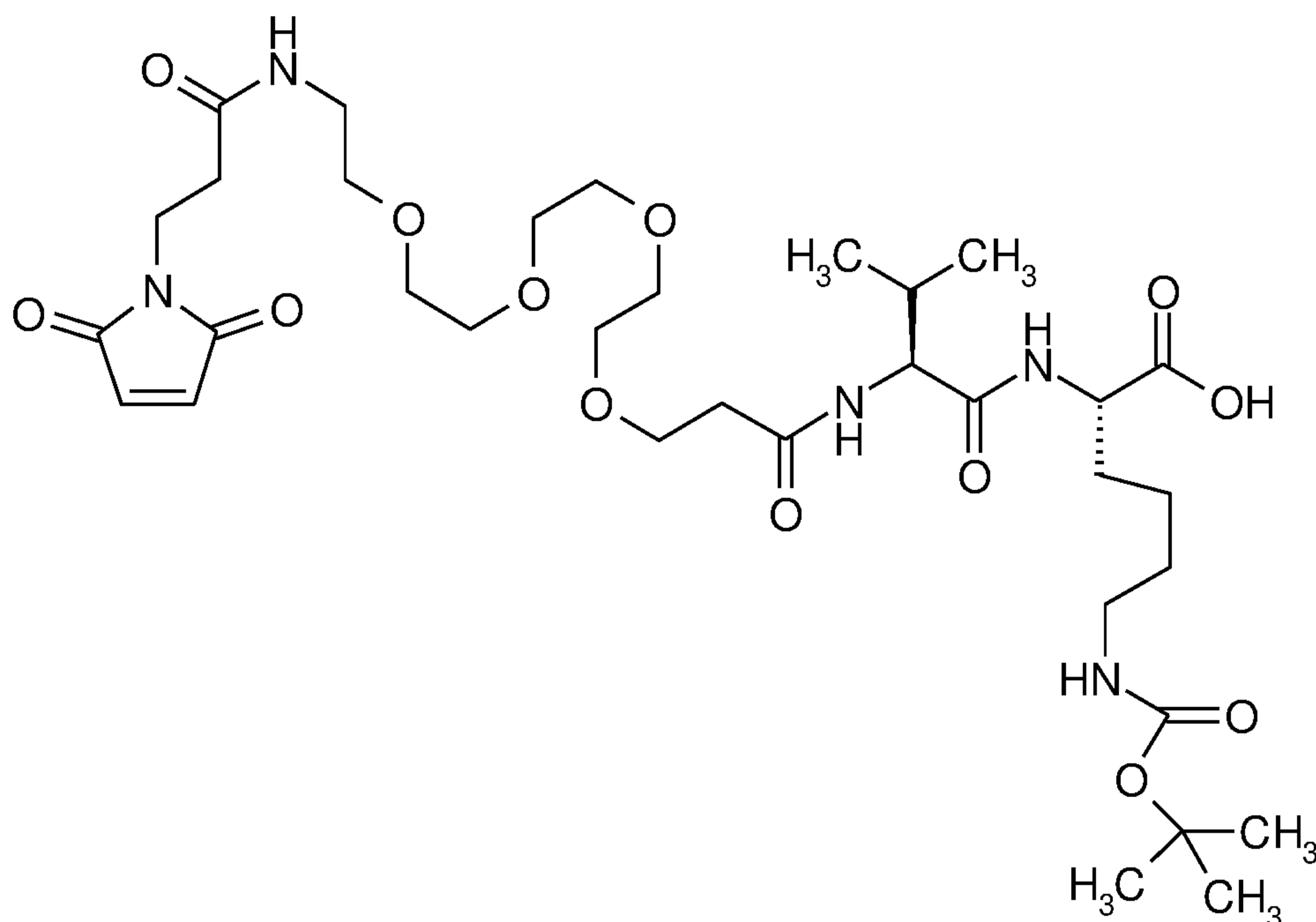
LC-MS (Method 1): R<sub>t</sub> = 0.53 min; MS (ESIpos): m/z = 346 [M+H]<sup>+</sup>.

30.0 mg (0.07 mmol) of L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine and 46.1 mg (0.07 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{27-[(2,5-dioxopyrrolidin-1-yl)oxy]-27-oxo-3,6,9,12,15,18,21,24-octaoxaheptacos-1-yl}propanamide were dissolved in 1.5 ml of DMF, and 6.8 mg (0.07 mmol) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10μ, flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 55.6 mg (90% of theory) of the title compound.

LC-MS (Method 1): R<sub>t</sub> = 0.77 min; MS (ESIpos): m/z = 920 [M+H]<sup>+</sup>.

#### Intermediate L41

N-[19-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine



600 mg (1.58 mmol) of N<sup>2</sup>-[(benzyloxy)carbonyl]-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were hydrogenated in 25.0 ml of water/ethanol/THF (1:1:0.5) using palladium on carbon (10%) at RT under standard pressure with hydrogen. The compound N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine is used without further purification in the next step of the synthesis.

10 LC-MS (Method 1): R<sub>t</sub> = 0.99 min; MS (ESIpos): m/z = 247 [M+H]<sup>+</sup>.

180.0 g (0.73 mmol) of N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were dissolved in 5.0 ml of DMF, and 74.0 mg (0.73 mmol) of triethylamine were added. 254.6 mg (0.73 mmol) of 2,5-dioxopyrrolidin-1-yl N-[(benzyloxy)carbonyl]-L-valinate and 74.0 mg (0.73 mmol) of triethylamine were then added. The reaction mixture was stirred at RT for 3.5 h. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprisil 250x30; 10μ, flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were then evaporated under reduced pressure and the residue was dried under high vacuum. This gave 294.1 mg (76% of theory) of the compound N-[(benzyloxy)carbonyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine.

25 LC-MS (Method 1): R<sub>t</sub> = 0.97 min; MS (ESIpos): m/z = 480 [M+H]<sup>+</sup>.

272.2 mg (0.57 mmol) of N-[(benzyloxy)carbonyl]-L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine were dissolved in 20.0 ml of ethyl acetate/ethanol/THF (1:1:1), 27.2 mg of palladium on activated carbon were added and the mixture was hydrogenated under standard pressure and at RT with hydrogen. The mixture was filtered through Celite<sup>(R)</sup> and the filter cake was washed thoroughly with ethyl acetate/ethanol/THF (1:1:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 182.0 mg (72% of theory) of the compound L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine.

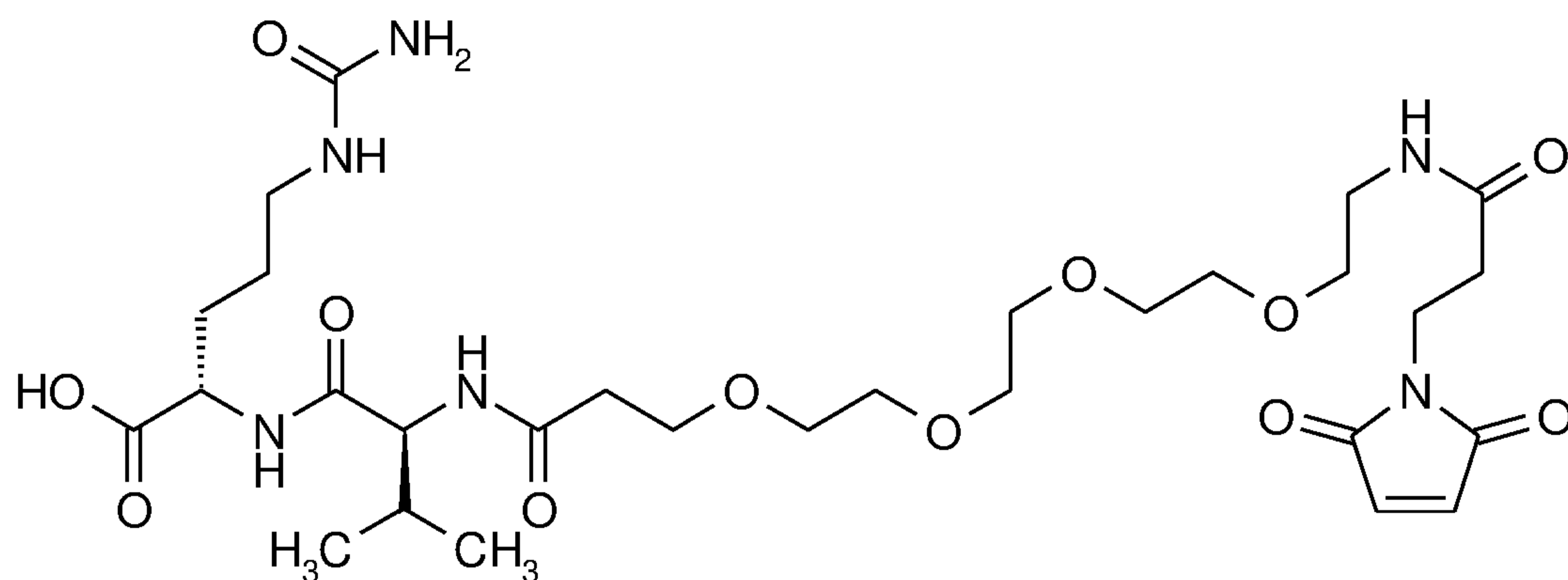
LC-MS (Method 1): R<sub>t</sub> = 0.53 min; MS (ESIpos): m/z = 346 [M+H]<sup>+</sup>.

30.0 mg (0.07 mmol) of L-valyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine and 34.3 mg (0.07 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide were dissolved in 1.5 ml of DMF, and 6.8 mg (0.07 mmol) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10μ, flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 40.6 mg (82% of theory) of the title compound.

LC-MS (Method 1): R<sub>t</sub> = 0.73 min; MS (ESIpos): m/z = 744 [M+H]<sup>+</sup>.

### Intermediate L42

N-[19-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithine



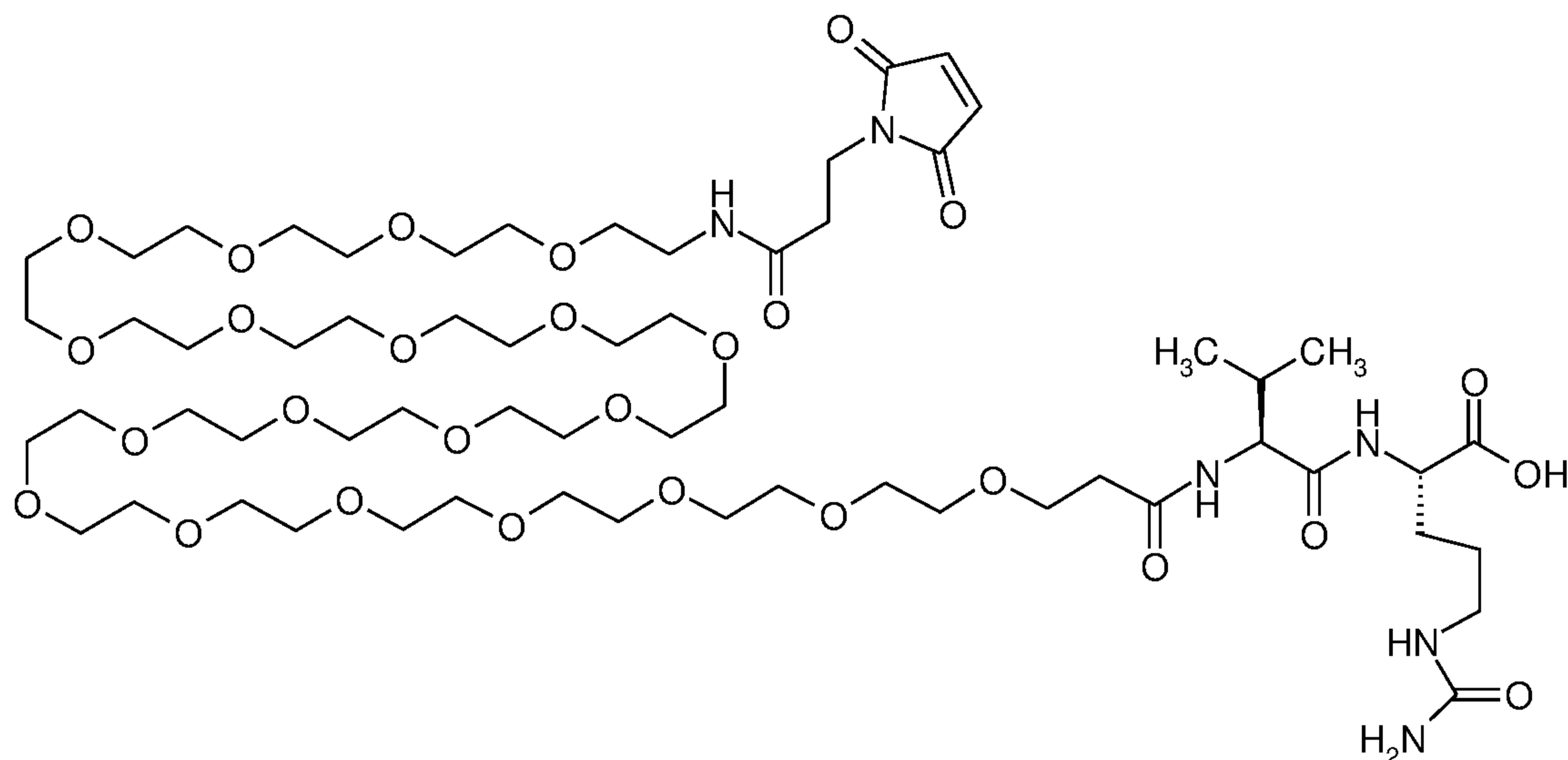
50.0 mg (0.18 mmol) of L-valyl-N<sup>5</sup>-carbamoyl-L-ornithine (Intermediate L37) were initially charged in DMF, and 93.6 mg (0.18 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide and 36.9 mg (0.37 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight. 21.9 mg (0.37 mmol) of HOAc were added and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 20.6 mg (14% of theory) of the title compound.

15

LC-MS (Method 1):  $R_t$  = 0.55 min; MS (ESIpos):  $m/z$  = 673 [M+H]<sup>+</sup>.

### Intermediate L43

20 N-[67-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-65-oxo-4,7,10,13,16,19,22,25,28,31,34,37,40,43,46,49,52,55,58,61-icosaoxa-64-azaheptahexacontan-1-oyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithine



11.3 mg (0.04 mmol) of L-valyl-N<sup>5</sup>-carbamoyl-L-ornithine  
 (Intermediate L37) were initially charged in DMF, and 50.0 mg  
 5 (0.04 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{63-  
 [(2,5-dioxopyrrolidin-1-yl)oxy]-63-oxo-  
 3,6,9,12,15,18,21,24,27,30,33,36,39,42,45,48,51,54,57,60-  
 icosaoxatrihexacont-1-yl}propanamide and 8.3 mg (0.08 mmol) of  
 triethylamine were added. The reaction mixture was stirred at  
 10 RT overnight. 4.9 mg (0.08 mmol) of HOAc were added and the  
 reaction mixture was purified directly by preparative RP-HPLC  
 (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min,  
 MeCN/water). The solvents were evaporated under reduced pressure  
 and the residue was dried under high vacuum. This gave 15.8 mg  
 15 (20% of theory) of the title compound.

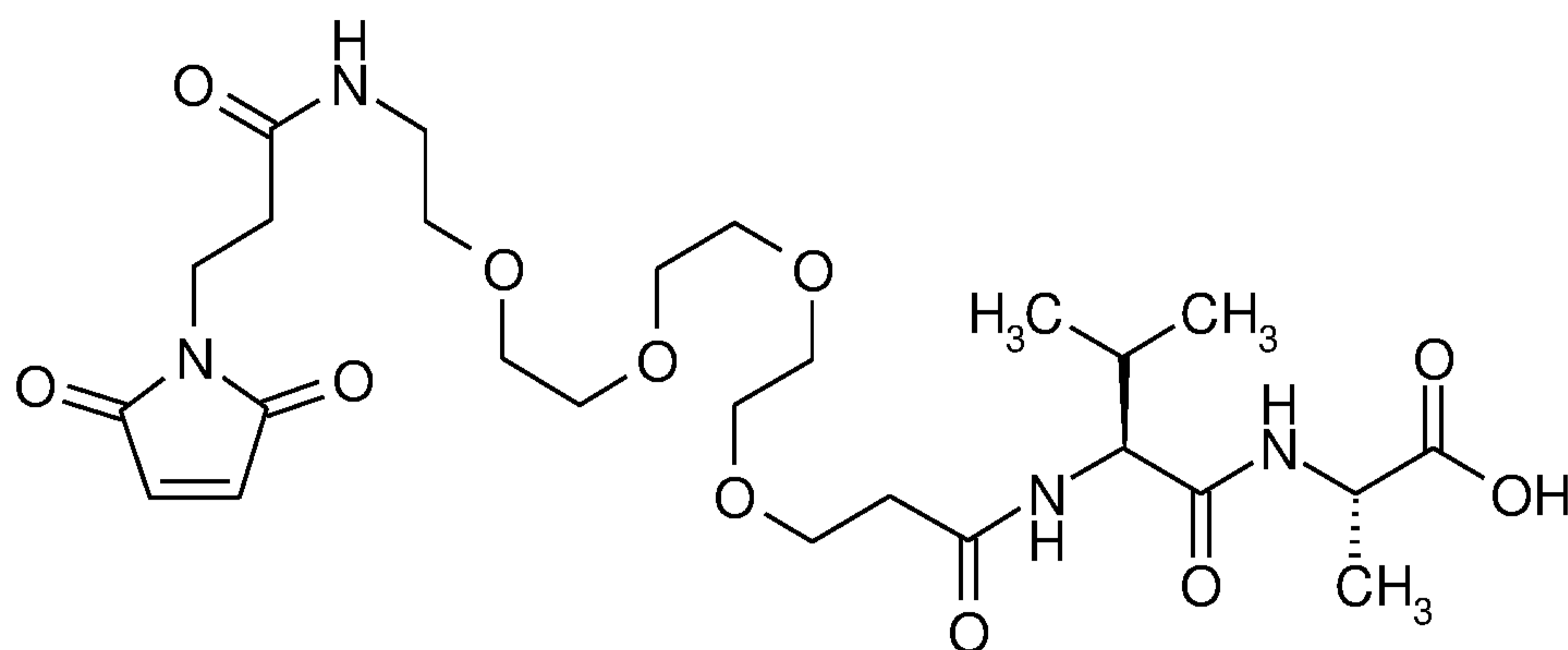
LC-MS (Method 4):  $R_t = 0.94$  min; MS (ESIpos):  $m/z = 1377$  [M+H]<sup>+</sup>.

#### Intermediate L44

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N-[19-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-  
 tetraoxa-16-azanonadecan-1-oyl]-L-valyl-L-alanine



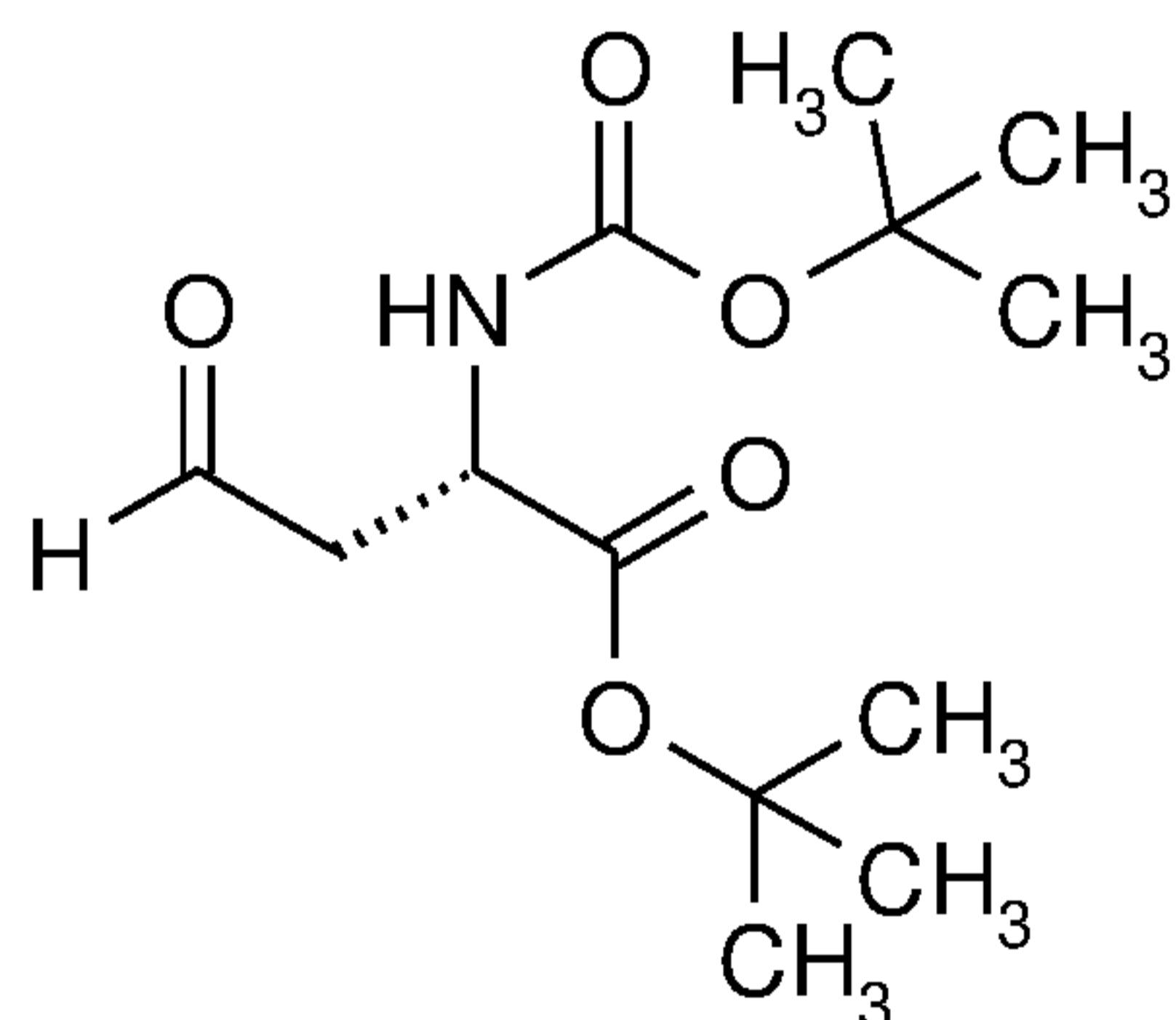


73.3 mg (0.39 mmol) of L-valyl-L-alanine were dissolved in 7.0 ml of DMF, and 200.0 mg (0.39 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide and 78.8 mg (0.78 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 103.3 mg (45% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.58 min; MS (ESIpos):  $m/z$  = 587 [M+H]<sup>+</sup>.

### Intermediate L45

tert-Butyl (2S)-2-[(tert-butoxycarbonyl)amino]-4-oxobutanoate



20

2.00 g (7.26 mmol) of tert-butyl N-(tert-butoxycarbonyl)-L-homoserinate were dissolved in 90 ml of dichloromethane, and 1.76 ml of pyridine and 4.62 g (10.90 mmol) of 1,1,1-triacetoxy-1 $\lambda^5$ ,2-benziodoxol-3(1H)-on (Dess-Martin periodinane) were

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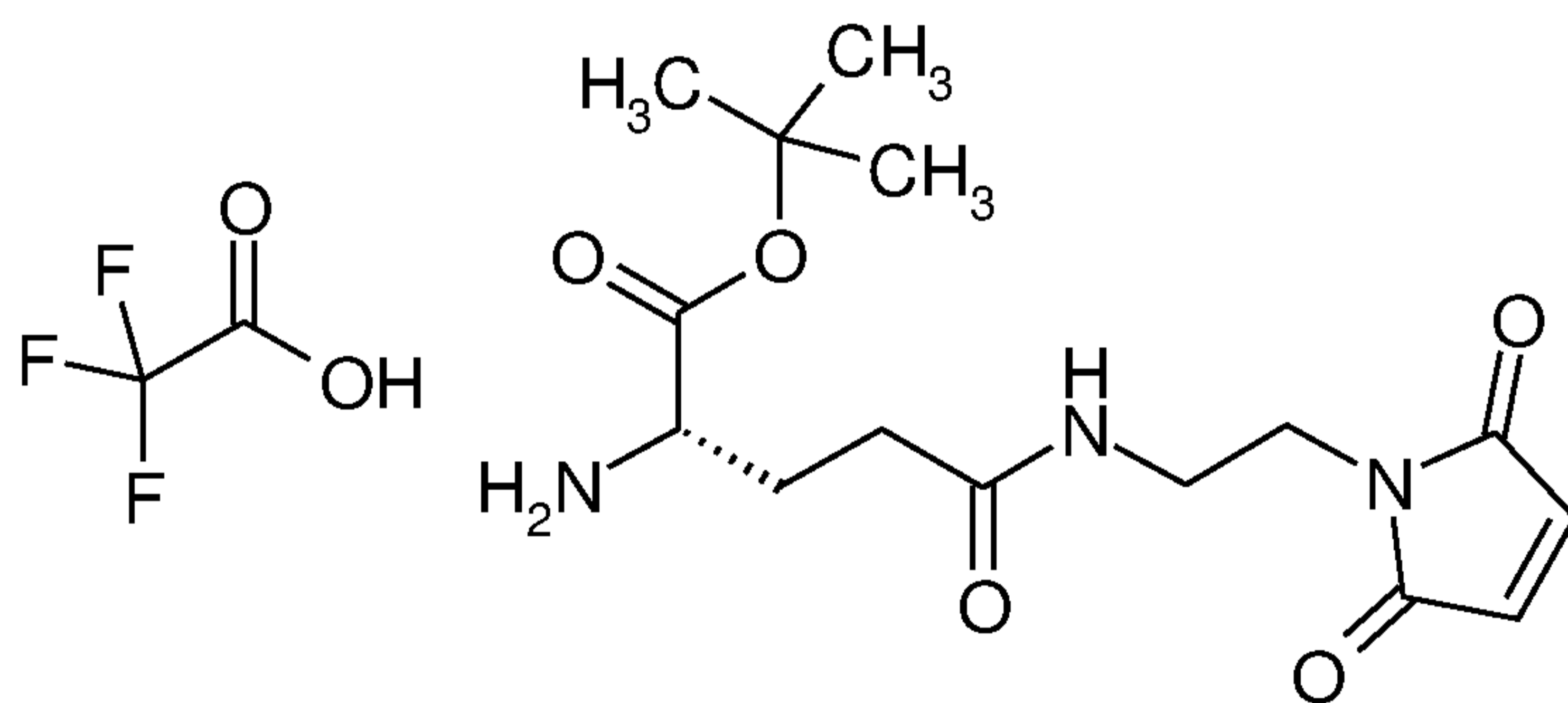
then added. The reaction was stirred at RT for 2 h and then diluted with 200 ml of dichloromethane and extracted twice with 10% strength sodium thiosulphate solution and then successively twice with 5% strength citric acid and twice with saturated sodium bicarbonate solution. The organic phase was separated off, dried over sodium sulphate and then dried under reduced pressure. 100 ml of diethyl ether and cyclohexane (v/v=1:1) were added to the residue, resulting in the formation of a white precipitate. This was filtered off with suction. The filtrate was concentrated on a rotary evaporator and dried under high vacuum, giving 1.74 g (88% of theory) of the target compound as a light-yellow oil.

LC-MS (Method 1):  $R_t = 0.85$  min; MS (ESIpos):  $m/z = 274$   $[M+H]^+$ .

$^1H$  NMR (400 MHz, DMSO- $d_6$ ):  $\delta$  [ppm] = 1.38 (s, 18H), 2.64-2.81 (m, 2H), 4.31-4.36 (m, 1H), 7.23 (d, 1H), 9.59 (s, 1H).

#### Intermediate L46

Trifluoroacetic acid / tert-butyl N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]-L-glutamate (1:1)



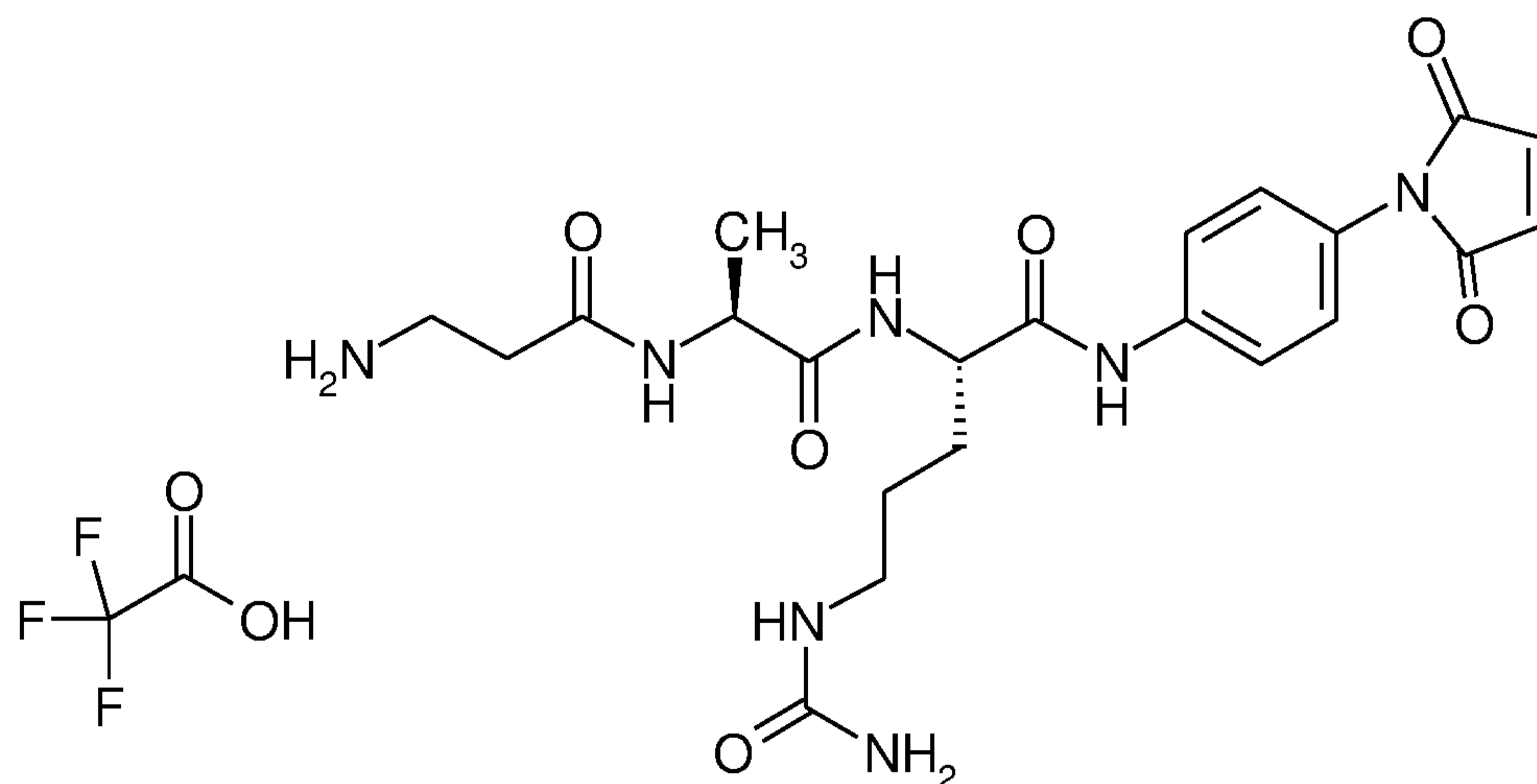
The title compound was prepared by first coupling 200 mg (0.79 mmol) of trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) with 263 mg (0.87 mmol) of (4S)-5-tert-butoxy-4-[(tert-butoxycarbonyl)amino]-5-oxopentanoic acid / trifluoroacetic acid (1:1) in the presence of EDC/HOBT and *N,N*-diisopropylethylamine and then deprotecting the amino group under gentle conditions by stirring for 1 h in 10% strength

trifluoroacetic acid in DCM at RT. Freeze-drying from acetonitrile/ water gave 85 mg (20% of theory) of the title compound over 2 steps.

5 LC-MS (Method 1):  $R_t = 0.37$  min; MS (ESIpos):  $m/z = 326$   $[M+H]^+$ .

#### Intermediate L47

10 Trifluoroacetic acid / beta-alanyl-L-alanyl-N<sup>5</sup>-carbamoyl-N-[4-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)phenyl]-L-ornithinamide (1:1)



15 The title compound was prepared by coupling Intermediate L8 with 2,5-dioxopyrrolidin-1-yl N-(tert-butoxycarbonyl)-beta-alaninate and subsequent deprotection with TFA.

20 LC-MS (Method 3):  $R_t = 1.36$  min; MS (ESIpos):  $m/z = 488$   $(M+H)^+$ .

#### Intermediate L48

25 Trifluoroacetic acid / (1R,2S)-2-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)

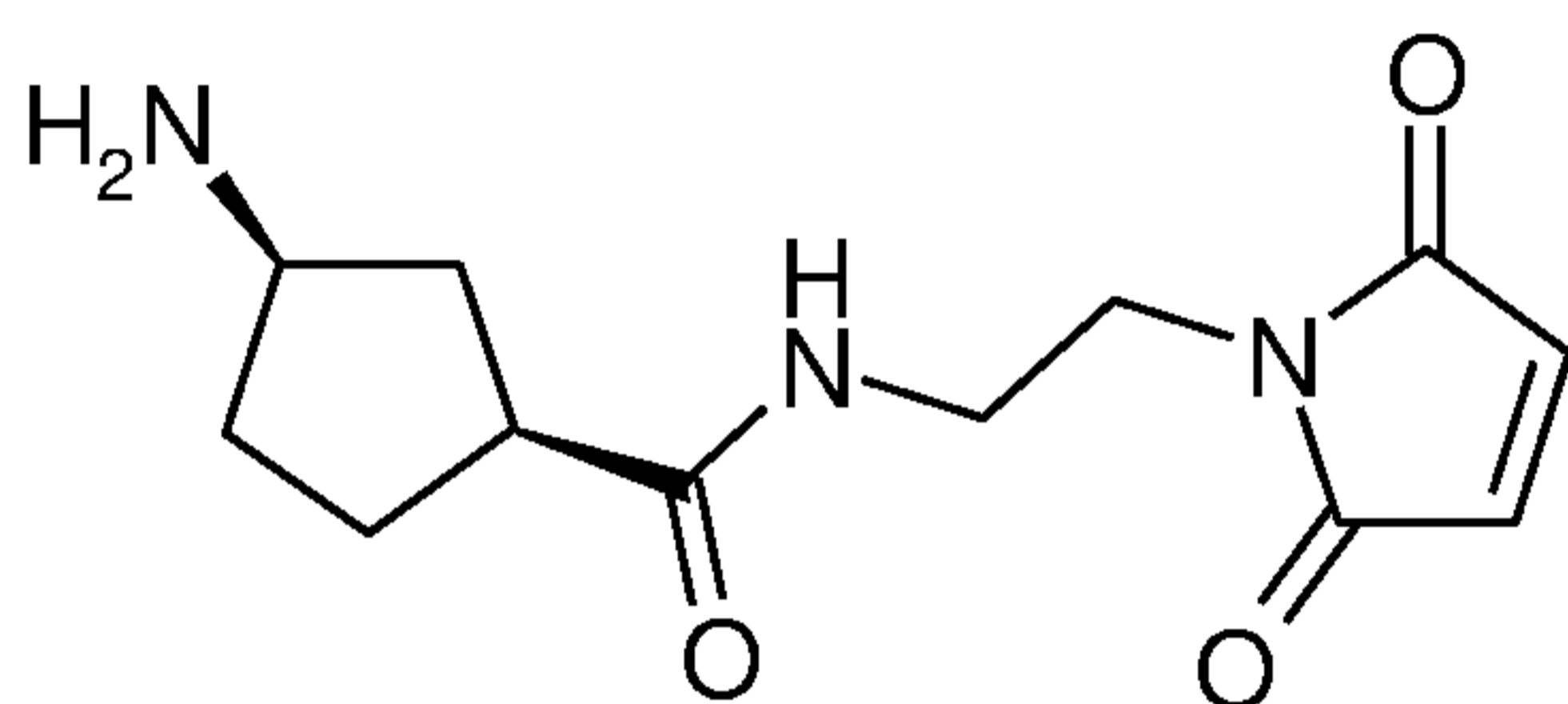
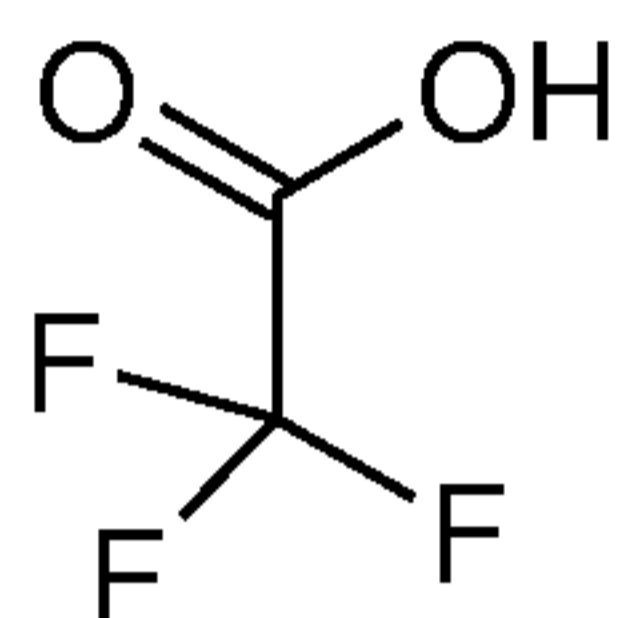


LC-MS (Method 1):  $R_t = 1.09$  min; MS (ESIpos):  $m/z = 593$  and  $595$  (M+H)<sup>+</sup>.

### Intermediate L50

5

Trifluoroacetic acid / (1S,3R)-3-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)



10

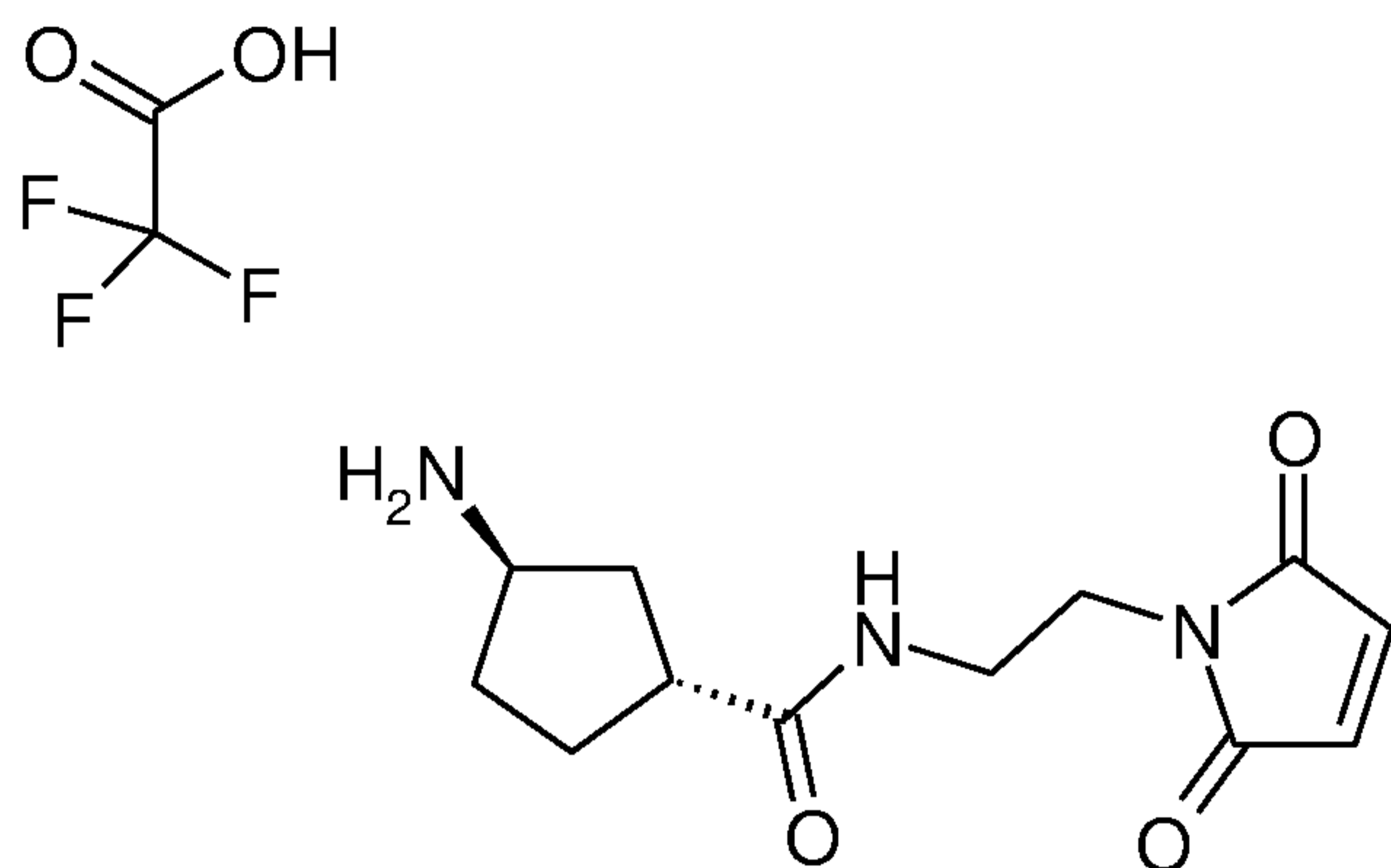
The title compound was prepared from commercially available (1S,3R)-3-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid and likewise commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) by coupling with HATU in the presence of *N,N*-diisopropylethylamine and subsequent deprotection with TFA.

HPLC (Method 11):  $R_t = 0.2$  min;

20 LC-MS (Method 3):  $R_t = 0.88$  min; MS (ESIpos):  $m/z = 252$  (M+H)<sup>+</sup>.

### Intermediate L51

25 Trifluoroacetic acid / (1R,3R)-3-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)

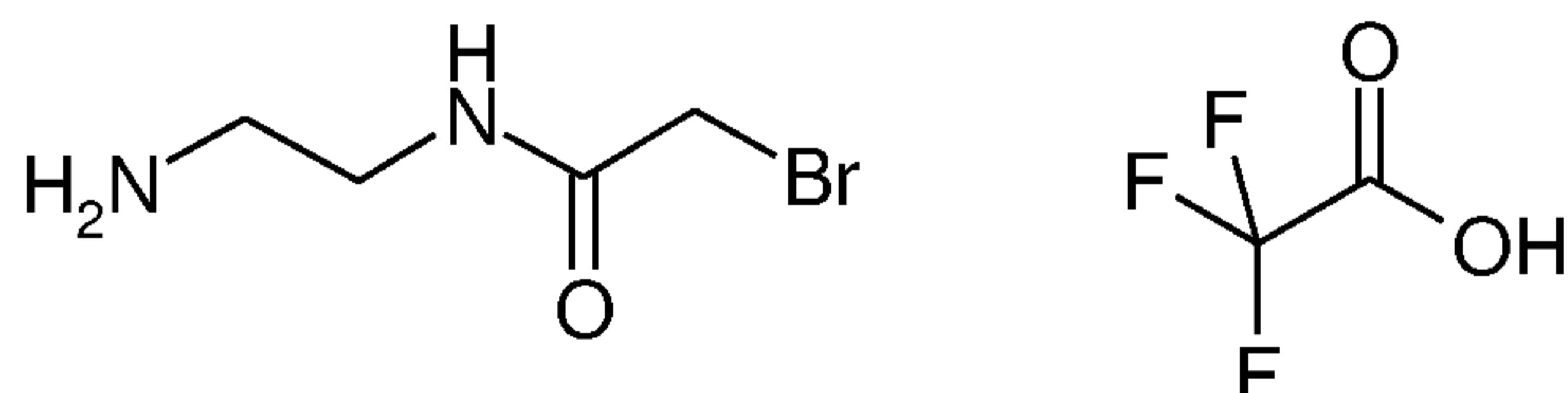


The title compound was prepared from commercially available (1R,3R)-3-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid and likewise commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) by coupling with HATU in the presence of *N,N*-diisopropylethylamine and subsequent deprotection with TFA.

10 LC-MS (Method 3):  $R_t = 0.98$  min; MS (ESIpos):  $m/z = 250$  (M-H)<sup>-</sup>.

### Intermediate L52

15 Trifluoroacetic acid / *N*-(2-aminoethyl)-2-bromoacetamide (1:1)



420 mg (2.62 mmol) of tert-butyl (2-aminoethyl)carbamate were taken up in 50 ml of dichloromethane, and 817 mg (3.15 mmol) of bromoacetic anhydride and 913  $\mu$ l (5.24 mmol) of *N,N*-diisopropylethylamine were added. The reaction was stirred at RT for 1 h and then dried under reduced pressure. The residue was purified by preparative HPLC.

25 This gave 577 mg of the protected intermediate which were then taken up in 50 ml of dichloromethane, and 10 ml of trifluoroacetic acid were added. After 1 h of stirring at RT, the reaction was concentrated under reduced pressure and the residue was lyophilized from acetonitrile/water. This gave 705

mg (65% of theory) of the title compound.

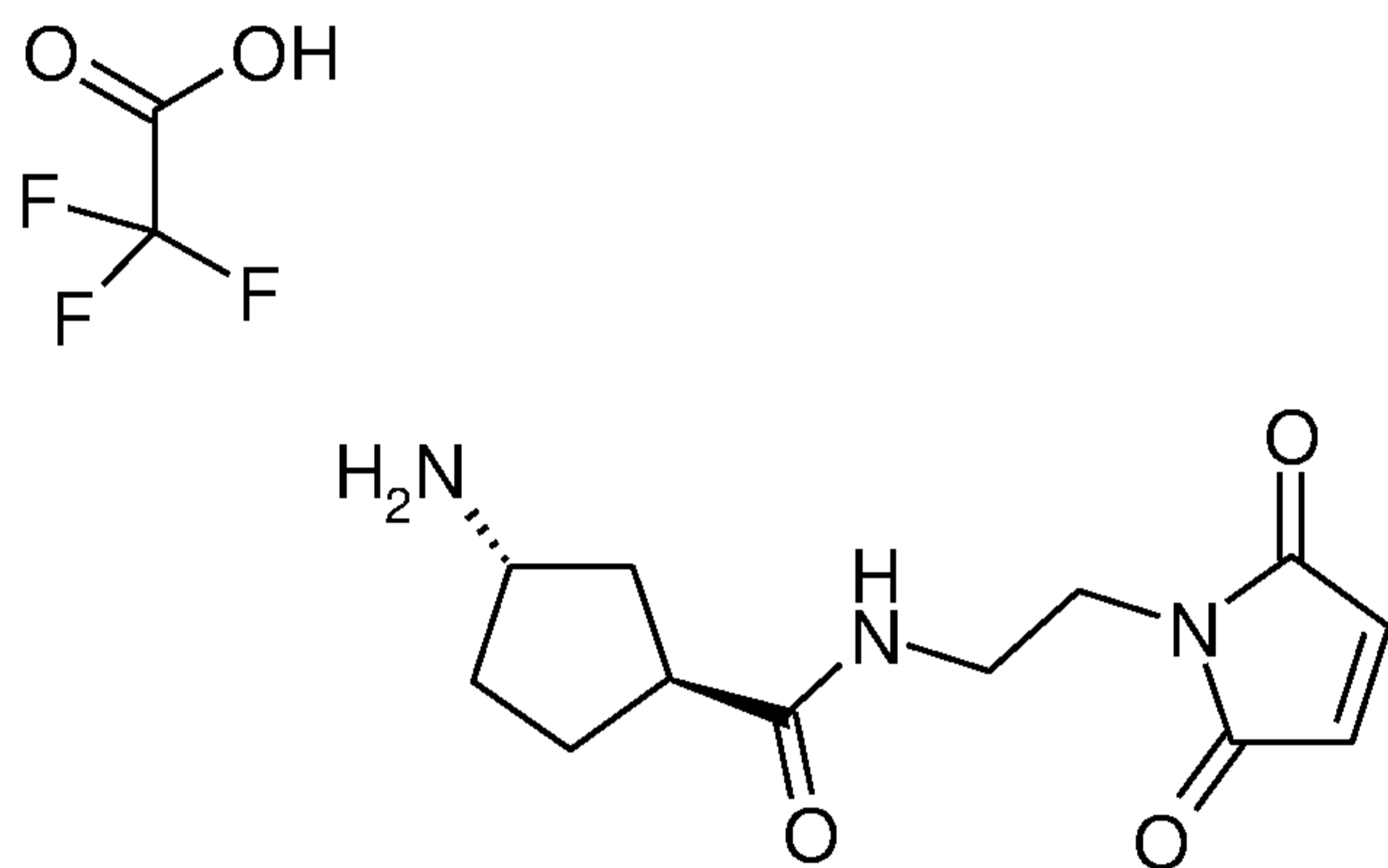
LC-MS (Method 3):  $R_t = 0.34$  min; MS (ESIpos):  $m/z = 181$  and  $183$  (M+H)<sup>+</sup>.

5

### Intermediate L53

Trifluoroacetic acid / (1S,3S)-3-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)

10



The title compound was prepared from commercially available (1S,3S)-3-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid and likewise commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) by coupling with HATU in the presence of *N,N*-diisopropylethylamine and subsequent deprotection with TFA.

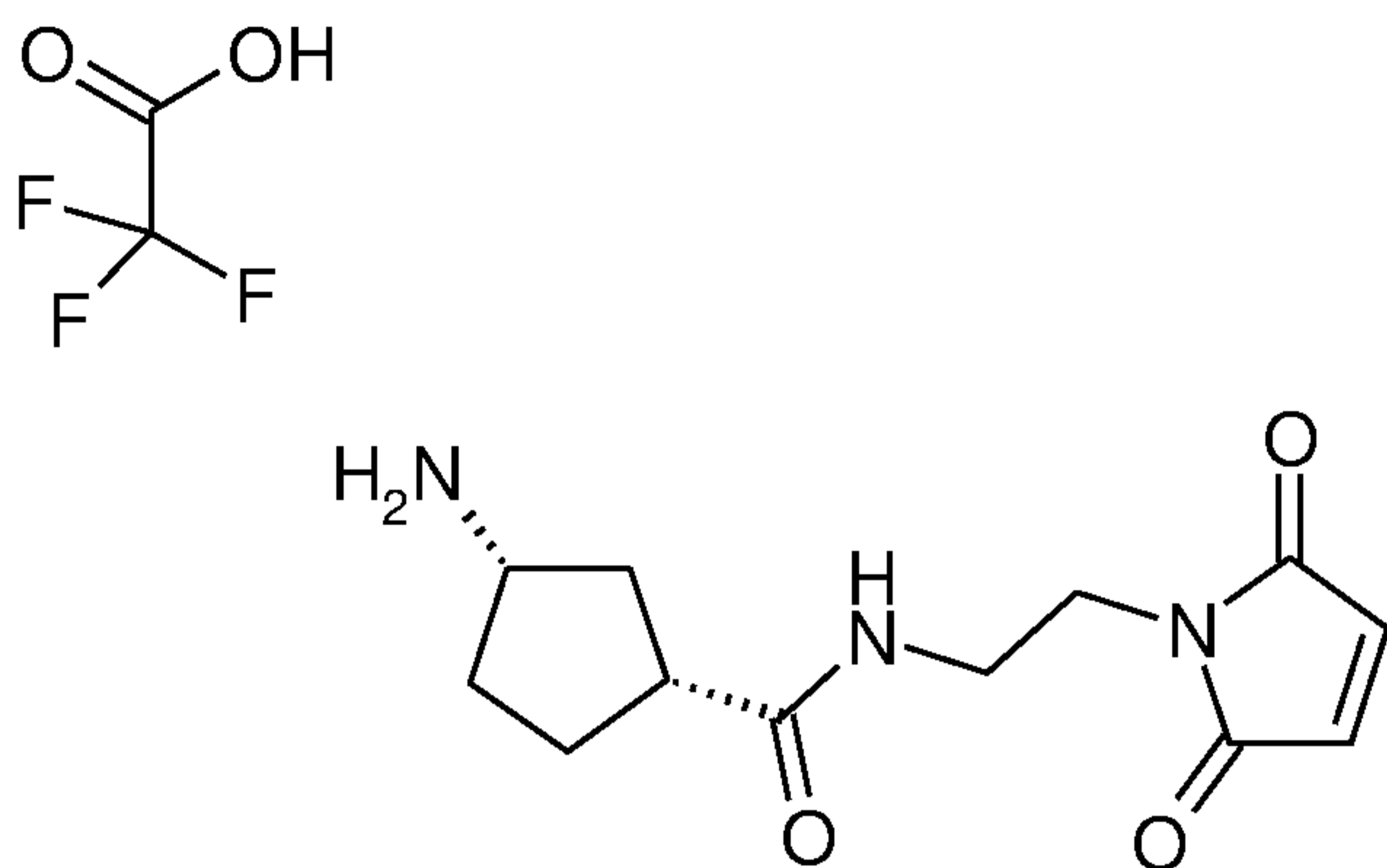
20 HPLC (Method 11):  $R_t = 0.19$  min;

LC-MS (Method 3):  $R_t = 0.88$  min; MS (ESIpos):  $m/z = 250$  (M-H)<sup>-</sup>.

### Intermediate L54

25

Trifluoroacetic acid / (1R,3S)-3-amino-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)

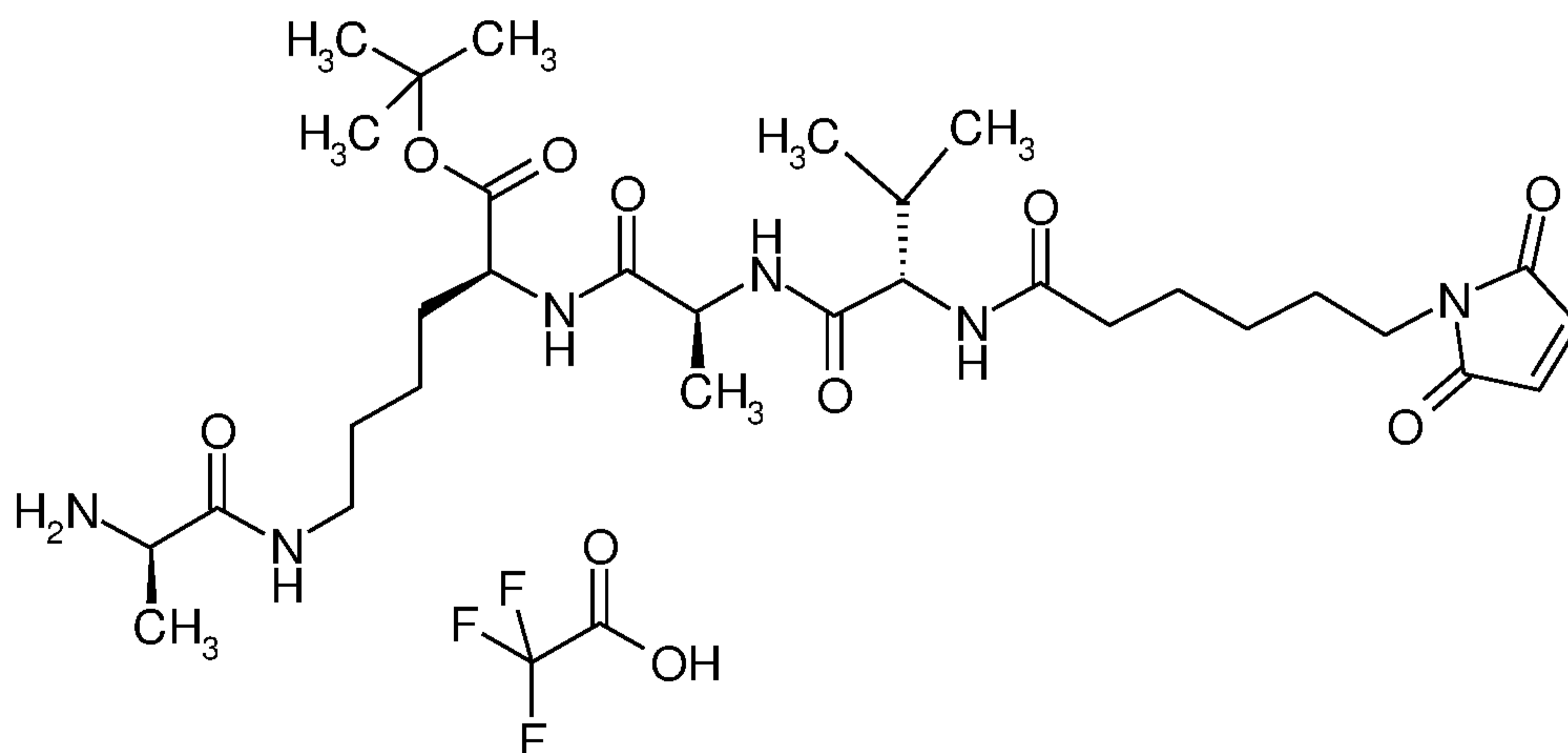


The title compound was prepared from commercially available (1R,3S)-3-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid and likewise commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) by coupling with HATU in the presence of *N,N*-diisopropylethylamine and subsequent deprotection with TFA.

10 LC-MS (Method 3):  $R_t = 0.89$  min; MS (ESIpos):  $m/z = 252$  (M+H)<sup>+</sup>.

### Intermediate L55

Trifluoroacetic acid / tert-butyl-N<sup>6</sup>-D-alanyl-N<sup>2</sup>-{N-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-hexanoyl]-L-valyl-L-alanyl}-L-lysinate (1:1)



20 The title compound was prepared by first coupling Intermediate L6 with *N*-(tert-butoxycarbonyl)-D-alanine in the presence of HATU, followed by deprotection at the amino group under gentle



conditions by stirring for 90 minutes in 5% strength trifluoroacetic acid in DCM at RT.

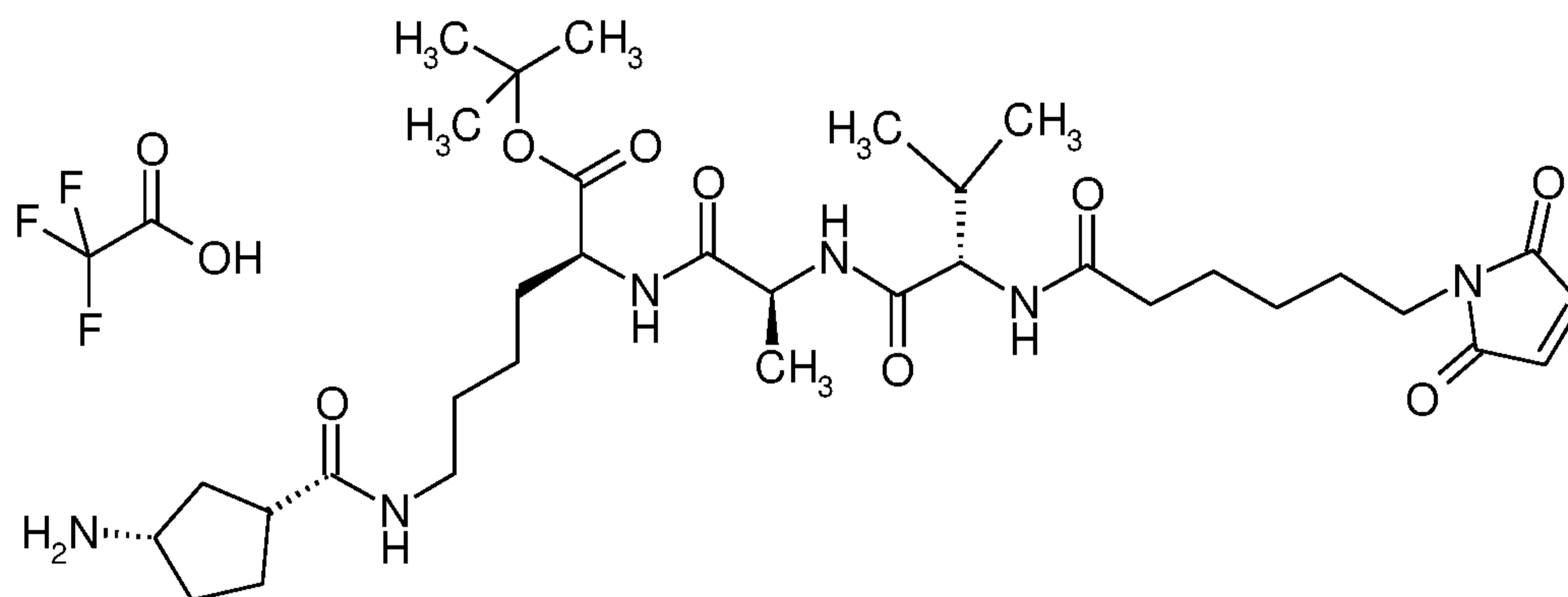
HPLC (Method 11):  $R_t = 1.35$  min;

5

LC-MS (Method 1):  $R_t = 0.67$  min; MS (ESIpos):  $m/z = 637$  (M+H)<sup>+</sup>.

### Intermediate L56

10 Trifluoroacetic acid / tert-butyl-N-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-L-alanyl-N<sup>6</sup>-{[(1R,3S)-3-aminocyclopentyl]carbonyl}-L-lysinate (1:1)



15

The title compound was prepared by first coupling Intermediate L6 with (1R,3S)-3-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid in the presence of HATU, followed by deprotection at the amino group under gentle conditions by stirring for 15 minutes in 25% strength trifluoroacetic acid in DCM at RT.

20

HPLC (Method 11):  $R_t = 1.4$  min;

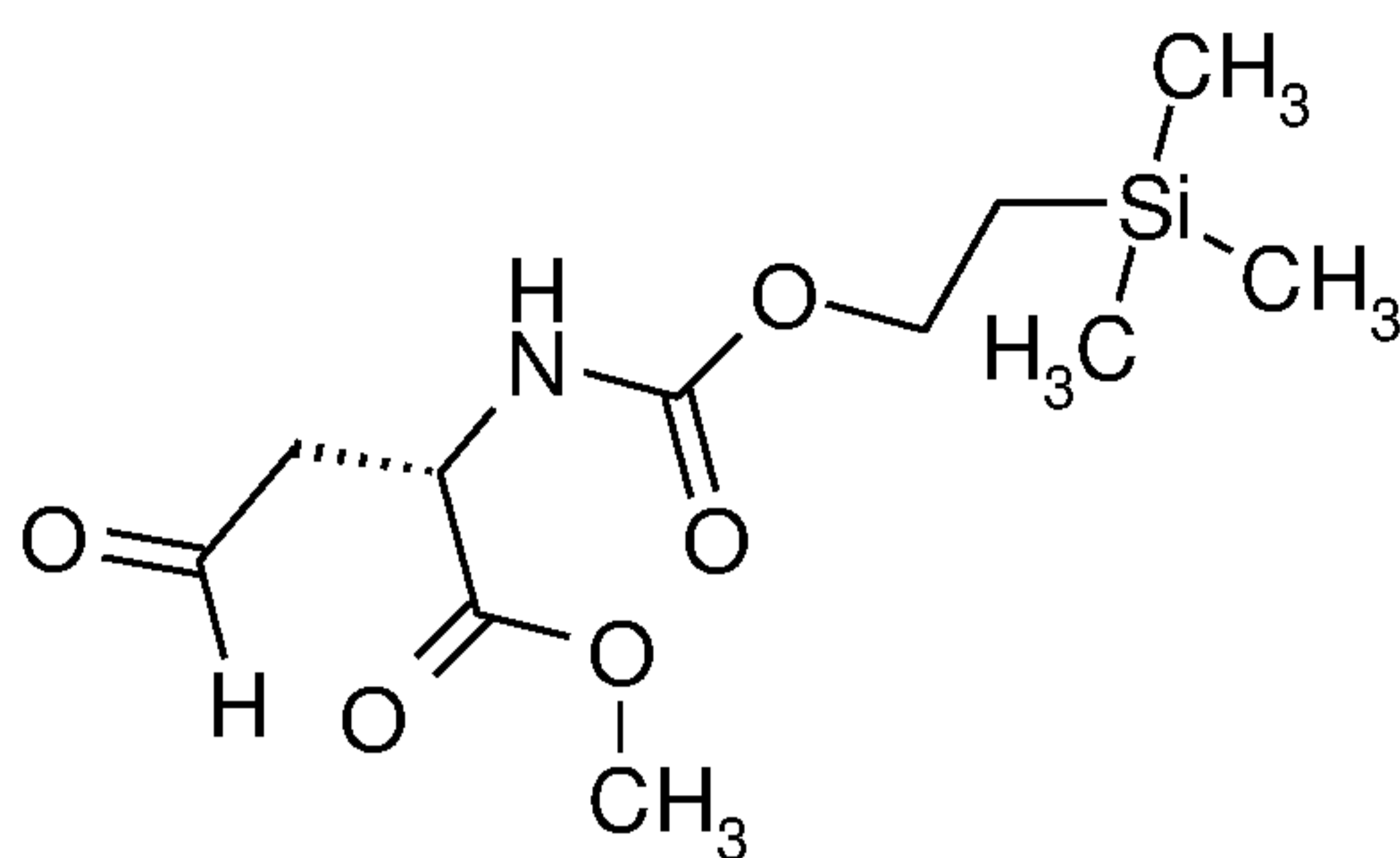
25

LC-MS (Method 1):  $R_t = 0.7$  min; MS (ESIpos):  $m/z = 677$  (M+H)<sup>+</sup>.

### Intermediate L57

Methyl (2S)-4-oxo-2-({[2-(trimethylsilyl)ethoxy]carbonyl}amino)butanoate

30



500.0 mg (2.72 mmol) of methyl L-asparaginate hydrochloride and 706.3 mg (2.72 mmol) of 2-(trimethylsilyl)ethyl 2,5-dioxopyrrolidine-1-carboxylate were initially charged in 5.0 ml of 1,4-dioxane, and 826.8 mg (8.17 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x40; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were then evaporated under reduced pressure and the residue was dried under high vacuum. This gave 583.9 mg (74% of theory) of the compound (3S)-4-methoxy-4-oxo-3-({[2-(trimethylsilyl)ethoxy]carbonyl}amino)butanoic acid.

LC-MS (Method 1):  $R_t$  = 0.89 min; MS (ESI<sub>neg</sub>):  $m/z$  = 290 (M-H)<sup>-</sup>.

592.9 mg of (3S)-4-methoxy-4-oxo-3-({[2-(trimethylsilyl)ethoxy]carbonyl}amino)butanoic acid were initially charged in 10.0 ml of 1,2-dimethoxyethane, the mixture was cooled to -15°C and 205.8 mg (2.04 mmol) of 4-methylmorpholine and 277.9 mg (2.04 mmol) of isobutyl chloroformate were added. The precipitate was filtered off after 15 min and twice with in each case 10.0 ml of 1,2-dimethoxyethane. The filtrate was cooled to -10°C, and 115.5 mg (3.05 mmol) of sodium borohydride dissolved in 10 ml of water were added with vigorous stirring. The phases were separated and the organic phase was washed in each case once with saturated sodium bicarbonate solution and saturated NaCl solution. The organic phase was dried over magnesium sulphate, the solvent was evaporated under reduced pressure and the residue was dried under high vacuum. This gave 515.9 mg (91% of theory) of the compound methyl N-([2-(trimethylsilyl)ethoxy]carbonyl)-L-homoserinate.

LC-MS (Method 1):  $R_t = 0.87$  min; MS (ESIpos):  $m/z = 278$  (M+H)<sup>+</sup>.

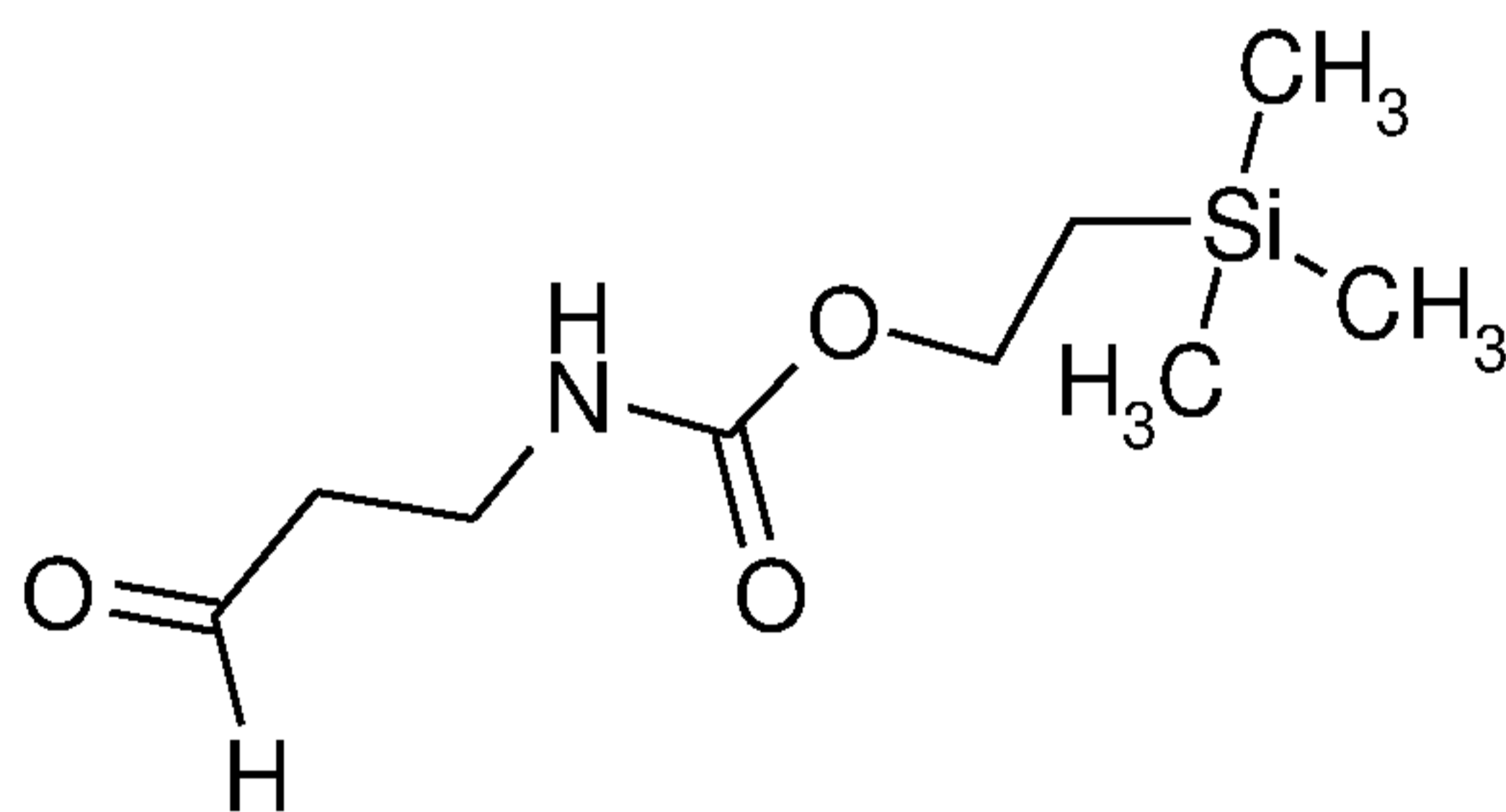
554.9 mg (2.00 mmol) of methyl N-{{[2-  
5 (trimethylsilyl)ethoxy]carbonyl}-L-homoserinate were initially  
charged in 30.0 ml of dichloromethane, and 1.27 g (3.0 mmol) of  
Dess-Martin periodinane and 474.7 mg (6.00 mmol) of pyridine  
were added. The mixture was stirred at RT overnight. After 4 h,  
the reaction was diluted with dichloromethane and the organic  
10 phase was washed in each case three times with 10% strength  
Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub> solution, 10% strength citric acid solution and saturated  
sodium bicarbonate solution. The organic phase was dried over  
magnesium sulphate and the solvent was evaporated under reduced  
pressure. This gave 565.7 mg (97% of theory) of the title  
15 compound.

<sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>):  $\delta$  [ppm] = 0.03 (s, 9H), 0.91 (m, 2H),  
2.70-2.79 (m, 1H), 2.88 (dd, 1H), 3.63 (s, 3H), 4.04 (m, 2H),  
4.55 (m, 1H), 7.54 (d, 1H), 9.60 (t, 1H).

20

### Intermediate L58

2-(Trimethylsilyl)ethyl (3-oxopropyl) carbamate



25

434.4 mg (5.78 mmol) of 3-amino-1-propanol and 1.50 g (5.78  
mmol) of 2-(trimethylsilyl)ethyl 2,5-dioxopyrrolidine-1-  
carboxylate were dissolved in 10.0 ml of dichloromethane, 585.3  
30 mg (5.78 mmol) of triethylamine were added and the mixture was  
stirred at RT overnight. The reaction mixture was diluted with  
dichloromethane and the organic phase was washed with water and  
saturated sodium bicarbonate solution and then dried over

magnesium sulphate. The solvent was evaporated under reduced pressure. The residue 2-(trimethylsilyl)ethyl (3-hydroxypropyl)carbamate (996.4 mg, 79% of theory) was dried under high vacuum and used without further purification in the next step of the synthesis.

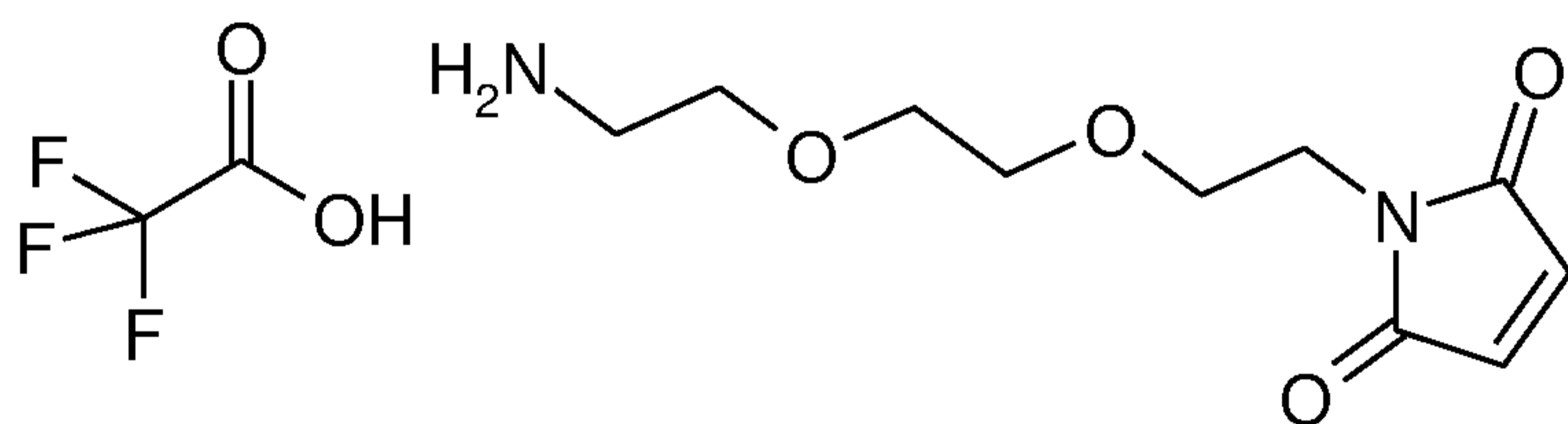
807.0 mg (3.68 mmol) of 2-(trimethylsilyl)ethyl (3-hydroxypropyl)carbamate were initially charged in 15.0 ml of chloroform and 15.0 ml of 0.05 N potassium carbonate/0.05 N sodium bicarbonate solution (1:1). 102.2 mg (0.37 mmol) of tetra-n-butylammonium chloride, 736.9 mg (5.52 mmol) of N-chlorosuccinimide and 57.5 mg (0.37 mmol) of TEMPO were then added and the reaction mixture was stirred vigorously at RT overnight. The reaction mixture was diluted with dichloromethane and the organic phase was washed with water and saturated NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was dried under high vacuum and used without further purification in the next step of the synthesis (890.3 mg).

20

### Intermediate L59

Trifluoroacetic acid / 1-{2-[2-(2-aminoethoxy)ethoxy]ethyl}-1H-pyrrole-2,5-dione (1:1)

25



300.0 mg (0.91 mmol) of tert-butyl (2-{2-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethoxy]ethoxy}ethyl)carbamate were initially charged in dichloromethane, 4.2 g (36.54 mmol) of TFA were added and the mixture was stirred at RT for 1 h (monitored by TLC: dichloromethane/methanol 10:1). The volatile components were evaporated under reduced pressure and the residue was co-distilled four times with dichloromethane. The residue was dried under high vacuum and used without further purification in the

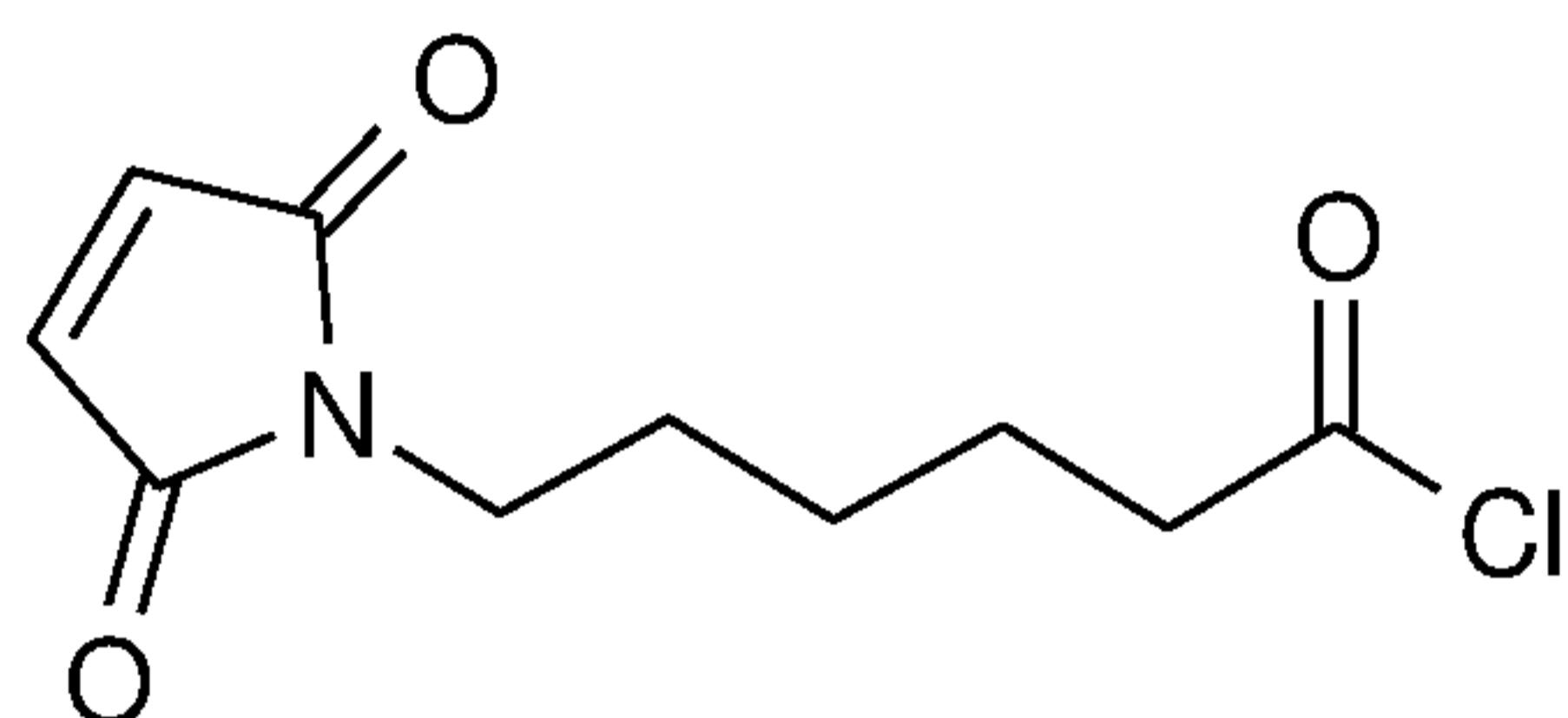
35

next step of the synthesis.

LC-MS (Method 1):  $R_t = 0.19$  min; MS (ESIpos):  $m/z = 229$  (M+H)<sup>+</sup>.

### 5 Intermediate L60

6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl chloride



10

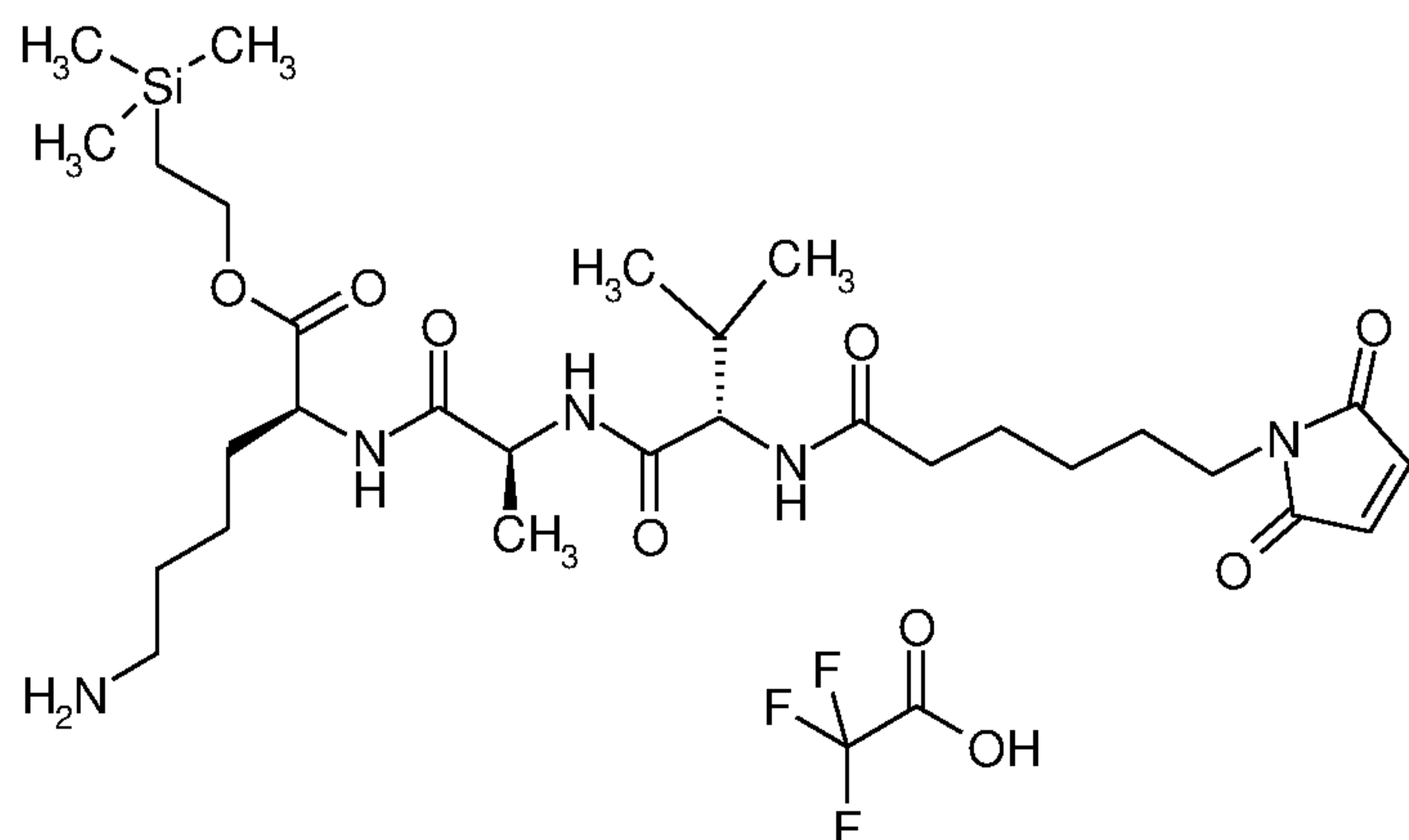
200.0 mg (0.95 mmol) of 6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoic acid were dissolved in 4.0 ml of dichloromethane, and 338.0 mg (2.84 mmol) of thionyl chloride were added. The reaction mixture was stirred at RT for 3 h, and 1 drop of DMF was then added. The mixture was stirred for another 1 h. The solvent was evaporated under reduced pressure and the residue was co-distilled three times with dichloromethane. The crude product was used without further purification in the next step of the synthesis.

20

### Intermediate L61

Trifluoroacetic acid / 2-(trimethylsilyl)ethyl-N-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-L-alanyl-L-lysinate (1:1)

25



First, the tripeptide derivative 2-(trimethylsilyl)ethyl-L-valyl-L-alanyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysinate was prepared from N<sup>2</sup>-[(benzyloxy) carbonyl]-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine  
5 according to classical methods of peptide chemistry (esterification with 2-(trimethylsilylethanol using EDCI/DMAP, hydrogenolysis, coupling with N-[(benzyloxy) carbonyl]-L-valyl-L-alanine in the presence of HATU and another hydrogenolysis). The title compound was prepared by coupling this partially  
10 protected peptide derivative with commercially available 6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoic acid in the presence of HATU and *N,N*-diisopropylethylamine. This was followed by deprotection at the amino group under gentle conditions by stirring for 2.5 hours in 5% strength trifluoroacetic acid in  
15 DCM at RT with retention of the ester protective group. Work-up and purification by preparative HPLC gave 438 mg of the title compound.

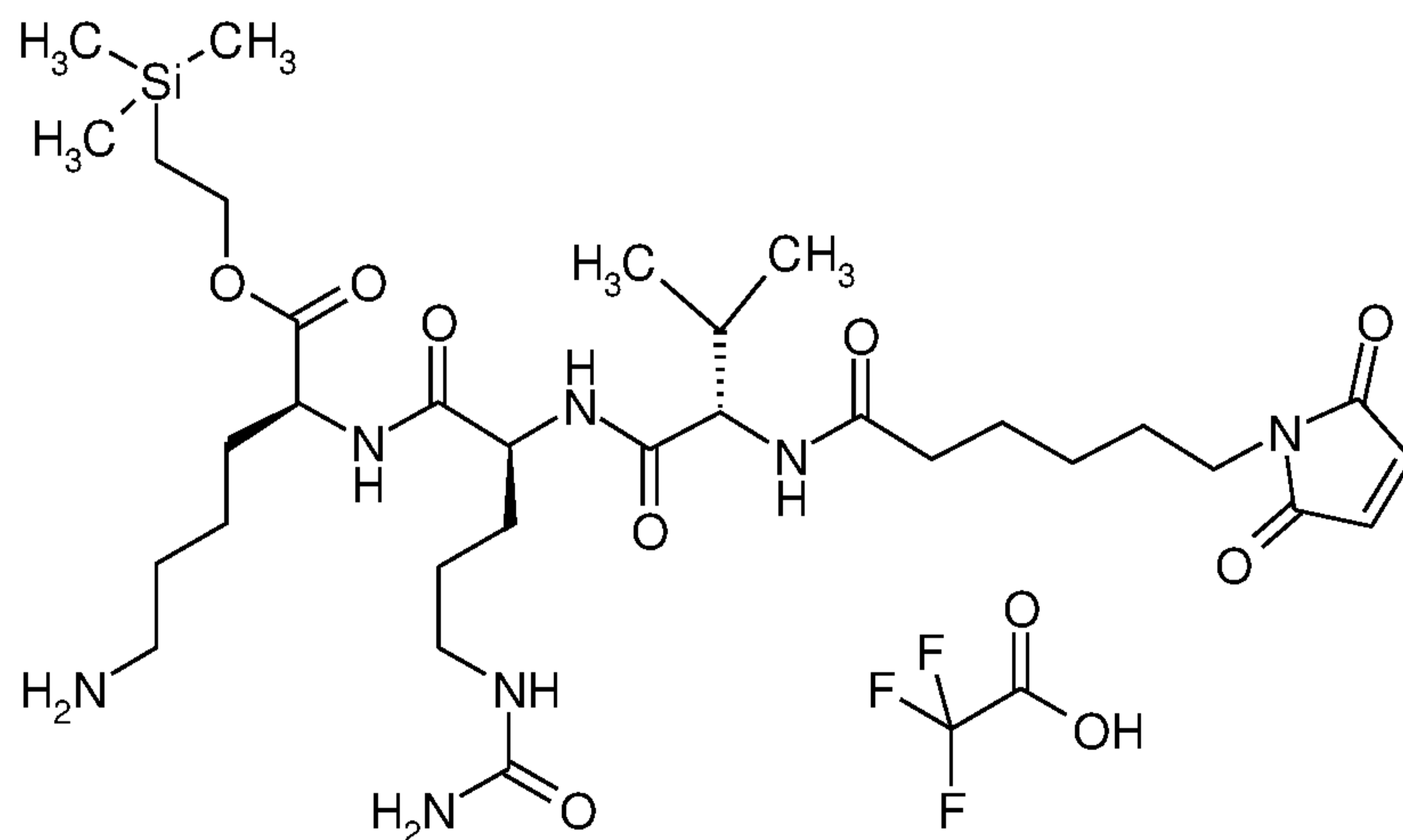
HPLC (Method 11):  $R_t = 1.69$  min;

20

LC-MS (Method 1):  $R_t = 0.78$  min; MS (ESIpos):  $m/z = 610$  (M+H)<sup>+</sup>.

### Intermediate L62

25 Trifluoroacetic acid / 2-(trimethylsilyl)ethyl-N-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithyl-L-lysinate (1:1)



First, 2-(trimethylsilyl)ethyl N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysinate was prepared from N<sup>2</sup>-[(benzyloxy)carbonyl]-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine according to classical methods of peptide chemistry. 148 mg (0.43 mmol) of this Intermediate were then coupled in the presence of 195 mg (0.51 mmol) of HATU and 149  $\mu$ l of *N,N*-diisopropylethylamine with 200 mg (0.43 mmol) of Intermediate L16. After concentration and purification of the residue by preparative HPLC, the protected intermediate was taken up in 20 ml of DCM and the tert-butoxycarbonyl protective group was removed by addition of 2 ml of trifluoroacetic acid and 1 h of stirring at RT. Concentration and lyophilization of the residue from acetonitrile/water gave 254 mg (63% of theory over 2 steps).

15

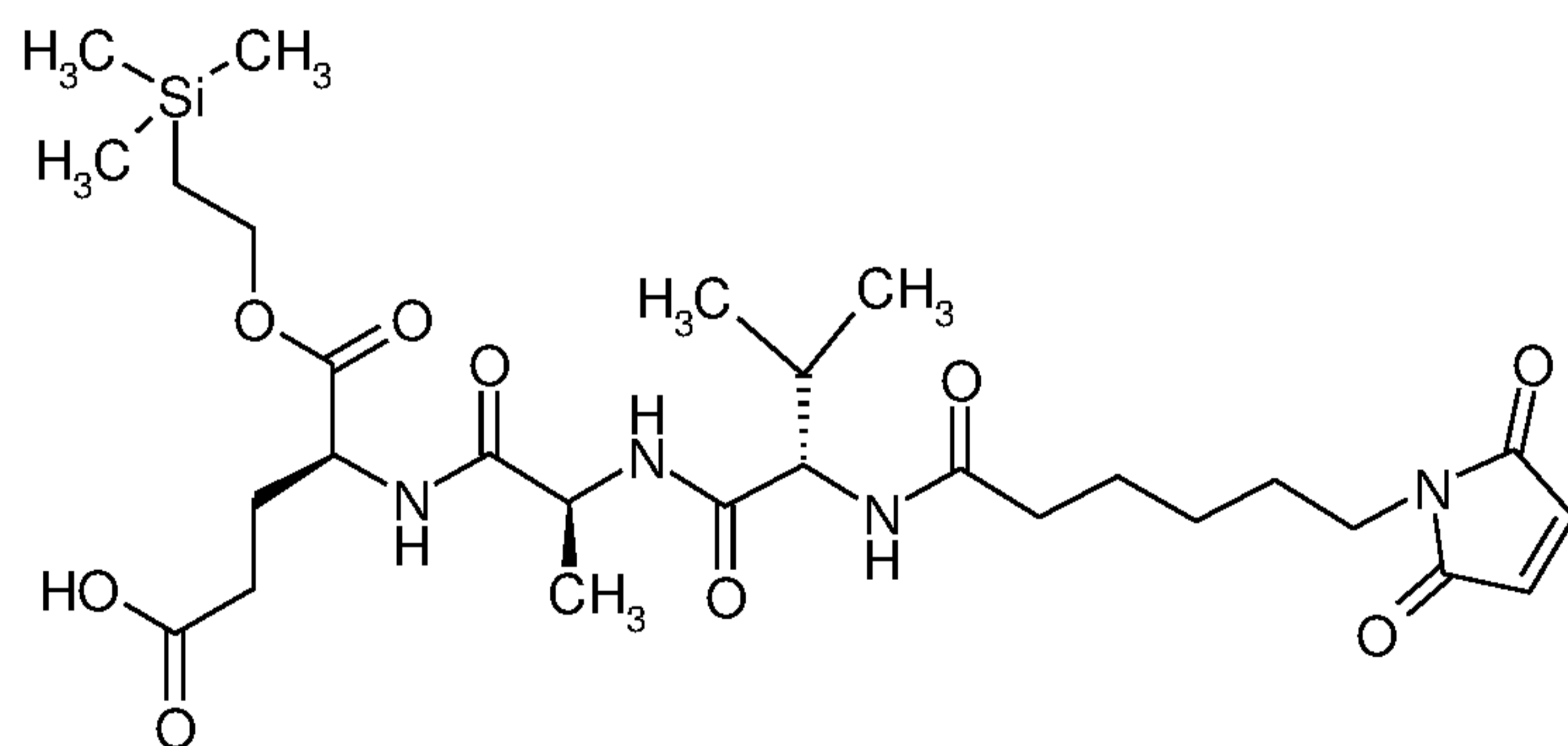
HPLC (Method 11):  $R_t = 1.51$  min;

LC-MS (Method 1):  $R_t = 0.68$  min; MS (ESIpos):  $m/z = 696$  (M+H)<sup>+</sup>.

### 20 Intermediate L63

(4S)-4-{[(2S)-2-{[(2S)-2-{[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]amino}-3-methylbutanoyl]amino}propanoyl]amino}-5-oxo-5-[2-(trimethylsilyl)ethoxy]pentanoic acid

25



First, the tripeptide derivative (4S)-4-{[(2S)-2-{[(2S)-2-amino-3-methylbutanoyl]amino}propanoyl]amino}-5-oxo-5-[2-(trimethylsilyl)ethoxy]pentanoic acid was prepared from (2S)-5-(benzyloxy)-2-[(tert-butoxycarbonyl)amino]-5-oxopentanoic acid according to classical methods of peptide chemistry

30

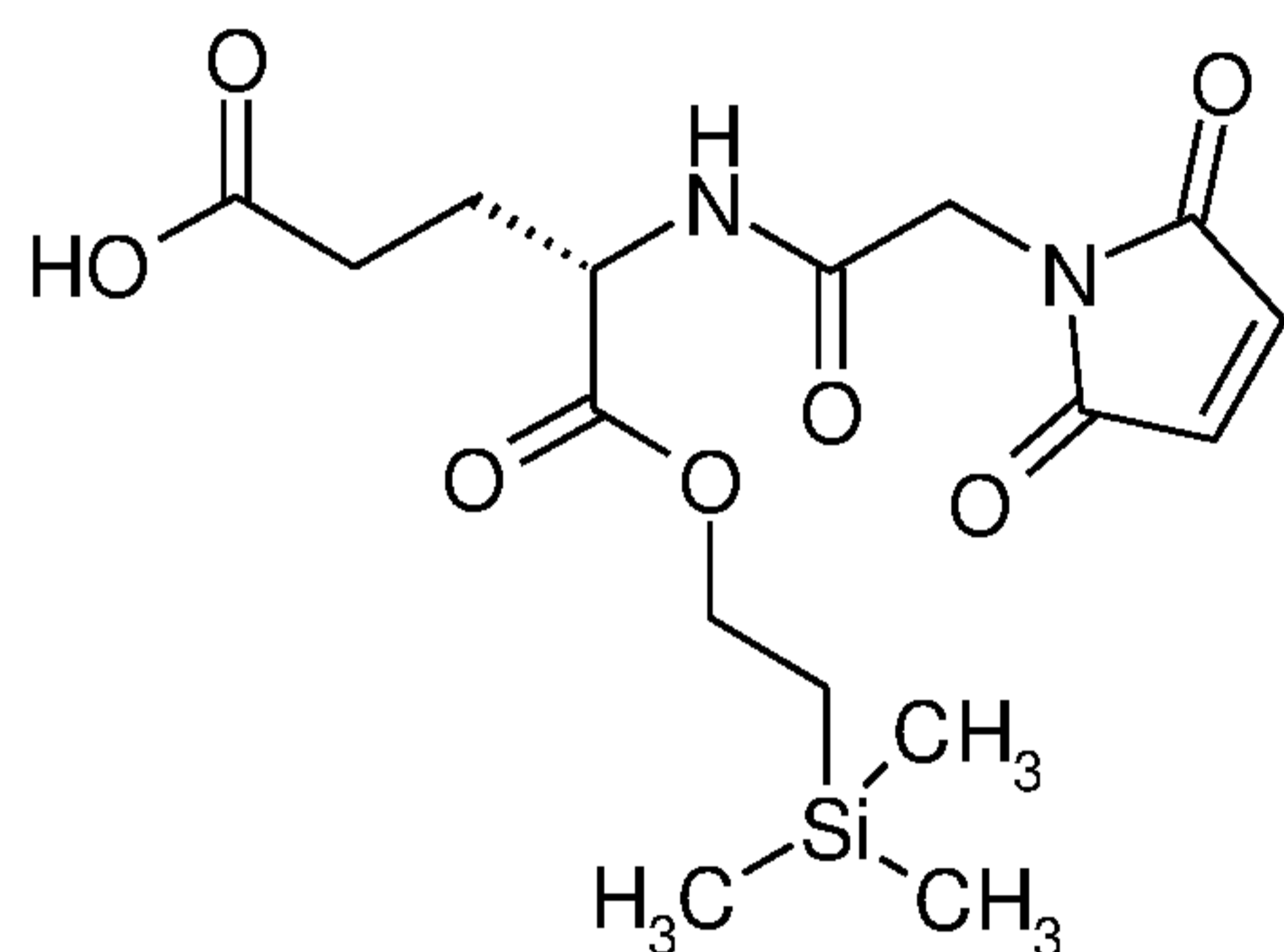
(esterification with 2-(trimethylsilylethanol using EDCI/DMAP, removal of the Boc protective group with trifluoroacetic acid, coupling with N-[(benzyloxy)carbonyl]-L-valyl-L-alanine in the presence of HATU and hydrogenolysis in methanol over 10% palladium on activated carbon). The title compound was prepared by coupling of this partially protected peptide derivative with commercially available 1-{6-[(2,5-dioxopyrrolidin-1-yl)oxy]-6-oxohexyl}-1H-pyrrole-2,5-dione. Work-up and purification by preparative HPLC gave 601 mg of the title compound.

10

LC-MS (Method 1):  $R_t = 0.96$  min; MS (ESIpos):  $m/z = 611$  (M+H)<sup>+</sup>.

#### Intermediate L64

15 (4S)-4-{[(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino}-5-oxo-5-[2-(trimethylsilyl)ethoxy]pentanoic acid



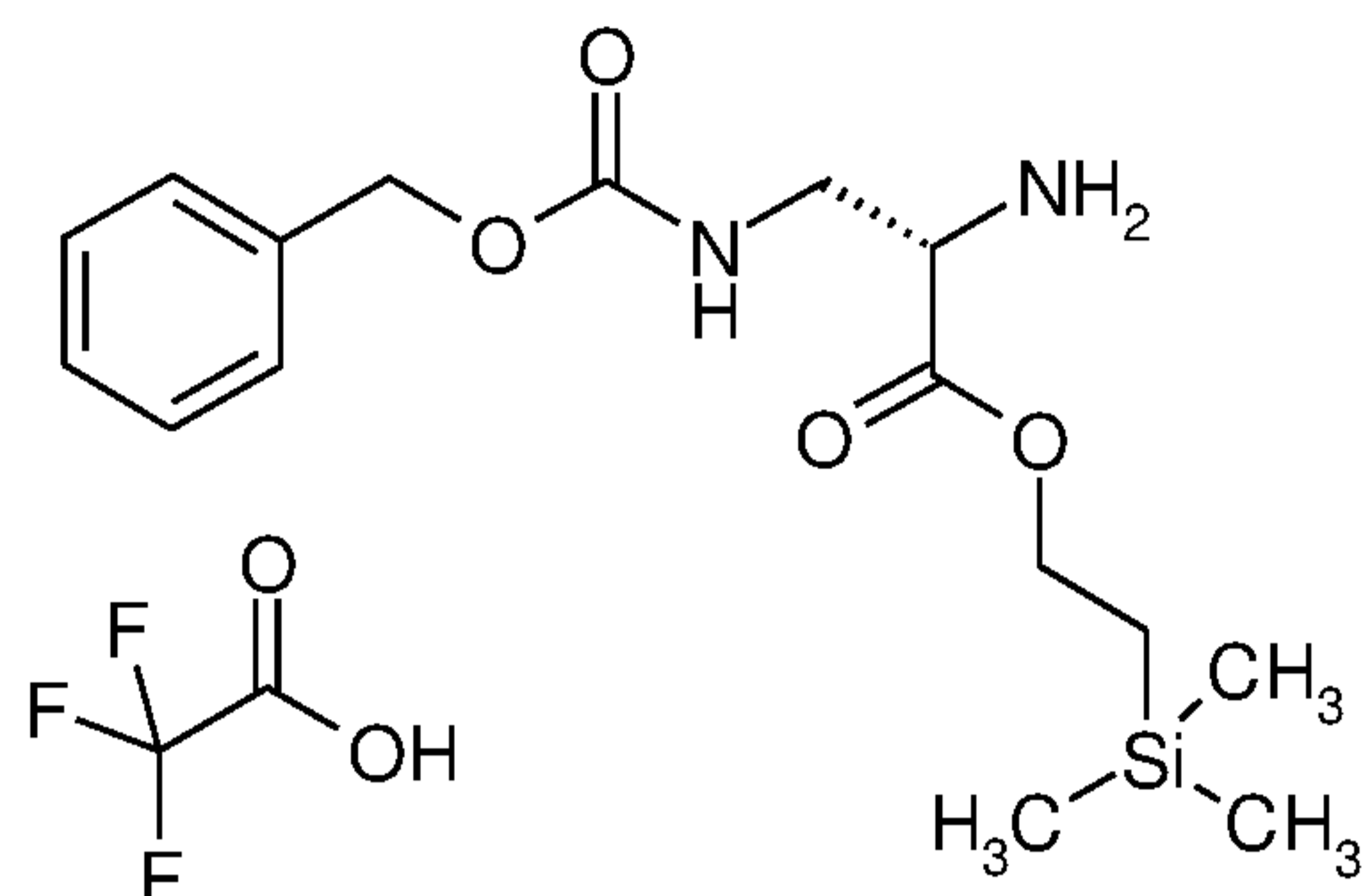
20 The title compound was prepared from (2S)-5-(benzyloxy)-2-[(tert-butoxycarbonyl)amino]-5-oxopentanoic acid according to classical methods of peptide chemistry (esterification with 2-(trimethylsilylethanol using EDCI/DMAP, removal of the Boc protective group with trifluoroacetic acid, hydrogenolytic cleavage of the benzyl ester in methanol over 10% palladium on activated carbon and coupling with 1-{2-[(2,5-dioxopyrrolidin-1-yl)oxy]-2-oxoethyl}-1H-pyrrole-2,5-dione in the presence of N,N-diisopropylethylamine).

25 LC-MS (Method 1):  $R_t = 0.84$  min; MS (ESIpos):  $m/z = 385$  (M+H)<sup>+</sup>.

#### Intermediate L65



Trifluoroacetic acid / 2-(trimethylsilyl)ethyl-3-  
 {[(benzyloxy)carbonyl]amino}-L-alaninate (1:1)



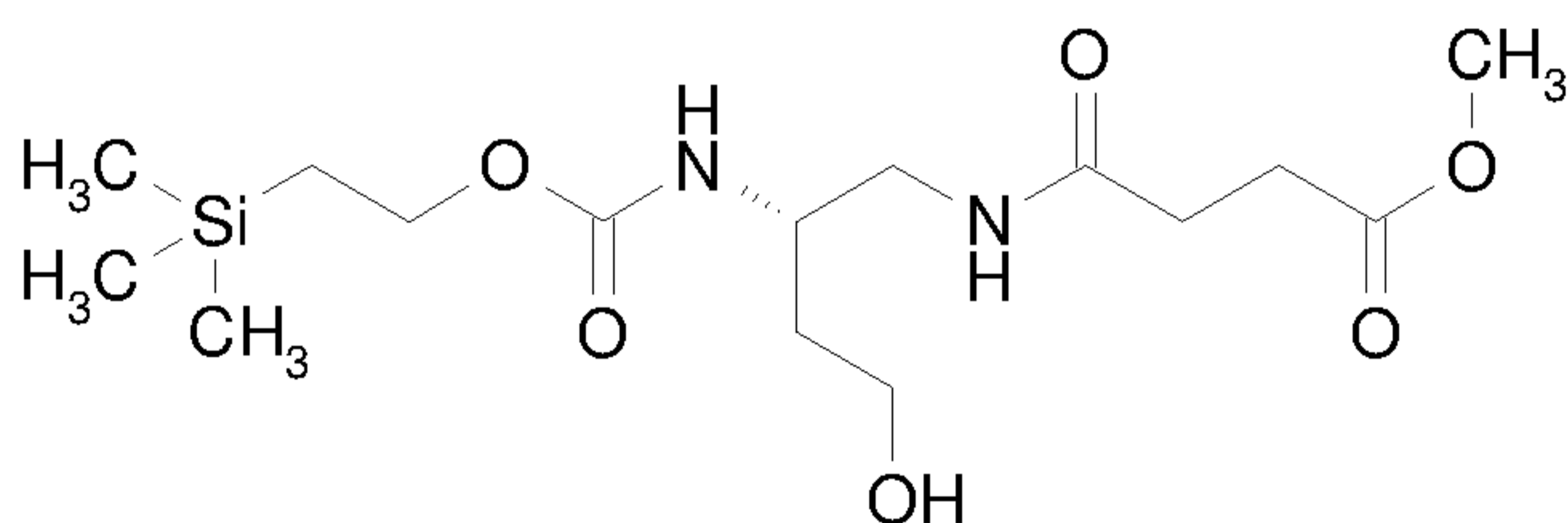
5

The title compound was prepared from 3-  
 {[(benzyloxy)carbonyl]amino}-N-(tert-butoxycarbonyl)-L-alanine  
 according to classical methods of peptide chemistry  
 10 (esterification with 2-(trimethylsilyl)ethanol using EDCI/DMAP  
 and removal of the Boc protective group with trifluoroacetic  
 acid. This gave 373 mg (79% of theory over 2 steps) of the title  
 compound.

15 LC-MS (Method 1):  $R_t = 0.72$  min; MS (ESIpos):  $m/z = 339$  (M+H)<sup>+</sup>.

### Intermediate L66

Methyl (8S)-8-(2-hydroxyethyl)-2,2-dimethyl-6,11-dioxo-5-oxa-  
 20 7,10-diaza-2-silatetradecan-14-oate



1000 mg (2.84 mmol) of (3S)-3-{[(benzyloxy)carbonyl]amino}-4-  
 25 [(tert-butoxycarbonyl)amino]butanoic acid were initially  
 charged in 10.0 ml of 1,2-dimethoxyethane, and 344.4 mg (3.4  
 mmol) of 4-methylmorpholine and 504 mg (3.69 mmol) of isobutyl  
 chloroformate were added. After 10 min of stirring at RT, the  
 reaction was cooled to 5°C and 161 mg (4.26 mmol) of sodium

borohydride dissolved in 3 ml of water were added a little at a time with vigorous stirring. After 1 h, the same amount of sodium borohydride was added again and the reaction was then slowly warmed to RT. 170 ml of water were added and the reaction was then extracted four times with in each case 200 ml of ethyl acetate. The phases were separated and the organic phase was washed once with citric acid and then with saturated sodium bicarbonate solution. The organic phase was dried over magnesium sulphate, the solvent was evaporated under reduced pressure and the residue was dried under high vacuum. This gave 760 mg (78% of theory) of the compound benzyl tert-butyl [(2S)-4-hydroxybutane-1,2-diyl]biscarbamate.

LC-MS (Method 1):  $R_t = 0.84$  min; MS (ESIpos):  $m/z = 339$  (M+H)<sup>+</sup>.

760 mg (2.16 mmol) of this intermediate dissolved in 13 ml of hydrogen chloride/dioxane were stirred at RT for 20 min. The reaction was then concentrated to 5 ml, and diethyl ether was added. The precipitate was filtered off and lyophilized from acetonitrile/water 1:1.

The product obtained in this manner was dissolved in 132 ml of DMF, and 345.5 mg (2.35 mmol) of 4-methoxy-4-oxobutanoic acid, 970 mg (2.55 mmol) of HATU and 1025  $\mu$ l of *N,N*-diisopropylethylamine were added. The mixture was stirred at RT for 5 min. The solvent was removed under reduced pressure and the residue that remained was purified by preparative HPLC. The appropriate fractions were combined and the acetonitrile was evaporated under reduced pressure. The aqueous phase that remained was extracted twice with ethyl acetate and the organic phase was then concentrated and dried under high vacuum.

The intermediate obtained in this manner was taken up in methanol and hydrogenated over 10% palladium on activated carbon at RT under hydrogen standard pressure for 1 h. The catalyst was then filtered off and the solvent was removed under reduced pressure.

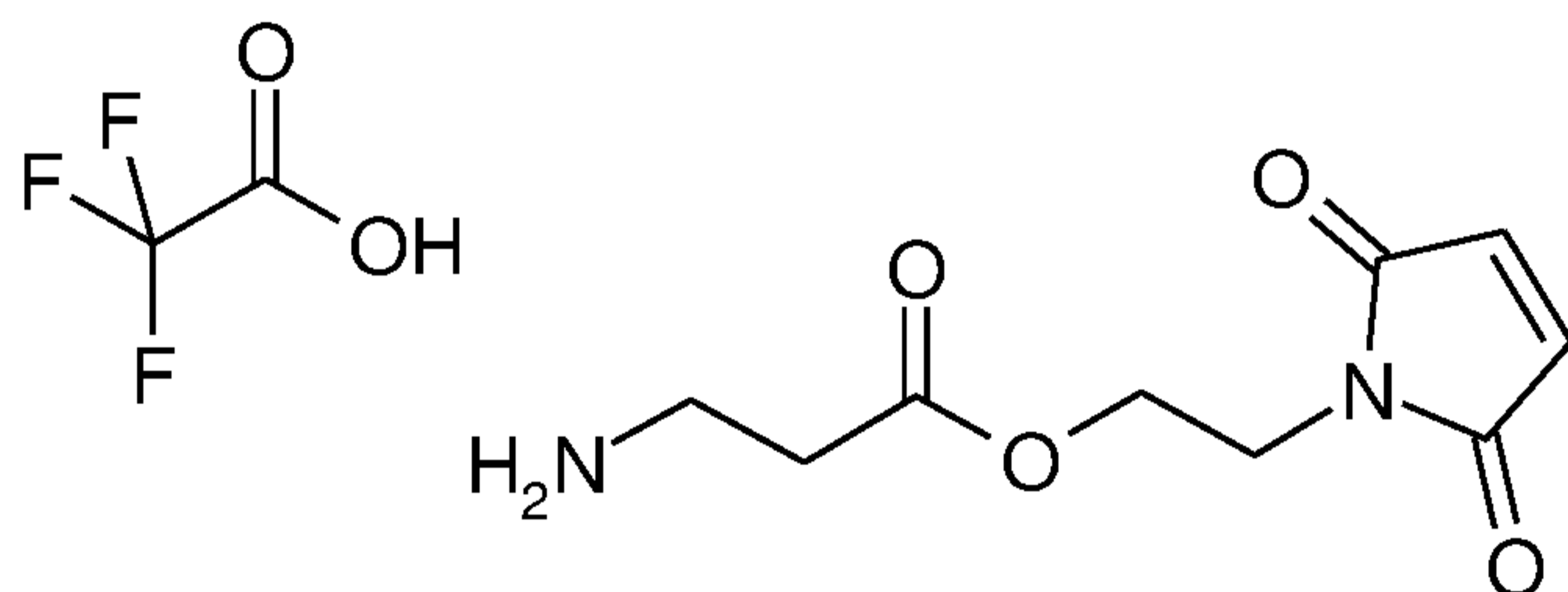
247 mg of this deprotected compound were taken up in 20 ml of

DMF, and 352 mg (1.36 mmol) of 1-([2-(trimethylsilyl)ethoxy]carbonyloxy)pyrrolidine-2,5-dione and 592  $\mu$ l of *N,N*-diisopropylethylamine were added. The reaction mixture was stirred at RT for 1 h and then concentrated, and the residue was purified by preparative HPLC. The solvents were then evaporated under reduced pressure and the residue was dried under high vacuum. This gave, over these 5 reaction steps, 218 mg of the title compound in a total yield of 21%.

10 LC-MS (Method 1):  $R_t$  = 0.74 min; MS (ESIpos):  $m/z$  = 363 (M+H)<sup>+</sup>.

### Intermediate L67

15 Trifluoroacetic acid / 2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl-beta-alaninate (1:1)



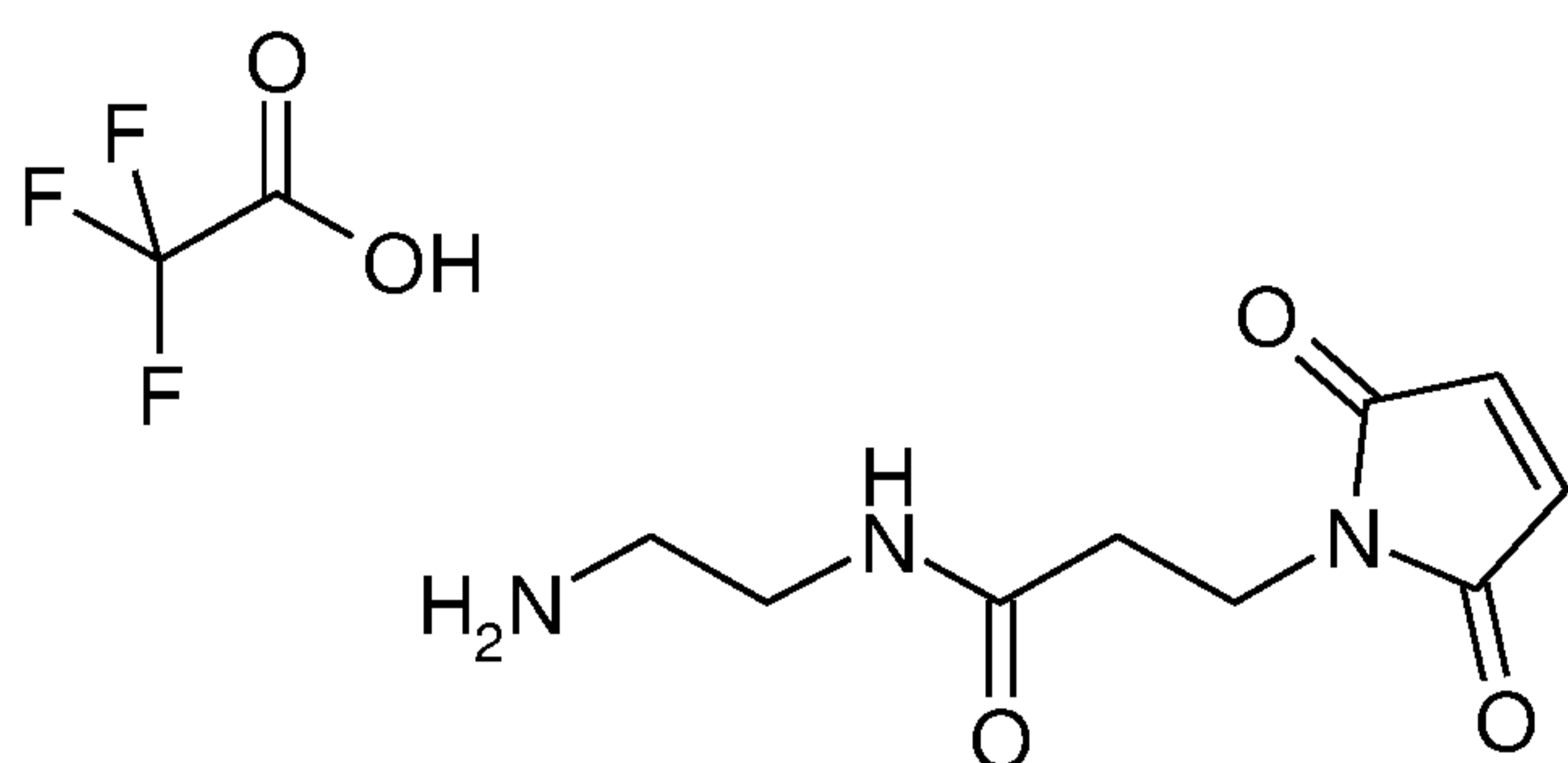
20 The title compound was prepared from 50 mg (0.354 mmol) of commercially available 1-(2-hydroxyethyl)-1H-pyrrole-2,5-dione by coupling with 134 mg (0.71 mmol) of *N*-(tert-butoxycarbonyl)-beta-alanine in 10 ml of dichloromethane in the presence of 1.5 equivalents of EDCI and 0.1 equivalent of 4-*N,N*-dimethylaminopyridine and subsequent deprotection with trifluoroacetic acid.

Yield: 56 mg (48% of theory over 2 stages)

30 LC-MS (Method 3):  $R_t$  = 1.15 min; MS (ESIpos):  $m/z$  = 213 (M+H)<sup>+</sup>.

### Intermediate L68

Trifluoroacetic acid / *N*-(2-aminoethyl)-2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)propanamide (1:1)



The title compound was prepared analogously to Intermediate L1  
 5 according to classical methods of peptide chemistry from  
 commercially available (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-  
 yl)propanoic acid and tert-butyl (2-aminoethyl)carbamate.

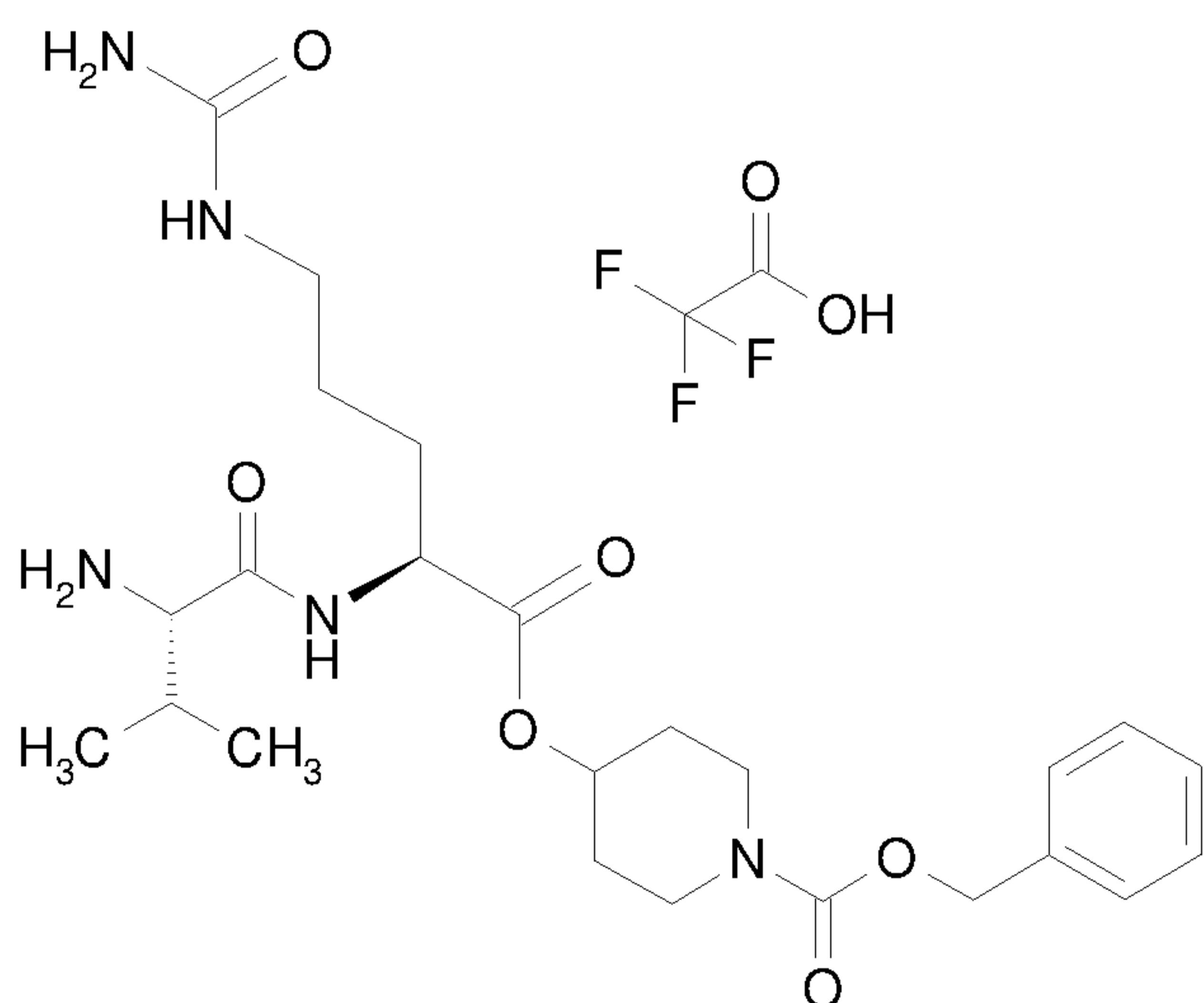
LC-MS (Method 1):  $R_t = 0.17$  min; MS (ESIpos):  $m/z = 212$  (M+H)<sup>+</sup>.

10

### Intermediate L69

Trifluoroacetic acid / 1-[(benzyloxy)carbonyl]piperidin-4-yl-L-  
 15 valyl-N<sup>5</sup>-carbamoyl-L-ornithinate (1:1)

15



The title compound was prepared by classical methods of peptide  
 chemistry from commercially available benzyl 4-  
 20 hydroxypiperidine-1-carboxylate by esterification with N<sup>2</sup>-(tert-  
 butoxycarbonyl)-N<sup>5</sup>-carbamoyl-L-ornithine using EDCI/DMAP,  
 subsequent Boc removal with TFA, followed by coupling with N-  
 [(tert-butoxy)carbonyl]-L-valine in the presence of HATU and

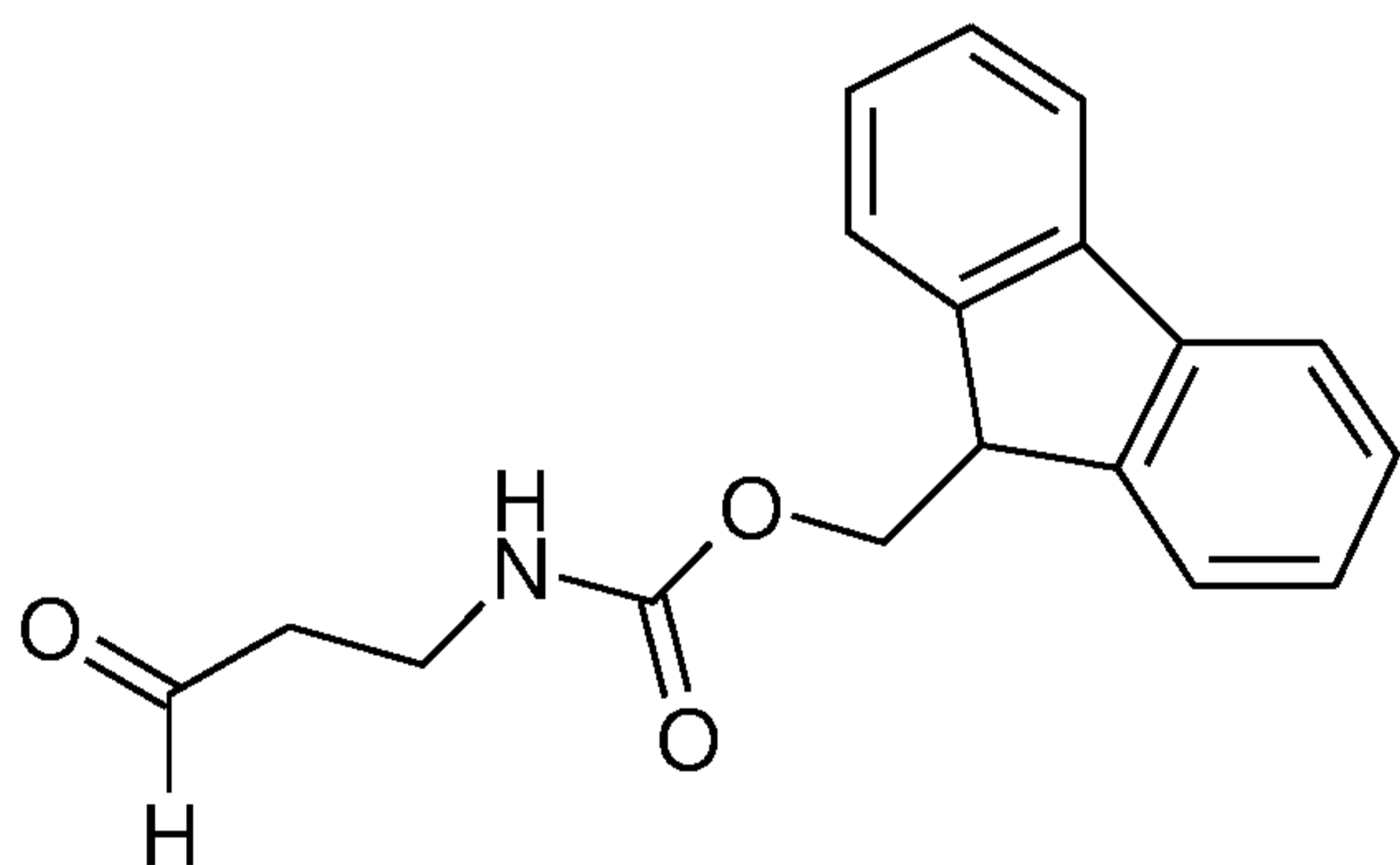
*N,N*-diisopropylethylamine and finally another Boc removal with TFA.

LC-MS (Method 1):  $R_t = 0.62$  min; MS (ESIpos):  $m/z = 492$  (M+H)<sup>+</sup>.

5

### Intermediate L70

9H-Fluoren-9-ylmethyl (3-oxopropyl) carbamate



10

1000.0 mg (3.36 mmol) of 9H-fluoren-9-ylmethyl (3-hydroxypropyl)carbamate were initially charged in 15.0 ml of chloroform and 15.0 ml of 0.05 N potassium carbonate/0.05 N sodium bicarbonate solution (1:1). 93.5 mg (0.34 mmol) of tetra-n-butylammonium chloride, 673.6 mg (5.04 mmol) of N-chlorosuccinimide and 52.5 mg (0.34 mmol) of TEMPO were then added and the reaction mixture was stirred vigorously at RT overnight. The reaction mixture was diluted with dichloromethane and the organic phase was washed with water and saturated NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was dried under high vacuum and purified on silica gel (mobile phase: cyclohexane/ethyl acetate 3:1-1:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 589.4 mg (58% of theory) of the title compound.

20

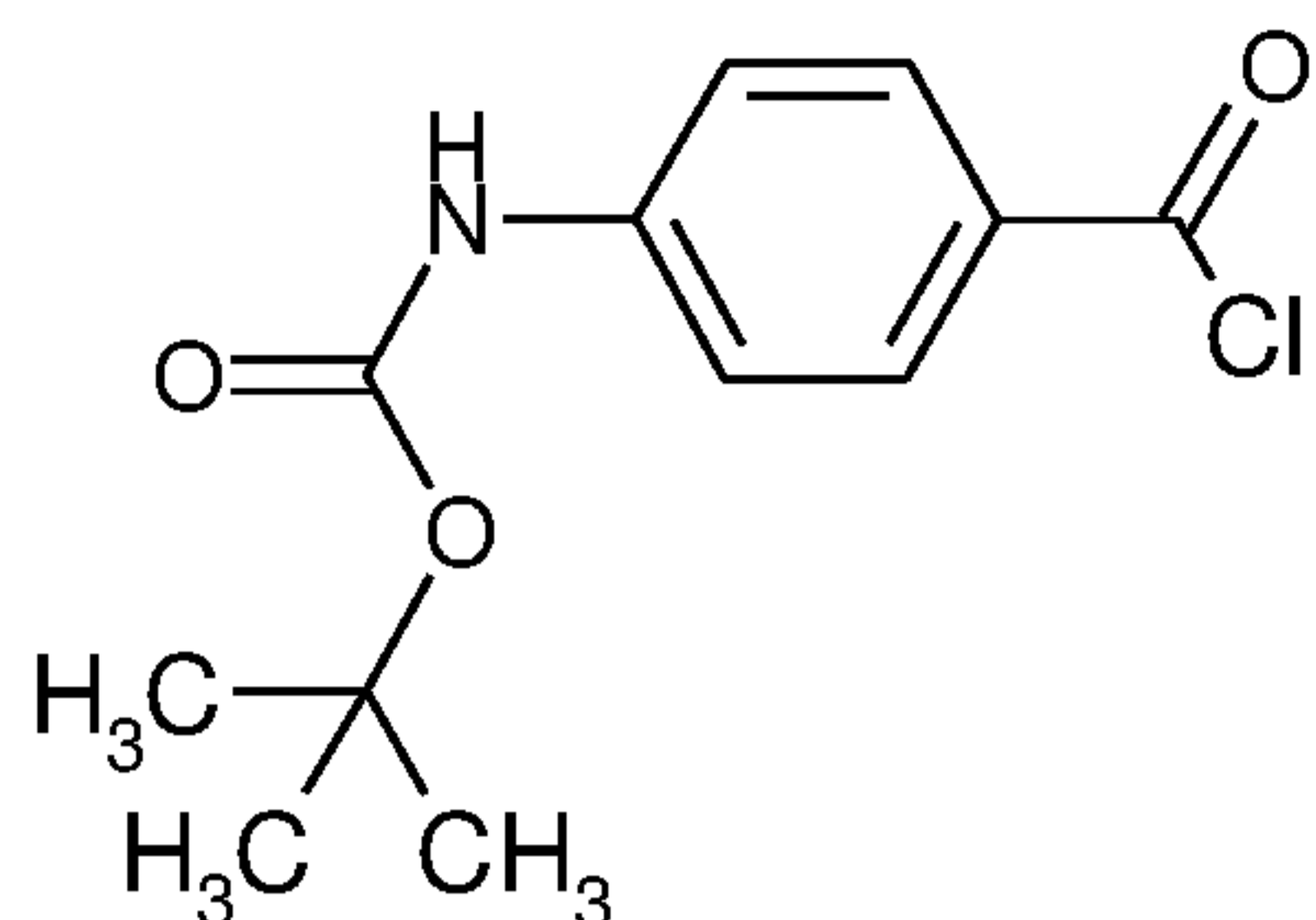
25

LC-MS (Method 6):  $R_t = 2.15$  min; MS (ESIpos):  $m/z = 296$  (M-H)<sup>+</sup>.

30

### Intermediate L71

tert-Butyl [4-(chlorocarbonyl)phenyl]carbamate



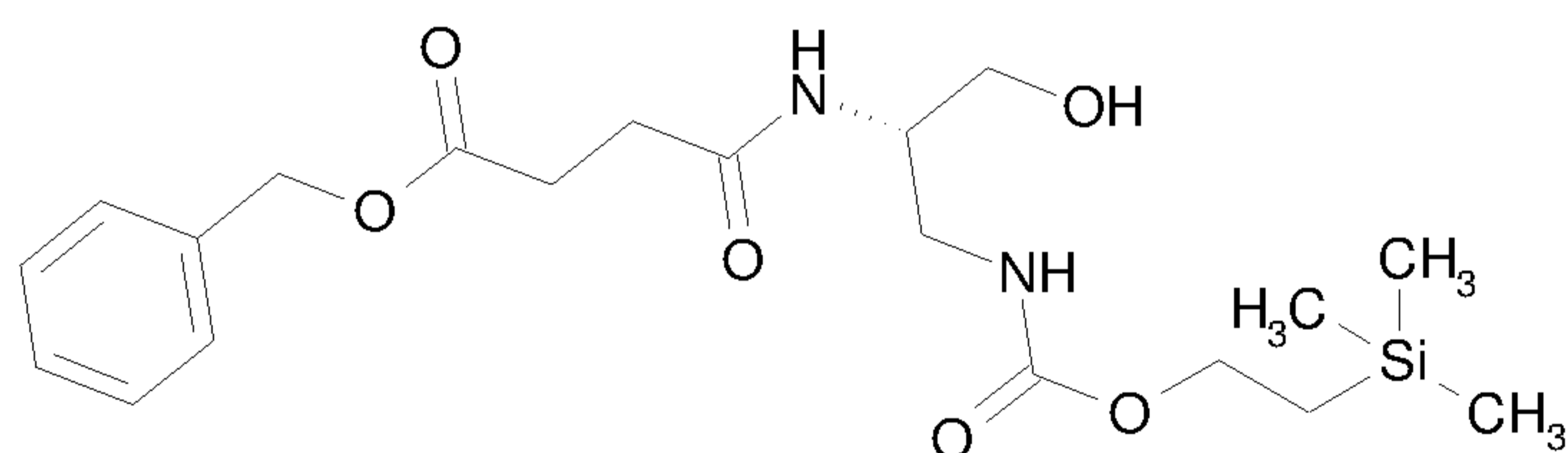
5 100.0 mg (0.42 mmol) of 4-[(tert-butoxycarbonyl)amino]benzoic acid were initially charged in 2.0 ml of dichloromethane, and 64.2 mg (0.51 mmol) of oxalyl chloride were added. The reaction mixture was stirred at RT for 30 min (monitored by TLC: dichloromethane/methanol). Another 192.6 mg (1.53 mmol) of  
 10 oxalyl chloride and 1 drop of DMF were then added and the mixture was stirred at RT for 1 h. The solvent was evaporated under reduced pressure and the residue was co-distilled repeatedly with dichloromethane. The residue was used without further purification in the next step of the synthesis.

15

### Intermediate L72

Benzyl (9S)-9-(hydroxymethyl)-2,2-dimethyl-6,11-dioxo-5-oxa-7,10-diaza-2-silatetradecan-14-oate

20



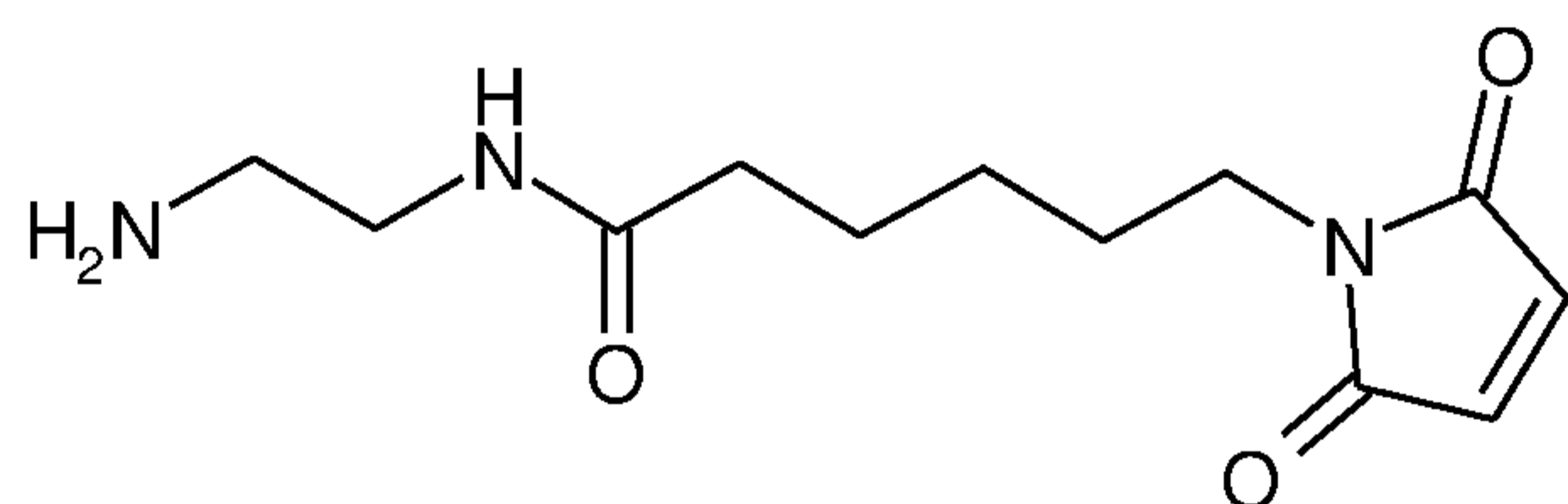
The title compound was prepared from commercially available benzyl tert-butyl [(2S)-3-hydroxypropan-1,2-diyl]biscarbamate  
 25 according to classical methods of peptide chemistry by hydrogenolytic removal of the Z protective group, subsequent coupling with 4-(benzyloxy)-4-oxobutanoic acid in the presence of EDCI/HOBT, followed by removal of the Boc protective group with TFA and finally by reaction with 1-({[2-  
 30 (trimethylsilyl)ethoxy]carbonyl}oxy)pyrrolidine-2,5-dione in

the presence of triethylamine.

LC-MS (Method 1):  $R_t = 0.94$  min; MS (ESIpos):  $m/z = 425$   $[M+H]^+$ .

## 5 Intermediate L73

N-(2-Aminoethyl)-6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanamide



10

395.5 mg (1.87 mmol) of 6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoic acid, 1.21 g (9.36 mmol) of *N,N*-diisopropylethylamine and 854.3 mg (2.25 mmol) of HATU were added to a solution of 300 mg (1.87 mmol) of *tert*-butyl (2-aminoethyl)carbamate in 20 ml of dimethylformamide. The reaction mixture was stirred at RT for 5 minutes. After concentration of the mixture, the residue was taken up in DCM and washed with water. The organic phase was washed with brine, dried over magnesium sulphate, filtered off and concentrated. This gave 408 mg (33%, purity 53%) of the title compound which were used without further purification.

LC-MS (Method 1):  $R_t = 0.75$  min; MS (ESIpos):  $m/z = 354$   $(M+H)^+$ .

25

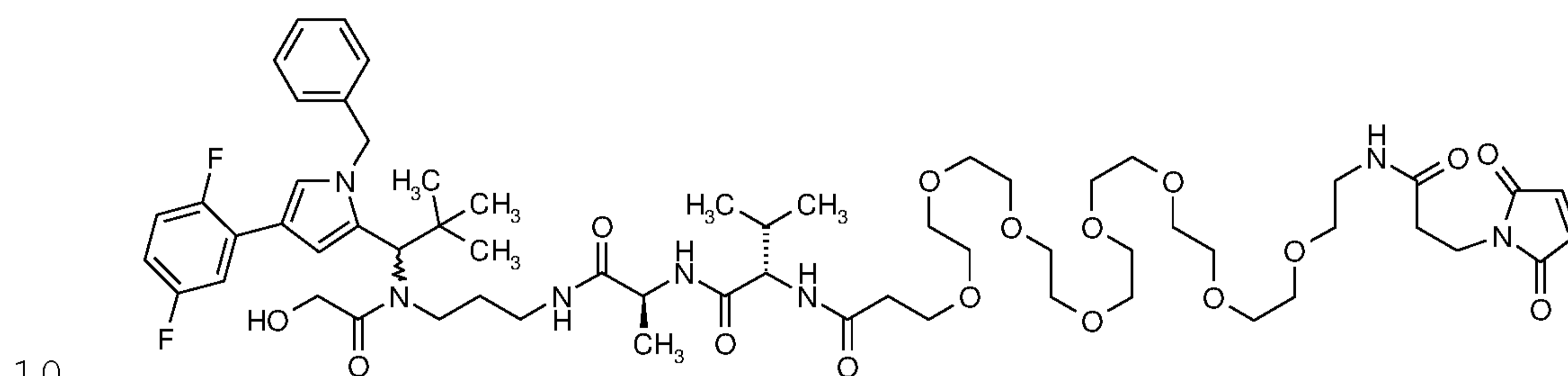
1 ml of TFA was added to a solution of *tert*-butyl (2-{{[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]amino}ethyl)carbamate (408 mg, 0.365 mmol) of in 7 ml of dichloromethane. The reaction mixture was stirred at RT for 0.5 h. The reaction mixture was concentrated under reduced pressure and the residue was co-distilled twice with dichloromethane. The residue was used further without further purification. This gave 384 mg (94%, purity 57%) of the title compound.

35

LC-MS (Method 1):  $R_t = 0.26$  min; MS (ESIpos):  $m/z = 254$  (M+H)<sup>+</sup>.

### Intermediate F82

5 N-[31-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-29-oxo-4,7,10,13,16,19,22,25-octaoxa-28-azahentriacontan-1-oyl]-L-valyl-N-{3-[[1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide



The synthesis of the title compound was carried out analogously to the synthesis of Intermediate F83. The racemic intermediates used were obtained analogously to the corresponding R-isomer intermediates.

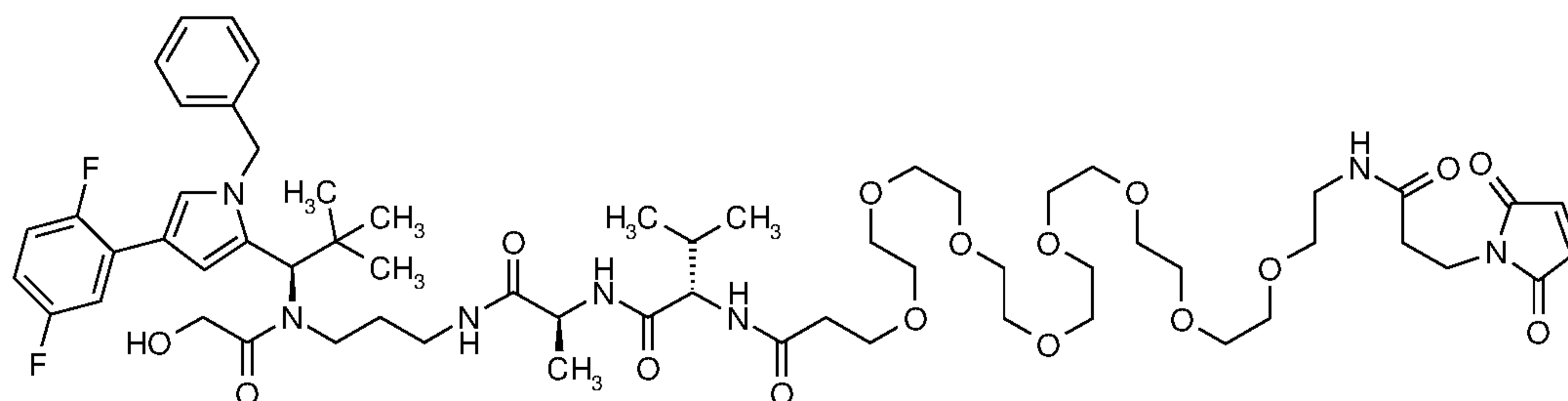
15 intermediates.

LC-MS (Method 2):  $R_t = 7.07$  min; MS (EIpos):  $m/z = 1236$  [M+Na]<sup>+</sup>.

### Intermediate F83

20 N-[31-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-29-oxo-4,7,10,13,16,19,22,25-octaoxa-28-azahentriacontan-1-oyl]-L-valyl-N-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide

25



30.0 mg (0.06 mmol) of N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-



(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide (Example 98) and 26.1 mg (0.06 mmol) of 2,5-dioxopyrrolidin-1-yl-N-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-alaninate were initially charged in 2.0 ml of DMF, and 19.4 mg (0.19 mmol) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight, and 11.5 mg (0.19 mmol) of HOAc were added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 41.9 mg (79% of theory) of the compound 9H-fluoren-9-ylmethyl [(2S)-1-({3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}amino)-1-oxopropan-2-yl]carbamate.

LC-MS (Method 1):  $R_t$  = 1.44 min; MS (ESIpos):  $m/z$  = 763 [M+H]<sup>+</sup>.

37.2 mg (0.05 mmol) of 9H-fluoren-9-ylmethyl [(2S)-1-({3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}amino)-1-oxopropan-2-yl]carbamate were dissolved in 1.5 ml of DMF, and 124.6 mg (1.46 mmol) of 2-aminoethanol were added. The reaction mixture was stirred at RT overnight. The reaction mixture was partitioned between ethyl acetate and water and the organic phase was washed twice with water and once with saturated NaCl solution. After drying over magnesium sulphate, the solvent was evaporated under reduced pressure and the residue was purified on silica gel (mobile phase: dichloromethane/methanol 10:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 14.2 mg (50% of theory) of the compound N-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide.

LC-MS (Method 1):  $R_t$  = 0.92 min; MS (ESIpos):  $m/z$  = 541 [M+H]<sup>+</sup>.

14.1 mg (0.03 mmol) of N-{3-[(1R)-1-[1-benzyl-4-(2,5-

difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide and 11.4 (0.03 mmol) of 2,5-dioxopyrrolidin-1-yl N-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-valinate were dissolved in 1.5 ml of DMF, and 7.9 mg (0.08 mmol) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight, and 4.7 mg (0.08 mmol) of HOAc were added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 15.9 mg (71% of theory) of the compound N-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-valyl-N-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide.

LC-MS (Method 1):  $R_t$  = 1.46 min; MS (ESIpos):  $m/z$  = 862 (M+H)<sup>+</sup>.

14.9 mg (0.02 mmol) of N-[(9H-fluoren-9-ylmethoxy)carbonyl]-L-valyl-N-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide were dissolved in 1.5 ml of DMF, and 44.2 mg (0.52 mmol) of 2-aminoethanol were added. The reaction mixture was stirred at RT overnight. The reaction mixture was partitioned between ethyl acetate and water and the organic phase was washed twice with water and once with saturated NaCl solution. After drying over magnesium sulphate, the solvent was evaporated under reduced pressure and the residue was purified on silica gel (mobile phase: dichloromethane/methanol 10:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 5.7 mg (52% of theory) of the compound L-valyl-N-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide.

LC-MS (Method 1):  $R_t$  = 0.92 min; MS (ESIpos):  $m/z$  = 640 (M+H)<sup>+</sup>.

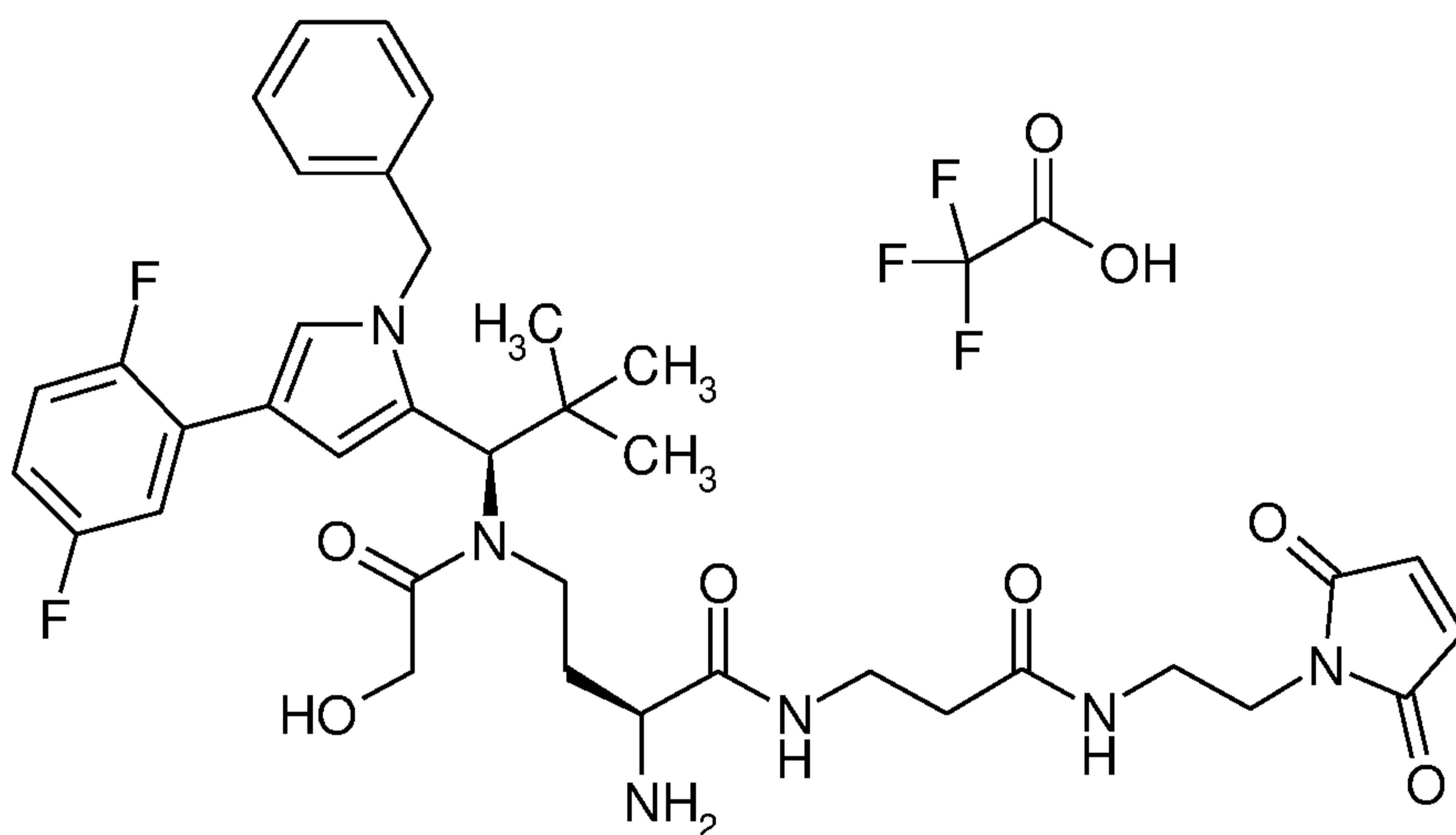
5.5 mg (8.6  $\mu$ mol) of L-valyl-N-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-

dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide and 6.5 mg (6.5  $\mu$ mol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{27-[(2,5-dioxopyrrolidin-1-yl)oxy]-27-oxo-3,6,9,12,15,18,21,24-octaoxaheptacos-1-yl}propanamide were dissolved in 1.0 ml of DMF, and 0.9 mg (8.6 mmol) of 4-methylmorpholine was added. The reaction mixture was stirred at RT overnight, and 0.8 mg (0.01 mmol) of HOAc was added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 7.7 mg (74% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 1.10 min; MS (ESIpos):  $m/z$  = 1214 (M+H)<sup>+</sup>.

#### Intermediate F84

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-(3-{[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]amino}-3-oxopropyl)butanamide (1:1)



25

First, 16.5 mg (0.02 mmol) of Intermediate C54 were taken up in 5 ml of DMF and reacted with 10.4 mg (0.041 mmol) of trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) in the presence of 11.7 mg (0.03 mmol) of HATU and 18  $\mu$ l

of *N,N*-diisopropylethylamine. After 5 min of stirring at RT, the mixture was concentrated and the residue was taken up in acetonitrile/water 1:1. The pH was adjusted to 2 with trifluoroacetic acid and the reaction was concentrated again.  
5 The residue that remained was purified by preparative HPLC. This gave 8 mg (42% of theory) of the protected intermediate.

LC-MS (Method 1):  $R_t = 1.38$  min; MS (EIpos):  $m/z = 929$  [M+H]<sup>+</sup>.

10 7.6 mg (0.008 mmol) of this intermediate were taken up in 3 ml of DMF, and 92 mg (0.82 mmol) of 1,4-diazabicyclo[2.2.2]octane were added. The reaction was treated in an ultrasonic bath for 1 h. 31  $\mu$ l of acetic acid were then added and the reaction was concentrated under high vacuum. The residue was purified by  
15 preparative HPLC. This gave 3 mg (45% of theory) of the title compound.

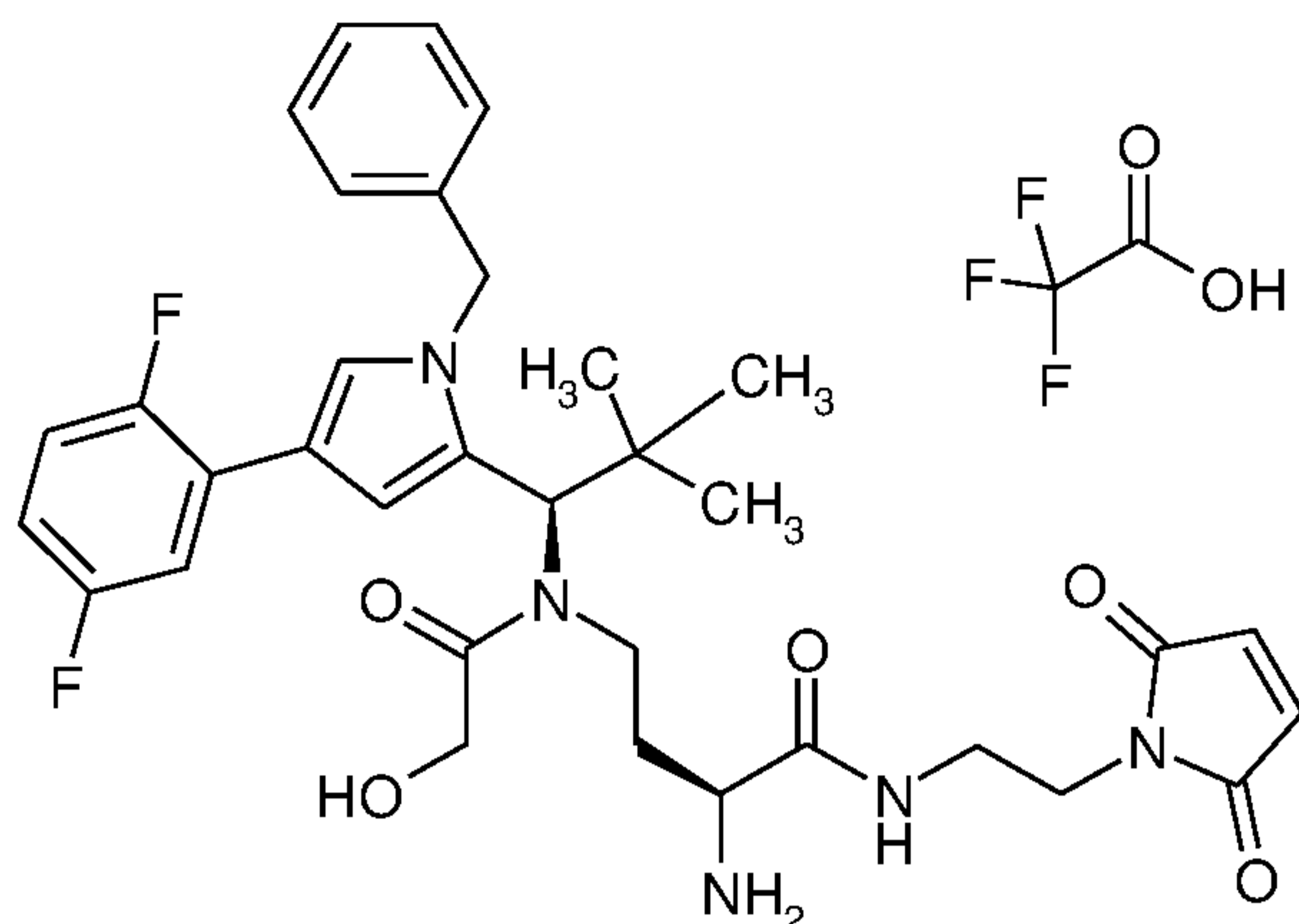
LC-MS (Method 1):  $R_t = 0.86$  min; MS (EIpos):  $m/z = 707$  [M+H]<sup>+</sup>.

20 <sup>1</sup>H NMR (500 MHz, DMSO-*d*<sub>6</sub>):  $\delta = 8.15$  (t, 1H), 7.9-8.1 (m, 4H), 7.6 (m, 1H), 7.5 (s, 1H), 7.15-7.35 (m, 6H), 6.9-7.0 (m, 3H), 6.85 (s, 1H), 5.6 (s, 1H), 4.9 and 5.2 (2d, 2H), 4.05 and 4.2 (2d, 2H), 3.1-3.2 (m, 4H), 2.15 (m, 2H), 0.7 and 1.45 (2m, 2H), 0.8 (s, 9H).

25

### Intermediate F85

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]butanamide (1:1)  
30



First, 10 mg (0.014 mmol) of Intermediate C53 were taken up in 3.4 ml of DMF and reacted with 7 mg (0.027 mmol) of trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) in the presence of 7.8 mg (0.02 mmol) of HATU and 12  $\mu$ l of *N,N*-diisopropylethylamine. After 15 min of stirring at RT, the mixture was concentrated and the residue was purified by preparative HPLC. This gave 6.6 mg (57% of theory) of the protected intermediate.

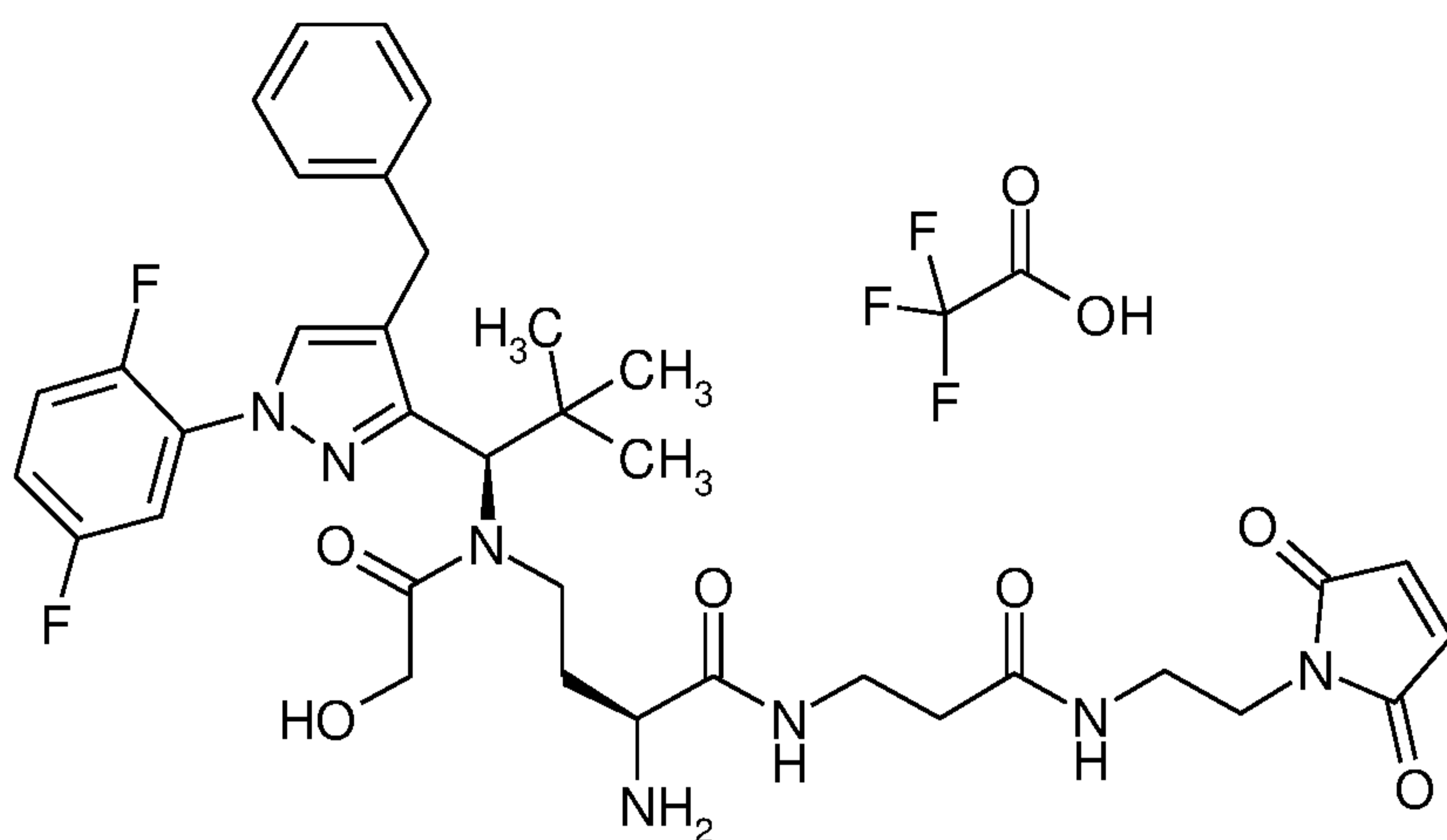
LC-MS (Method 1):  $R_t$  = 1.4 min; MS (EIpos):  $m/z$  = 858  $[M+H]^+$ .

6.6 mg (0.008 mmol) of this intermediate were taken up in 2 ml of DMF, and 86 mg (0.77 mmol) of 1,4-diazabicyclo[2.2.2]octane were added. The reaction was treated in an ultrasonic bath for 2 h. 44  $\mu$ l of acetic acid were then added and the reaction was concentrated under high vacuum. The residue was purified by preparative HPLC. This gave 3.3 mg (53% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.88 min; MS (EIpos):  $m/z$  = 636  $[M+H]^+$ .

### Intermediate F86

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]-N-(3-{[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]amino}-3-oxopropyl)butanamide (1:1)

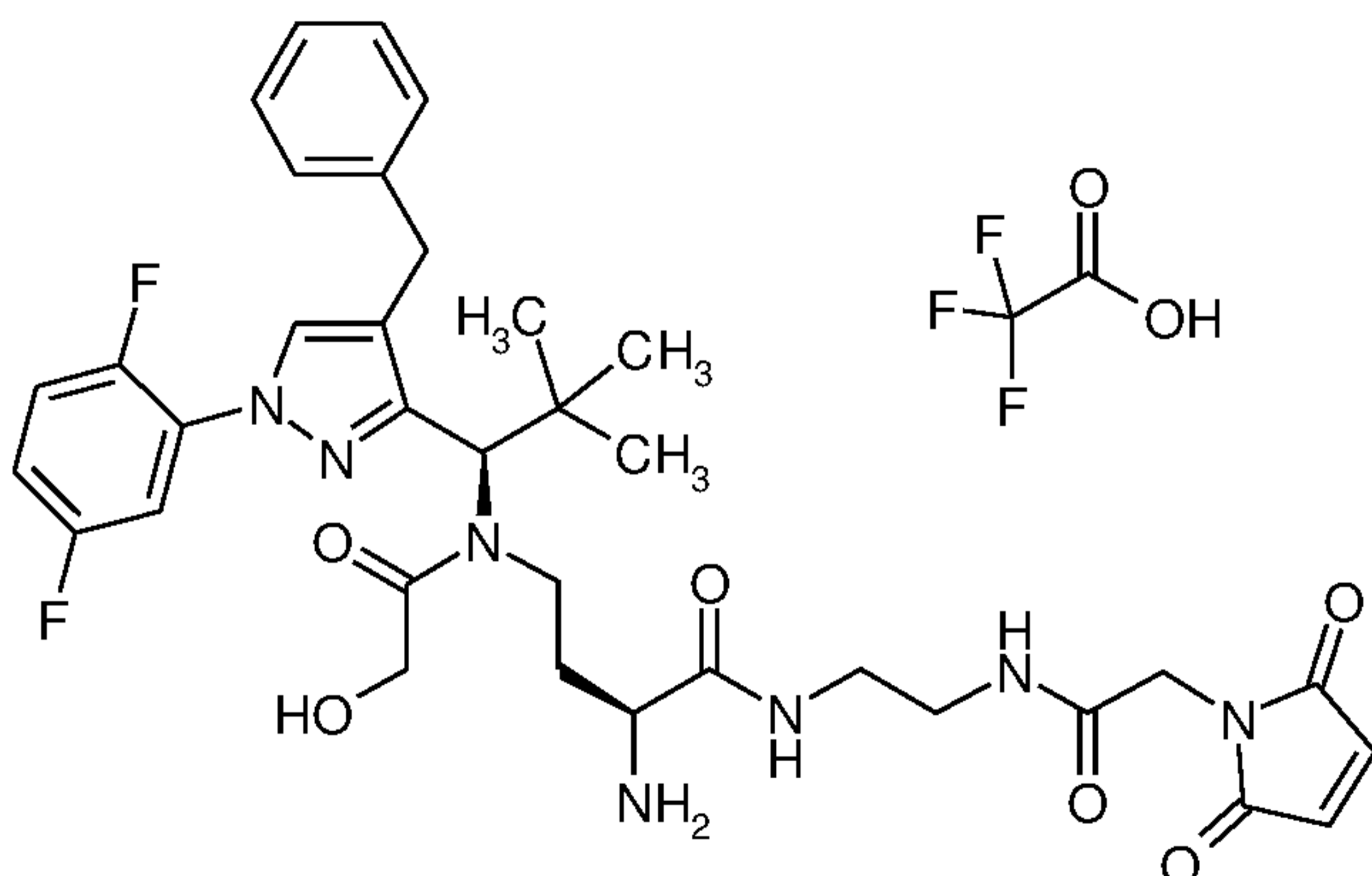


The title compound was prepared from 8 mg (0.012 mmol) of  
 5 Intermediate C51 by reaction with 4.5 mg (0.017 mmol) of  
 trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione  
 (1:1) in the presence of 5.8 mg (0.015 mmol) of HATU and 10  $\mu$ l  
 of *N,N*-diisopropylethylamine and subsequent deprotection with  
 trifluoroacetic acid. This gave 7 mg (78% of theory over 2  
 10 steps).

LC-MS (Method 1):  $R_t$  = 0.83 min; MS (EIpos):  $m/z$  = 708  $[M+H]^+$ .

### Intermediate F87

15 Trifluoroacetic acid / (2*S*)-2-amino-4-[(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-*N*-(3-{[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]amino}-3-oxopropyl)butanamide  
 20 (1:1)

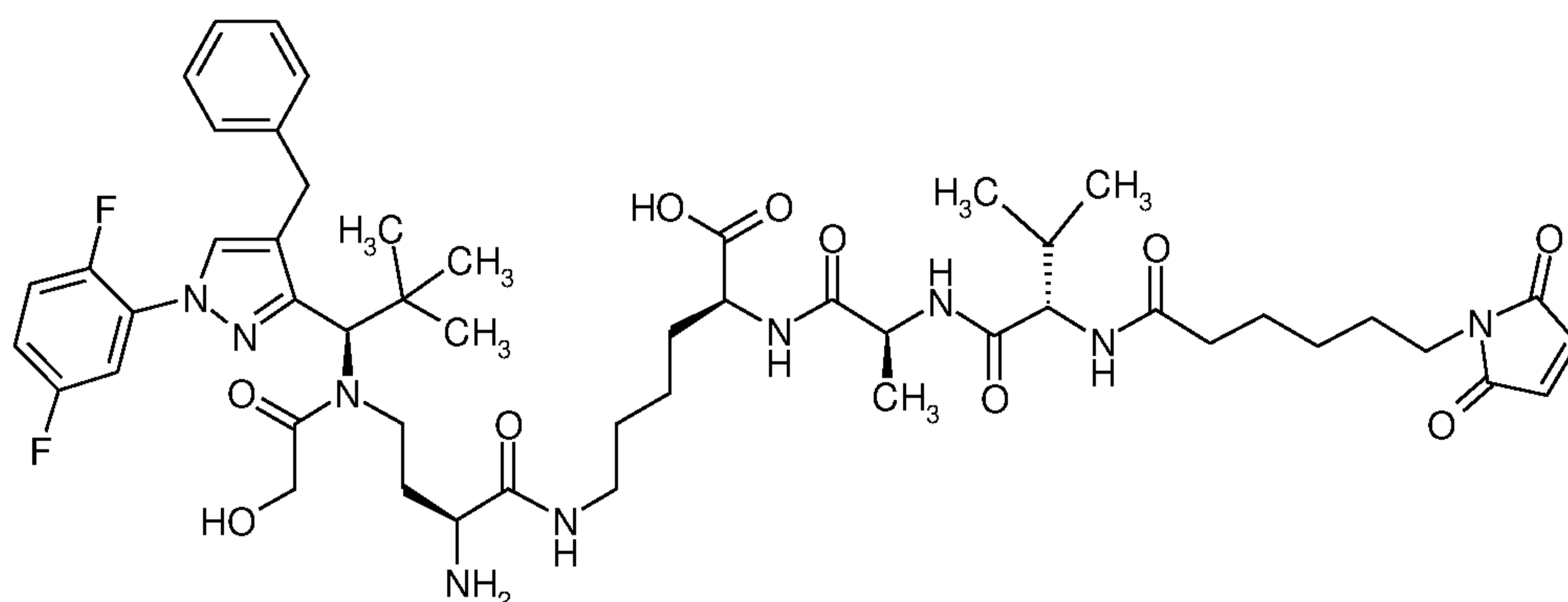


The title compound was prepared analogously to Intermediate F2 from 16 mg (0.025 mmol) of Intermediate C49 by reaction with 24 mg (0.076 mmol) of Intermediate L1 in the presence of EDCI/HOBT and *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid. This gave 3 mg of the title compound (14% of theory over 2 steps).

LC-MS (Method 1):  $R_t = 0.88$  min; MS (EIpos):  $m/z = 694$   $[M+H]^+$ .

10

### Intermediate F88



15 The compound was prepared analogously to Intermediate F8.

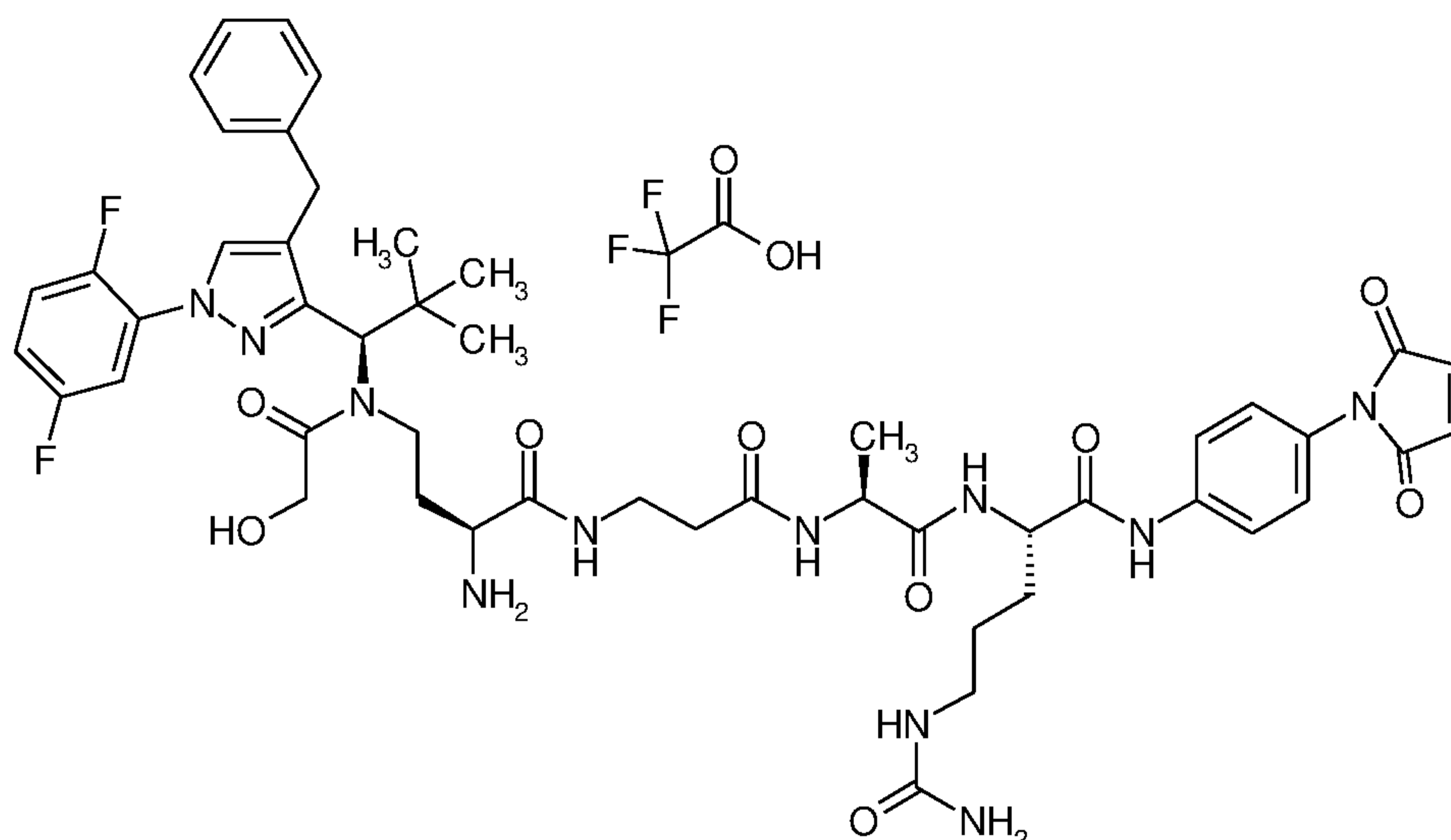
LC-MS (Method 5):  $R_t = 2.97$  min; MS (EIpos):  $m/z = 1006$   $[M+H]^+$ .

### Intermediate F89

20

Trifluoroacetic acid / *N*-{(2*S*)-2-amino-4-[(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1*H*-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl-L-alanyl-*N*<sup>5</sup>-carbamoyl-*N*-[4-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)phenyl]-L-ornithinamide (1:1)

25

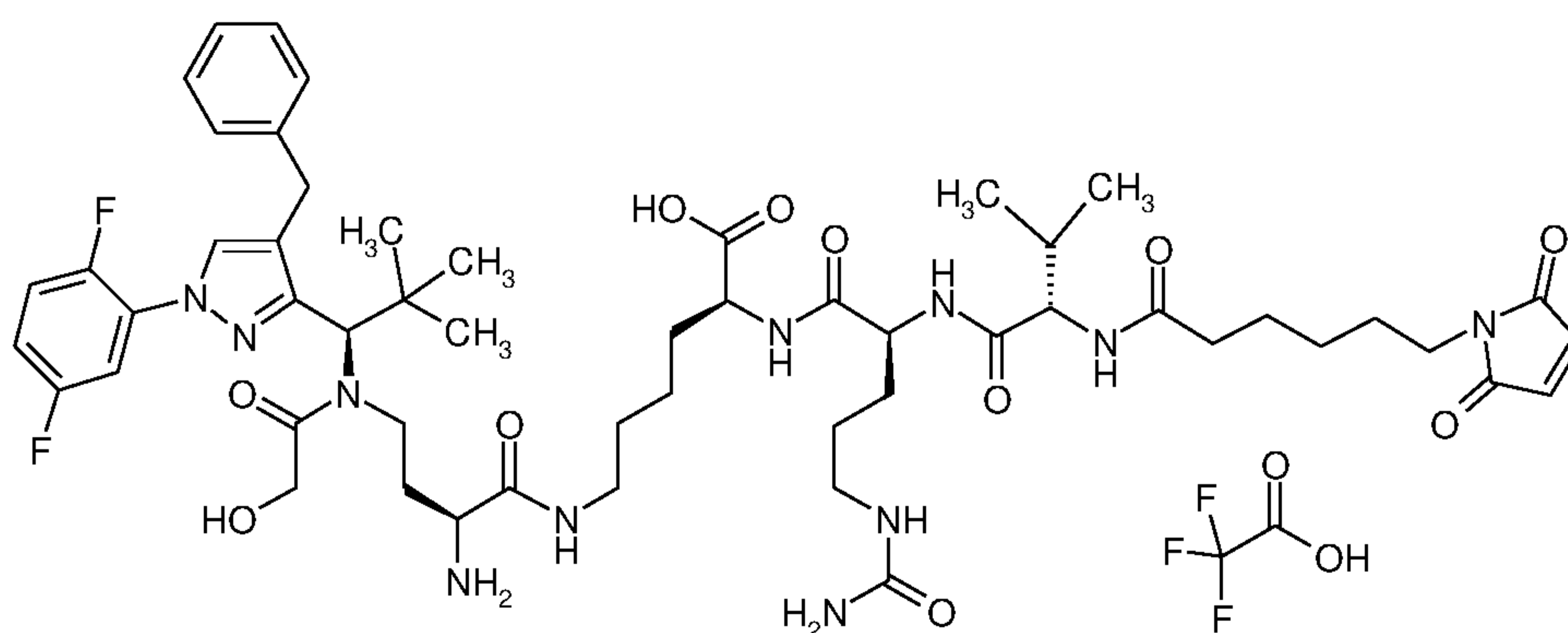


The title compound was prepared from 8 mg (0.012 mmol) of Intermediate C51 by reaction with 7.4 mg (0.014 mmol) of Intermediate L8 in the presence of 5.8 mg (0.015 mmol) of HATU and 10  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid. This gave 10 mg (78% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t = 0.87$  min; MS (EIpos):  $m/z = 984$   $[M+H]^+$ .

### Intermediate F90

15 *N*-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-*N*<sup>5</sup>-carbamoyl-L-ornithyl-*N*<sup>6</sup>-{(2*S*)-2-amino-4-[(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-lysine / trifluoroacetic acid (1:1)



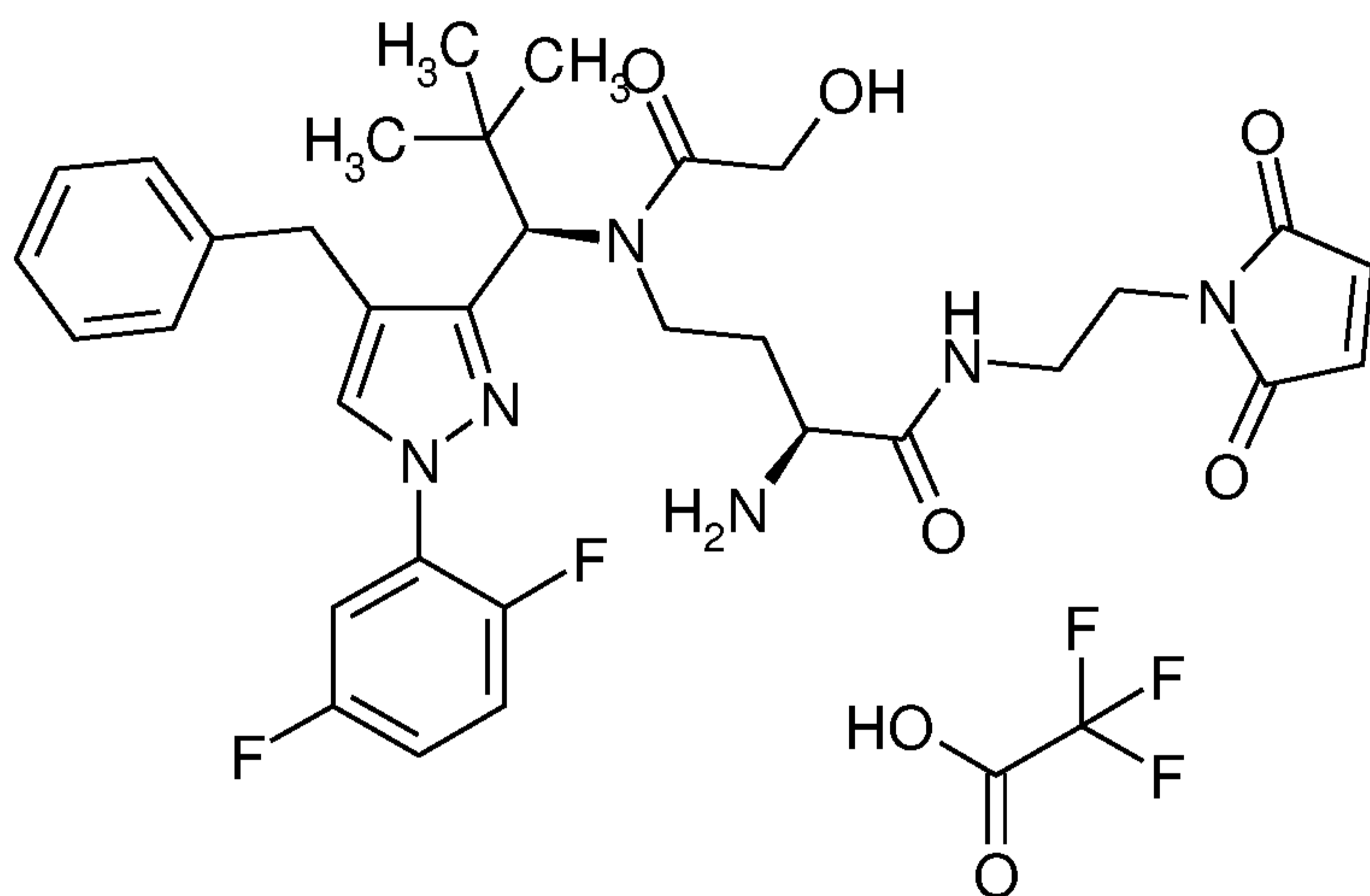


The title compound was prepared from 11 mg (0.018 mmol) of Intermediate C49 by reaction with 13.7 mg (0.018 mmol) of Intermediate L17 in the presence of 34 mg (0.089 mmol) of HATU and 19  $\mu$ l of *N,N*-diisopropylethylamine and subsequent  
 5 deprotection with trifluoroacetic acid. This gave 7.5 mg (35% of theory over 2 steps).

LC-MS (Method 8):  $R_t$  = 6.78 min; MS (EIpos):  $m/z$  = 1092  $[M+H]^+$ .

### 10 Intermediate F91

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-[2-(2,5-dioxo-2,5-dihydro-  
 15 1H-pyrrol-1-yl)ethyl]butanamide (1:1)

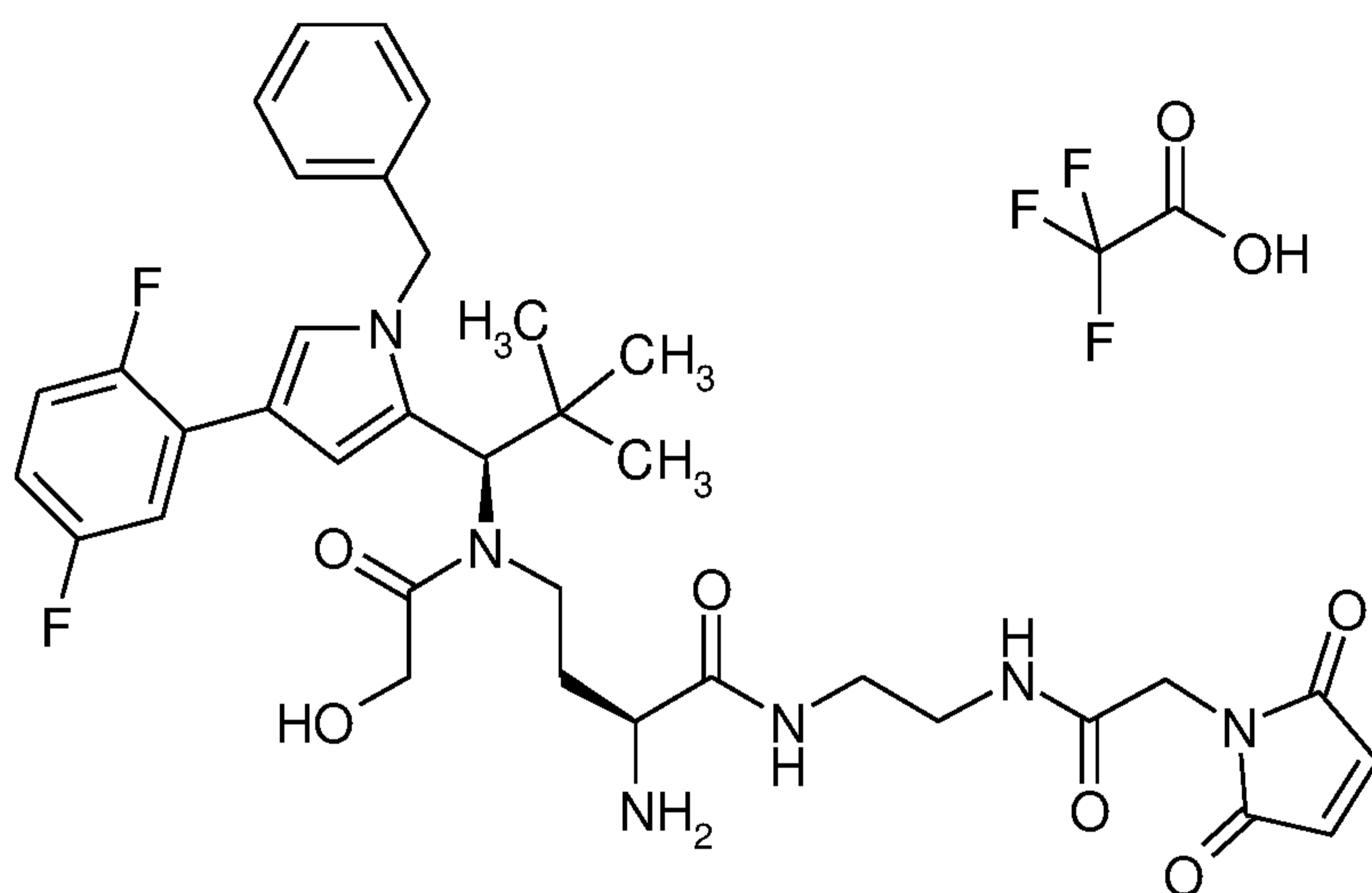


9.3 mg (0.01 mmol) of tert-butyl [(2S)-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-1-  
 20 { [2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]amino }-1-oxobutan-2-yl]carbamate were dissolved in 2 ml of dichloromethane, and 740 mg (6.49 mmol, 0.50 ml) of trifluoroacetic acid were added and the mixture was  
 25 stirred at RT for 1.5 h. The reaction mixture was then concentrated and the residue was taken up in acetonitrile and water and lyophilized. This gave 9.2 mg (96% of theory) of the target compound.

LC-MS (Method 1):  $R_t = 0.88$  min; MS (EIpos):  $m/z = 637$   $[M+H]^+$ .

### Intermediate F104

5 Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]-N-(2-[[2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino}ethyl)butanamide (1:1)



10

10 mg (0.014 mmol) of Intermediate C53 were dissolved in 3.3 ml of DMF, and 8.5 mg (0.027 mmol) of Intermediate L1, 7.8 mg (0.02 mmol) of HATU and 12  $\mu$ l of *N,N*-diisopropylethylamine were added.  
 15 The reaction was stirred at RT for 15 min and then concentrated. The residue was purified by preparative HPLC giving, after lyophilization, 5.6 mg (38% of theory) of the protected intermediate.

20 LC-MS (Method 1):  $R_t = 1.32$  min; MS (ESIpos):  $m/z = 915$   $(M+H)^+$ .

5.6 mg (0.006 mmol) of this intermediate were taken up in 2 ml of DMF, and 69 mg (0.61 mmol) of 1,4-diazabicyclo[2.2.2]octane were added. The reaction was treated in an ultrasonic bath for  
 25 2 h. 35  $\mu$ l of acetic acid were then added and the reaction was concentrated under high vacuum. The residue was purified by preparative HPLC. This gave 2.4 mg (48% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.84$  min; MS (EIpos):  $m/z = 693$  [M+H]<sup>+</sup>.

HPLC (Method 11):  $R_t = 1.91$  min;

5 Alternatively, the title compound was also prepared from  
Intermediate C58. 15 mg (0.023 mmol) of Intermediate C58 were  
initially reacted with 11 mg (0.036 mmol) of Intermediate L1 in  
the presence of 13 mg (0.034 mmol) of HATU and 10  $\mu$ l of *N,N*-  
diisopropylethylamine. After 60 min of stirring at RT, the  
10 mixture was concentrated and the residue was purified by  
preparative HPLC. This gave 12.3 mg (63% of theory) of the  
protected intermediate.

LC-MS (Method 1):  $R_t = 1.3$  min; MS (EIpos):  $m/z = 837$  [M+H]<sup>+</sup>.

15

In the second step, this intermediate was dissolved in 3 ml of  
2,2,2-trifluoroethanol. 12 mg (0.088 mmol) of zinc chloride were  
added, and the reaction was stirred at 50°C for 2 h. 26 mg (0.088  
mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid and 2 ml of  
20 a 0.1% strength aqueous trifluoroacetic acid solution were then  
added. The reaction was purified by preparative HPLC.  
Concentration of the appropriate fractions and lyophilization  
of the residue from acetonitrile/water gave 8.1 mg (68% of  
theory) of the title compound.

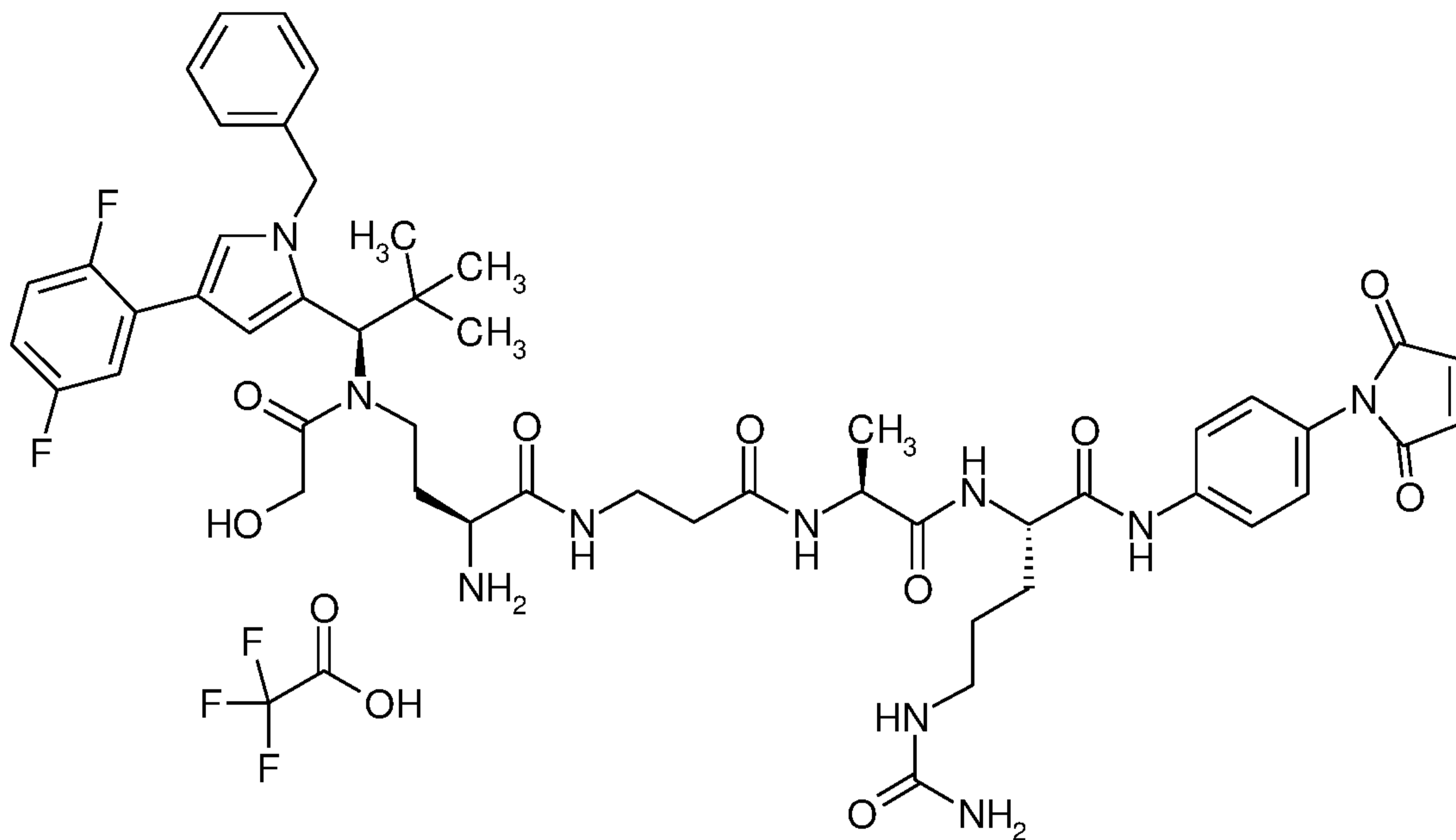
25

LC-MS (Method 1):  $R_t = 0.89$  min; MS (ESIpos):  $m/z = 693$  (M+H)<sup>+</sup>.

### Intermediate F106

30 Trifluoroacetic acid / N-{(2*S*)-2-amino-4-[(1*R*)-1-[1-benzyl-4-  
(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-  
dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl-L-  
alanyl-N<sup>5</sup>-carbamoyl-N-[4-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-  
yl)phenyl]-L-ornithinamide (1:1)

35



The title compound was prepared analogously to Intermediate F104 from Intermediate C53 and Intermediate L47.

5

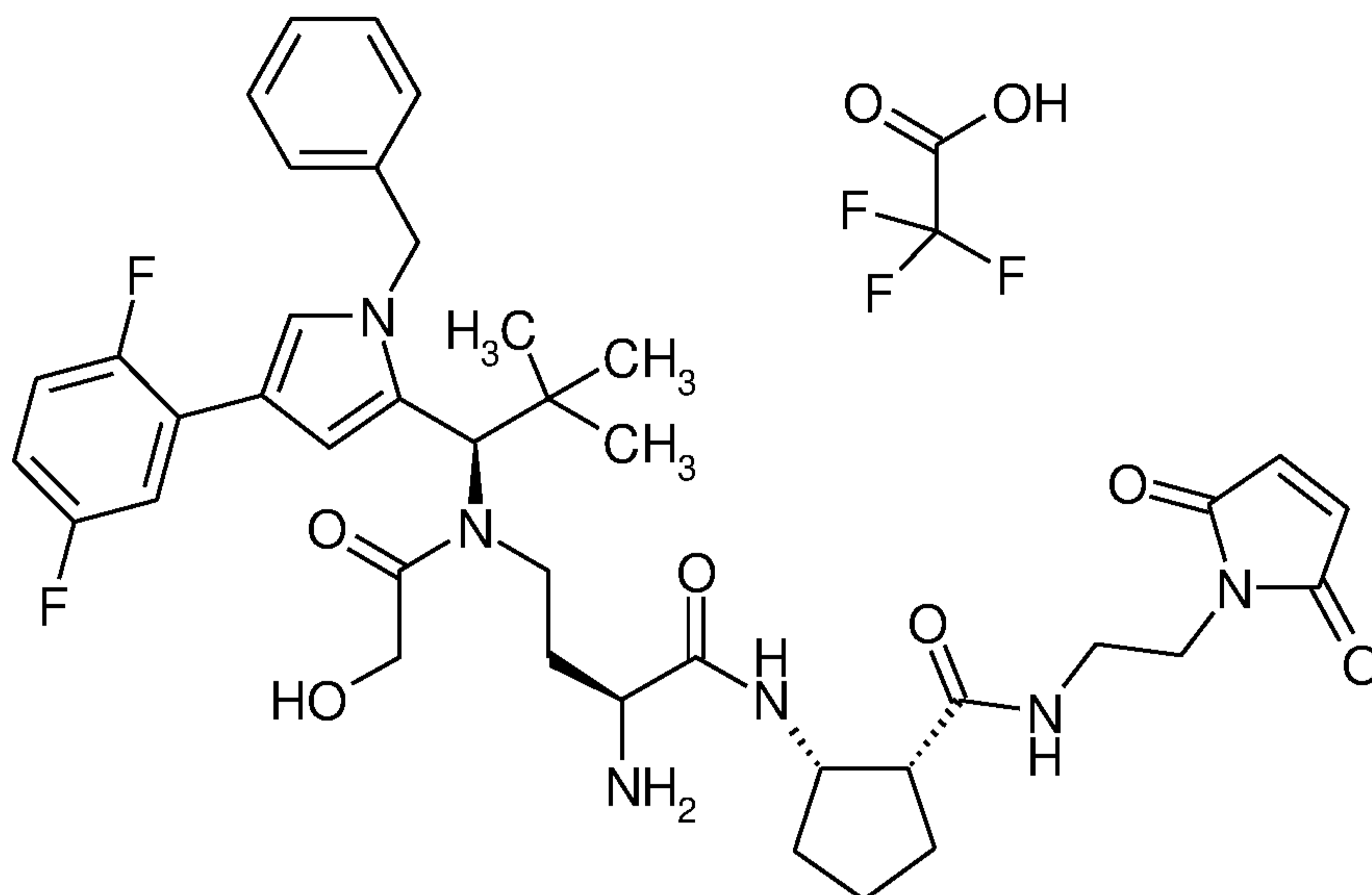
HPLC (Method 11):  $R_t = 1.85$  min;

LC-MS (Method 1):  $R_t = 0.86$  min; MS (ESIpos):  $m/z = 983$  (M+H)<sup>+</sup>.

10 **Intermediate F108**

Trifluoroacetic acid / (1R,2S)-2-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)

15

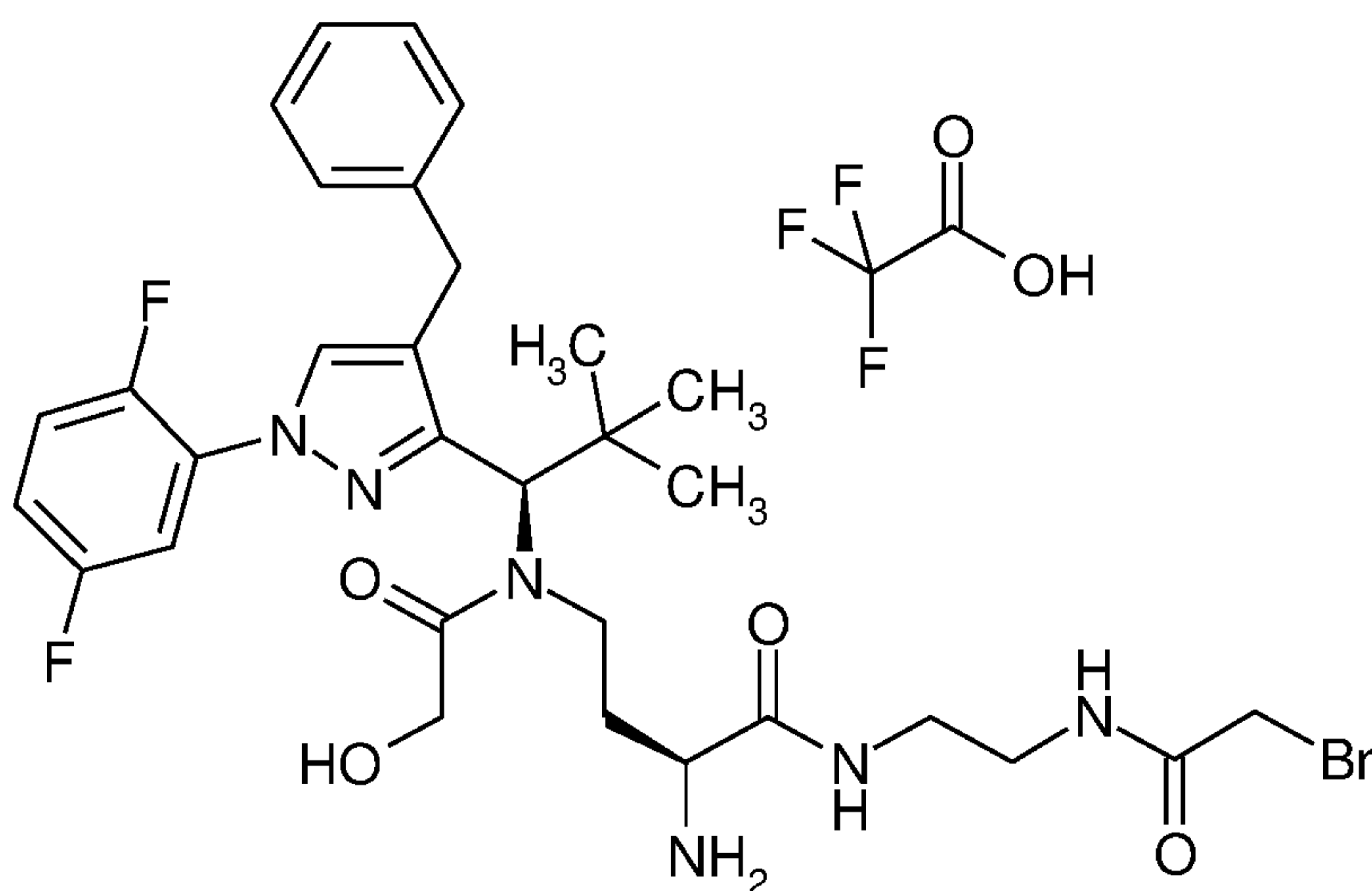


The title compound was prepared analogously to Intermediate F104 from 20 mg (0.027 mmol) of Intermediate C53 and 24 mg (0.054 mmol) of Intermediate L48. This gave 3 mg (14% of theory over 2 steps).

LC-MS (Method 1):  $R_t = 0.93$  min; MS (EIpos):  $m/z = 747$   $[M+H]^+$ .

#### 10 Intermediate F109

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-{2-[(bromoacetyl)amino]ethyl}butanamide (1:1)



17 mg (0.026 mmol) of Intermediate C57 were taken up in 3 ml of DMF and reacted with 7 mg (0.027 mmol) of commercially available 1-(2-bromoacetoxy)pyrrolidine-2,5-dione in the presence of 14  $\mu$ l of *N,N*-diisopropylethylamine. After 15min of stirring at RT, the mixture was concentrated and the residue was purified by preparative HPLC. This gave 7 mg (33% of theory) of this intermediate.

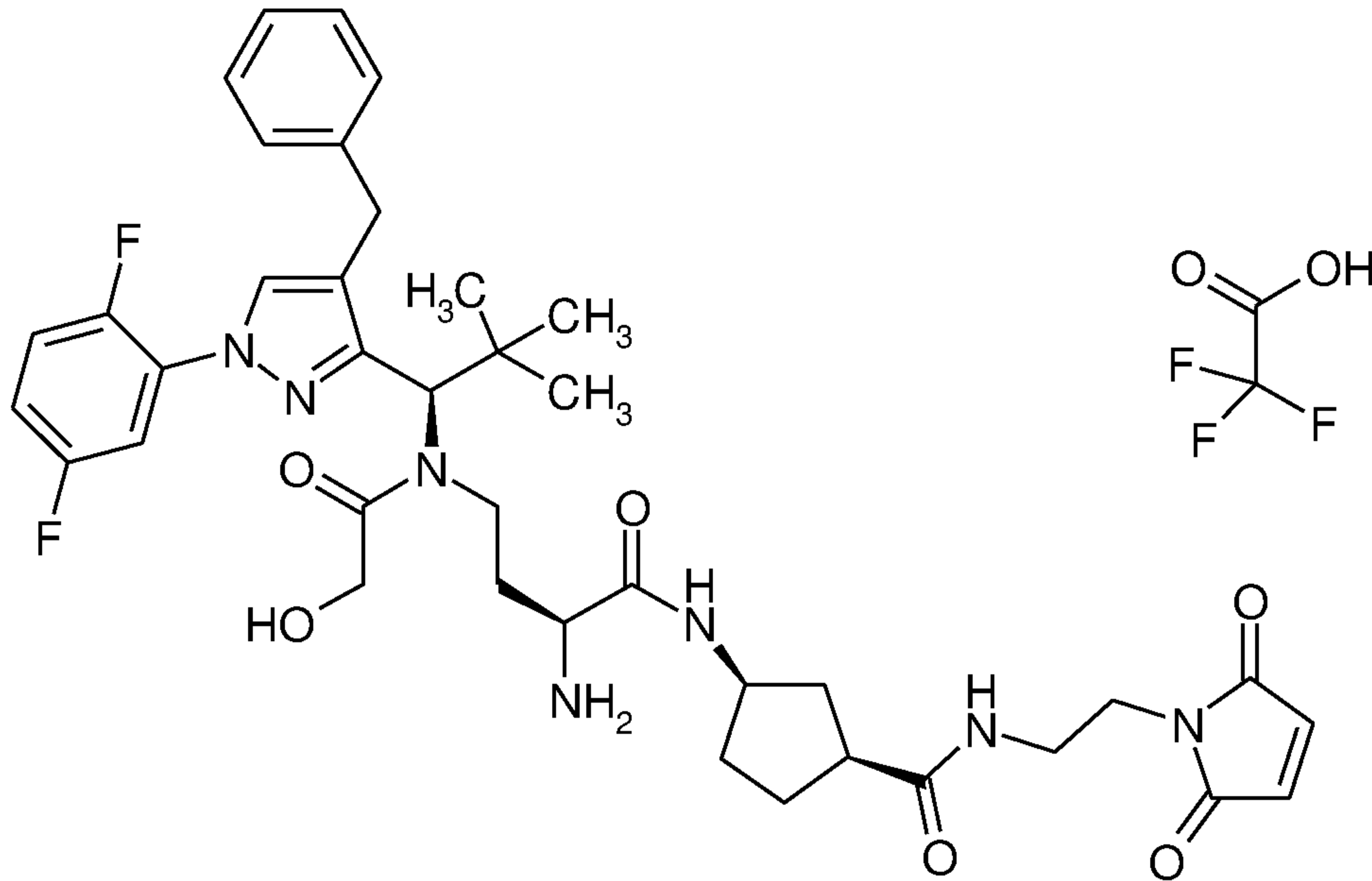
LC-MS (Method 1):  $R_t = 1.29$  min; MS (ESIpos):  $m/z = 777$  and  $779$  (M+H)<sup>+</sup>.

This intermediate was taken up in 1 ml of dichloromethane and deprotected with 1 ml of trifluoroacetic acid. After concentration and lyophilization from acetonitrile/water, 6 mg (88% of theory) of the title compound were obtained.

LC-MS (Method 1):  $R_t = 0.86$  min; MS (ESIpos):  $m/z = 677/679$  (M+H)<sup>+</sup>.

## 20 Intermediate F112

Trifluoroacetic acid / (1*S*,3*R*)-3-({(2*S*)-2-amino-4-[(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1*H*-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-*N*-[2-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)

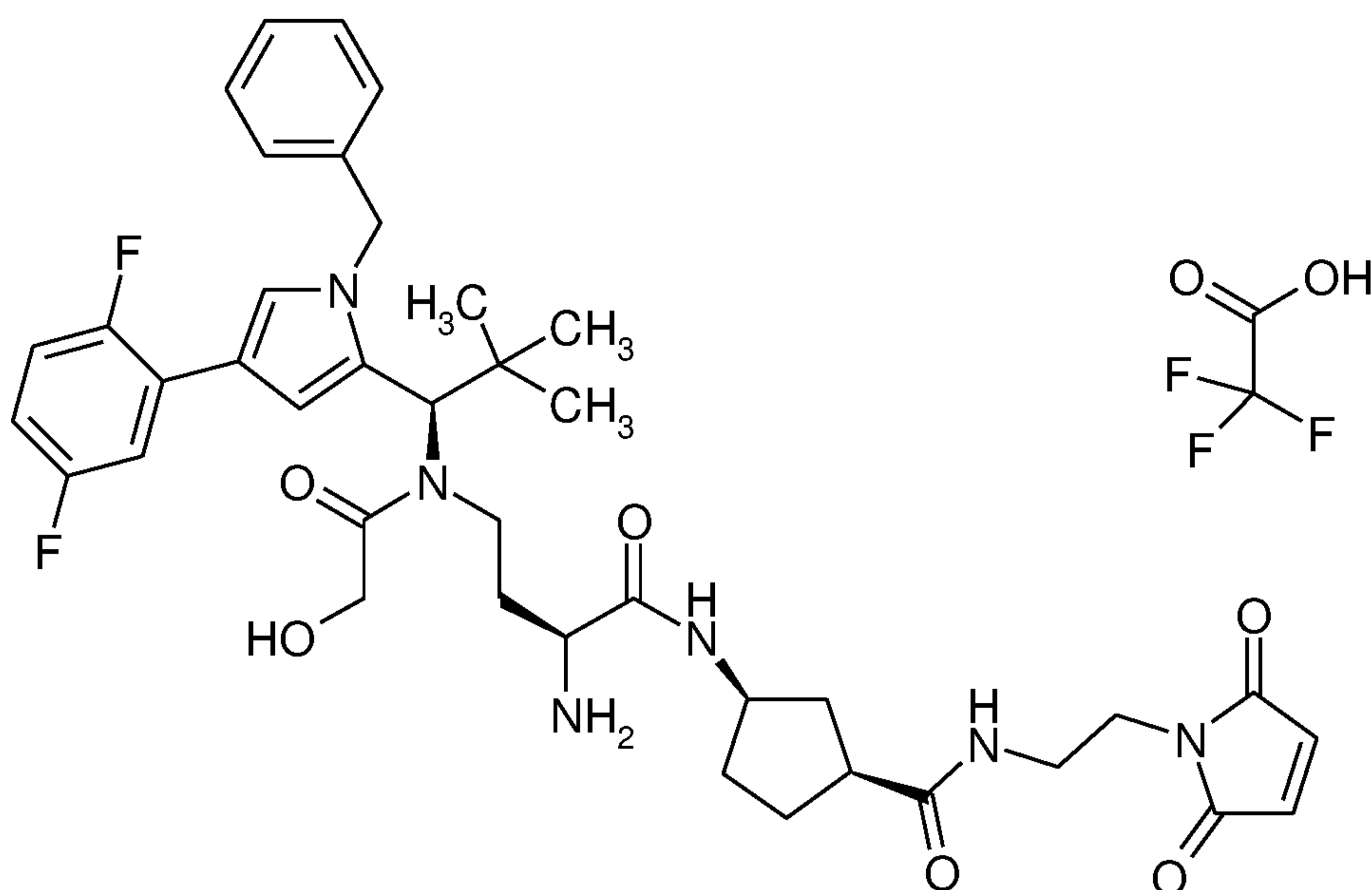


The title compound was prepared from 15 mg (0.024 mmol) of Intermediate C49 by reaction with 18 mg (0.049 mmol) of Intermediate L50 in the presence of 14 mg (0.037 mmol) of HATU and 21  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid. This gave 12 mg (51% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t$  = 0.89 min; MS (EIpos):  $m/z$  = 748  $[M+H]^+$ .

### Intermediate F113

15 Trifluoroacetic acid / (1*S*,3*R*)-3-({(2*S*)-2-amino-4-[(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-*N*-[2-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)



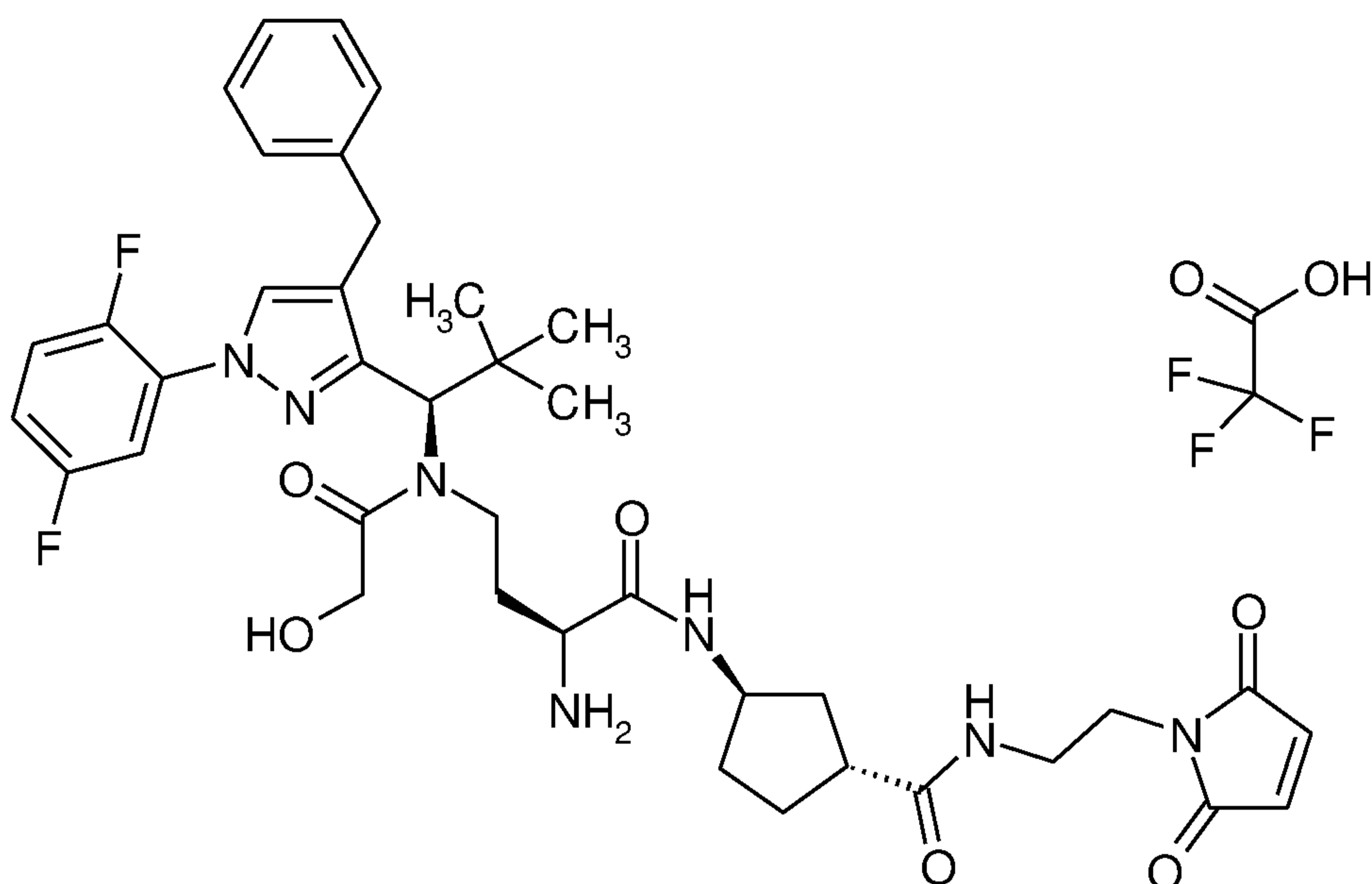
The title compound was prepared from 15 mg (0.019 mmol) of Intermediate C53 by reaction with 14 mg (0.038 mmol) of Intermediate L50 in the presence of 11 mg (0.029 mmol) of HATU and 17  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with 133 mg of DABCO in 2 ml of DMF. Purification by HPLC gave 4 mg (24% of theory over 2 steps).

10 LC-MS (Method 5):  $R_t$  = 2.77 min; MS (EIpos):  $m/z$  = 747 [M+H]<sup>+</sup>.

### Intermediate F115

15 Trifluoroacetic acid / (1R,3R)-3-({(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)



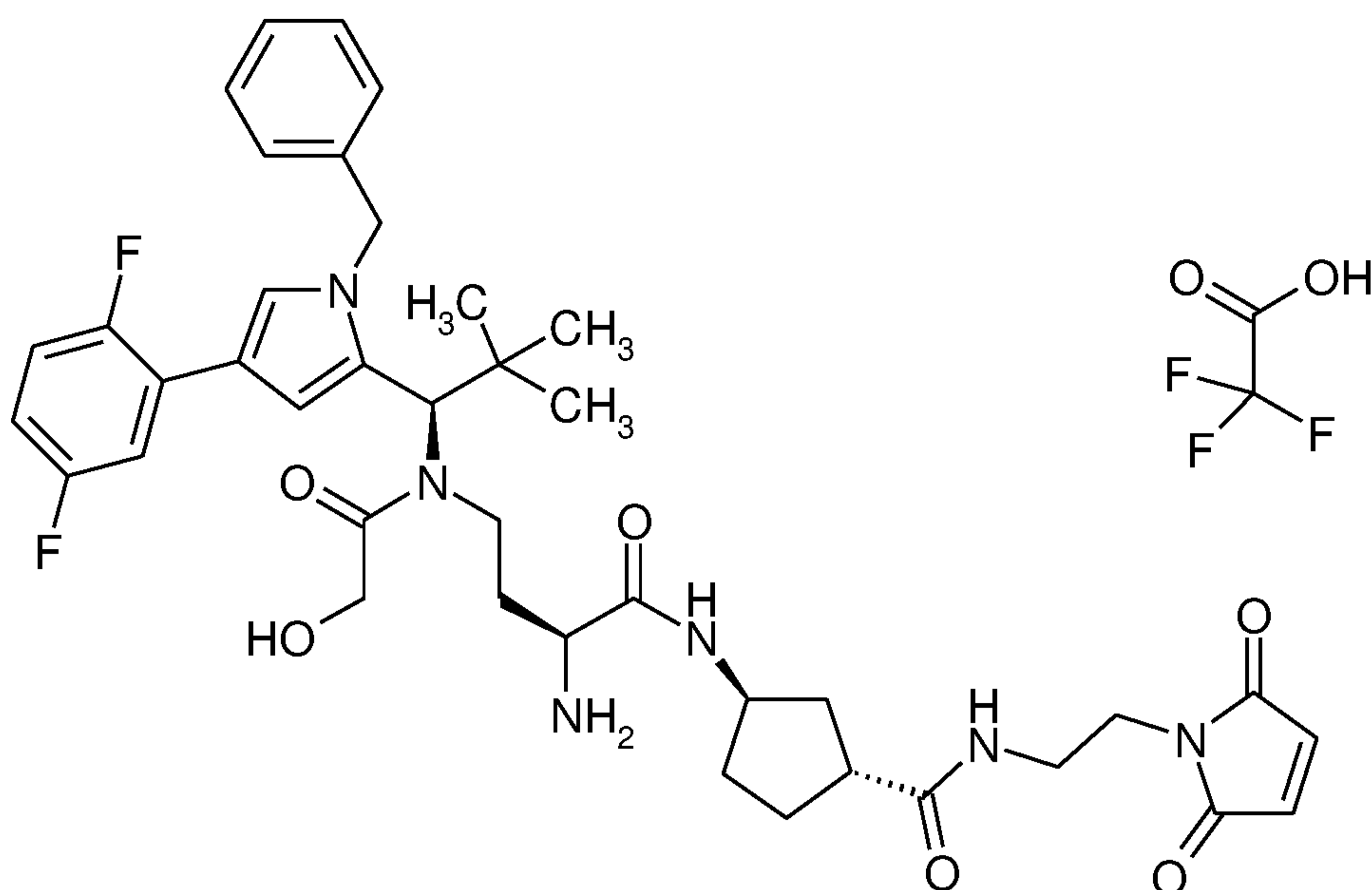


The title compound was prepared from 15 mg (0.024 mmol) of Intermediate C49 by reaction with 18 mg (0.047 mmol) of Intermediate L51 in the presence of 13 mg (0.035 mmol) of HATU and 21  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid. This gave 12 mg (51% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t$  = 0.87 min; MS (EIpos):  $m/z$  = 748  $[M+H]^+$ .

### Intermediate F116

15 Trifluoroacetic acid / (1R,3R)-3-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)

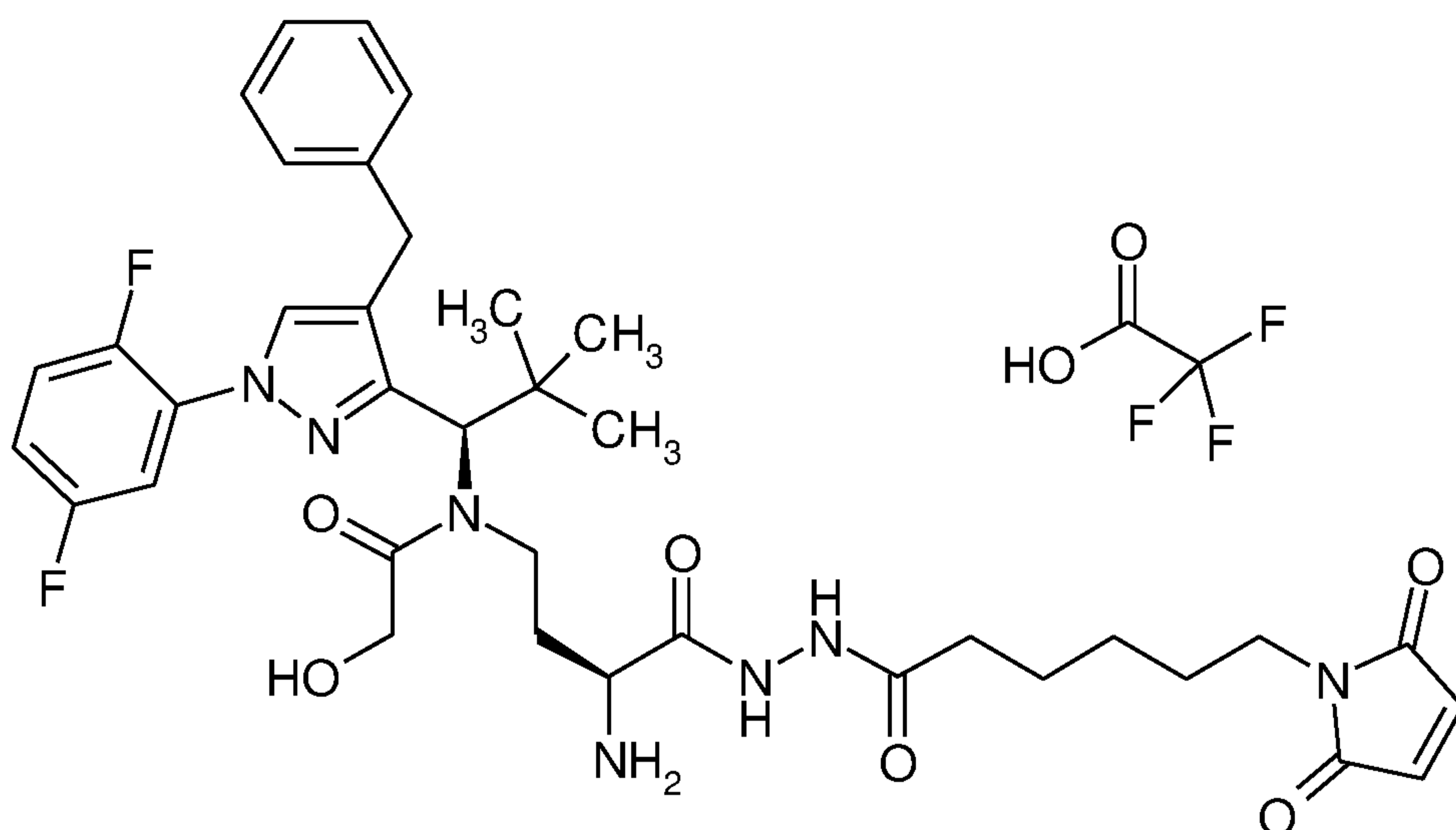


The title compound was prepared from 11 mg (0.014 mmol) of Intermediate C51 by reaction with 11 mg (0.028 mmol) of Intermediate L51 in the presence of 8 mg (0.021 mmol) of HATU and 12  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with 87 mg of DABCO in 2 ml of DMF. Purification by HPLC gave 3.3 mg (28% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t$  = 0.92 min; MS (EIpos):  $m/z$  = 747  $[M+H]^+$ .

### Intermediate F117

15 Trifluoroacetic acid / *N*-[(3*S*)-3-amino-4-{2-[6-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)hexanoyl]hydrazino}-4-oxobutyl]-*N*-{(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1*H*-pyrazol-3-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide (1:1)



The title compound was prepared according to classical methods of peptide chemistry from Intermediate C49. First, C49 was  
 5 coupled with 9H-fluoren-9-ylmethyl hydrazinecarboxylate in the presence of HATU. The Fmoc protective group was then removed with piperidine in DMF and the hydrazide obtained was coupled with 6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoic acid in the presence of HATU. In the last step, the Boc protective group  
 10 was removed with TFA in dichloromethane.

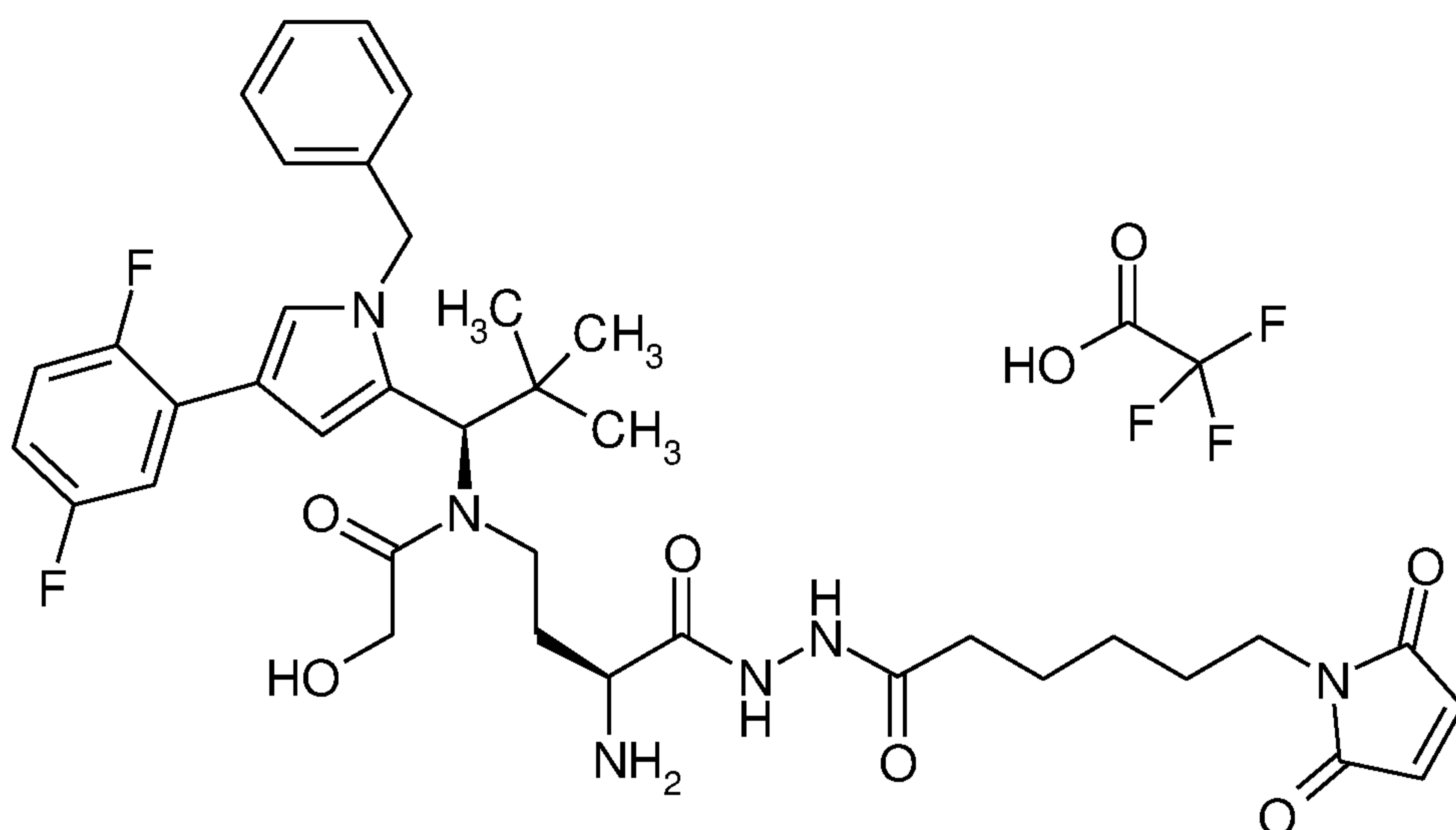
LC-MS (Method 1):  $R_t = 0.93$  min; MS (EIpos):  $m/z = 722$   $[M+H]^+$ .

### Intermediate F118

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Trifluoroacetic acid / N-[(3S)-3-amino-4-{2-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]hydrazino}-4-oxobutyl]-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide (1:1)

20



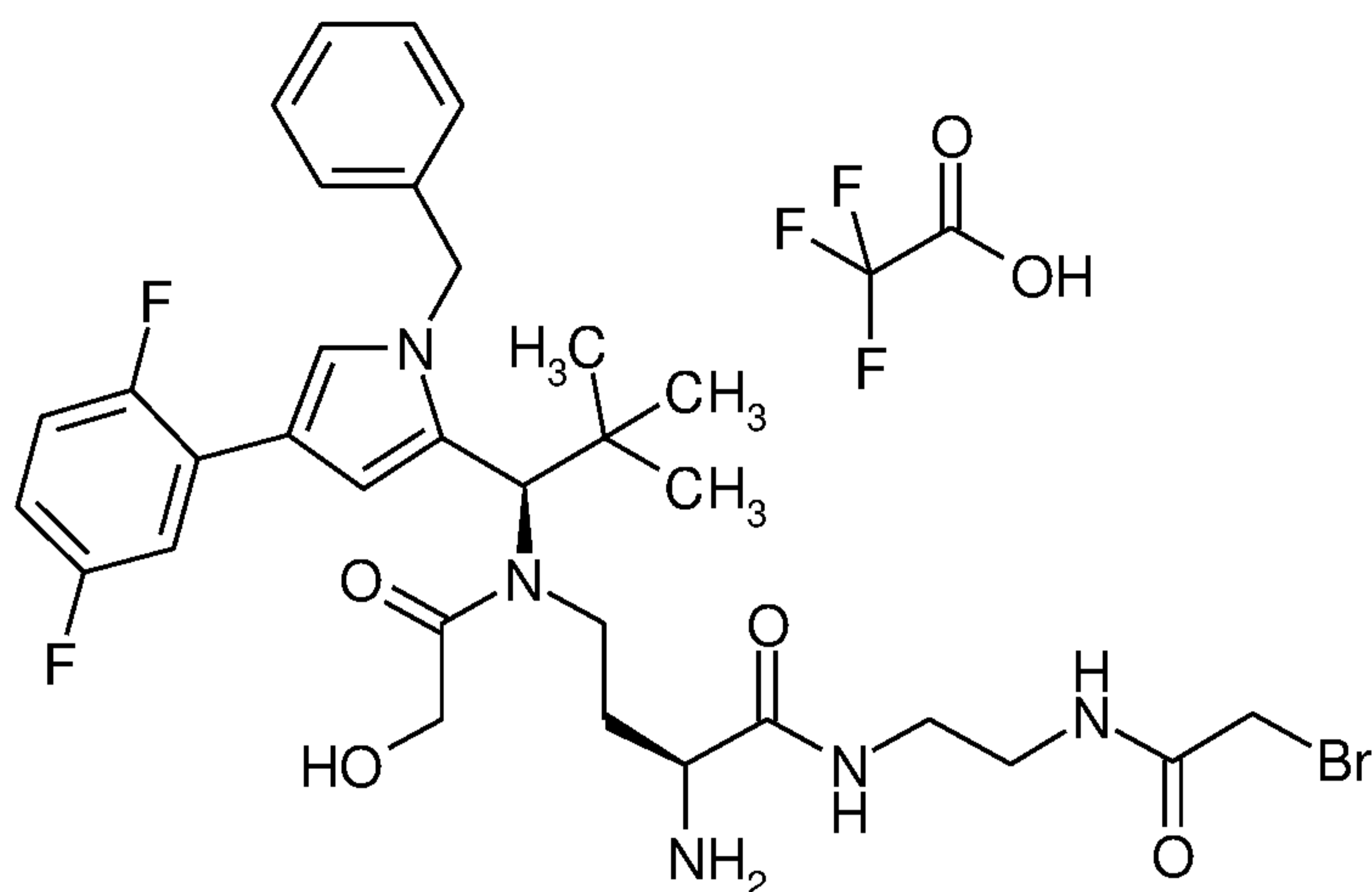
In the first step, the title compound was prepared analogously to Intermediate F3 from 15 mg (0.019 mmol) of Intermediate C53  
 5 by coupling with commercially available 6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanehydrazide in the presence of HATU. The Fmoc protective group was then removed with 142 mg of DABCO in DMF. Purification by HPLC gave 3 mg (19% of theory) of the title compound.

10

LC-MS (Method 1):  $R_t = 0.90$  min; MS (EIpos):  $m/z = 721$   $[M+H]^+$ .

### Intermediate F119

15 Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-{2-[(bromoacetyl)amino]ethyl}butanamide (1:1)



29 mg (0.044 mmol) of Intermediate C58 were taken up in 3.4 ml of DMF, and 36 mg (0.087 mmol) of Intermediate L52, 25 mg (0.065 mmol) of HATU and 19  $\mu$ l of *N,N*-diisopropylethylamine were added. After 60 min of stirring at RT, the mixture was concentrated and the residue was purified by preparative HPLC. This gave 26.4 mg (73% of theory) of the intermediate.

10 LC-MS (Method 1):  $R_t$  = 1.34 min; MS (ESIpos):  $m/z$  = 820 and 822 (M+H)<sup>+</sup>.

This intermediate was dissolved in 3 ml of 2,2,2-trifluoroethanol. 6.5 mg (0.048 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 4 h. 13.9 mg (0.048 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid and 2 ml of a 0.1% strength aqueous trifluoroacetic acid solution were added. The reaction was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 14.4 mg (58% of theory) of the title compound.

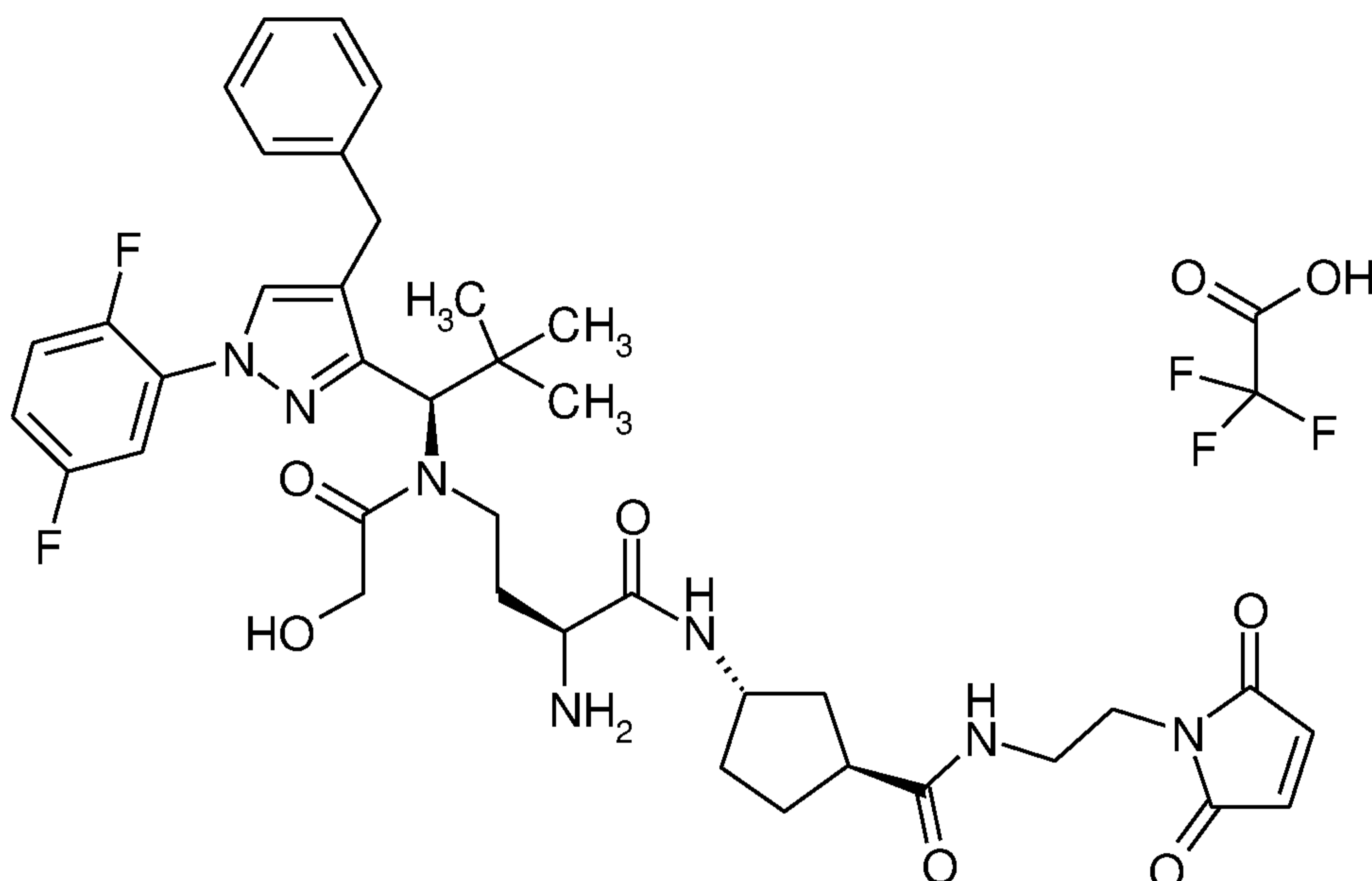
LC-MS (Method 1):  $R_t$  = 0.88 min; MS (ESIpos):  $m/z$  = 676 and 678 (M+H)<sup>+</sup>.

25

### Intermediate F121

Trifluoroacetic acid / (1*S*,3*S*)-3-({(2*S*)-2-amino-4-[(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1*H*-pyrazol-3-yl]-2,2-

dimethylpropyl} (glycoloyl) amino]butanoyl} amino) -N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl) ethyl] cyclopentanecarboxamide (1:1)



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The title compound was prepared from 10 mg (0.016 mmol) of Intermediate C49 by reaction with 11.5 mg (0.031 mmol) of Intermediate L53 in the presence of 9 mg (0.024 mmol) of HATU and 14  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid. This gave 9 mg (61% of theory over 2 steps).

10

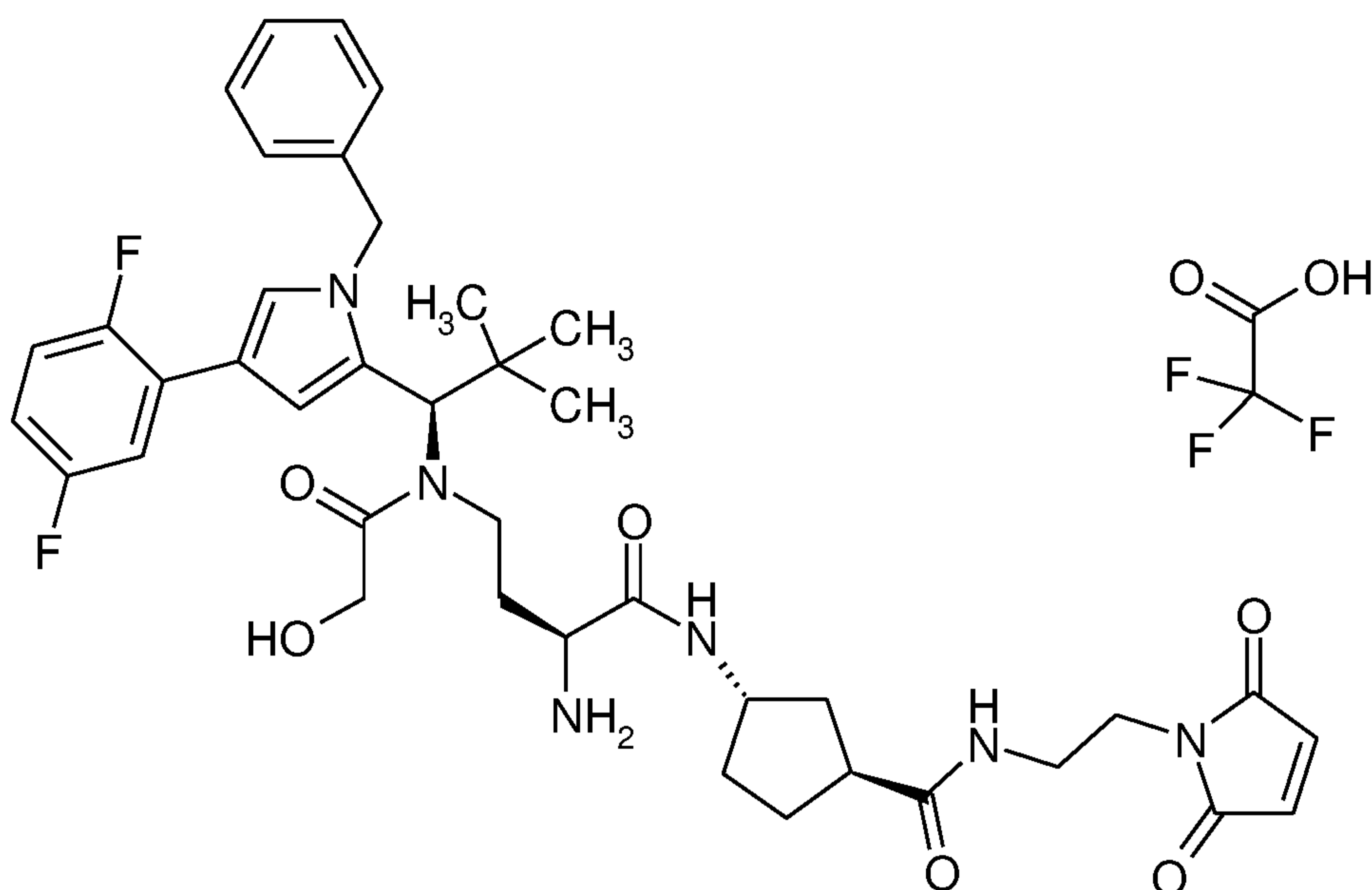
LC-MS (Method 1):  $R_t$  = 0.84 min; MS (EIpos):  $m/z$  = 748  $[M+H]^+$ .

15

### Intermediate F122

Trifluoroacetic acid / (1*S*,3*S*)-3-({(2*S*)-2-amino-4-[(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl} amino) -N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl) ethyl] cyclopentanecarboxamide (1:1)

20

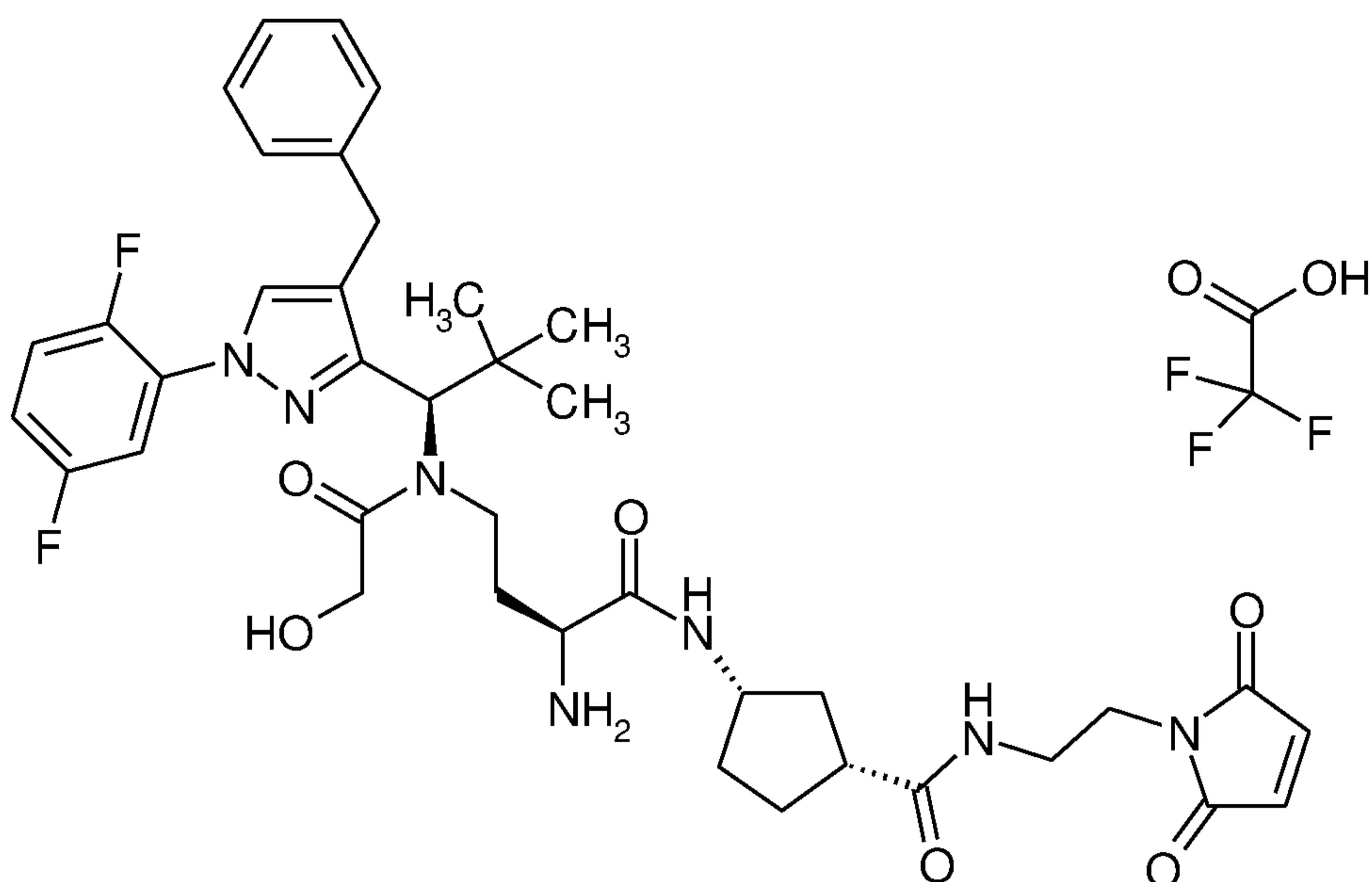


The title compound was prepared from 15 mg (0.019 mmol) of Intermediate C53 by reaction with 14 mg (0.038 mmol) of Intermediate L53 in the presence of 11 mg (0.029 mmol) of HATU and 17  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with 202 mg of DABCO in 3 ml of DMF. Purification by HPLC gave 4 mg (24% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t$  = 0.87 min; MS (EIpos):  $m/z$  = 747 [M+H]<sup>+</sup>.

#### Intermediate F124

15 Trifluoroacetic acid / (1R,3S)-3-({(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)



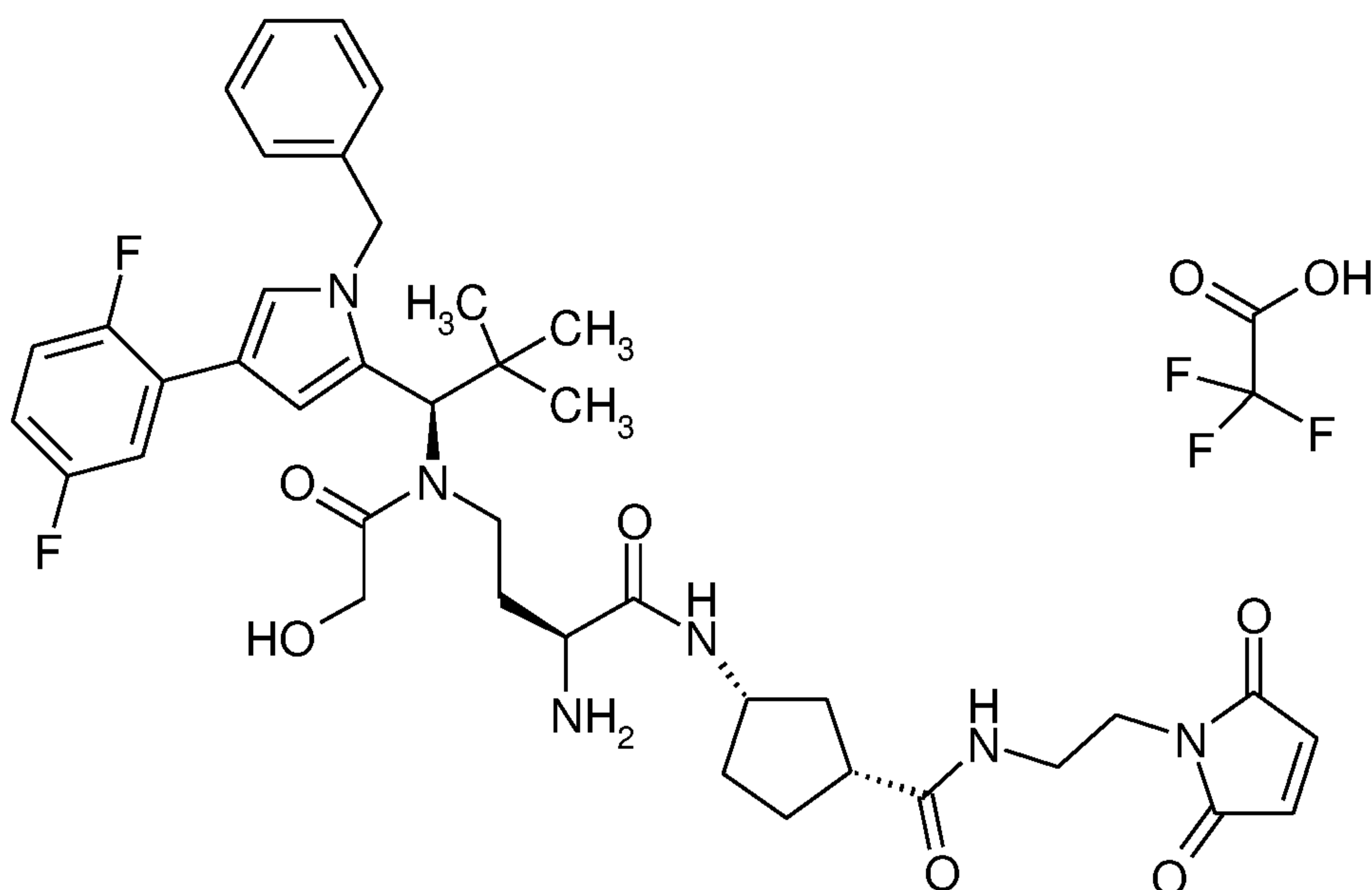
The title compound was prepared from 10 mg (0.016 mmol) of Intermediate C49 by reaction with 11.5 mg (0.031 mmol) of Intermediate L54 in the presence of 9 mg (0.024 mmol) of HATU and 14  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid. This gave 9 mg (66% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t$  = 0.84 min; MS (EIpos):  $m/z$  = 748  $[M+H]^+$ .

### Intermediate F125

15 Trifluoroacetic acid / (1*R*,3*S*)-3-({(2*S*)-2-amino-4-[(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-*N*-[2-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)ethyl]cyclopentanecarboxamide (1:1)





The title compound was prepared from 15 mg (0.019 mmol) of Intermediate C53 by reaction with 14 mg (0.038 mmol) of Intermediate L54 in the presence of 11 mg (0.029 mmol) of HATU and 17  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with 127 mg of DABCO in 3 ml of DMF. Purification by HPLC gave 3 mg (17% of theory over 2 steps).

10 LC-MS (Method 4):  $R_t$  = 1.08 min; MS (EIpos):  $m/z$  = 769  $[M+Na]^+$ .

### Intermediate F126

15 *N*-(Bromoacetyl)-*L*-valyl-*L*-alanyl-*N*<sup>6</sup>-{(2*S*)-2-amino-4-[(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1*H*-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-*L*-lysine / trifluoroacetic acid (1:1)



12 mg (0.015 mmol) of Intermediate C59 were dissolved in 2.4 ml of DMF, and 14.6 mg (0.046 mmol) of Intermediate L1, 6 mg (0.031 mmol) of 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide hydrochloride, 5.9 mg (0.039 mmol) of 1-hydroxy-1*H*-benzotriazole hydrate and 8  $\mu$ l of *N,N*-diisopropylethylamine were added. After 1 h of stirring at RT, the mixture was concentrated and the residue was purified by preparative HPLC. This gave 11 mg (70% of theory) of this intermediate.

10

LC-MS (Method 1):  $R_t = 1.34$  min; MS (ESIpos):  $m/z = 942$  (M+H)<sup>+</sup>.

11 mg (0.011 mmol) of this intermediate were taken up in 2 ml of DMF, and 123 mg (1.1 mmol) of 1,4-diazabicyclo[2.2.2]octane were added. The reaction was treated in an ultrasonic bath for 2 h. 63  $\mu$ l of acetic acid were then added and the reaction was concentrated under high vacuum. The residue was purified by preparative HPLC. This gave 2 mg (22% of theory) of the title compound.

20

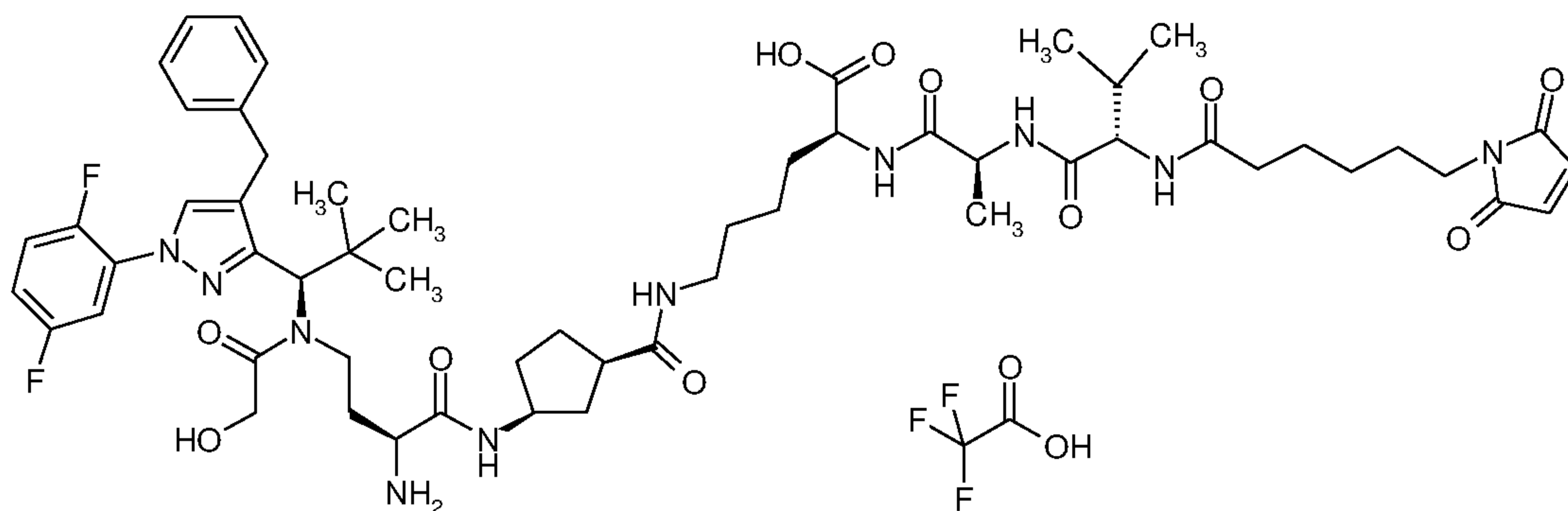
LC-MS (Method 1):  $R_t = 0.89$  min; MS (EIpos):  $m/z = 721$  [M+H]<sup>+</sup>.

HPLC (Method 11):  $R_t = 1.95$  min;

### 25 Intermediate F129

*N*-[6-(2,5-Dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)hexanoyl]-*L*-valyl-*L*-alanyl-*N*<sup>6</sup>-{[(1*R*,3*S*)-3-({(2*S*)-2-amino-4-[(1*R*)-1-[4-benzyl-1-(2,5-difluorophenyl)-1*H*-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)cyclopentyl]carbonyl}-*L*-lysine / trifluoroacetic acid

30

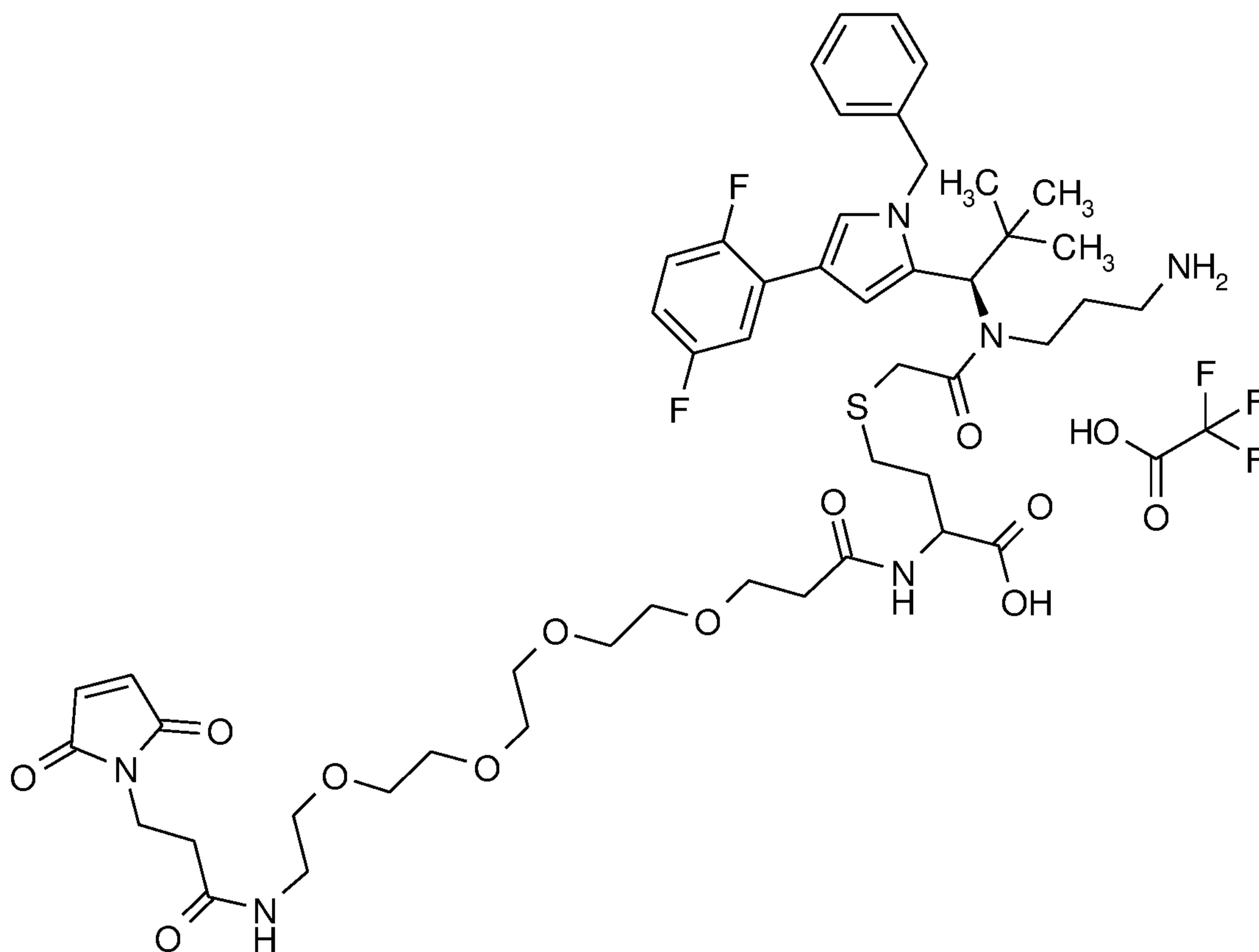


The title compound was prepared analogously to Intermediate F128 from 10 mg (0.016 mmol) of Intermediate C49 by reaction with 19 mg (0.024 mmol) of Intermediate L56 in the presence of 12 mg (0.031 mmol) of HATU and 14  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with trifluoroacetic acid. This gave 13.5 mg (70% of theory over 2 steps).

10 LC-MS (Method 1):  $R_t$  = 0.9 min; MS (EIpos):  $m/z$  = 1117  $[M+H]^+$ .

### Intermediate F142

15 R/S-{2-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-homocysteine / trifluoroacetic acid (1:1)



20.0 mg (23.7  $\mu\text{mol}$ ) of R/S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-homocysteine / trifluoroacetic acid (1:1) and 13.4 mg (26.04 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide were dissolved in 1.0 ml of DMF, and 4.8 mg (47.34  $\mu\text{mol}$ ) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight. 3.6 mg (0.06 mmol) of acetic acid were added and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 12.4 mg (44% of theory) of the compound R/S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]homocysteine.

LC-MS (Method 1):  $R_t = 1.30$  min; MS (ESIpos):  $m/z = 1129$  (M+H)<sup>+</sup>.

10.0 mg (8.85  $\mu$ mol) of R/S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-homocysteine were dissolved in trifluoroethanol, and 3.1 mg (22.71  $\mu$ mol) of zinc dichloride were added. The reaction mixture was stirred at 50°C overnight. 3.9 mg (0.01 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred briefly and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was lyophilized with a little water. This gave 7.6 mg (78% of theory) of the title compound.

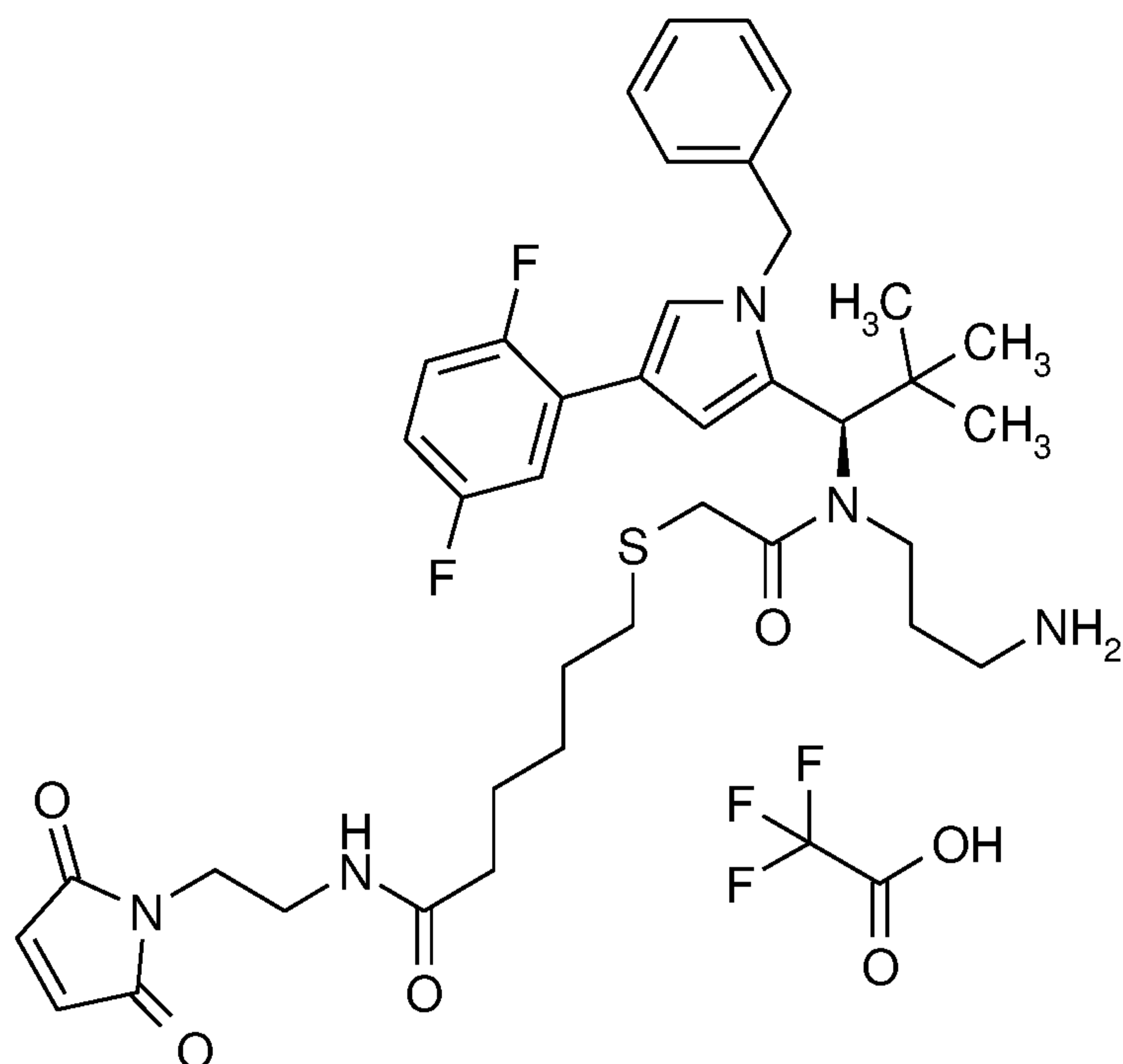
LC-MS (Method 1):  $R_t = 0.94$  min; MS (ESIpos):  $m/z = 983$  (M+H)<sup>+</sup>.

<sup>1</sup>H NMR (400 MHz, DMSO-d<sub>6</sub>):  $\delta$  [ppm] = 0.50 (m, 1H), 0.81 (s, 9H), 1.49 (m, 1H), 1.89 (m, 1H), 2.05 (m, 1H), 2.29-2.43 (m, 4H), 2.45-2.55 (m, 2H), 2.58-2.74 (m, 2H), 3.10-3.20 (m, 2H), 3.21-3.40 (m, 2H), 3.42-3.54 (m, 16H), 3.55-3.65 (m, 4H), 4.28 (m, 1H), 4.91 (dd, 1H), 5.18 (dd, 1H), 5.60 (s, 1H), 6.95 (m, 1H), 7.00 (s, 2H), 7.15-7.38 (m, 7H), 7.53 (s, 1H), 7.68 (m, 1H), 8.00 (m, 2H).

### Intermediate F143

Trifluoroacetic acid / 6-({2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulphanyl)-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]hexanamide (1:1)

35



30.0 mg (0.05 mmol) of 2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate and 13.5 mg (0.07 mmol) of 6-(acetylsulphanyl)hexanoic acid were initially charged in 2.0 ml of methanol with a drop of water. 23.0 mg (0.17 mmol) of potassium carbonate were added. The reaction mixture was stirred at 50°C for 4 h. Ethyl acetate was added to the reaction mixture. The organic phase was washed with saturated NaCl solution and dried over magnesium sulphate. The solvent was evaporated under reduced pressure. The residue was purified by preparative RP-HPLC (column: Reprosil 125x30; 10µ, flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 54.2 mg (90% of theory) of the compound 11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaicosan-20-oic acid.

20

LC-MS (Method 1):  $R_t = 1.49$  min; MS (ESIpos):  $m/z = 1106$  (M+H)<sup>+</sup>.

54.0 mg (0.07 mmol) of 11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-

dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaicosan-20-oic acid and 16.7 mg (0.09 mmol) of 1-(2- $\beta$ -aminoethyl)-1H-pyrrole-2,5-dione hydrochloride (1:1) were initially charged in 3.0 ml of acetonitrile, and 75.0 mg (0.58 mmol) of *N,N*-diisopropylethylamine were added. 60.0 mg (0.09 mmol) of T3P (50% in acetonitrile) were added and the mixture was stirred at RT overnight. The reaction was quenched with water and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 42.8 mg (68% of theory) of the compound 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl){[(6-([2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]amino)-6-oxohexyl)sulphanyl]acetyl}amino)propyl]carbamate.

LC-MS (Method 1):  $R_t$  = 1.48 min; MS (ESIpos):  $m/z$  = 866 (M+H)<sup>+</sup>.

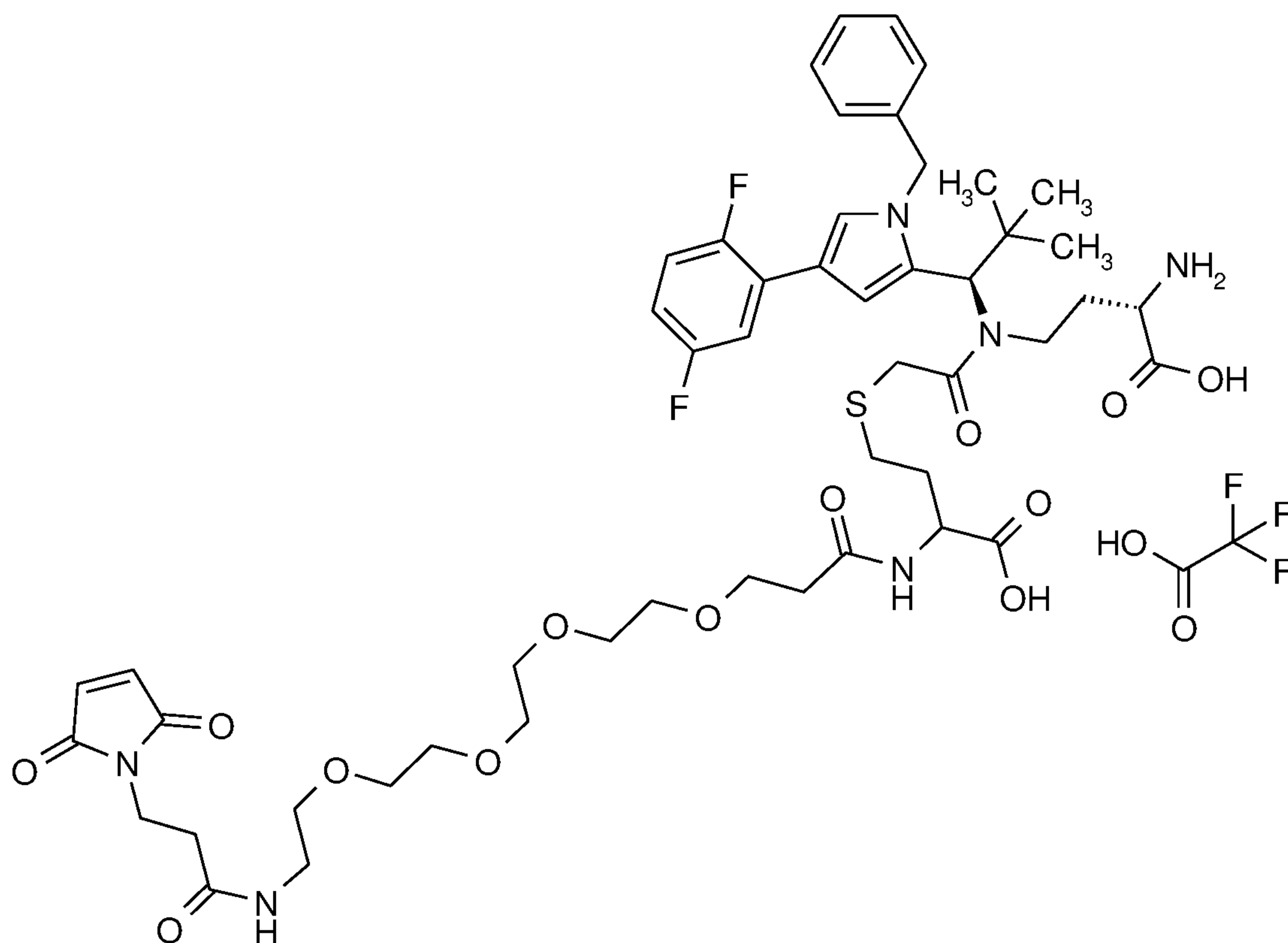
20.0 mg (0.02 mmol) of 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl){[(6-([2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]amino)-6-oxohexyl)sulphanyl]acetyl}amino)propyl]carbamate were dissolved in 2.0 ml of trifluoroethanol, and 4.7 mg (0.04 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C overnight, and 10.1 mg (0.04 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid were then added and the mixture was stirred for 10 min. Water (0.1% TFA) was added and the reaction mixture was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 9.2 mg (48% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.98 min; MS (ESIpos):  $m/z$  = 722 (M+H)<sup>+</sup>.

#### Intermediate F146



R/S-[2-([(3S)-3-Amino-3-carboxypropyl]{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)-2-oxoethyl]-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]homocysteine / trifluoroacetic acid (1:1)



10 25.0 mg (28.12  $\mu\text{mol}$ ) of R/S-[(8S)-11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-8-carboxy-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl]homocysteine (Intermediate C12) and 15.9 mg (30.93  $\mu\text{mol}$ ) of

15 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide were dissolved in 2.0 ml of DMF, and 11.4 mg (112.48  $\mu\text{mol}$ ) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight. 7.6 mg (0.13 mmol) of acetic acid were added and the reaction mixture was purified directly

20 by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum.

This gave 23.9 mg (59% of theory) of the compound R/S-[(8S)-11-  
{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
dimethylpropyl}-8-carboxy-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-  
diazasilatridecan-13-yl]-N-[19-(2,5-dioxo-2,5-dihydro-1H-  
5 pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-  
oyl]homocysteine.

LC-MS (Method 1):  $R_t = 1.26$  min; MS (ESIpos):  $m/z = 1173$  (M+H)<sup>+</sup>.

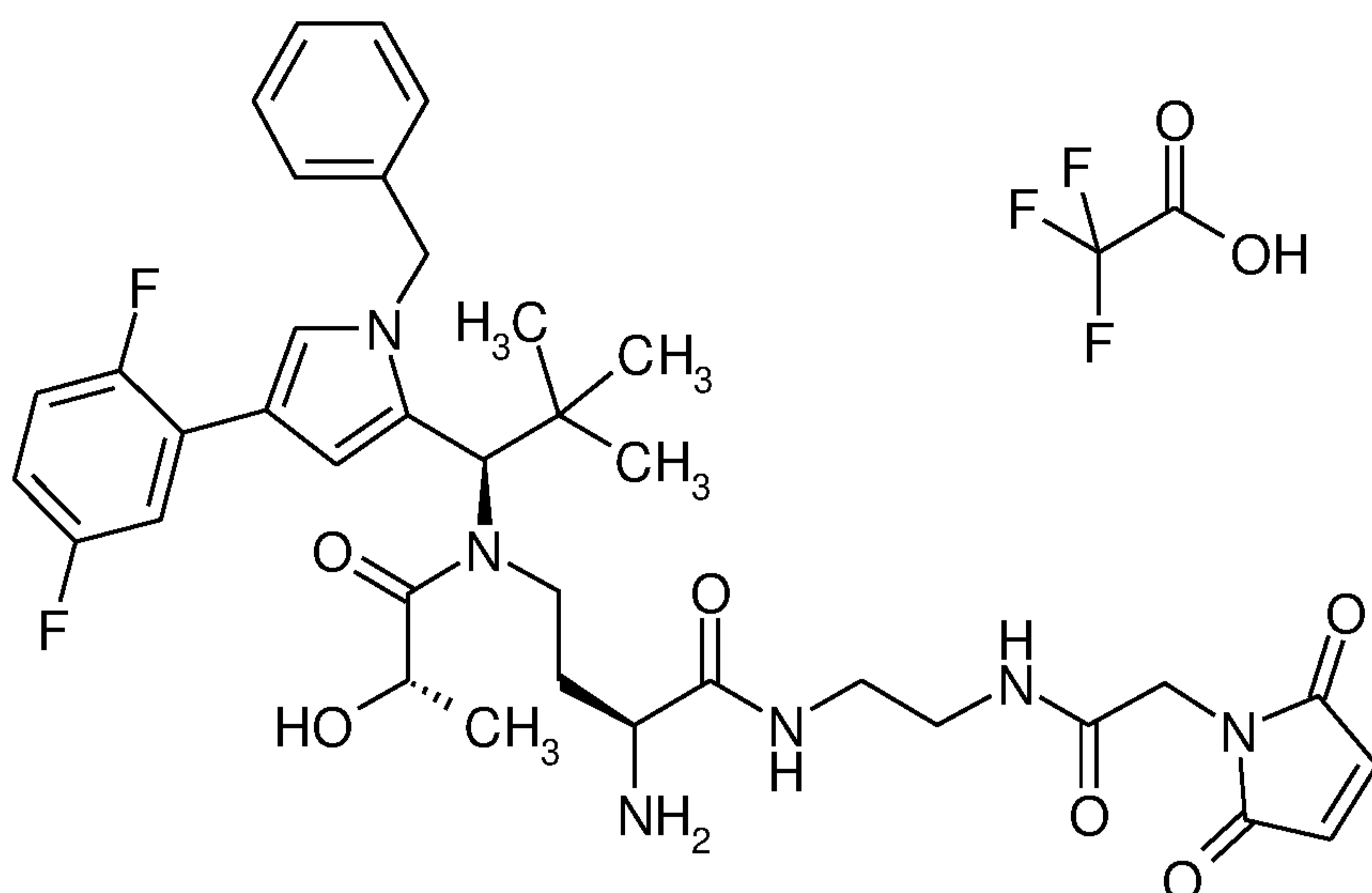
10 11.8 mg (8.23  $\mu$ mol) of R/S-[(8S)-11-[(1R)-1-[1-benzyl-4-(2,5-  
difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]-8-carboxy-  
2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diazasilatridecan-13-  
yl]-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-  
4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]homocysteine were  
15 dissolved in trifluoroethanol, and 1.7 mg (12.35  $\mu$ mol) of zinc  
dichloride were added. The reaction mixture was stirred at 50°C  
overnight. 3.6 mg (0.01 mmol) of ethylenediamine-N,N,N',N'-  
tetraacetic acid were added, the reaction mixture was stirred  
briefly and water (0.1% TFA) was then added. Purification was  
20 carried out directly by preparative RP-HPLC (column: Reprosil  
250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The  
solvents were evaporated under reduced pressure and the residue  
was dried under high vacuum. This gave 5.8 mg (62% of theory)  
of the title compound.

25

LC-MS (Method 4):  $R_t = 1.20$  min; MS (ESIpos):  $m/z = 1029$  (M+H)<sup>+</sup>.

### Intermediate F153

30 Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-  
(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl][(2S)-  
2-hydroxypropanoyl]amino)-N-(2-[[2,5-dioxo-2,5-dihydro-1H-  
pyrrol-1-yl]acetyl]amino)ethyl)butanamide (1:1)



The synthesis was carried out analogously to Intermediate F104 from Intermediate C60.

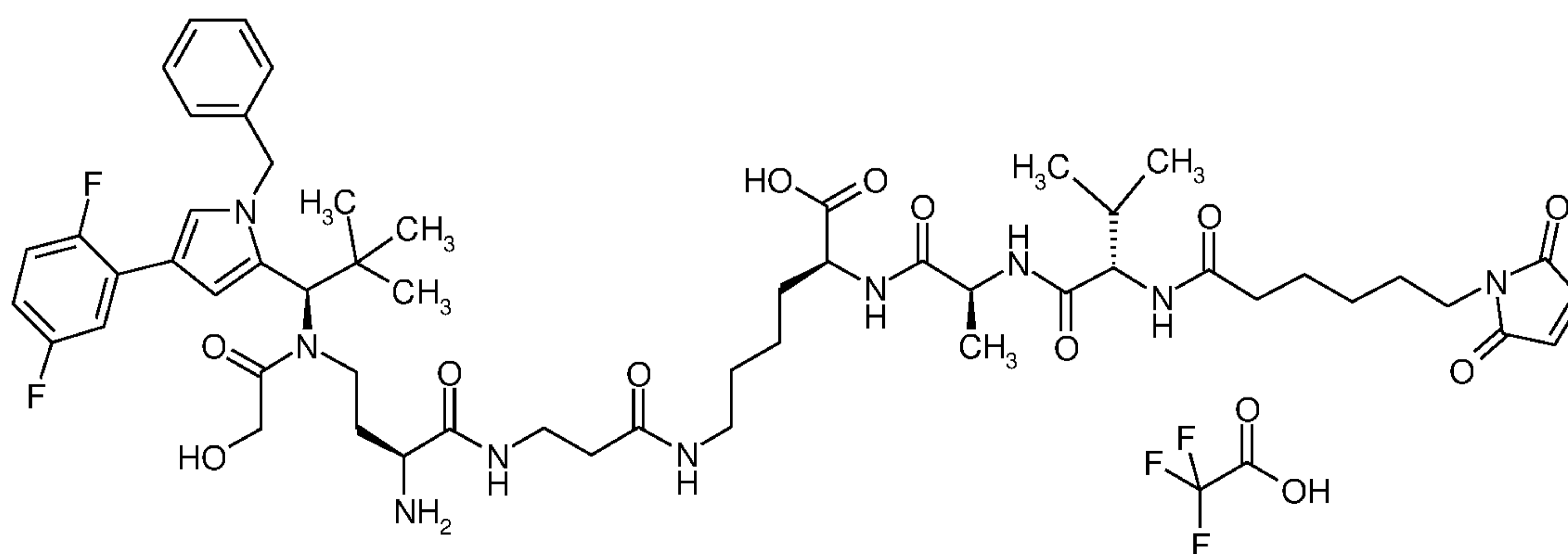
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LC-MS (Method 1):  $R_t = 1.1$  min; MS (ESIpos):  $m/z = 707$  (M+H)<sup>+</sup>.

### Intermediate F155

10 N<sup>6</sup>-(N-{(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)-N<sup>2</sup>-{N-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-L-alanyl}-L-lysine / trifluoroacetic acid (1:1)

15



The title compound was prepared by coupling of 14 mg (0.019 mmol) of Intermediate C61 with 15 mg (0.021 mmol) of Intermediate L61 in the presence of 8.7 mg (0.023 mmol) of HATU and 17  $\mu$ l of

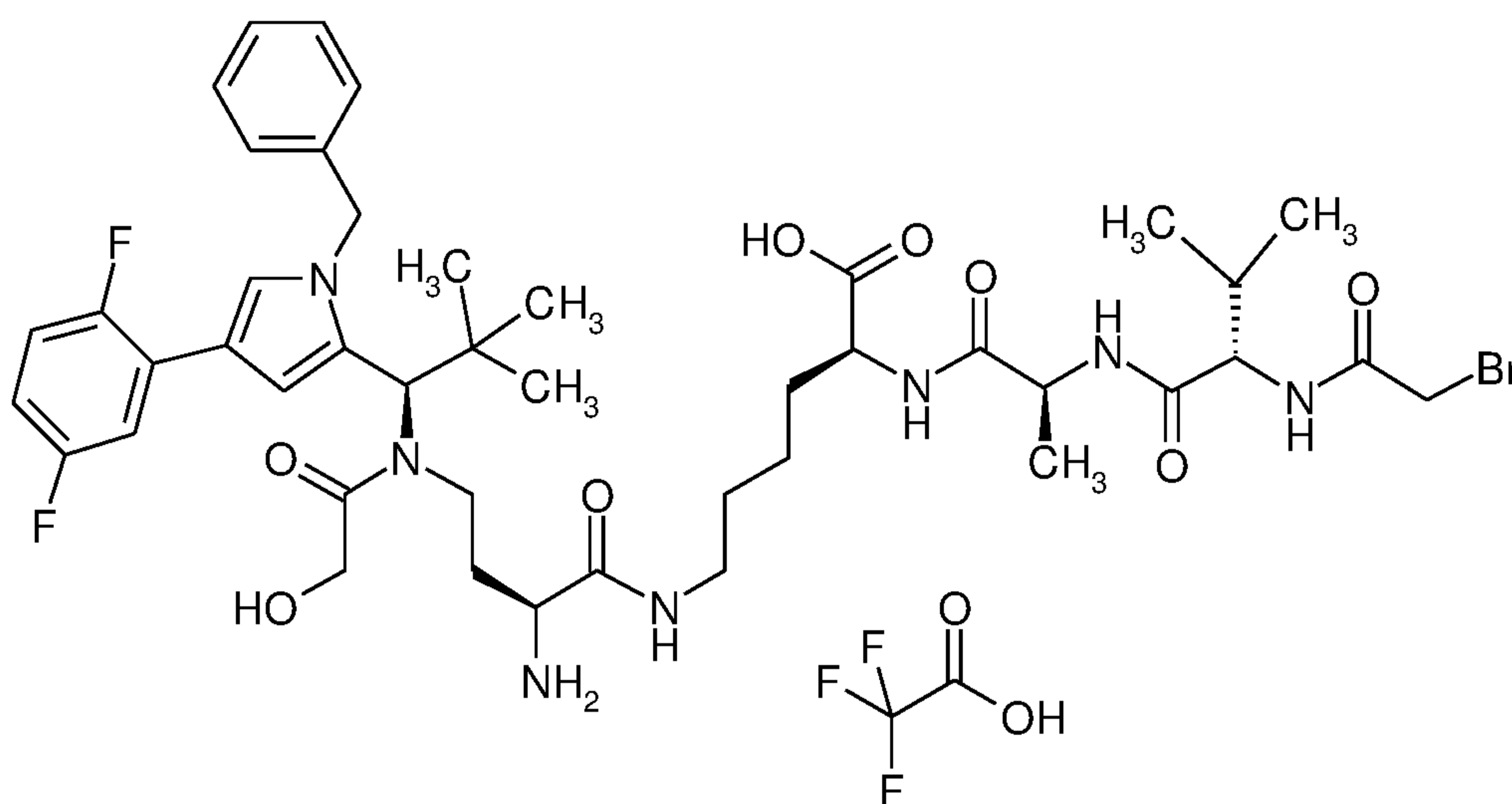
20 *N,N*-diisopropylethylamine and subsequent deprotection with zinc

chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 13 mg (59% of theory over 2 steps) of the title compound.

5 LC-MS (Method 1):  $R_t = 0.86$  min; MS (ESIpos):  $m/z = 1076$  (M+H)<sup>+</sup>.

### Intermediate F156

10 N-(Bromoacetyl)-L-valyl-L-alanyl-N<sup>6</sup>-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-lysine /  
trifluoroacetic acid (1:1)



15 First, the tripeptide derivative 2-(trimethylsilyl)ethyl-L-valyl-L-alanyl-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysinate was prepared from N<sup>2</sup>-[(benzyloxy)carbonyl]-N<sup>6</sup>-(tert-butoxycarbonyl)-L-lysine according to classical methods of peptide chemistry  
20 (esterification with 2-(trimethylsilylethanol using EDCI/DMAP, hydrogenolysis, coupling with N-[(benzyloxy)carbonyl]-L-valyl-L-alanine in the presence of HATU and another hydrogenolysis).

25 84 mg (0.163 mmol) of this Intermediate were taken up in 2.5 ml of DMF, and 58 mg (0.244 mmol) of 1-(2-bromoacetoxy)pyrrolidine-2,5-dione were added. After 10 min of stirring at RT, the mixture was concentrated, the residue was taken up in acetonitrile/water 1:1 and the mixture was adjusted with trifluoroacetic acid to

pH 2 and purified by preparative HPLC. After concentration of the appropriate fractions, the residue was taken up in 15 ml of a 5% strength trifluoroacetic acid solution in DCM and stirred at RT for 2 h. The mixture was then concentrated with slight cooling and the residue was lyophilized from acetonitrile/water 1:1. 53 mg (50% of theory) of this intermediate were obtained over 2 steps.

LC-MS (Method 1):  $R_t = 0.72$  min; MS (ESIpos):  $m/z = 537$  and  $539$  (M+H)<sup>+</sup>.

For the synthesis of the title compound, 18 mg (0.027 mmol) of this intermediate were taken up in 4 ml of DMF, and 16 mg (0.025 mmol) of Intermediate C61 and 19 mg of HATU and 9  $\mu$ l of *N,N*-diisopropylethylamine were added. After 5 min of stirring at RT, a few drops of trifluoroacetic acid were added and the reaction was purified by preparative HPLC. After concentration of the appropriate fractions and lyophilization from acetonitrile/water 1:1, the intermediate obtained was dissolved in 3 ml of 2,2,2-trifluoroethanol. Following addition of 4.8 mg (0.035 mmol) of zinc chloride, the reaction was stirred at 50°C for 2.5 h. 10 mg (0.035 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid were then added, and the reaction was diluted with acetonitrile/water and filtered. Purification was carried out by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 3.2 mg (13% of theory) of the title compound over 2 steps.

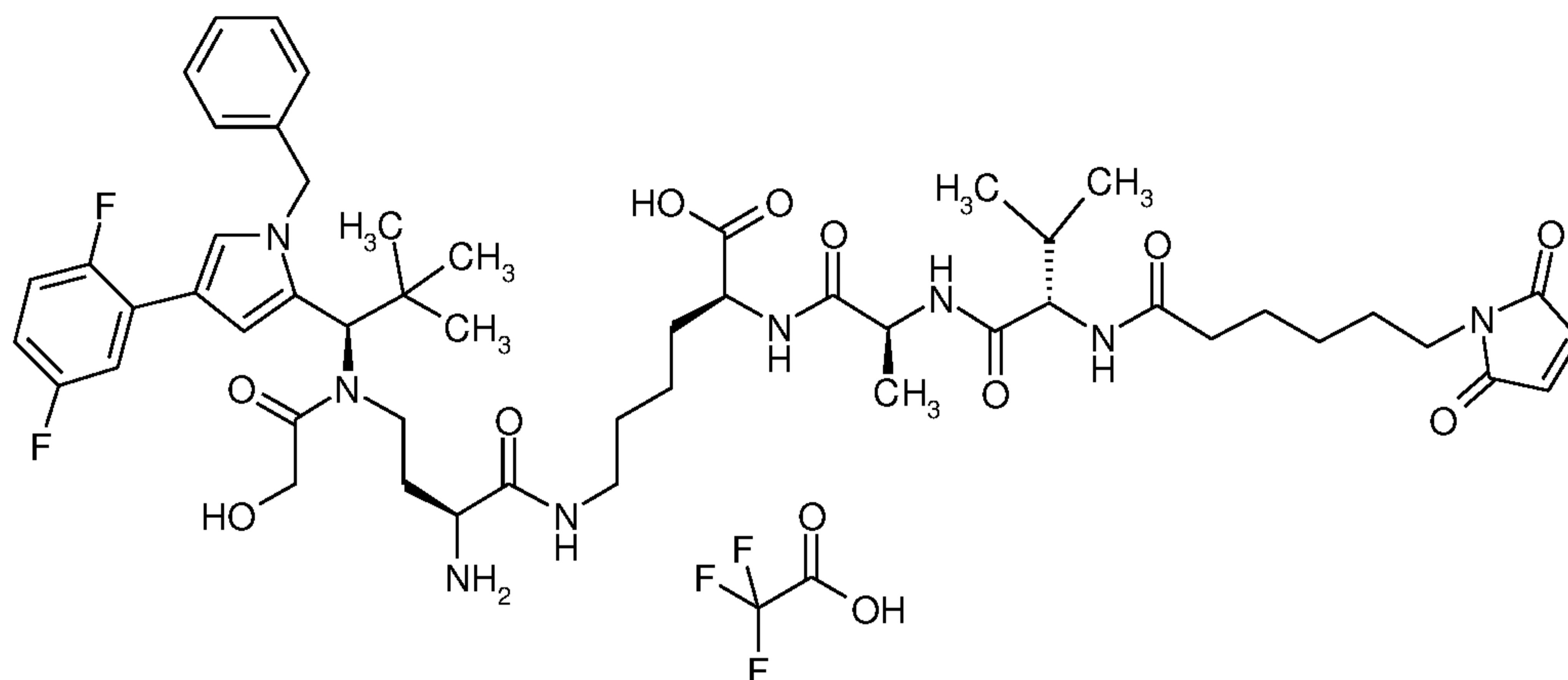
HPLC (Method 11):  $R_t = 1.94$  min;

LC-MS (Method 5):  $R_t = 2.79$  min; MS (ESIpos):  $m/z = 932$  and  $934$  (M+H)<sup>+</sup>.

### 35 Intermediate F163

*N*-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-*L*-valyl-*L*-alanyl-*N*6-[(2*S*)-2-amino-4-[(1*R*)-1-[1-benzyl-4-(2,5-

difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino] butanoyl}-L-lysine /  
trifluoroacetic acid (1:1)



5

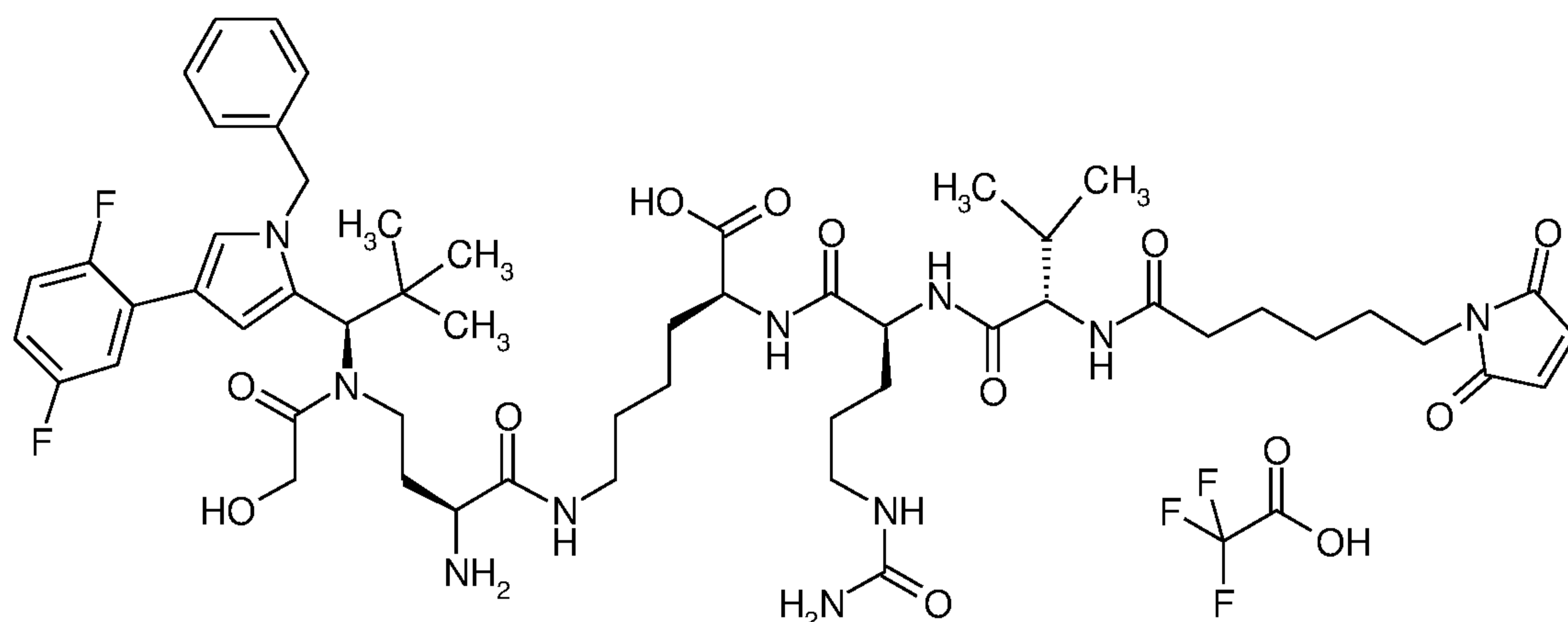
The title compound was prepared by coupling of 37 mg (0.056 mmol) of Intermediate C58 and 41 mg (0.056 mmol) of Intermediate L61 in the presence of HATU and subsequent deblocking with zinc chloride as described for Intermediate F119. This gave 12 mg (19% of theory over 2 steps) of the title compound.

HPLC (Method 11):  $R_t = 1.49$  min;

15 LC-MS (Method 1):  $R_t = 0.89$  min; MS (ESIpos):  $m/z = 1005$  (M+H)<sup>+</sup>.

### Intermediate F164

N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-  
20 N5-carbamoyl-L-ornithyl-N6-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-lysine /  
trifluoroacetic acid (1:1)



The title compound was prepared analogously to Intermediate F155 by coupling of 20 mg (0.030 mmol) of Intermediate C58 with 27 mg (0.033 mmol) of Intermediate L62 in the presence of HATU and *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol.

HPLC (Method 11):  $R_t = 1.92$  min;

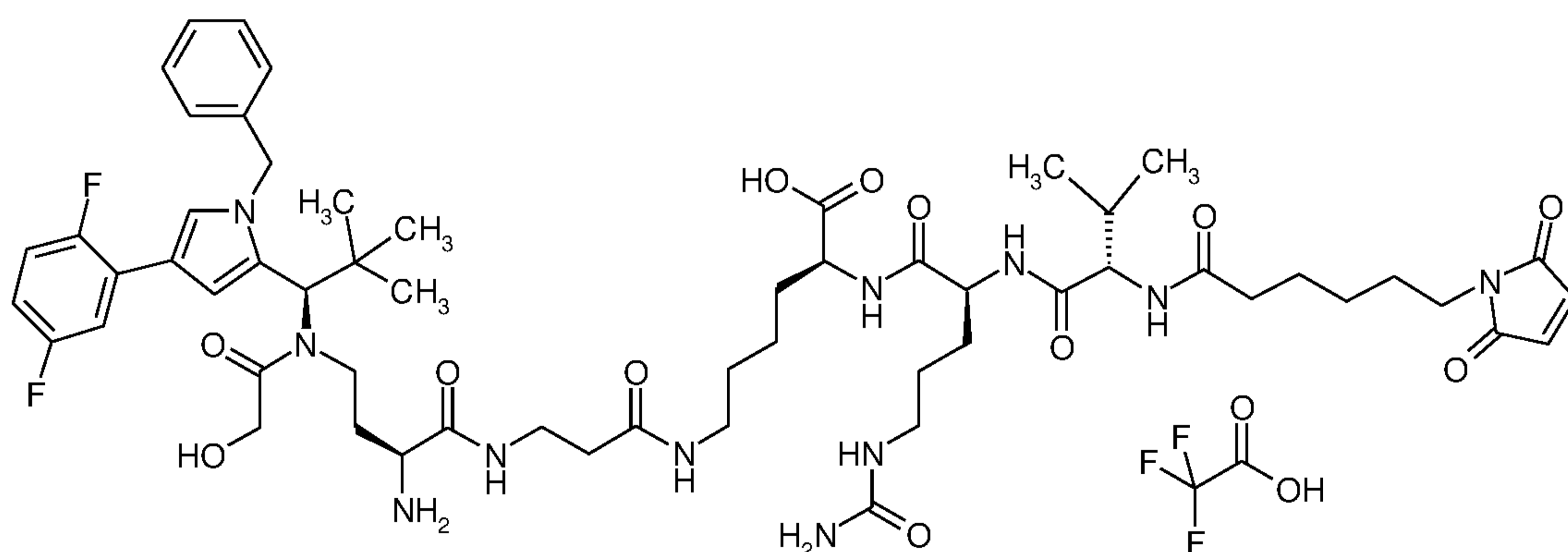
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LC-MS (Method 1):  $R_t = 0.87$  min; MS (ESIpos):  $m/z = 1091$  (M+H)<sup>+</sup>.

### Intermediate F165

15  $N^6$ - (N- { (2S)-2-Amino-4- [ { (1R)-1- [1-benzyl-4- (2,5-difluorophenyl)-1H-pyrrol-2-yl] -2,2-dimethylpropyl } (glycoloyl) amino] butanoyl } -beta-alanyl) - $N^2$ - { N- [6- (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl) hexanoyl] -L-valyl - $N^5$ -carbamoyl-L-ornithyl } -L-lysine / trifluoroacetic acid (1:1)

20



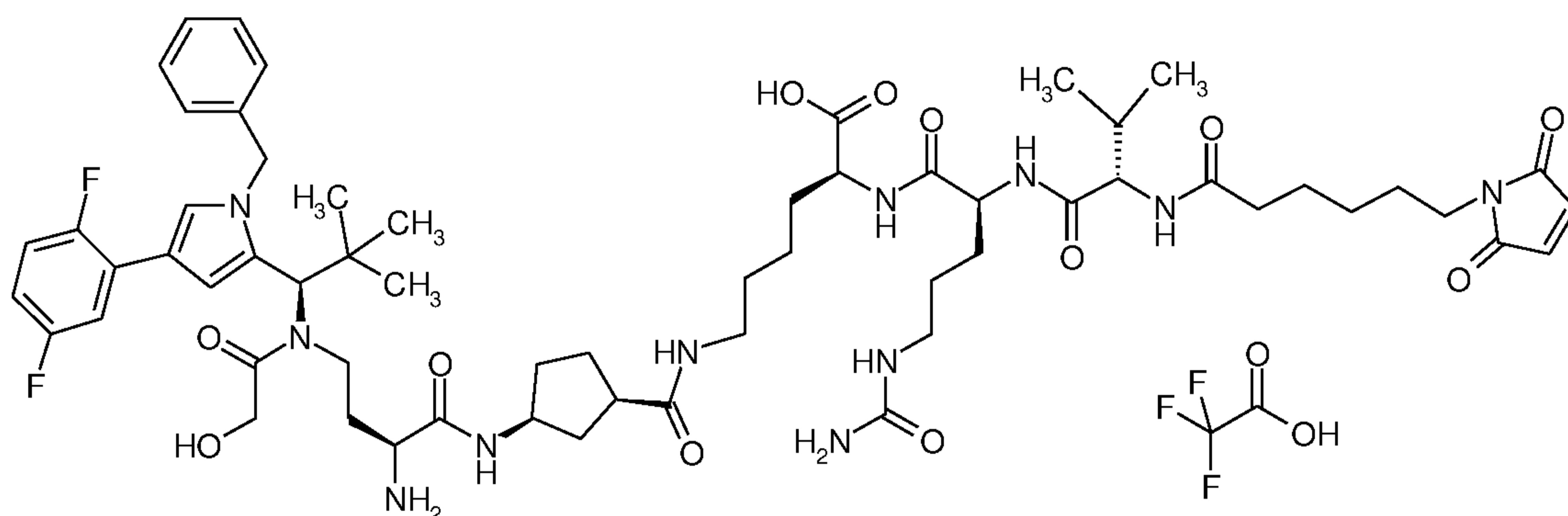
The title compound was prepared analogously to Intermediate F155

by coupling of 15 mg (0.021 mmol) of Intermediate C61 with 18 mg (0.023 mmol) of Intermediate L62 in the presence of HATU and subsequent deprotection with zinc chloride in trifluoroethanol.

5 LC-MS (Method 1):  $R_t = 0.88$  min; MS (ESIpos):  $m/z = 1162$  (M+H)<sup>+</sup>.

### Intermediate F166

10 N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithyl-N<sup>6</sup>-{[(1R,3S)-3-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)cyclopentyl]carbonyl}-L-lysine / trifluoroacetic acid (1:1)



15

First, trifluoroacetic acid / benzyl (1R, 3S)-3-aminocyclopentanecarboxylate (1:1) was prepared from commercially available (1R, 3S)-3-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid according to classical methods of peptide chemistry by esterification with benzyl alcohol using EDCI/DMAP and subsequent removal of the tert-butoxycarbonyl protective group with TFA in DCM.

25 51 mg (0.076 mmol) of this intermediate were taken up in 6 ml of DMF and coupled with 50 mg (0.076 mmol) of Intermediate C58 in the presence of HATU and *N,N*-diisopropylethylamine. After purification by preparative HPLC, the intermediate was taken up in methanol and hydrogenated over 10% palladium on activated carbon at RT under hydrogen standard pressure for 2 h. The catalyst was then filtered off, the solvent was removed under reduced pressure and the product was purified by preparative

30



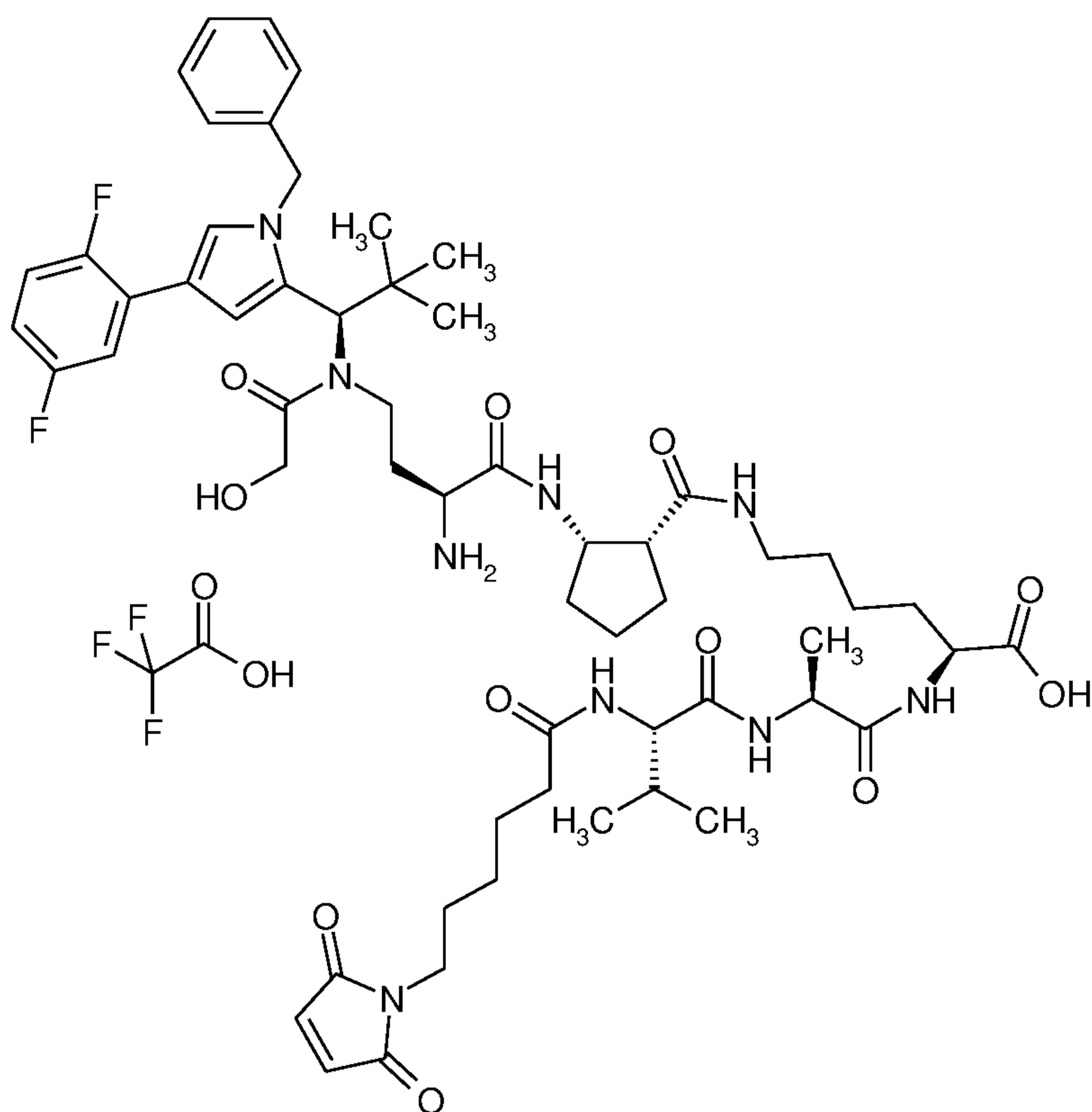
HPLC. Lyophilization from dioxane gave 21 mg (34% of theory over 2 steps) of (1R,3S)-3-{[(2S)-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-2-([2-(trimethylsilyl)ethoxy]carbonyl)amino)butanoyl]amino}cyclopentanecarboxylic acid.

The title compound was prepared analogously to Intermediate F155 by coupling of 10.5 mg (0.013 mmol) of this intermediate with 11.4 mg (0.014 mmol) of Intermediate L62 in the presence of HATU and subsequent deprotection with zinc chloride in trifluoroethanol. Purification by preparative HPLC gave 8.6 mg (48% of theory over 2 steps) of the title compound.

LC-MS (Method 1):  $R_t = 0.88$  min; MS (ESIpos):  $m/z = 1203$  (M+H)<sup>+</sup>.

#### Intermediate F168

N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-L-alanyl-N<sup>6</sup>-{[(1R,2S)-2-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl)amino)cyclopentyl]carbonyl}-L-lysine / trifluoroacetic acid (1:1)



First, trifluoroacetic acid / benzyl (1R,2S)-2-aminocyclopentanecarboxylate (1:1) was prepared from commercially available (1R,2S)-2-[(tert-butoxycarbonyl)amino]cyclopentanecarboxylic acid according to classical methods of peptide chemistry by esterification with benzyl alcohol using EDCI/DMAP and subsequent removal of the tert-butoxycarbonyl protective group with TFA in DCM.

102 mg (0.305 mmol) of this intermediate were taken up in 12 ml of DMF and coupled with 100 mg (0.152 mmol) of Intermediate C58 in the presence of HATU and *N,N*-diisopropylethylamine. After purification by preparative HPLC, the intermediate was taken up in methanol and hydrogenated over 10% palladium on activated carbon at RT under hydrogen standard pressure for 2 h. The catalyst was then filtered off, the solvent was removed under reduced pressure and the product was purified by preparative HPLC. Lyophilization from acetonitrile/water 1:1 gave 70 mg (59% of theory over 2 steps) of (1R,2S)-2-[[[(2S)-4-[[[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-2-({[2-(trimethylsilyl)ethoxy]carbonyl}amino)butanoyl]amino]cyclopent

anecarboxylic acid.

The title compound was then prepared by coupling of 20 mg (0.013 mmol) of this intermediate with 16.6 mg (0.023 mmol) of Intermediate L61 in the presence of 9.5 mg (0.025 mmol) of HATU and 18  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 9.3 mg (30% of theory over 2 steps) of the title compound.

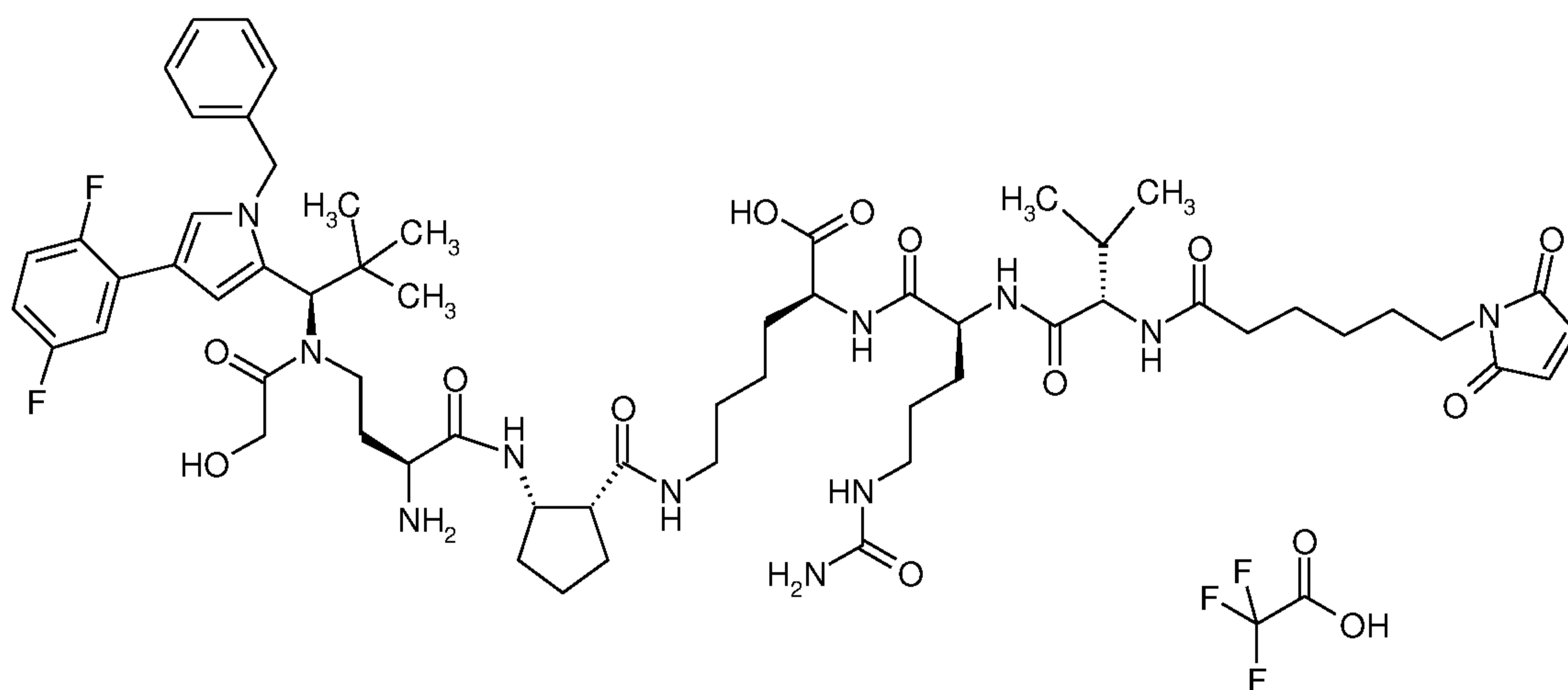
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LC-MS (Method 1):  $R_t = 0.98$  min; MS (ESIpos):  $m/z = 1116$  (M+H)<sup>+</sup>.

### Intermediate F169

15 N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithyl-N<sup>6</sup>-{[(1R,2S)-2-({(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)cyclopentyl]carbonyl}-L-lysine / trifluoroacetic acid (1:1)

20



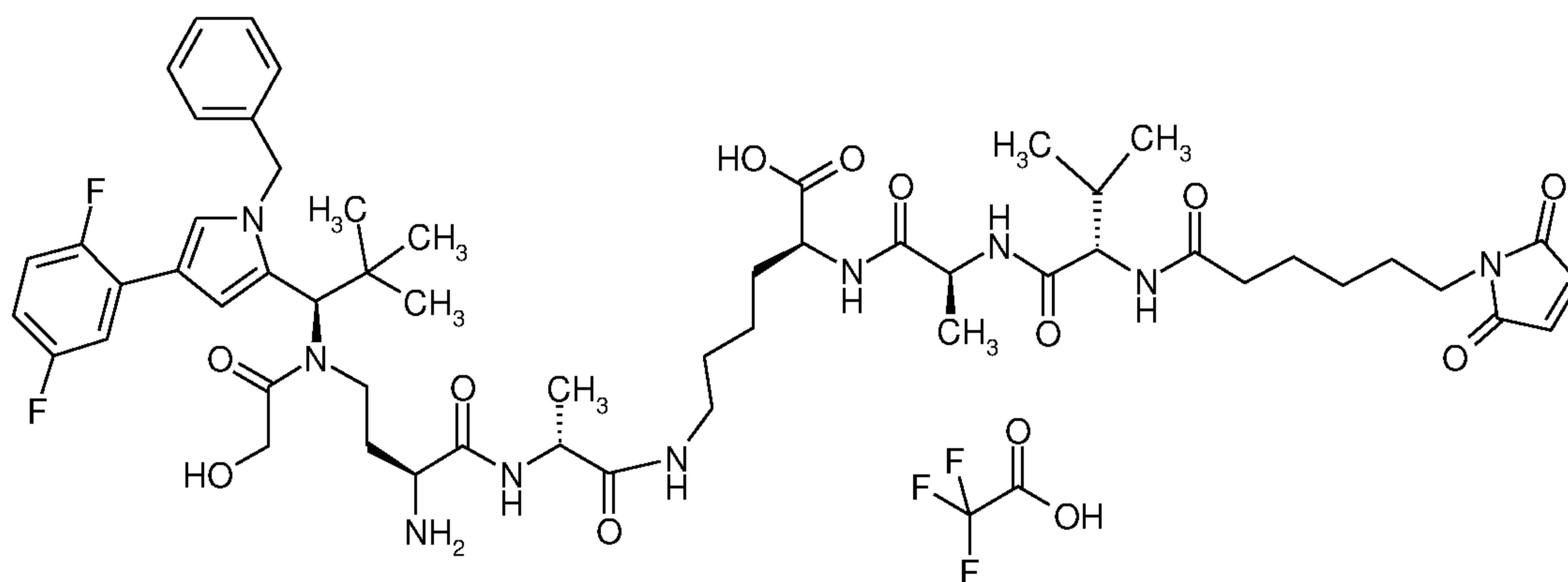
The synthesis of the title compound was carried out analogously to Intermediate F168 from Intermediates C58 and L62.

25

LC-MS (Method 1):  $R_t = 0.91$  min; MS (ESIpos):  $m/z = 1202$  (M+H)<sup>+</sup>.

### Intermediate F171

N<sup>6</sup>- (N- { (2S) -2-Amino-4- [ { (1R) -1- [1-benzyl-4- (2, 5-  
 difluorophenyl) -1H-pyrrol-2-yl] -2, 2-  
 dimethylpropyl } (glycoloyl) amino] butanoyl } -D-alanyl) -N<sup>2</sup>- { N- [6-  
 (2, 5-dioxo-2, 5-dihydro-1H-pyrrol-1-yl) hexanoyl] -L-valyl-L-  
 5 alanyl} -L-lysine / trifluoroacetic acid (1:1)

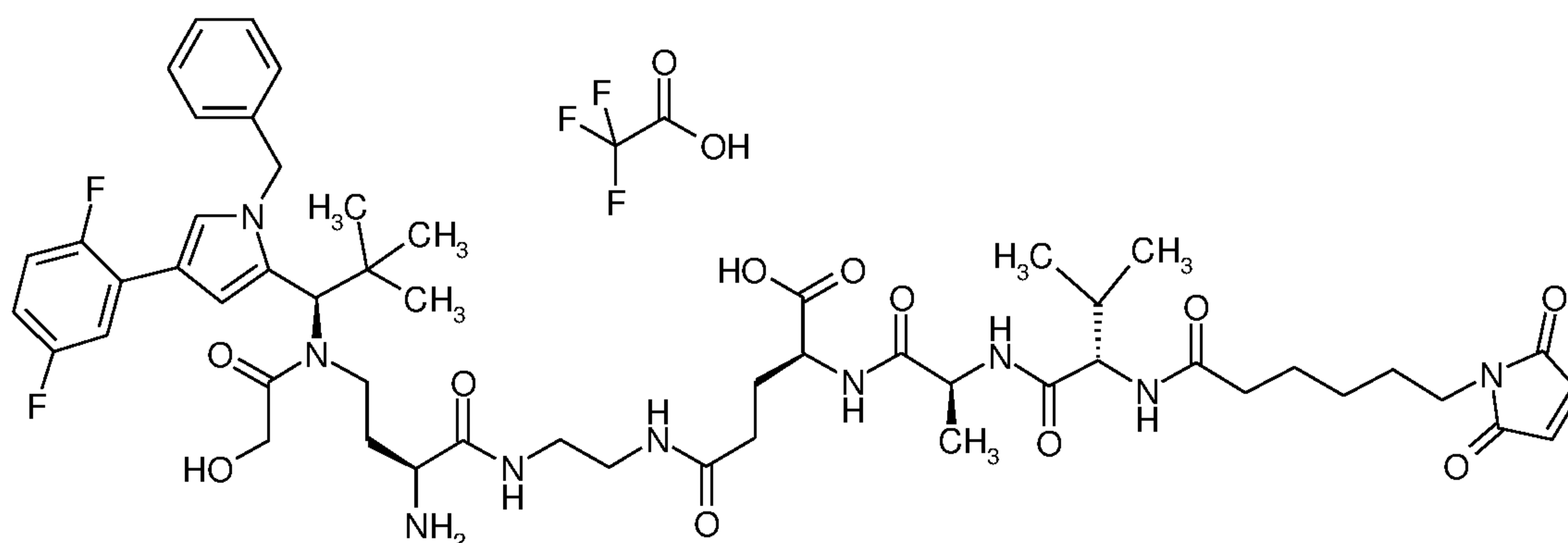


The synthesis of the title compound was carried out analogously  
 10 to Intermediate F155 from Intermediates C62 and L61.

LC-MS (Method 1): R<sub>t</sub> = 0.93 min; MS (ESIpos): m/z = 1076 (M+H)<sup>+</sup>.

### Intermediate F173

15 N- [6- (2, 5-Dioxo-2, 5-dihydro-1H-pyrrol-1-yl) hexanoyl] -L-valyl-  
 L-alanyl-N- [2- ( { (2S) -2-amino-4- [ { (1R) -1- [1-benzyl-4- (2, 5-  
 difluorophenyl) -1H-pyrrol-2-yl] -2, 2-  
 dimethylpropyl } (glycoloyl) amino] butanoyl } amino) ethyl] -L-  
 20 glutamine / trifluoroacetic acid (1:1)



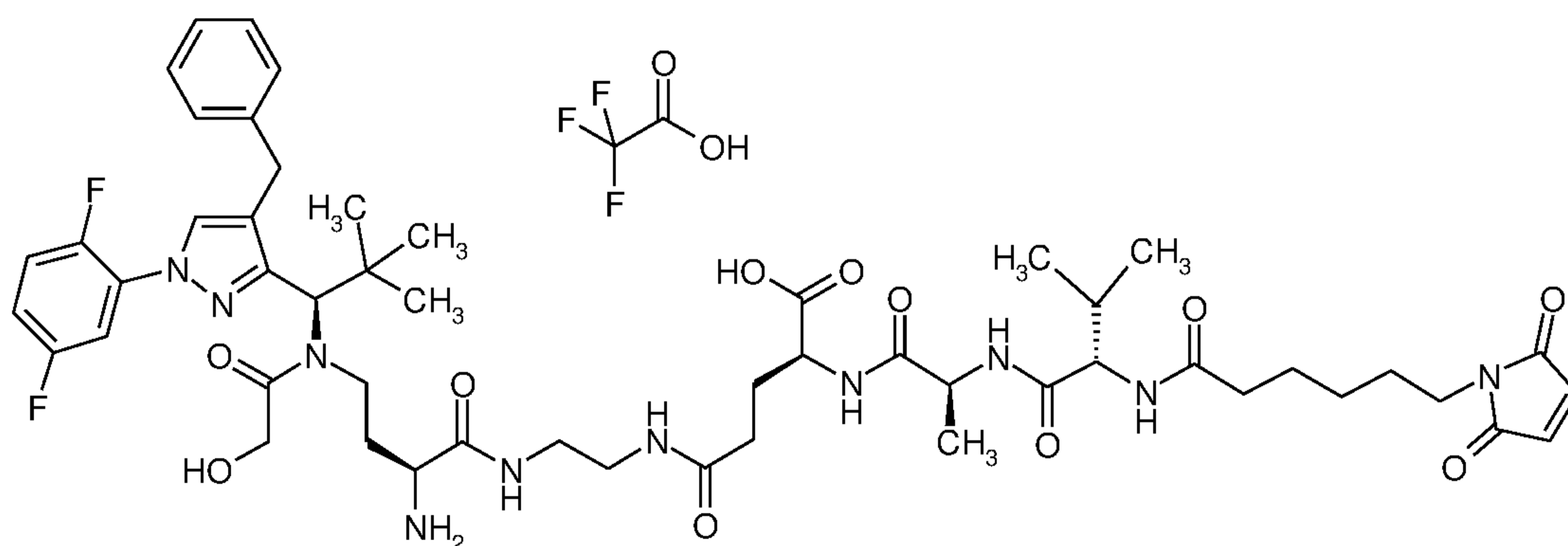
The title compound was prepared from 15 mg (0.018 mmol) of

Intermediate C64 by coupling with 12 mg (0.02 mmol) of Intermediate L63 in the presence of 7.7 mg (0.02 mmol) of HATU and 16  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 12 mg (58% of theory over 2 steps) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.91 min; MS (EIpos):  $m/z$  = 1048  $[M+H]^+$ .

### 10 Intermediate F174

N-[6-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoyl]-L-valyl-L-alanyl-N-[2-((2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]-L-glutamine / trifluoroacetic acid (1:1)

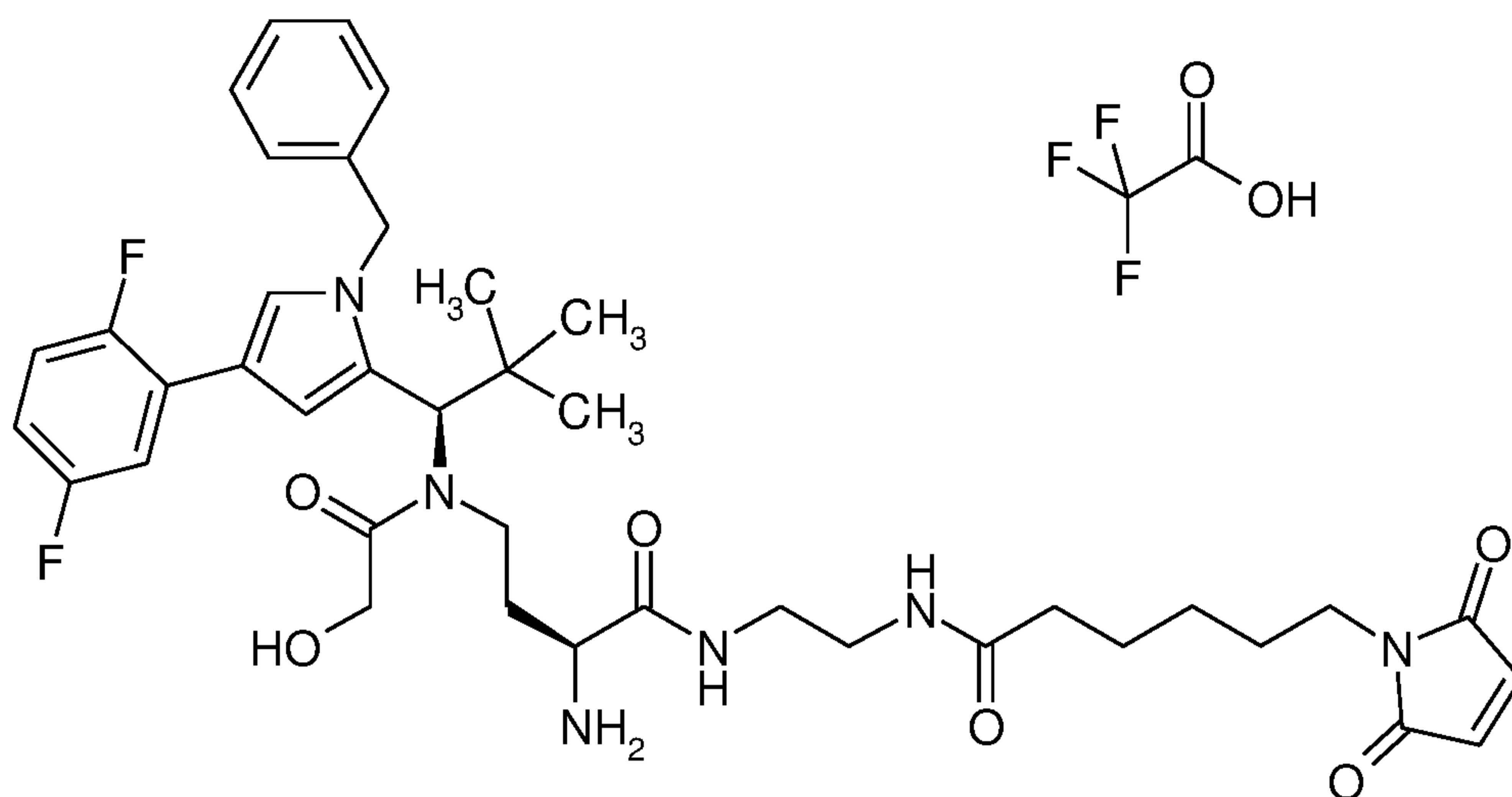


20 The title compound was prepared analogously to Intermediate F172 from Intermediates C57 and L63.

LC-MS (Method 1):  $R_t$  = 0.9 min; MS (EIpos):  $m/z$  = 1049  $[M+H]^+$ .

### 25 Intermediate F175

Trifluoroacetic acid / N-[2-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]-6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanamide (1:1)

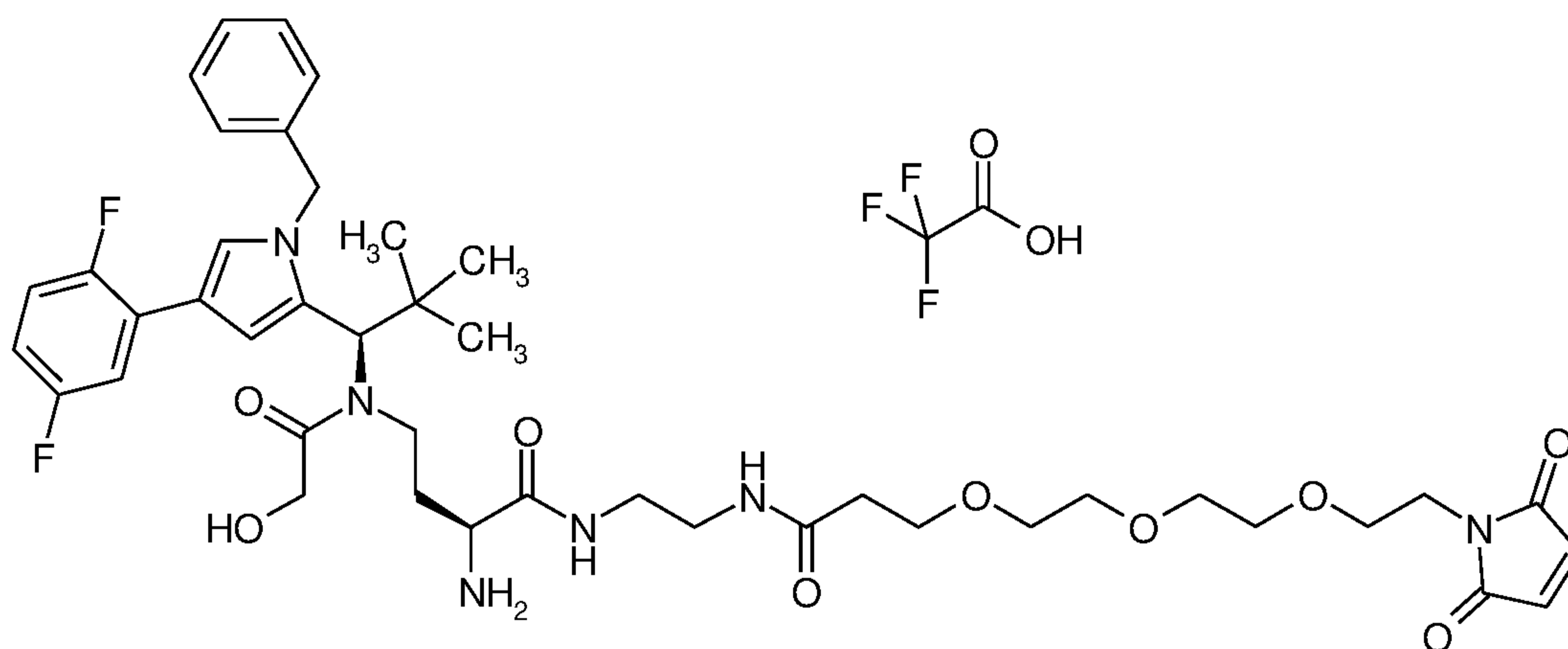


The title compound was prepared by coupling of 11 mg (0.013 mmol) of Intermediate C64 with 3.4 mg (0.016 mmol) of 6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanoic acid in the presence of 6.7 mg (0.018 mmol) of HATU and 9  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 8 mg (69% of theory over 2 steps) of the title compound.

LC-MS (Method 1):  $R_t$  = 1.35 min; MS (EIpos):  $m/z$  = 893  $[M+H]^+$ .

### Intermediate F176

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]-N-[1-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-12-oxo-3,6,9-trioxa-13-azapentadecan-15-yl]butanamide (1:1)



The title compound was prepared by coupling of 5 mg (0.006 mmol) of Intermediate C64 with 2 mg (0.007 mmol) of 3-(2-{2-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethoxy]ethoxy}ethoxy)propanoic acid, the preparation of which is described under Intermediate L15, in the presence of 3.5 mg (0.009 mmol) of HATU and 4  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 2 mg (35% of theory over 2 steps) of the title compound.

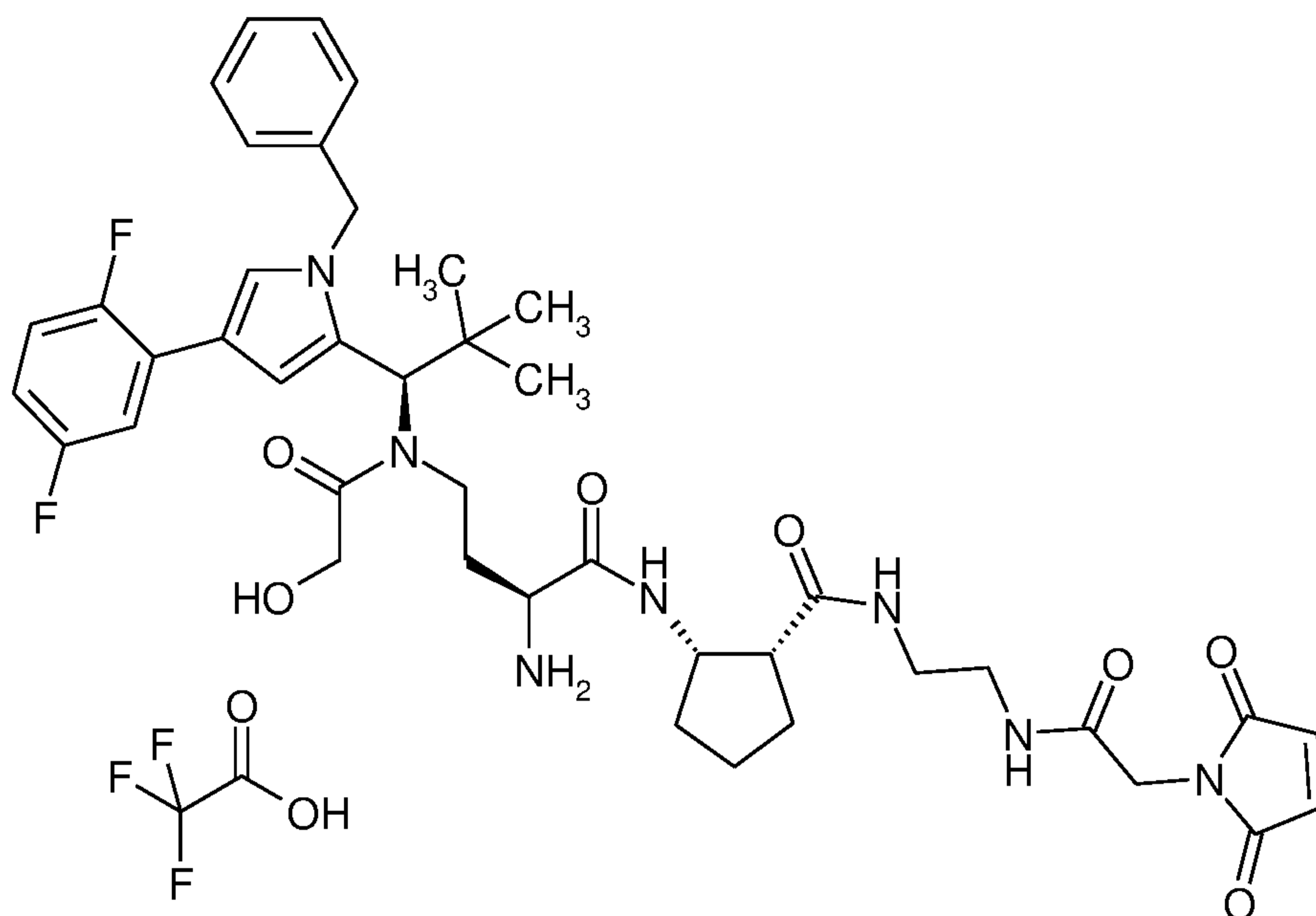
LC-MS (Method 1):  $R_t$  = 0.86 min; MS (EIpos):  $m/z$  = 839  $[M+H]^+$ .

15

### Intermediate F177

Trifluoroacetic acid / (1*R*,2*S*)-2-({(2*S*)-2-amino-4-[(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-*N*-(2-{[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino}ethyl)cyclopentanecarboxamide (1:1)

20



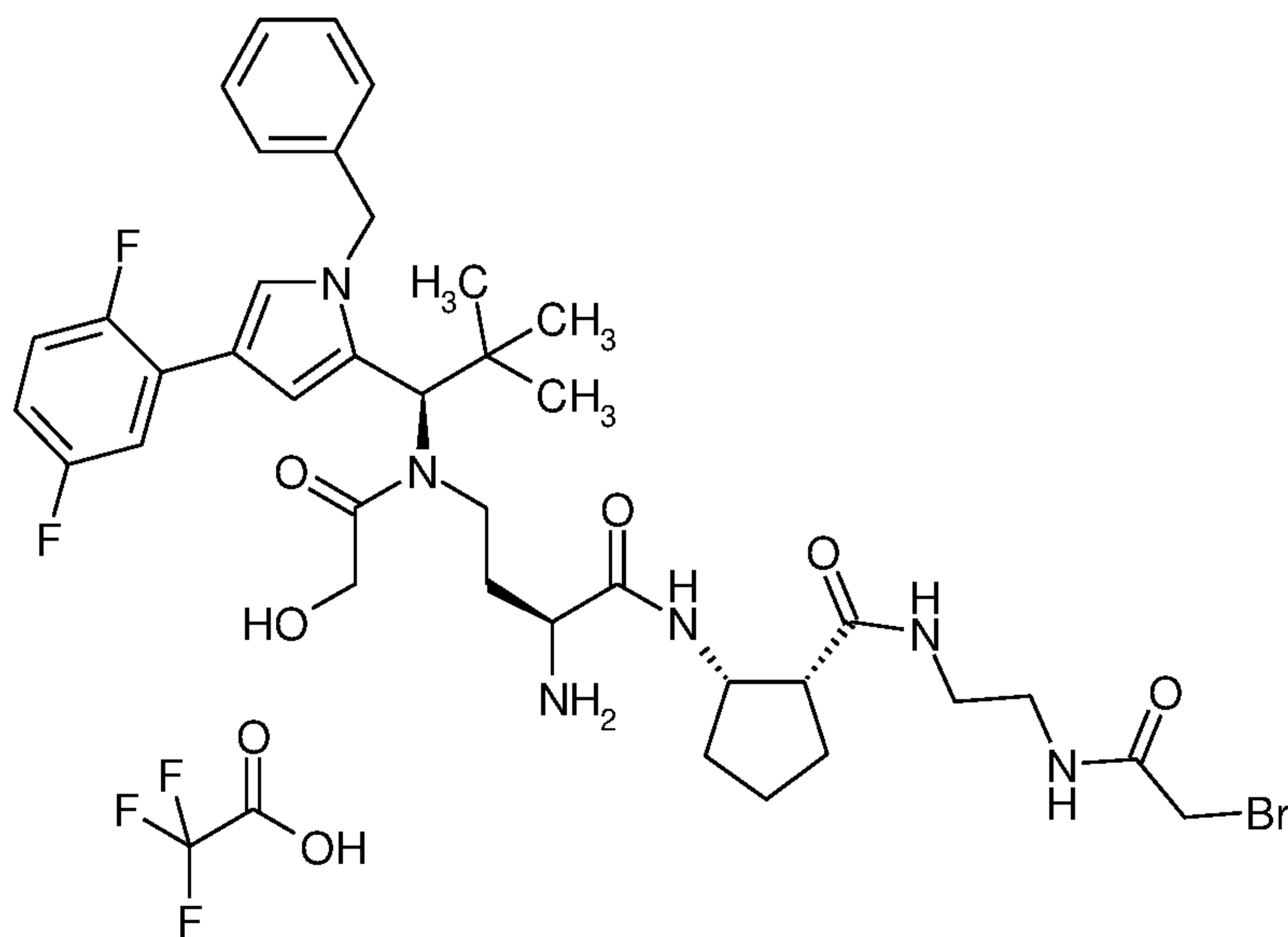
The title compound was prepared analogously to Intermediate F168 using, instead of Intermediate L61, the Intermediate L1.

5

LC-MS (Method 1):  $R_t = 0.86$  min; MS (EIpos):  $m/z = 804$   $[M+H]^+$ .

### Intermediate F178

10 Trifluoroacetic acid / (1R,2S)-2-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl)amino)-N-{2-[(bromoacetyl)amino]ethyl}cyclopentanecarboxamide (1:1)



15

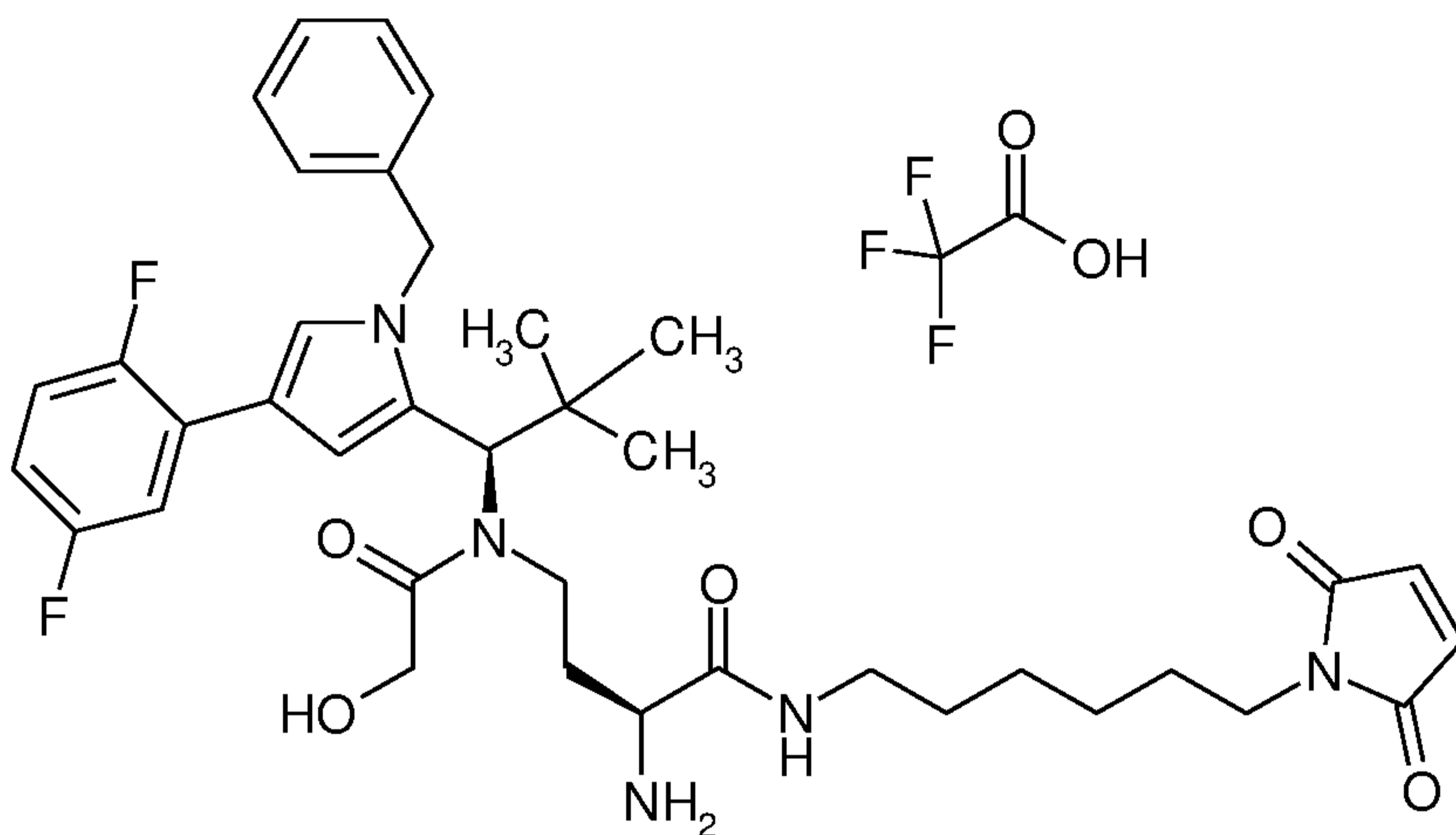


The title compound was prepared analogously to Intermediate F177 using, instead of Intermediate L1, the Intermediate L52.

- 5 LC-MS (Method 1):  $R_t = 0.89$  min; MS (EIpos):  $m/z = 787$  and  $789$   $[M+H]^+$ .

### Intermediate F179

- 10 Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-[6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexyl]butanamide (1:1)



15

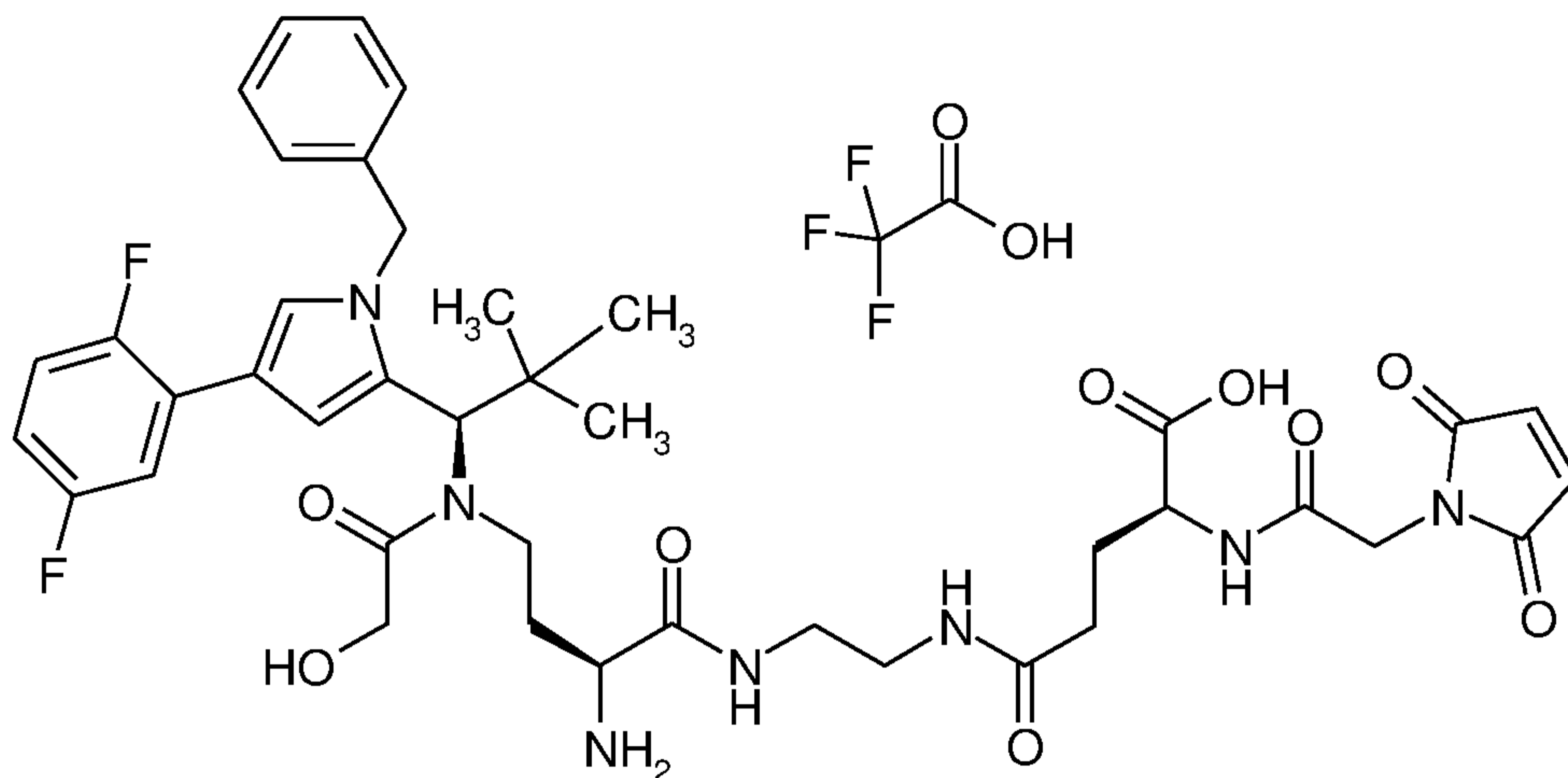
- The title compound was prepared by coupling of 15 mg (0.023 mmol) of Intermediate C58 with 6 mg (0.025 mmol) of 1-(6-aminohexyl)-1H-pyrrole-2,5-dione in the presence of 13 mg (0.034 mmol) of HATU and 16  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 8.5 mg (46% of theory over 2 steps) of the title compound.

25

- LC-MS (Method 6):  $R_t = 2.22$  min; MS (EIpos):  $m/z = 692$   $[M+H]^+$ .

### Intermediate F180

N-[2-((2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl)amino)ethyl]-N2-[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]-L-glutamine /  
 5 trifluoroacetic acid (1:1)



The title compound was prepared by coupling of 9.6 mg (0.012  
 10 mmol) of Intermediate C64 with 5 mg (0.013 mmol) of Intermediate  
 L64 in the presence of 7 mg (0.018 mmol) of HATU and 6  $\mu$ l of  
*N,N*-diisopropylethylamine and subsequent deprotection with zinc  
 chloride in trifluoroethanol as described for Intermediate F119.  
 Purification by preparative HPLC gave 3.1 mg (28% of theory over  
 15 2 steps) of the title compound.

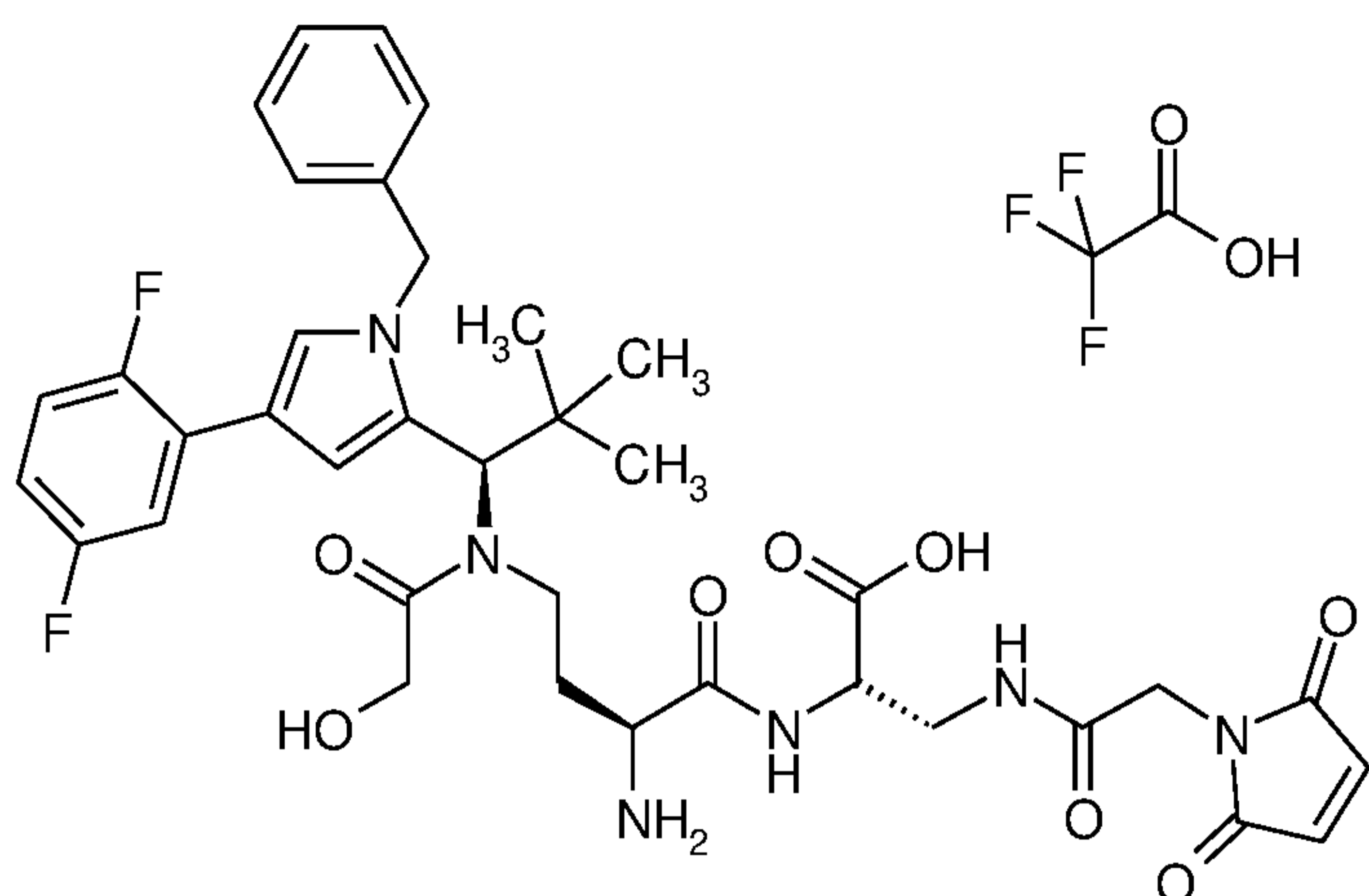
LC-MS (Method 1):  $R_t$  = 0.85 min; MS (EIpos):  $m/z$  = 822  $[M+H]^+$ .

### Intermediate F192

20

N-((2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl)-  
 3-[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino-L-  
 alanine / trifluoroacetic acid (1:1)

25



60 mg (0.091 mmol) of Intermediate C58 were taken up in 8 ml of DMF and coupled with 45 mg (0.100 mmol) of Intermediate L65 in the presence of 42 mg (0.11 mmol) of HATU and 64  $\mu$ l of *N,N*-diisopropylethylamine. After purification by preparative HPLC, the intermediate was taken up in 10 ml of ethanol and hydrogenated over 10% palladium on activated carbon at RT under hydrogen standard pressure for 45 min. The catalyst was then filtered off, the solvent was removed under reduced pressure and the product was purified by preparative HPLC. Lyophilization from acetonitrile/water 1:1 gave 24.5 mg (31% of theory over 2 steps) of 2-(trimethylsilyl)ethyl 3-amino-N-[(2S)-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-2-((2-(trimethylsilyl)ethoxy)carbonyl)amino)butanoyl]-L-alaninate.

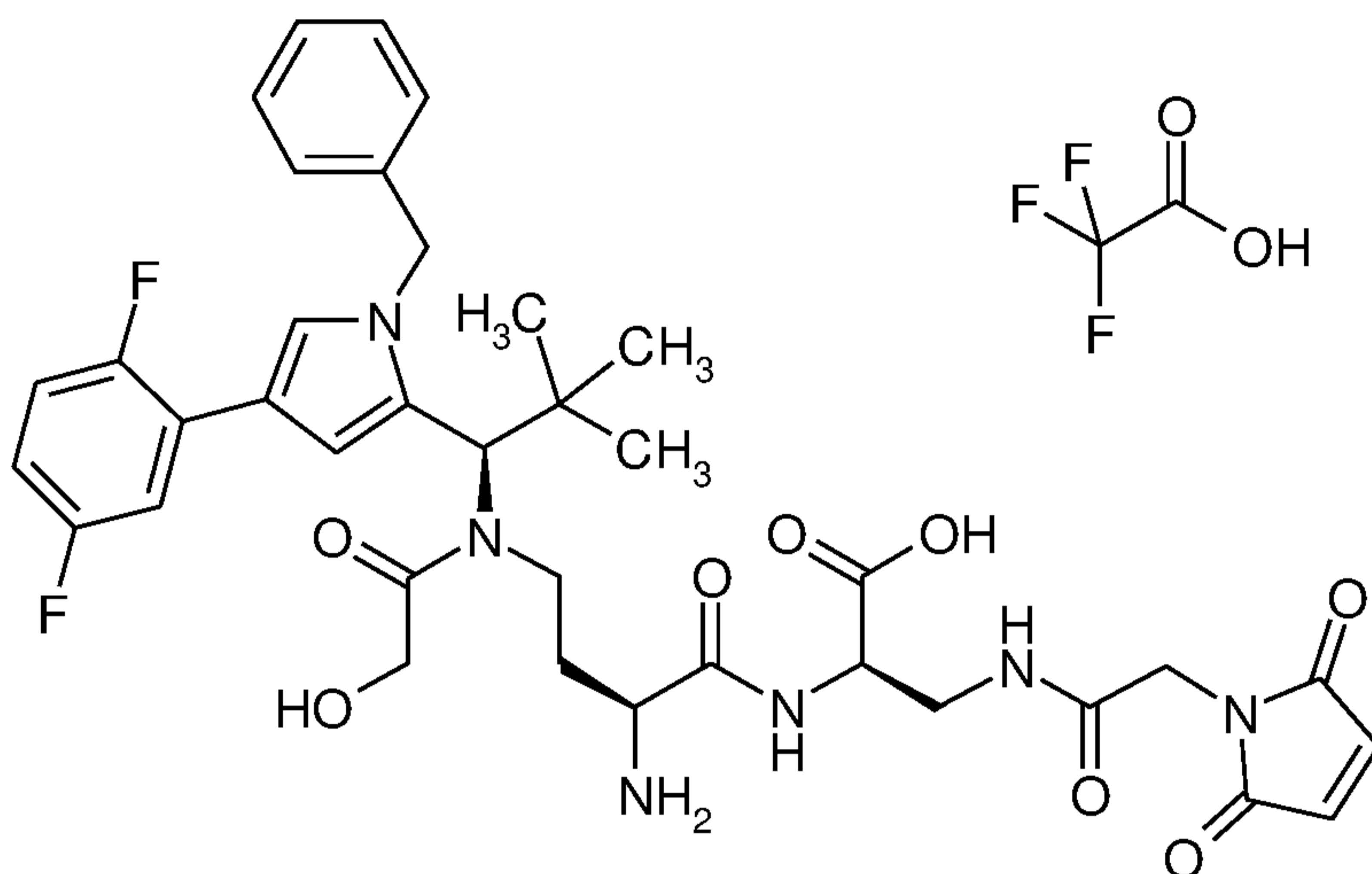
LC-MS (Method 1):  $R_t$  = 1.17 min; MS (EIpos):  $m/z$  = 844  $[M+H]^+$ .

The title compound was then prepared by coupling of 10 mg (0.012 mmol) of this intermediate with 2 mg (0.013 mmol) of commercially available (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetic acid intermediate in the presence of 5.4 mg (0.014 mmol) of HATU and 8  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 3.5 mg (33% of theory over 2 steps) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.81 min; MS (ESIpos):  $m/z$  = 737  $(M+H)^+$ .

**Intermediate F193**

N-{(2S)-2-Amino-4-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl]-3-[[{(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino}-D-alanine / trifluoroacetic acid (1:1)

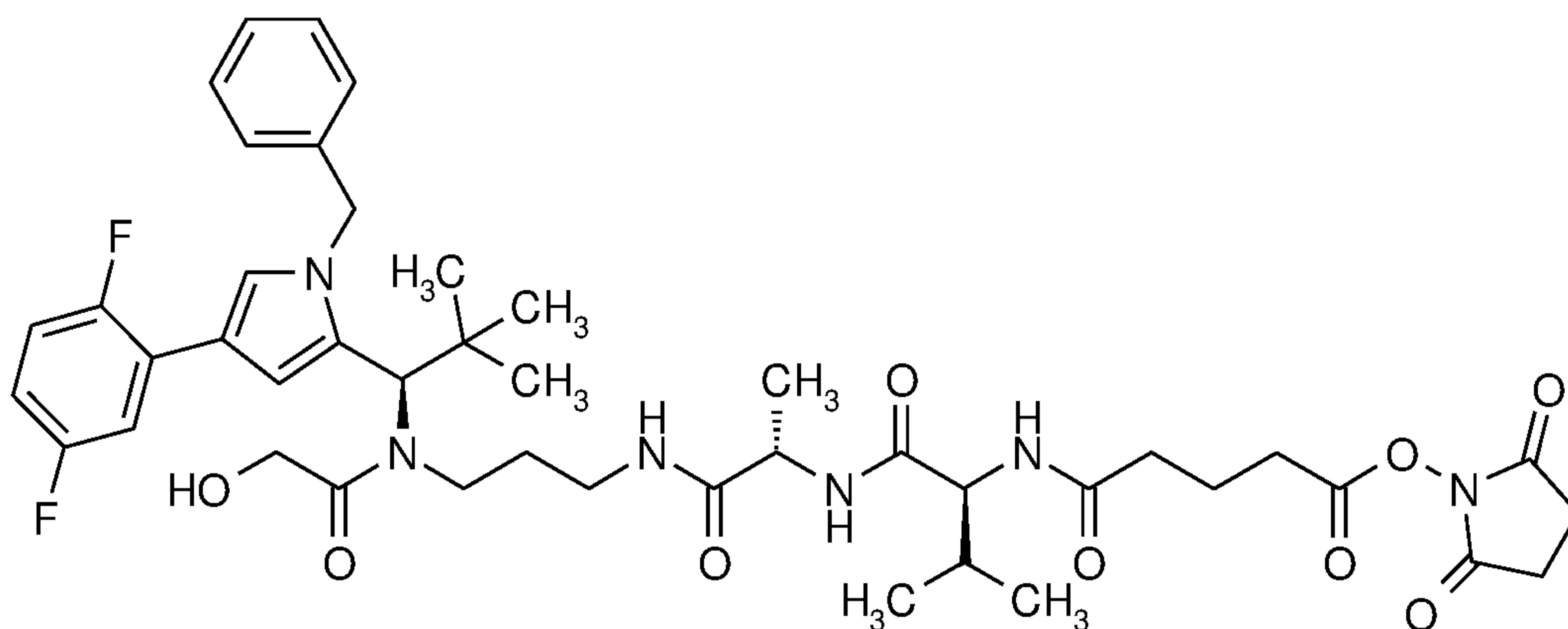


10 The synthesis of the title compound was carried out analogously to Intermediate F192 from 3-[[{(benzyloxy)carbonyl]amino}-N-(tert-butoxycarbonyl)-D-alanine / N-cyclohexylcyclohexanamine (1:1).

15 LC-MS (Method 1):  $R_t = 0.87$  min; MS (ESIpos):  $m/z = 737$  (M+H)<sup>+</sup>.

**Intermediate F194**

N-{5-[(2,5-Dioxopyrrolidin-1-yl)oxy]-5-oxopentanoyl}-L-valyl-  
 20 N-{3-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propyl}-L-alaninamide

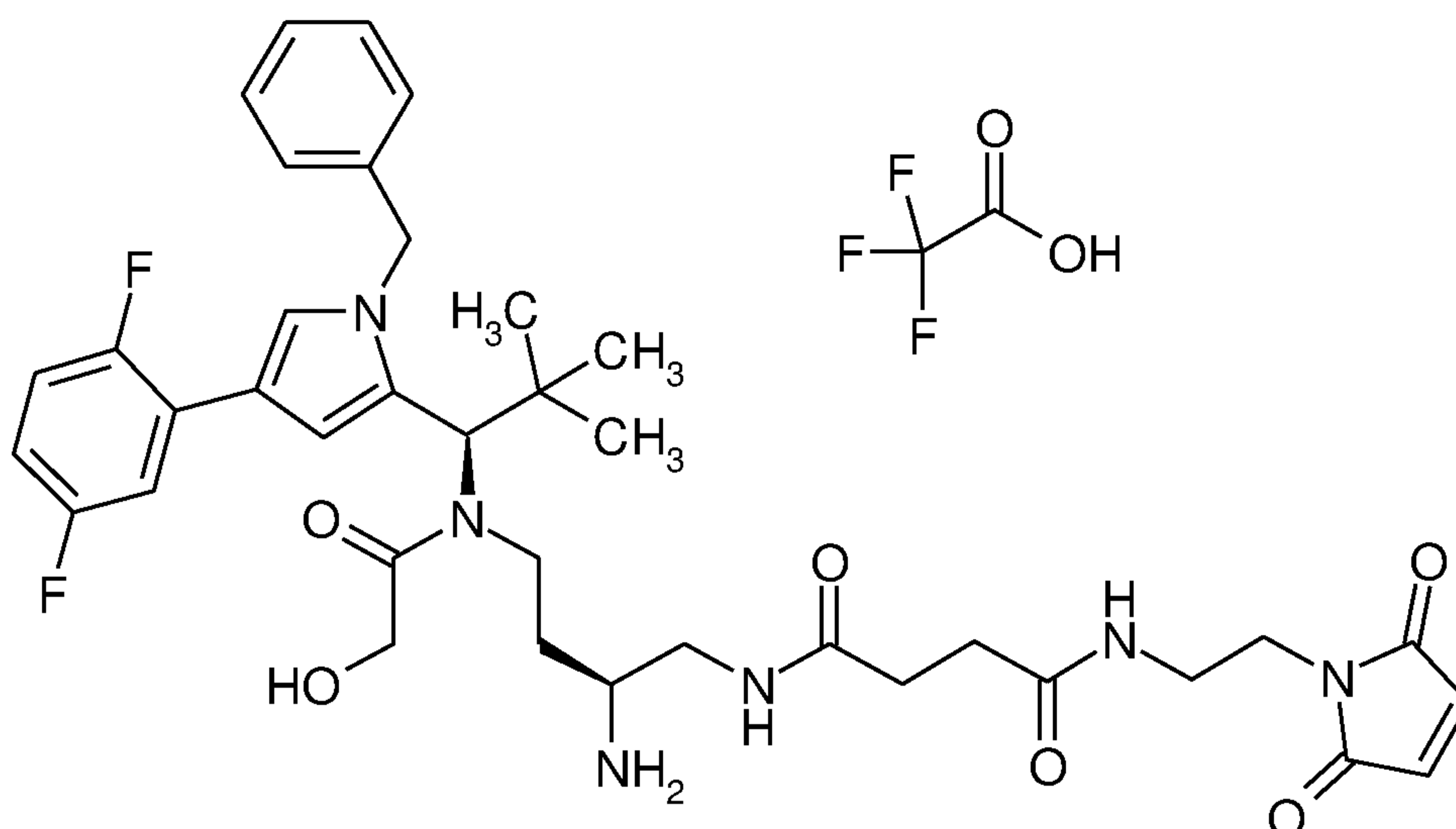


The title compound was prepared from Example 98 first by coupling with N-[(benzyloxy)carbonyl]-L-valyl-L-alanine in the presence of HATU and *N,N*-diisopropylethylamine. In the next step, the Z protective group was removed by hydrogenating for 1 hour over 10% palladium on activated carbon at RT under hydrogen standard pressure and then converting the deprotected intermediate as described for Intermediate F58 by reaction with 1,1'-[(1,5-dioxopentane-1,5-diyl)bis(oxy)]dipyrrolidine-2,5-dione into the title compound.

LC-MS (Method 1):  $R_t = 1.19$  min; MS (ESIpos):  $m/z = 851$   $[M+H]^+$ .

### 15 Intermediate F195

Trifluoroacetic acid / N-[(2*S*)-2-amino-4-[(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]butyl]-N'-[2-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)ethyl]succinamide (1:1)



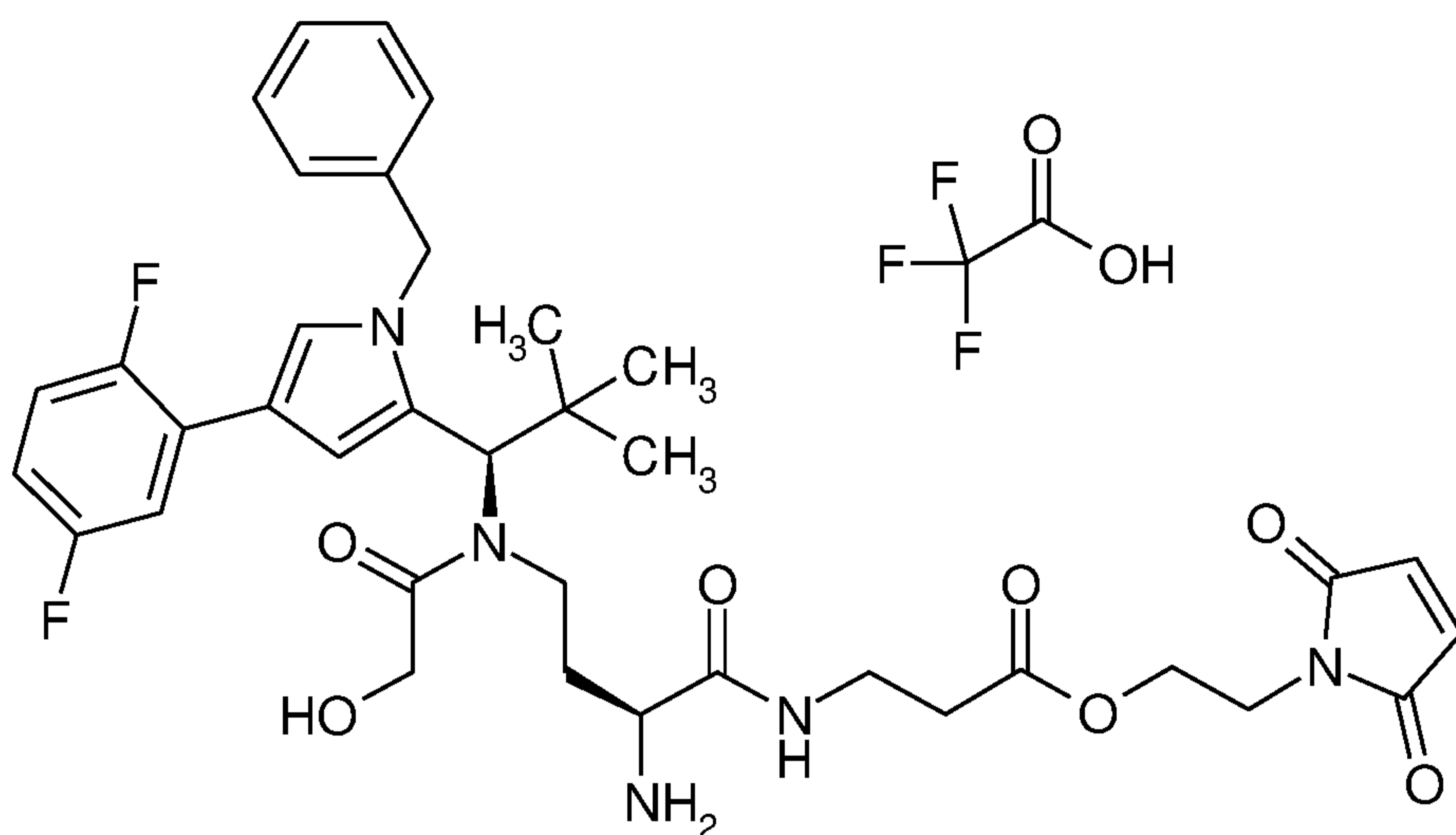
The title compound was prepared by coupling of 26 mg (0.035 mmol) of Intermediate C65 with 18 mg (0.07 mmol) of commercially available trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) in 8 ml of DMF in the presence of 40 mg (0.1054 mmol) of HATU and 61  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 16 mg (43% of theory over 2 steps) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.85 min; MS (ESIpos):  $m/z$  = 721 ( $M+H$ )<sup>+</sup>.

<sup>1</sup>H NMR (500 MHz, DMSO-*d*<sub>6</sub>):  $\delta$  = 7.99 (t, 1H), 7.95 (t, 1H), 7.6-7.75 (m, 4H), 7.5 (s, 1H), 7.2-7.4 (m, 6H), 6.8-7.0 (m, 4H), 5.63 (s, 1H), 4.9 and 5.2 (2d, 2H), 4.26 and 4.0 (2d, 2H), 3.3-3.6 (m, 4H), 3.15-3.25 (m, 3H), 2.85-3.0 (m, 2H), 2.2-2.3 (m, 4H), 0.64 and 1.49 (2m, 2H), 0.81 (s, 9H).

### Intermediate F196

Trifluoroacetic acid / 2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl-N-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alaninate (1:1)



First, 15 mg (0.023 mmol) of Intermediate C58 were taken up in 4 ml of DMF and reacted with 8.2 mg (0.025 mmol) of Intermediate L67 in the presence of 13.0 mg (0.034 mmol) of HATU and 16  $\mu$ l of *N,N*-diisopropylethylamine. After 30 min of stirring at RT, the mixture was concentrated and the residue was purified by preparative HPLC. After combination of the appropriate fractions and evaporation of the solvent, the residue was lyophilized from acetonitrile/water 1:1. This gave 4.3 mg (20% of theory) of the protected intermediate.

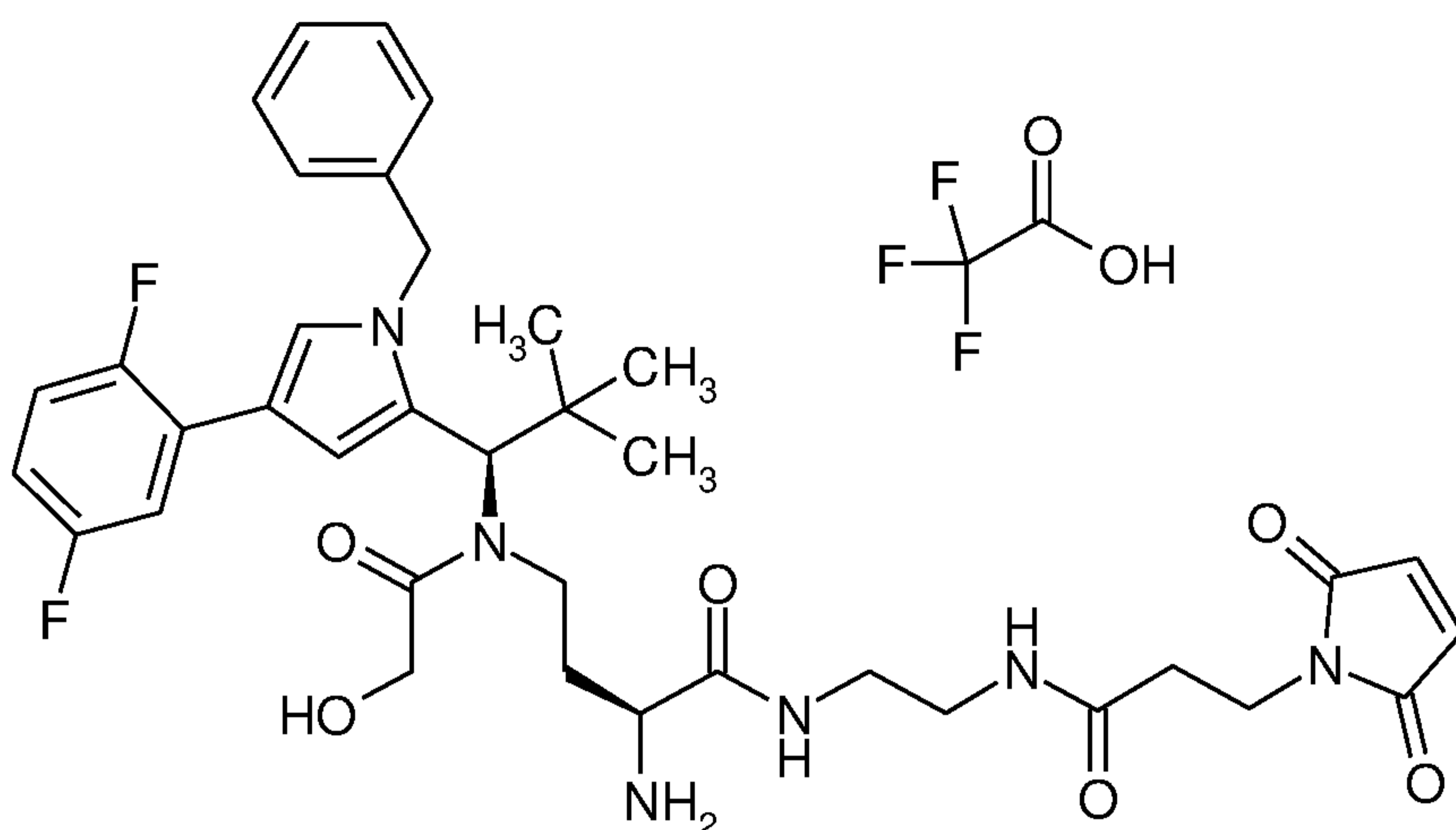
LC-MS (Method 1):  $R_t$  = 1.35 min; MS (EIpos):  $m/z$  = 852  $[M+H]^+$ .

4.3 mg (4.5  $\mu$ mol) of the intermediate were dissolved in 1 ml of trifluoroethanol and deprotected with 3.65 mg (27  $\mu$ mol) zinc chloride as described for Intermediate F119. Purification by preparative HPLC gave 1 mg (25% of theory over 2 steps) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.88 min; MS (ESIpos):  $m/z$  = 708  $(M+H)^+$

### Intermediate F204

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-(2-{[3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)propanoyl]amino}ethyl)butanamide (1:1)



25 mg (0.038 mmol) of Intermediate C58 were initially reacted  
 5 with 16.5 mg (75% pure) (0.038 mmol) of Intermediate L68 in the  
 presence of 17 mg (0.046 mmol) of HATU and 20  $\mu$ l of *N,N*-  
 diisopropylethylamine. After 60 min of stirring at RT, the  
 mixture was concentrated and the residue was purified by  
 preparative HPLC. This gave 18.3 mg (56% of theory) of the  
 10 protected intermediate.

RGM

LC-MS (Method 1):  $R_t$  = 1.32 min; MS (EIpos):  $m/z$  = 851  $[M+H]^+$ .  
 15

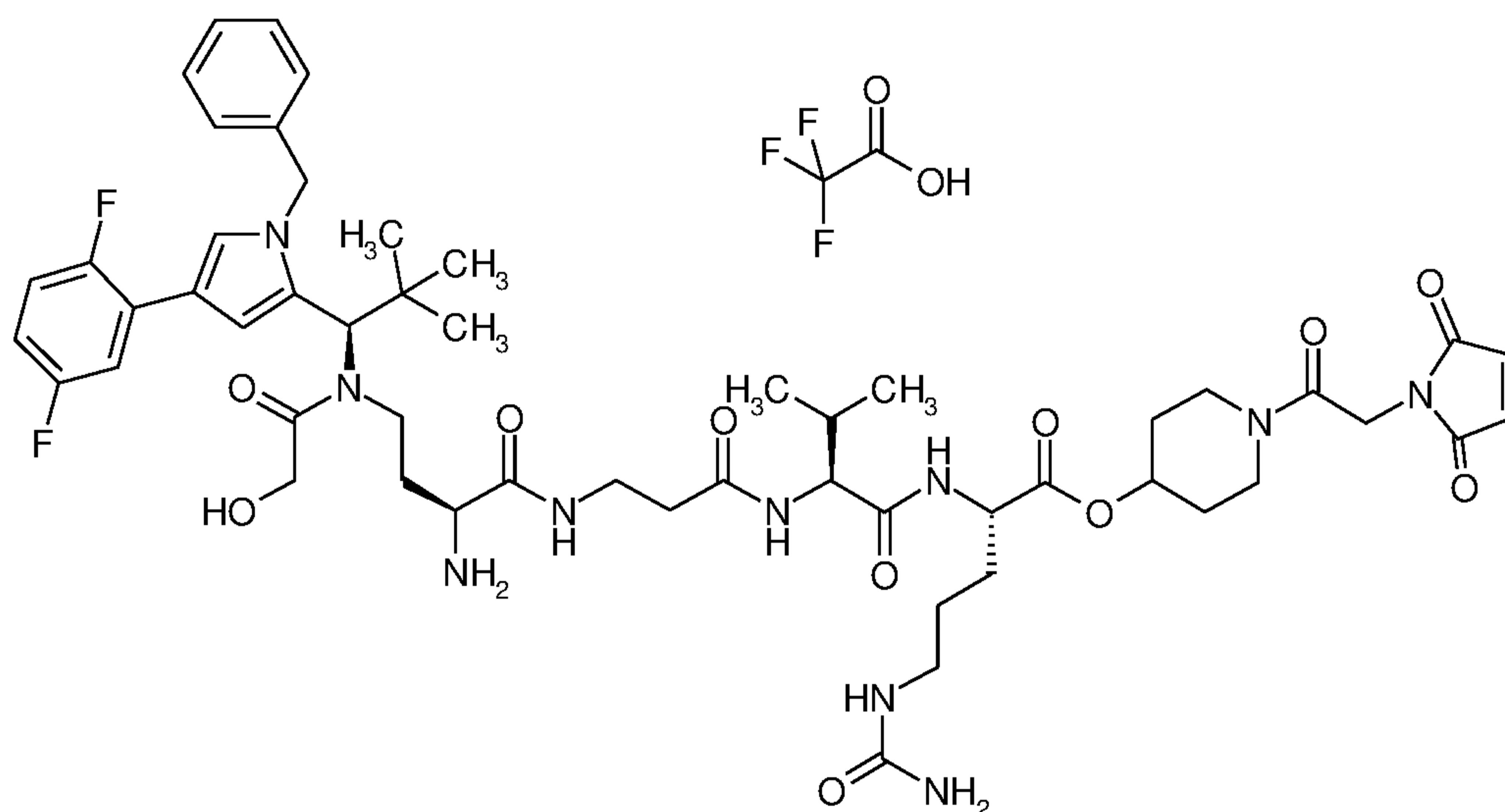
In the second step, this intermediate was dissolved in 3 ml of  
 2,2,2-trifluoroethanol. 12 mg (0.086 mmol) of zinc chloride were  
 added, and the reaction was stirred at 50°C for 2 h. 25 mg (0.086  
 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid and 2 ml of  
 20 a 0.1% strength aqueous trifluoroacetic acid solution were then  
 added. The reaction was purified by preparative HPLC.  
 Concentration of the appropriate fractions and lyophilization  
 of the residue from acetonitrile/water gave 11 mg (62% of theory)  
 of the title compound.

25 LC-MS (Method 1):  $R_t$  = 0.85 min; MS (ESIpos):  $m/z$  = 707  $(M+H)^+$ .

**Intermediate F205**



Trifluoroacetic acid / 1-[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]piperidin-4-yl N-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl]-beta-alanyl-L-valyl-L-alanine (1:1)

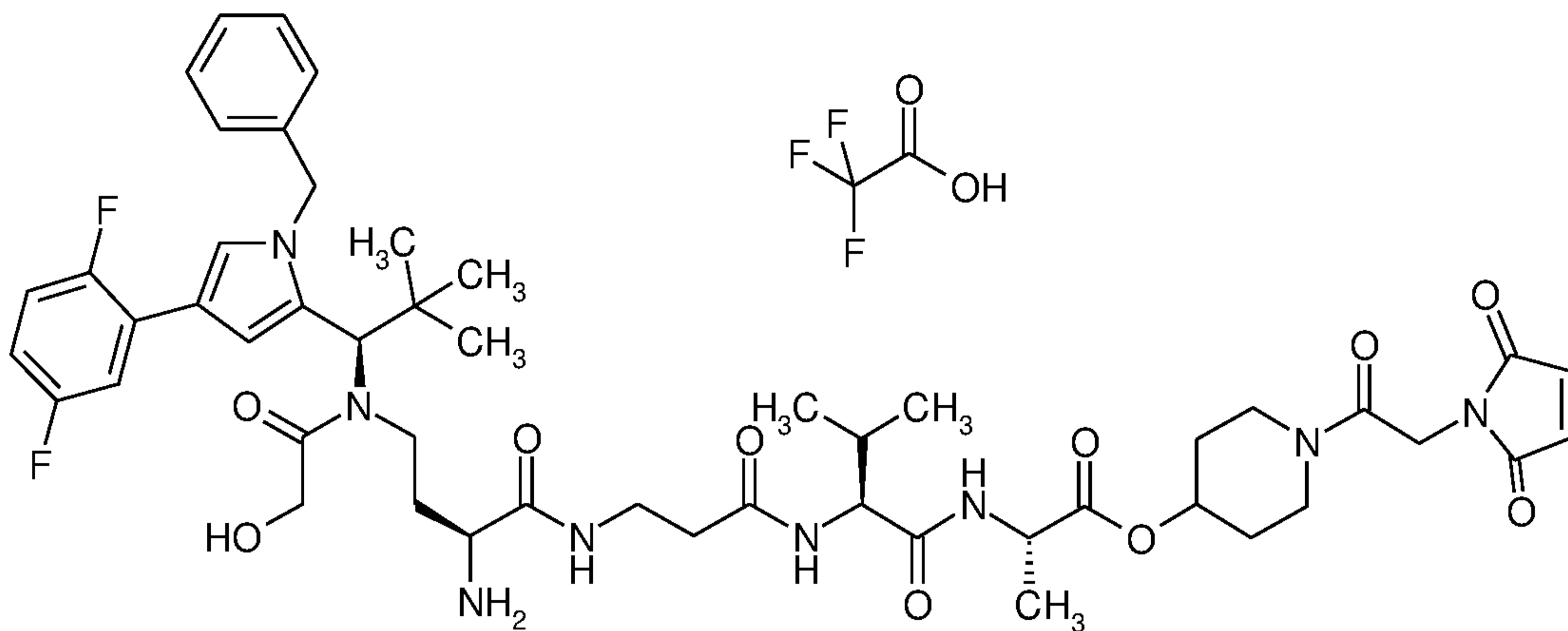


The synthesis was carried out by coupling of 25 mg (0.034 mmol) of Intermediate C61 and 29 mg (0.041 mmol) of Intermediate L69 in the presence of HATU and *N,N*-diisopropylethylamine, followed by hydrogenation with palladium on activated carbon (10%) under standard pressure, then coupling with (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetic acid in the presence of HATU and *N,N*-diisopropylethylamine and finally removal of the 2-(trimethylsilyl)ethoxycarbonyl protective group with zinc chloride. HPLC purification gave 11 mg (26% of theory over 4 steps).

LC-MS (Method 1):  $R_t = 0.86$  min; MS (ESIpos):  $m/z = 1061$  (M+H)<sup>+</sup>.

### Intermediate F206

Trifluoroacetic acid / 1-[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]piperidin-4-yl N-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl]-beta-alanyl-L-valyl-L-alanine (1:1)



The synthesis was carried out analogously to Intermediate F205.

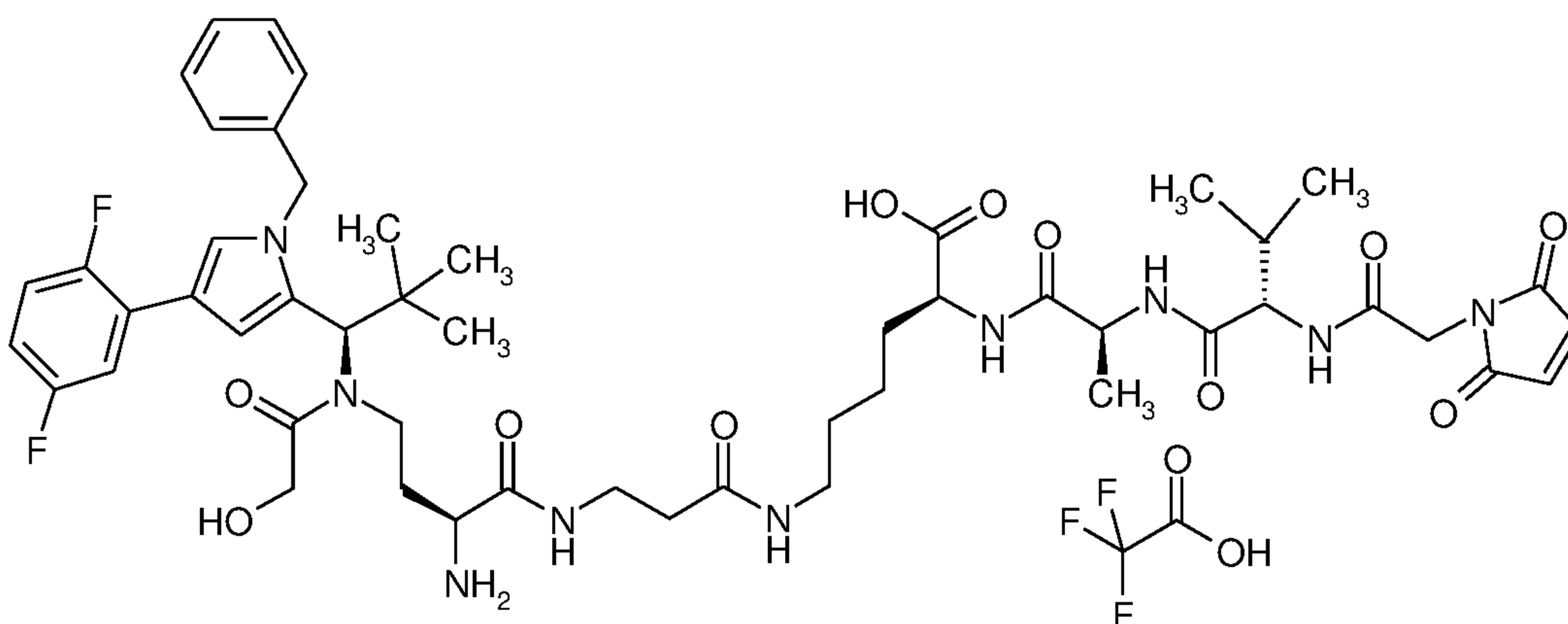
5

LC-MS (Method 1):  $R_t = 0.86$  min; MS (ESIpos):  $m/z = 975$  (M+H)<sup>+</sup>.

### Intermediate F207

10 N<sup>6</sup>- (N- { (2S) -2-Amino-4- [ { (1R) -1- [1-benzyl-4- (2,5-  
difluorophenyl) -1H-pyrrol-2-yl] -2,2-  
dimethylpropyl } (glycoloyl) amino] butanoyl } -beta-alanyl) -N<sup>2</sup>- { N-  
[ (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl) acetyl] -L-valyl-L-  
alanyl } -L-lysine / trifluoroacetic acid (1:1)

15



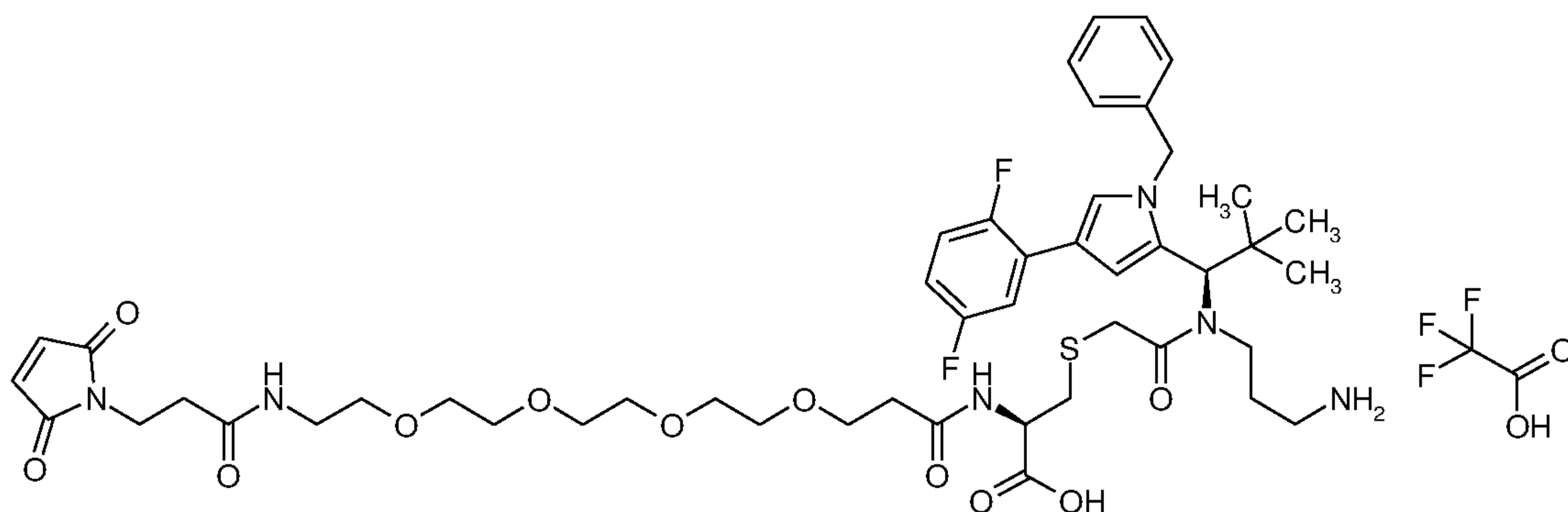
The title compound was prepared analogously to Intermediate F155.

20

LC-MS (Method 1):  $R_t = 0.81$  min; MS (ESIpos):  $m/z = 1020$  (M+H)<sup>+</sup>.

Intermediate F209

R-{2-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-cysteine / trifluoroacetic acid (1:1)



10

93.9 mg (0.78 mmol) of L-cysteine were suspended in a solution of 93.0 mg (1.11 mmol) of sodium bicarbonate and 0.9 ml of water. 70.0 mg (0.11 mmol) of 2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate (Intermediate C70), dissolved in 6.0 ml of isopropanol, and 202.3 mg (1.33 mmol) of 1,8-diazabicyclo[5.4.0]undec-7-ene were added. The reaction mixture was stirred at 50°C for 90 min. Water (0.1% TFA) was added, and the reaction was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water; 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 53.9 mg (59% of theory) of the compound R-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1).

15

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LC-MS (Method 1):  $R_t$  = 1.24 min; MS (ESIpos):  $m/z$  = 717 (M+H)<sup>+</sup>.

30

86.0 mg (0.1 mmol) of R-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-

cysteine / trifluoroacetic acid (1:1) and 58.5 mg (0.11 mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide were dissolved in 4.0 ml of DMF, and 20.9 mg (0.21 mmol) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight. 15.5 mg (0.26 mmol) of HOAc were added and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water; 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 68.6 mg (59% of theory) of the compound R-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-cysteine.

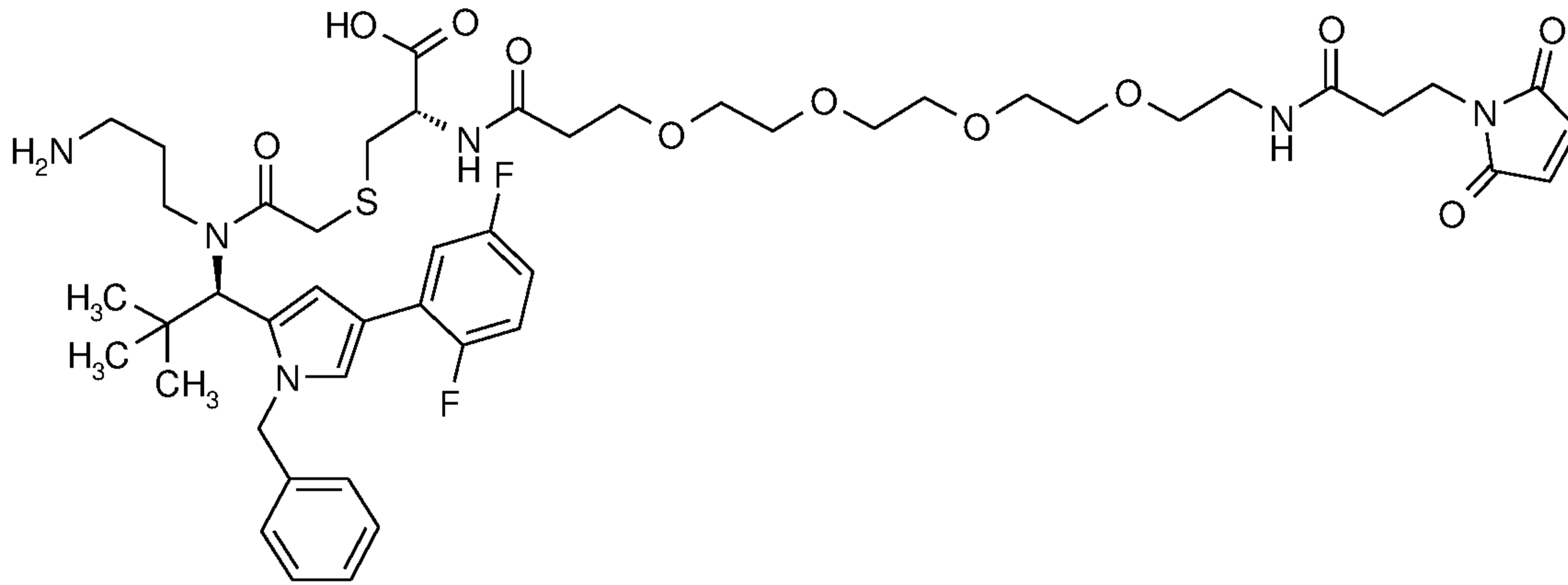
LC-MS (Method 6):  $R_t$  = 2.88 min; MS (ESIpos):  $m/z$  = 1115 (M+H)<sup>+</sup>.

46.4 mg (0.04 mmol) of R-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-cysteine were dissolved in 2.0 ml of trifluoroethanol, and 17.0 mg (0.13 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C overnight. Another 8.5 mg (0.07 mmol) of zinc dichloride were added, and the mixture was stirred at 50°C overnight. 36.5 mg (0.13 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 19.4 mg (43% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.94 min; MS (ESIpos):  $m/z$  = 971 (M+H)<sup>+</sup>.

**Intermediate F210**

S-{2-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-D-cysteine / trifluoroacetic acid (1:1)



10

The title compound was prepared analogously to the synthesis of Intermediate F209 using D-cysteine.

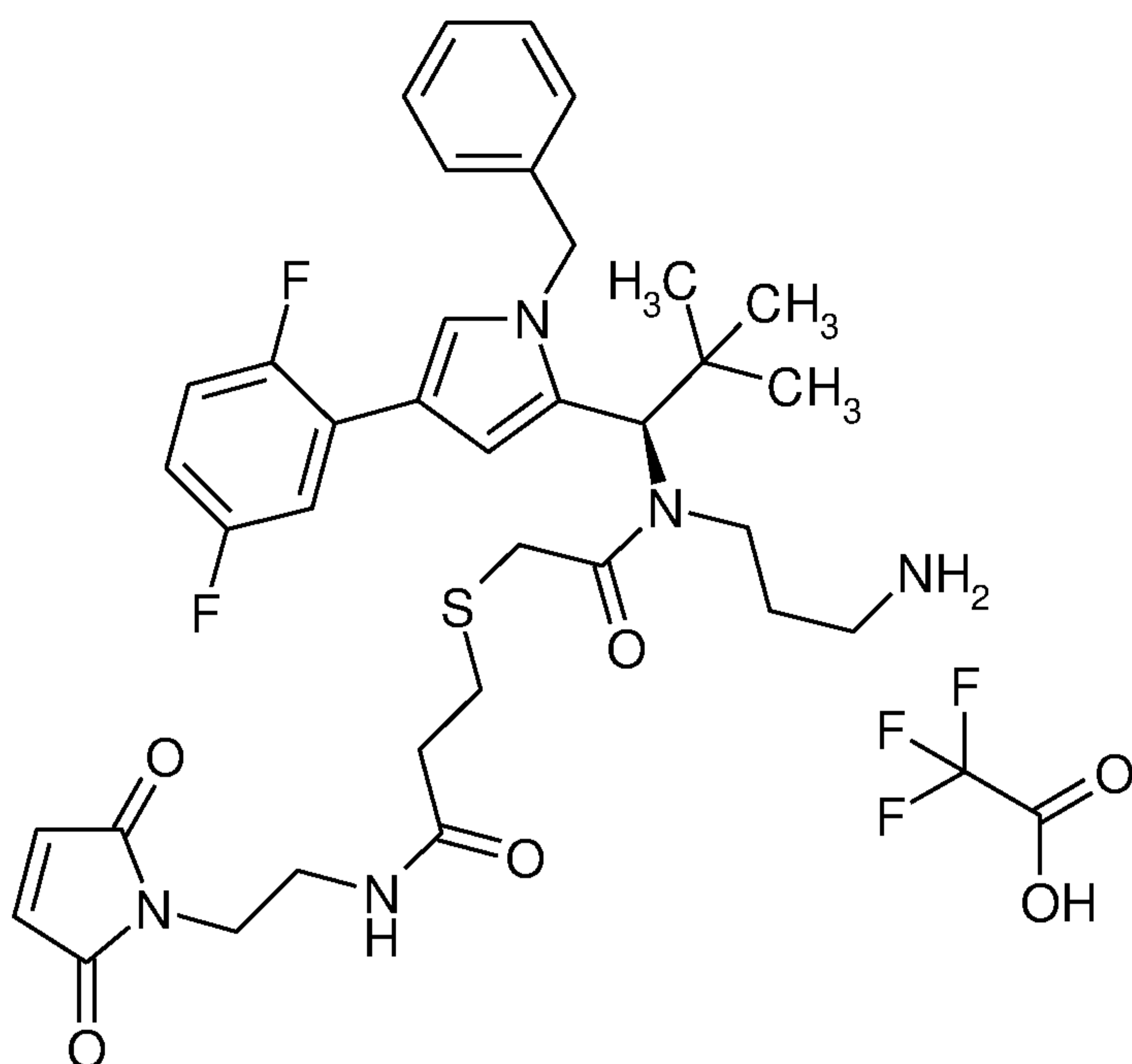
LC-MS (Method 1):  $R_t = 0.91$  min; MS (ESIpos):  $m/z = 971$  (M+H)<sup>+</sup>.

15

**Intermediate F211**

Trifluoroacetic acid / 3-({2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulphanyl)-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]propanamide (1:1)

20



30.0 mg (0.05 mmol) of 2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl} carbamate (Intermediate C70) were initially charged together with 5.5 mg (0.05 mmol) of 3-sulphonylpropanoic acid in 0.5 ml of methanol with a drop of water. 23.0 mg (0.17 mmol) of potassium carbonate were then added, and the reaction mixture was stirred at 50°C for 4 h. Ethyl acetate was added and the organic phase was washed once with water and once with saturated NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was used without further purification in the next step of the synthesis. This gave 30.3 mg (86% of theory) of the compound 11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid.

LC-MS (Method 1):  $R_t = 1.39$  min; MS (ESIpos):  $m/z = 702$  (M+H)<sup>+</sup>.

30.0 mg (0.04 mol) of 11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid and 9.8 mg (0.06 mmol) of 1-(2-aminoethyl)-1H-

pyrrole-2,5-dione hydrochloride (1:1) were initially charged in 2.0 ml of acetonitrile, and 44.2 mg (0.34 mmol) of *N,N*-diisopropylethylamine were added. 35.4 mg (0.06 mmol) T3P (50 % in ethyl acetate) were added, and the reaction mixture was stirred at RT overnight. Water was added, and purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 22.0 mg (63% of theory) of the compound 2-(trimethylsilyl)ethyl [3-((1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-dimethylpropyl){[(3-{[2-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)ethyl]amino}-3-oxopropyl)sulphanyl]acetyl}amino)propyl]carbamate.

LC-MS (Method 1):  $R_t$  = 1.41 min; MS (ESIpos):  $m/z$  = 824 (M+H)<sup>+</sup>.

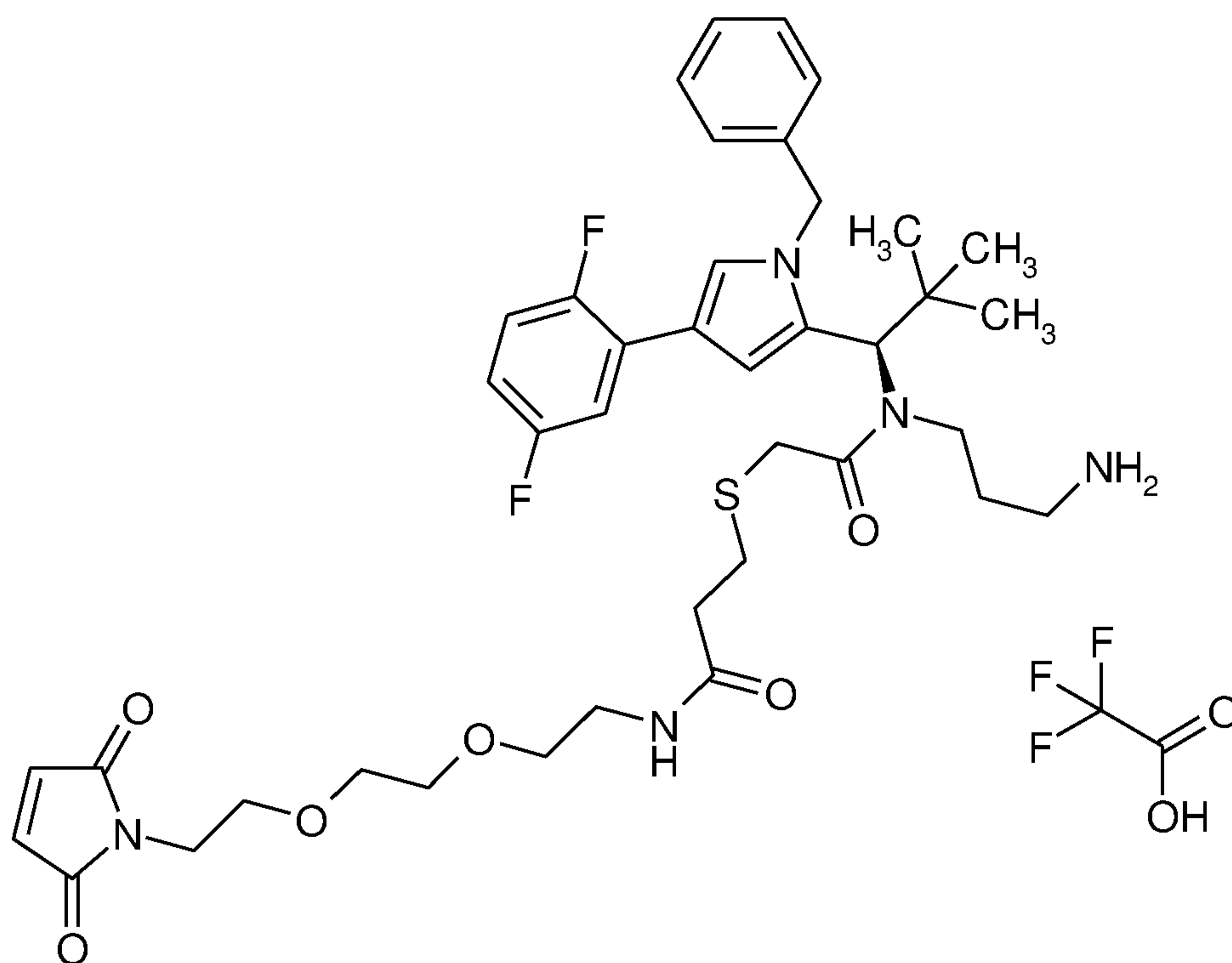
22.0 mg (0.03 mol) of 2-(trimethylsilyl)ethyl [3-((1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-dimethylpropyl){[(3-{[2-(2,5-dioxo-2,5-dihydro-1*H*-pyrrol-1-yl)ethyl]amino}-3-oxopropyl)sulphanyl]acetyl}amino)propyl]carbamate were dissolved in 1.0 ml of trifluoroethanol, and 9.1 mg (0.07 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C for 5 h. 19.5 mg (0.07 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 15.0 mg (71% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.88 min; MS (ESIpos):  $m/z$  = 680 (M+H)<sup>+</sup>.

### 35 Intermediate F212

Trifluoroacetic acid / *N*-(3-aminopropyl)-*N*-{(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-dimethylpropyl}-1-

(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-10-oxo-3,6-dioxa-13-thia-9-azapentadecan-15-amide (1:1)



5

28.8 mg (0.04 mmol) of 11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid (Intermediate C69) were initially charged together with 18.3 mg (0.05 mmol) of trifluoroacetic acid / 1-((2-((2-aminoethoxy)ethoxy)ethyl)-1H-pyrrole-2,5-dione (1:1) (Intermediate L59) in 1.9 ml of acetonitrile. 42.4 mg (0.33 mmol) of *N,N*-diisopropylethylamine were then added, and 33.9 mg (0.05 mmol) of T3P (50% in ethyl acetate) were added dropwise. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 10.7 mg (26% of theory) of the compound 2-(trimethylsilyl)ethyl [16-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-1-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-10,15-dioxo-3,6-dioxa-13-thia-9,16-diazanonadecan-19-yl]carbamate.

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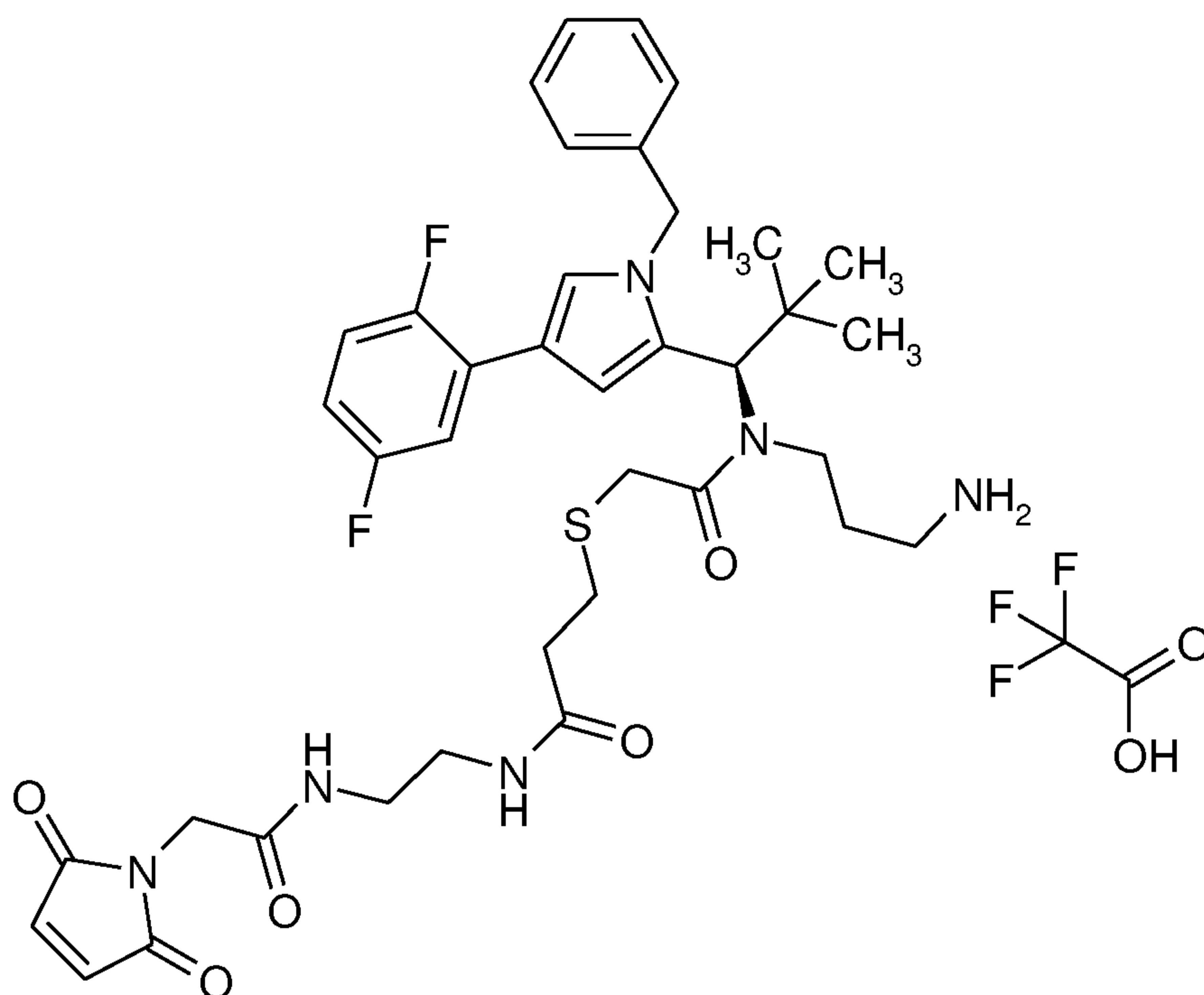
LC-MS (Method 1):  $R_t = 1.44$  min; MS (ESIpos):  $m/z = 812$  (M+H)<sup>+</sup>.

10.7 mg (0.01 mol) of 2-(trimethylsilyl)ethyl [16-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-1-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-10,15-dioxo-3,6-dioxo-13-thia-9,16-diazanonadecan-19-yl]carbamate were dissolved in 0.8 ml of trifluoroethanol, and 8.0 mg (0.06 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C for 5 h. 17.1 mg (0.06 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum.

LC-MS (Method 1):  $R_t = 1.03$  min; MS (ESIpos):  $m/z = 768$  (M+H)<sup>+</sup>.

### Intermediate F213

20 Trifluoroacetic acid / 3-({2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulphonyl)-N-(2-{[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino}ethyl)propanamide  
25 (1:1)



27.5 mg (0.04 mmol) of 11-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid (Intermediate C69) were initially charged together with 15.9 mg (0.05 mmol) of trifluoroacetic acid / N-(2-aminoethyl)-2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetamide (1:1) (Intermediate L1) in 1.8 ml of acetonitrile. 32.4 mg (0.31 mmol) of *N,N*-diisopropylethylamine were then added, and 32.4 mg (0.05 mmol) of T3P (50% in ethyl acetate) were added dropwise. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 11.9 mg (35% of theory) of the compound 2-(trimethylsilyl)ethyl [13-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]-1-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-2,7,12-trioxo-10-thia-3,6,13-triazahexadecan-16-yl]carbamate.

LC-MS (Method 1):  $R_t$  = 1.39 min; MS (ESIpos):  $m/z$  = 881 (M+H)<sup>+</sup>.

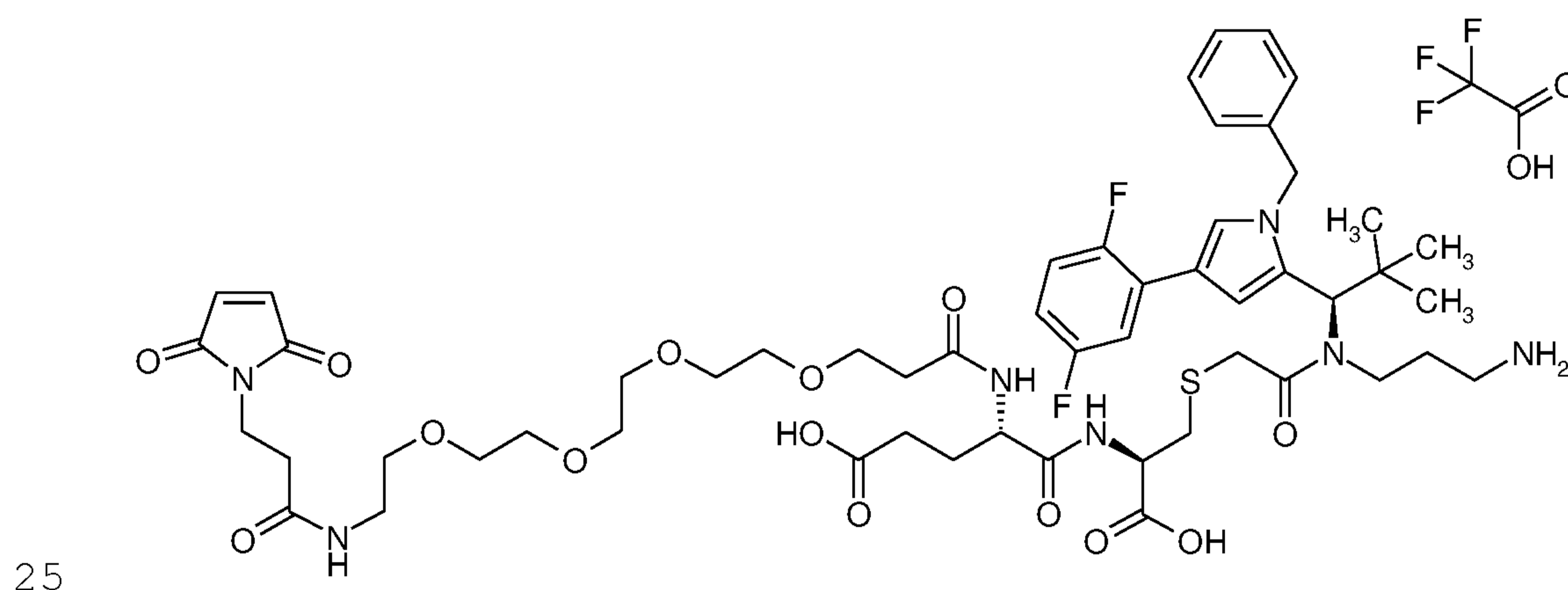
11.9 mg (0.01 mol) of 2-(trimethylsilyl)ethyl-[13-[(1R)-1-[1-

benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-1-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-2,7,12-trioxo-10-thia-3,6,13-triazahexadecan-16-yl]carbamate were dissolved in 1.0 ml of trifluoroethanol, and 5.5 mg (0.04 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C overnight. 11.8 mg (0.04 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 7.4 mg (60% of theory) of the title compound.

LC-MS (Method 5):  $R_t$  = 2.75 min; MS (ESIpos):  $m/z$  = 737 (M+H)<sup>+</sup>.

#### Intermediate F214

N-[19-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-alpha-glutamyl-S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-L-cysteine / trifluoroacetic acid (1:1)



111.7 mg (0.30 mmol) of (2S)-5-(benzyloxy)-2-[(benzyloxy)carbonyl]amino}-5-oxopentanoic acid were initially charged in 3.0 ml of DMF, and 46.1 (0.30 mmol) of HOBt, 96.6 mg (0.30 mmol) of TBTU and 38.9 mg (0.30 mmol) of N,N-diisopropylethylamine were added. The reaction mixture was

30

stirred at RT for 10 min. 250.0 mg (0.30 mmol) of S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1) (Intermediate C71) dissolved in 116.3 mg (0.9 mmol) of *N,N*-diisopropylethylamine and 3.0 ml of DMF were then added. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 257.0 mg (80% of theory) of the compound (16R)-11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-16-(((2S)-5-(benzyloxy)-2-(((benzyloxy)carbonyl)amino)-5-oxopentanoyl)amino)-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid.

LC-MS (Method 1):  $R_t$  = 1.55 min; MS (ESIpos):  $m/z$  = 1071 (M+H)<sup>+</sup>.

Under argon, 24.6 mg (0.11 mmol) of palladium(II) acetate were initially charged in 5.0 ml of dichloromethane, and 33.2 mg (0.33 mmol) of triethylamine and 254.3 mg (2.19 mmol) of triethylsilane were added. The reaction mixture was stirred at RT for 5 min, and 234.1 mg (0.22 mmol) of (16R)-11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-16-(((2S)-5-(benzyloxy)-2-(((benzyloxy)carbonyl)amino)-5-oxopentanoyl)amino)-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid dissolved in 5.0 ml of dichloromethane were added. The reaction mixture was stirred at RT overnight. The reaction mixture was filtered through a cardboard filter and the filter cake was washed with dichloromethane. The solvent was evaporated under reduced pressure without heating. The residue was purified by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 177.5 mg (85% of theory) of the compound L-alpha-glutamyl-S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-

1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1).

5 LC-MS (Method 1):  $R_t = 1.07$  min; MS (ESIpos):  $m/z = 846$  (M+H)<sup>+</sup>.

20.0 mg (20.83  $\mu$ mol) L-alpha-glutamyl-S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1) were initially charged together with 11.8 mg (22.91  $\mu$ mol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide in 1.5 ml of DMF, and 6.3 mg (62.49  $\mu$ mol) of 4-methylmorpholine were added. 15 The reaction mixture was stirred at RT overnight, and 4.4 mg (0.07 mmol) of acetic acid were then added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried 20 under high vacuum. This gave 19.1 mg (74% of theory) of the compound N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-alpha-glutamyl-S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine. 25

LC-MS (Method 1):  $R_t = 1.24$  min; MS (ESIpos):  $m/z = 1244$  (M+H)<sup>+</sup>.

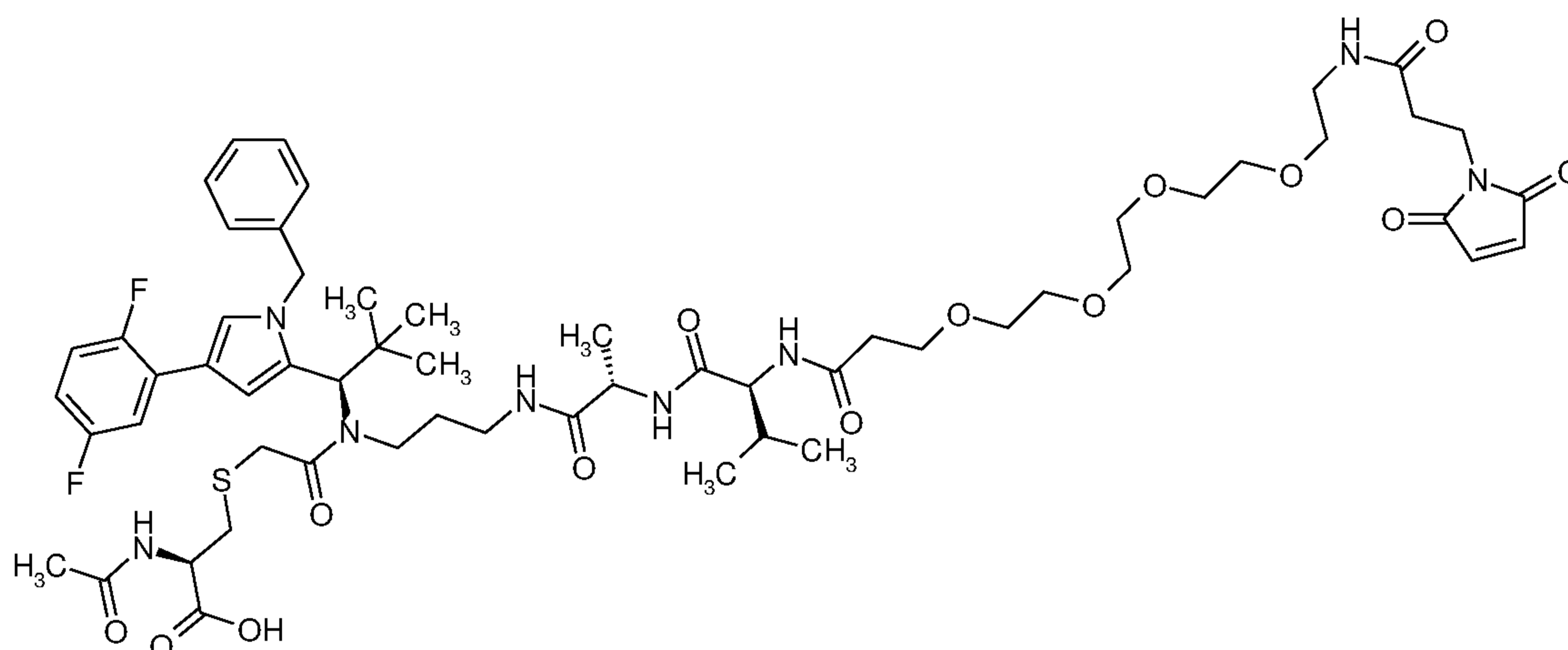
17.5 mg (14.06  $\mu$ mol) of N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-alpha-glutamyl-S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine were dissolved in 1.5 ml of trifluoroethanol, and 11.5 mg (84.37  $\mu$ mol) of zinc 35 dichloride were added. The reaction mixture was stirred at 50°C for 4 h. 24.7 mg (0.08 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was

carried out directly by preparative RP-HPLC (column: Reprisil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 10.8 mg (63% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.89 min; MS (ESIpos):  $m/z$  = 1100 (M+H)<sup>+</sup>.

### Intermediate F215

N-[19-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-valyl-N-{3-[(2R)-2-acetamido-2-carboxyethyl]sulphonyl}acetyl}{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]propyl}-L-alaninamide



14.9 mg (0.02 mmol) of N-acetyl-S-[2-([3-(L-alanyl-amino)propyl]{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)-2-oxoethyl]-L-cysteine / trifluoroacetic acid (1:1) (Example 229) and 7.1 mg (0.02 mmol) of 2,5-dioxopyrrolidin-1-yl-N-[(benzyloxy)carbonyl]-L-valinate were initially charged in 1.0 ml of DMF, and 5.7 mg (0.06 mmol) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight, and 4.5 mg (0.08 mmol) of acetic acid were then added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprisil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue

was dried under high vacuum. This gave 13.3 mg (78% of theory) of the compound N-[(benzyloxy)carbonyl]-L-valyl-N-{3-[[[(2R)-2-acetamido-2-carboxyethyl]sulphonyl]acetyl]}{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]propyl}-L-alaninamide.

LC-MS (Method 1):  $R_t = 1.24$  min; MS (ESIpos):  $m/z = 919$  (M+H)<sup>+</sup>.

11.1 mg (0.01 mmol) of N-[(benzyloxy)carbonyl]-L-valyl-N-{3-[[[(2R)-2-acetamido-2-carboxyethyl]sulphonyl]acetyl]}{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]propyl}-L-alaninamide were dissolved in 5.0 ml of ethanol, 1.0 mg of palladium on activated carbon (10%) was added and the mixture was hydrogenated at RT and standard pressure overnight. The reaction mixture was filtered through Celite and the filter cake was washed with an ethanol/THF/water mixture. The solvents were evaporated under reduced pressure. The residue was purified by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was lyophilized. This gave 7.5 mg (69% of theory) of the compound L-valyl-N-{3-[[[(2R)-2-acetamido-2-carboxyethyl]sulphonyl]acetyl]}{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]propyl}-L-alaninamide / trifluoroacetic acid (1:1).

LC-MS (Method 1):  $R_t = 0.86$  min; MS (ESIpos):  $m/z = 785$  (M+H)<sup>+</sup>.

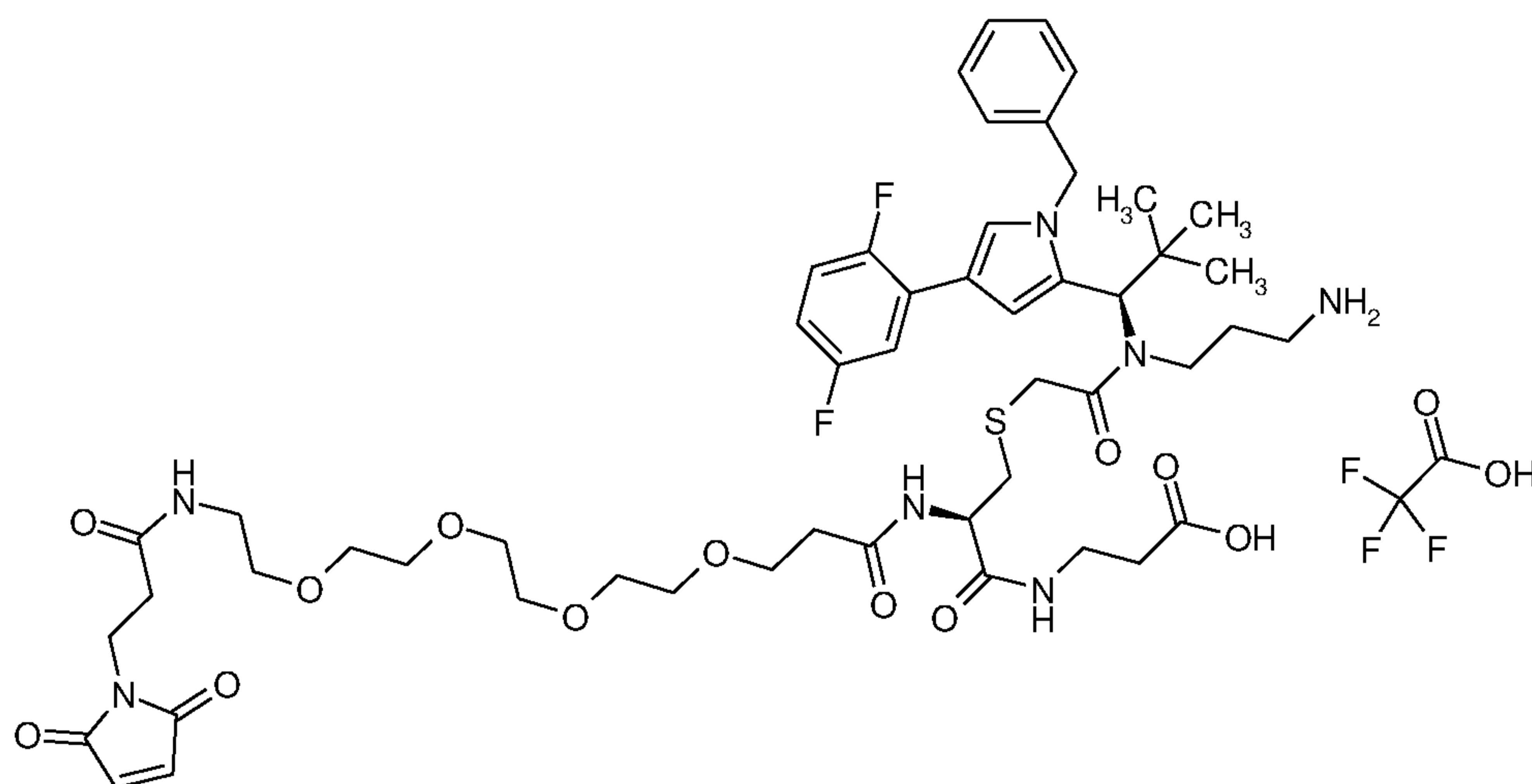
7.3 mg (8.12  $\mu$ mol) of L-valyl-N-{3-[[[(2R)-2-acetamido-2-carboxyethyl]sulphonyl]acetyl]}{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]propyl}-L-alaninamide / trifluoroacetic acid (1:1) were initially charged together with 4.6 mg (8.93  $\mu$ mol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide in 0.5 ml of DMF, and 2.5 mg (24.36  $\mu$ mol) of 4-methylmorpholine were added. The reaction mixture was stirred

at RT overnight, and 4.4 mg (0.03 mmol) of acetic acid were then added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under  
 5 reduced pressure and the residue was dried under high vacuum. This gave 4.9 mg (50% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 1.07 min; MS (ESIpos):  $m/z$  = 1183 (M+H)<sup>+</sup>.

### 10 Intermediate F216

S-{2-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-N-[19-  
 (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-  
 15 tetraoxa-16-azanonadecan-1-oyl]-L-cysteinyl-beta-alanine /  
 trifluoroacetic acid (1:1)



20 Under argon, 30.2 mg (0.06 mmol) of N,N'-  
 bis[(benzyloxy)carbonyl]-L-cystine were initially charged in  
 2.0 ml of water and 2.0 ml of isopropanol, and 56.7 mg (0.20  
 mmol) of TCEP were added. The reaction mixture was stirred at  
 RT for 30 min. 50.0 mg (0.08 mmol) of 2-(trimethylsilyl)ethyl  
 25 {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-  
 2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate  
 (Intermediate C70), dissolved in 2.0 ml of isopropanol, and  
 122.2 mg (0.48 mmol) of 1,8-diazabicyclo[5.4.0]undec-7-ene were  
 then added, and the reaction mixture was stirred at 50°C for 7



h. Another 122.2 mg (0.48 mmol) of 1,8-diazabicyclo[5.4.0]undec-7-ene were then added, and the reaction mixture was stirred at 50°C for 1 h. The mixture was diluted with ethyl acetate and the organic phase was extracted with water and saturated sodium bicarbonate solution and washed with saturated NaCl solution. The organic phase was dried over magnesium sulphate and the solvent was evaporated under reduced pressure. The residue was purified by preparative RP-HPLC (column: Reprosil 250x30; 10µ, flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 43.1 mg (64% of theory) of the compound S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[(benzyloxy)carbonyl]-L-cysteine.

LC-MS (Method 1):  $R_t = 1.46$  min; MS (ESIpos):  $m/z = 851$  (M+H)<sup>+</sup>.

16.5 mg (0.05 mmol) of 4-methylbenzenesulphonic acid / benzyl beta-alaninate (1:1) were initially charged together with 14.0 mg (0.11 mmol) of *N,N*-diisopropylethylamine in 1.5 ml of acetonitrile. The reaction mixture was stirred at RT for 3 min, and 30.8 mg (0.04 mmol) of S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[(benzyloxy)carbonyl]-L-cysteine dissolved in 1.5 ml of acetonitrile, 23.4 mg (0.18 mmol) of *N,N*-diisopropylethylamine and 29.9 mg (0.05 mmol) of T3P (50% in ethyl acetate) were then added. The reaction mixture was stirred at RT overnight. Water was added, and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10µ, flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. The compound obtained was benzyl S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[(benzyloxy)carbonyl]-L-cysteinyll-beta-alaninate.

LC-MS (Method 1):  $R_t = 1.59$  min; MS (ESIpos):  $m/z = 1012$  (M+H)<sup>+</sup>.

43.8 mg (43.3  $\mu$ mol) of benzyl S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[(benzyloxy)carbonyl]-L-cysteinyl-beta-alaninate were dissolved in 8.0 ml of ethanol, 4.4 mg of palladium on activated carbon (10%) were added and the mixture was hydrogenated at RT and standard pressure overnight. The reaction mixture was filtered through a cardboard filter and the filter cake was washed with ethanol. The solvent was evaporated under reduced pressure. Two more times, the residue was treated as just described. The residue was purified by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 4.9 mg (50% of theory) of the compound S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteinyl-beta-alanine / trifluoroacetic acid (1:1).

LC-MS (Method 1):  $R_t = 1.08$  min; MS (ESIpos):  $m/z = 788$  (M+H)<sup>+</sup>.

14.5 mg (16.1  $\mu$ mol) of S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteinyl-beta-alanine / trifluoroacetic acid (1:1) were initially charged together with 9.1 mg (17.7  $\mu$ mol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide in 1.0 ml of DMF, and 4.9 mg (48.2  $\mu$ mol) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight, and 3.4 mg (0.06 mmol) of acetic acid were then added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 4.9 mg (50%

of theory) of the compound S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteinyl-beta-alanine / trifluoroacetic acid (1:1).

5

LC-MS (Method 1):  $R_t = 1.28$  min; MS (ESIpos):  $m/z = 1186$  (M+H)<sup>+</sup>.

14.1 mg (11.9  $\mu$ mol) of S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-cysteinyl-beta-alanine / trifluoroacetic acid (1:1) were dissolved in 1.5 ml of trifluoroethanol, and 9.7 mg (71.3  $\mu$ mol) of zinc dichloride were added. The reaction mixture was stirred at 50°C for 3 h. Another 9.7 mg (71.3  $\mu$ mol) of zinc dichloride were added, and the reaction mixture was stirred at 50°C for 3 h. Another 9.7 mg (71.3  $\mu$ mol) of zinc dichloride were added, and the reaction mixture was stirred at 70°C for 4 h. 20.8 mg (0.07 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was lyophilized. This gave 6.2 mg (44% of theory) of the title compound.

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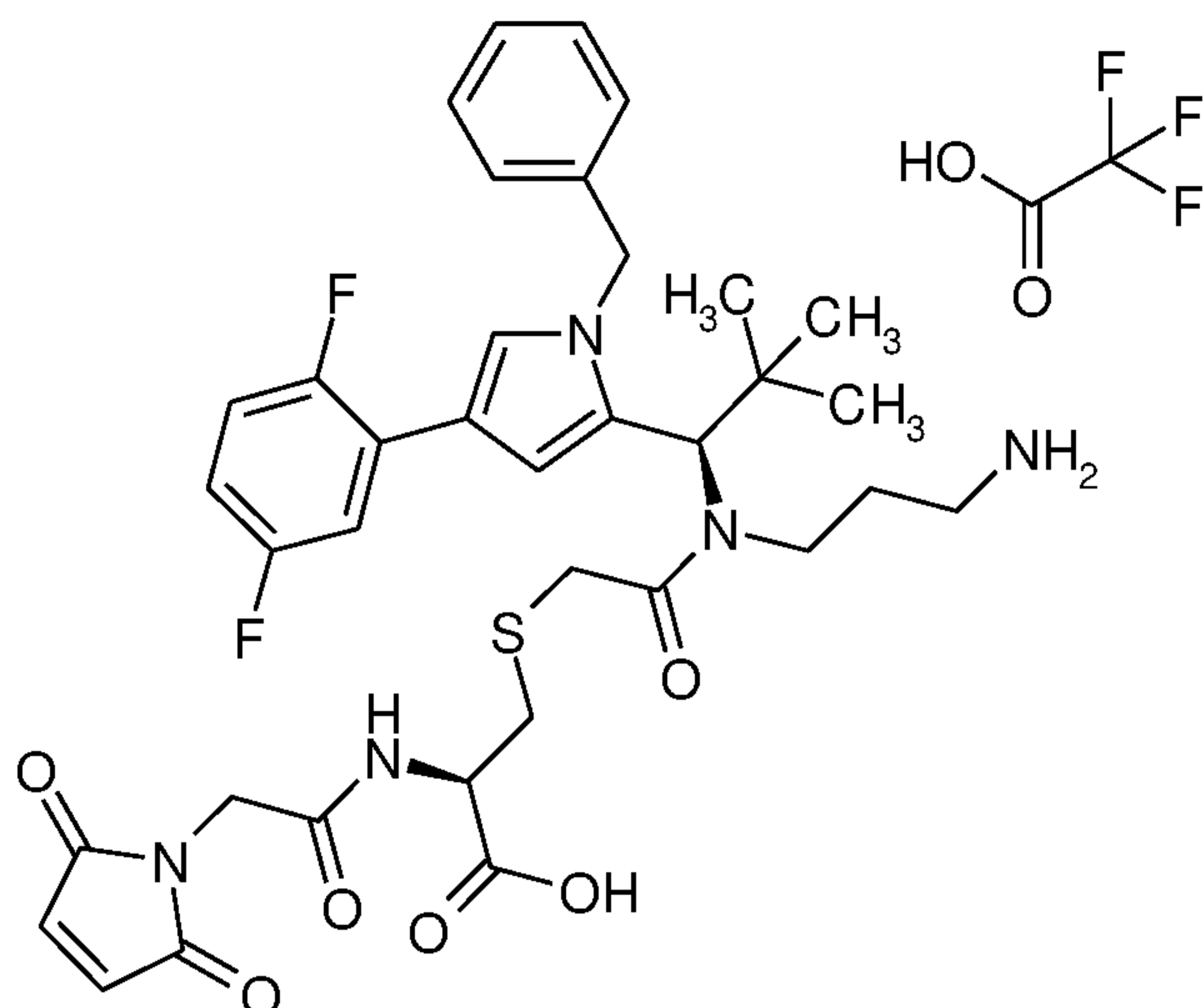
25

LC-MS (Method 1):  $R_t = 0.82$  min; MS (ESIpos):  $m/z = 1042$  (M+H)<sup>+</sup>.

### 30 Intermediate F217

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S-{2-[(3-Aminopropyl)((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino]-2-oxoethyl}-N-[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]-L-cysteine / trifluoroacetic acid (1:1)



Under argon, 7.5 mg (0.05 mmol) of (2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetic acid were initially charged in 1.5 ml of DMF,  
 5 and 7.5 mg (0.05 mmol) of HOBT, 15.5 mg (0.05 mmol) of TBTU and 6.2 mg (0.05 mmol) of *N,N*-diisopropylethylamine were added. The reaction mixture was stirred at RT for 10 min. 40.0 mg (0.05 mmol) of *S*-(11-((1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-*L*-cysteine  
 10 trifluoroacetic acid (1:1) (Intermediate C71), dissolved in 1.5 ml of DMF, and 18.7 mg (0.14 mmol) of *N,N*-diisopropylethylamine were then added, and the reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 11.2 mg (25% of theory) of the compound *S*-(11-((1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-*N*-[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]-*L*-cysteine.  
 20

LC-MS (Method 1):  $R_t$  = 1.37 min; MS (ESIpos):  $m/z$  = 854 ( $M+H$ )<sup>+</sup>.  
 25

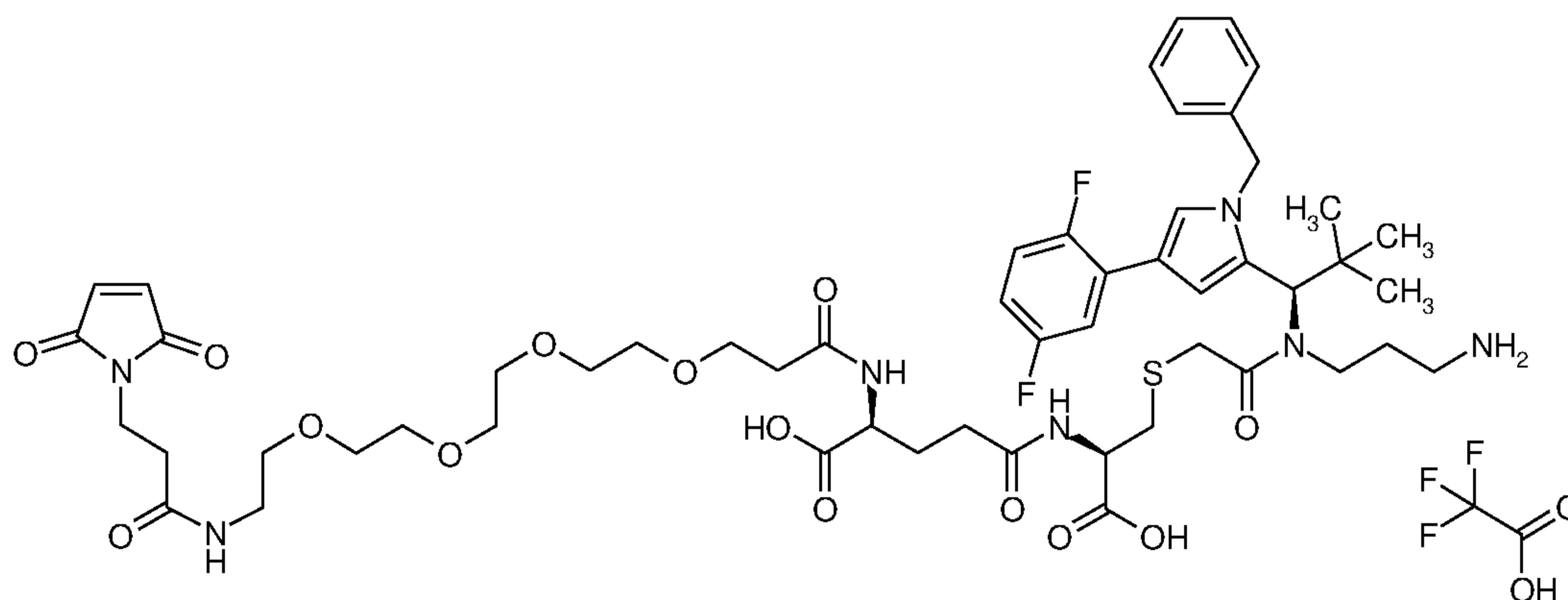
10.9 mg (12.8  $\mu$ mol) of *S*-(11-((1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-*N*-

[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]-L-cysteine were dissolved in 2.0 ml of trifluoroethanol, and 10.4 mg (76.6  $\mu$ mol) zinc dichloride were added. The reaction mixture was stirred at 50°C for 4 h. 22.4 mg (0.08 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprisil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was lyophilized. This gave 7.5 mg (65% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.92 min; MS (ESIpos):  $m/z$  = 710 (M+H)<sup>+</sup>.

### 15 Intermediate F218

*N*-[19-(2,5-Dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L- $\gamma$ -glutamyl-S-{2-[(3-aminopropyl){(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-L-cysteine / trifluoroacetic acid (1:1)



25 Under argon, 22.9 mg (0.06 mmol) of (4*S*)-5-(benzyloxy)-4-[[ (benzyloxy)carbonyl]amino]-5-oxopentanoic acid were initially charged in 2.0 ml of DMF, and 9.4 mg (0.05 mmol) of HOBT, 19.8 mg (0.06 mmol) of TBTU and 8.0 mg (0.06 mmol) of *N,N*-diisopropylethylamine were added. The reaction mixture was stirred at RT for 10 min. 51.2 mg (0.06 mmol) of *S*-(11-[(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-

dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine (Intermediate C71), dissolved in 1.0 ml of DMF, and 23.9 mg (0.19 mmol) of *N,N*-diisopropylethylamine were then added, and the reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 11.2 mg (25% of theory) of the compound (16R)-11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-16-{[(4S)-5-(benzyloxy)-4-[(benzyloxy)carbonyl]amino]-5-oxopentanoyl]amino}-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid.

LC-MS (Method 1):  $R_t$  = 1.52 min; MS (ESIpos):  $m/z$  = 1070 (M+H)<sup>+</sup>.

Under argon, 3.9 mg (0.02 mmol) of palladium(II) acetate were initially charged in 1.0 ml of dichloromethane, and 5.3 mg (0.05 mmol) of triethylamine and 254.3 mg (2.19 mmol) of triethylsilane were added. The reaction mixture was stirred at RT for 5 min, and 18.6 mg (0.02 mmol) of (16R)-11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-16-{[(4S)-5-(benzyloxy)-4-[(benzyloxy)carbonyl]amino]-5-oxopentanoyl]amino}-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid dissolved in 1.0 ml of dichloromethane were added. The solvent was evaporated under reduced pressure without heating. The residue was taken up in acetonitrile, filtered through a syringe filter and purified by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 11.0 mg (66% of theory) of the compound L-gamma-glutamyl-S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1).

LC-MS (Method 1):  $R_t = 1.14$  min; MS (ESIpos):  $m/z = 846$  (M+H)<sup>+</sup>.

15.0 mg (15.6  $\mu\text{mol}$ ) of L-gamma-glutamyl-S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine / trifluoroacetic acid (1:1) were initially charged together with 8.8 mg (17.2  $\mu\text{mol}$ ) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-yl}propanamide in 1.0 ml of DMF, and 4.7 mg (46.9  $\mu\text{mol}$ ) of 4-methylmorpholine were added. The reaction mixture was stirred at RT overnight, and 3.3 mg (0.06 mmol) of

acetic acid were then added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 14.2 mg (70% of theory) of the compound N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-gamma-glutamyl-S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine.

LC-MS (Method 4)  $R_t = 1.24$  min; MS (ESIpos):  $m/z = 1244$  (M+H)<sup>+</sup>.

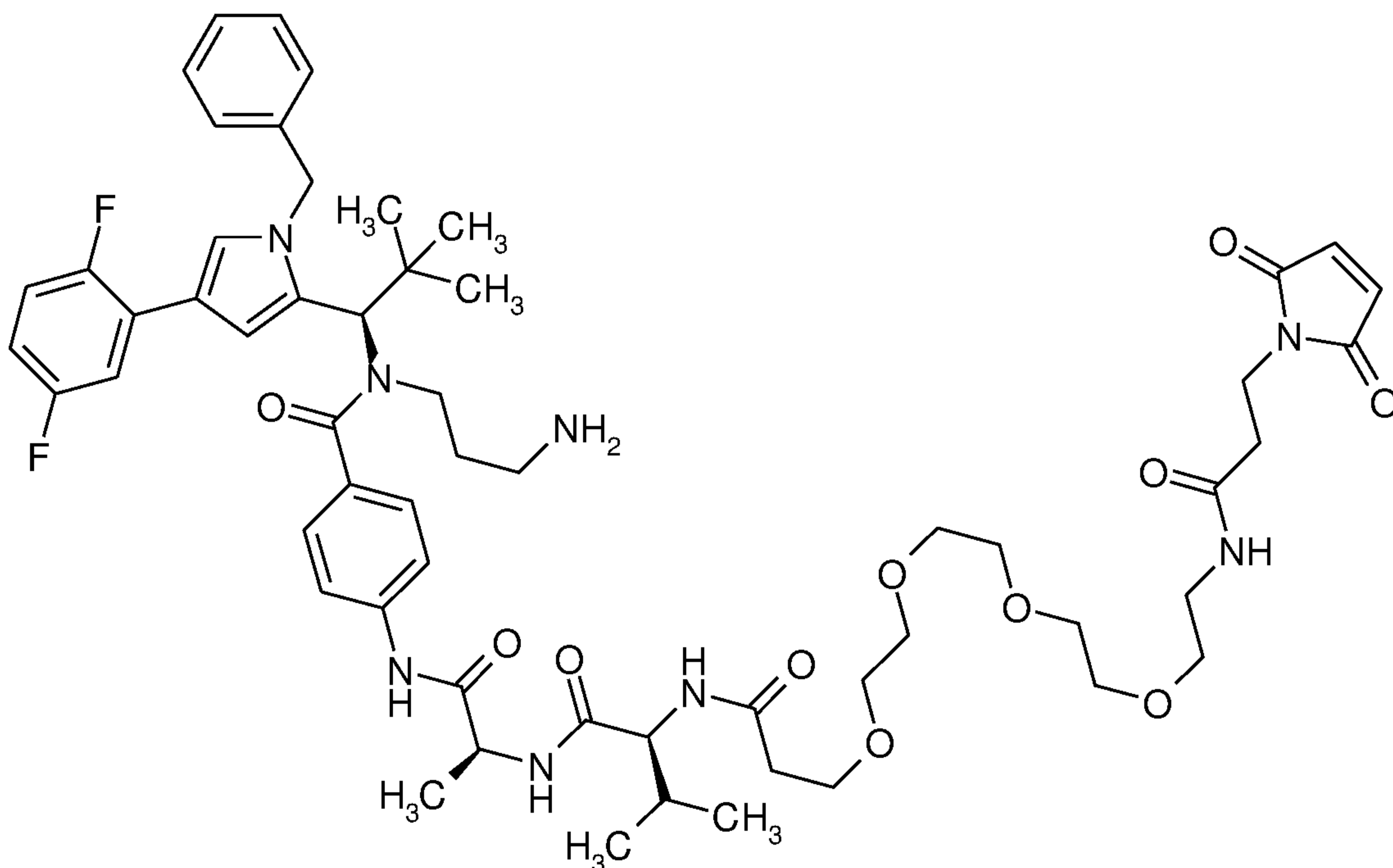
13.8 mg (11.1  $\mu\text{mol}$ ) of N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-gamma-glutamyl-S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine were dissolved in 2.0 ml of trifluoroethanol, and 9.1 mg (66.5  $\mu\text{mol}$ ) zinc dichloride were added. The reaction mixture was stirred at 50°C for 4 h. 19.4 mg (0.07 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The

solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 6.9 mg (50% of theory) of the title compound.

5 LC-MS (Method 1):  $R_t = 0.89$  min; MS (ESIpos):  $m/z = 1100$  (M+H)<sup>+</sup>.

### Intermediate F235

10 Trifluoroacetic acid / N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-valyl-N-{4-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}carbamoyl]phenyl}-L-alaninamide (1:1)



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120.0 mg (0.22 mmol) of 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino)propyl]carbamate (see synthesis of  
 20 Intermediate C11) and 52.1 mg (0.28 mmol) of 4-nitrobenzoyl chloride were dissolved in 8.0 ml of dichloromethane, and 28.4 mg (0.28 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and the residue was purified by preparative RP-  
 25 HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min,



MeCN/water, 0.1% TFA). This gave 97.7 mg (64% of theory) of the compound 2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(4-nitrobenzoyl)amino]propyl}carbamate.

5

LC-MS (Method 1):  $R_t = 1.54$  min; MS (ESIpos):  $m/z = 705$  (M+H)<sup>+</sup>.

97.0 mg (0.14 mmol) of 2-(trimethylsilyl)ethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(4-nitrobenzoyl)amino]propyl}carbamate were dissolved in 5.0 ml of ethanol, 9.7 mg of palladium on activated carbon (10%) were added and the mixture was hydrogenated at standard pressure for 5 h. The reaction mixture was filtered through a cardboard filter and the filter cake was washed with ethanol. The solvent was evaporated under reduced pressure. The residue was used without further purification in the next step of the synthesis. This gave 87.4 mg (88% of theory) of the compound 2-(trimethylsilyl)ethyl {3-[(4-aminobenzoyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]propyl}carbamate.

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LC-MS (Method 1):  $R_t = 1.47$  min; MS (ESIpos):  $m/z = 675$  (M+H)<sup>+</sup>.

59.3 mg (0.09 mmol) of 2-(trimethylsilyl)ethyl {3-[(4-aminobenzoyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]propyl}carbamate and 25.5 mg (0.11 mmol) of N-[(benzyloxy)carbonyl]-L-alanine were initially charged together with 68.1 mg (0.53 mmol) of N,N-diisopropylethylamine in 5.0 ml of acetonitrile. 72.7 mg (0.11 mmol) of T3P (50% in ethyl acetate) were added slowly. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 52.2 mg (68% of theory) of the compound benzyl [(2S)-1-{[4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[3-({[2-

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(trimethylsilyl)ethoxy]carbonyl}amino)propyl]carbamoylethyl]phenyl]amino}-1-oxopropan-2-yl]carbamate.

LC-MS (Method 1):  $R_t = 1.48$  min; MS (ESIpos):  $m/z = 880$  (M+H)<sup>+</sup>.

5

23.9 mg (0.03 mmol) of benzyl [(2S)-1-{{4-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)}[3-({[2-

(trimethylsilyl)ethoxy]carbonyl}amino)propyl]carbamoylethyl]phenyl]amino}-1-oxopropan-2-yl]carbamate were dissolved in 3.0 ml of ethyl acetate, 2.4 mg of palladium on activated carbon (10%) were added and the mixture was hydrogenated at standard pressure for 2 h. The reaction mixture was filtered through a paper filter and the filter cake was washed with ethyl acetate. The solvent was evaporated under reduced pressure. The residue was used without further purification in the next step of the synthesis. This gave 20.1 mg (90% of theory) of the compound 2-(trimethylsilyl)ethyl [3-([4-(L-alanyl)amino]benzoyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)propyl]carbamate.

20

LC-MS (Method 1):  $R_t = 1.13$  min; MS (ESIpos):  $m/z = 746$  (M+H)<sup>+</sup>.

20.0 mg (0.03 mmol) of 2-(trimethylsilyl)ethyl [3-([4-(L-alanyl)amino]benzoyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)propyl]carbamate were initially charged together with 14.9 mg (0.04 mmol) of 2,5-dioxopyrrolidin-1-yl N-[(benzyloxy)carbonyl]-L-valinate in 2.0 ml of DMF, and 5.4 mg (0.05 mmol) of 4-methylmorpholine were added. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave the compound N-[(benzyloxy)carbonyl]-L-valyl-N-[4-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)}[3-({[2-(trimethylsilyl)ethoxy]carbonyl}amino)propyl]carbamoylethyl]phenyl]-L-alaninamide.

35

LC-MS (Method 1):  $R_t = 1.49$  min; MS (ESIpos):  $m/z = 979$  (M+H)<sup>+</sup>.

17.0 mg (17.4  $\mu$ mol) of N-[(benzyloxy)carbonyl]-L-valyl-N-[4-  
5 ([(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
dimethylpropyl])[3-([2-  
(trimethylsilyl)ethoxy]carbonyl)amino)propyl]carbamoyl)phenyl]  
-L-alaninamide were dissolved in 2.5 ml of ethyl acetate, 1.7  
mg of palladium on activated carbon (10%) were added and the  
10 mixture was hydrogenated at standard pressure overnight. The  
reaction mixture was filtered through a paper filter and the  
filter cake was washed with ethyl acetate. The solvent was  
evaporated under reduced pressure and the residue was dried  
under high vacuum. This gave 15.3 mg (60% of theory) of the  
15 compound L-valyl-N-[4-([(1R)-1-[1-benzyl-4-(2,5-  
difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl])[3-([2-  
(trimethylsilyl)ethoxy]carbonyl)amino)propyl]carbamoyl)phenyl]  
-L-alaninamide.

20 LC-MS (Method 1):  $R_t = 1.15$  min; MS (ESIpos):  $m/z = 845$  (M+H)<sup>+</sup>.

15.3 mg (0.01 mmol) of L-valyl-N-[4-([(1R)-1-[1-benzyl-4-(2,5-  
difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl])[3-([2-  
(trimethylsilyl)ethoxy]carbonyl)amino)propyl]carbamoyl)phenyl]  
25 -L-alaninamide were initially charged together with 7.9 mg (0.02  
mmol) of 3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-N-{15-[(2,5-  
dioxopyrrolidin-1-yl)oxy]-15-oxo-3,6,9,12-tetraoxapentadec-1-  
yl}propanamide in 2.4 ml of DMF, and 1.9 mg (0.02 mmol) of 4-  
methylmorpholine were added. The reaction mixture was stirred  
30 at RT overnight, and 1.4 mg (0.02 mmol) of acetic acid were then  
added. The reaction mixture was purified directly by preparative  
RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min,  
MeCN/water). The solvents were evaporated under reduced pressure  
and the residue was dried under high vacuum. This gave 11.7 mg  
35 (70% of theory) of the compound N-[19-(2,5-dioxo-2,5-dihydro-  
1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-  
oyl]-L-valyl-N-[4-([(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
1H-pyrrol-2-yl]-2,2-dimethylpropyl])[3-([2-

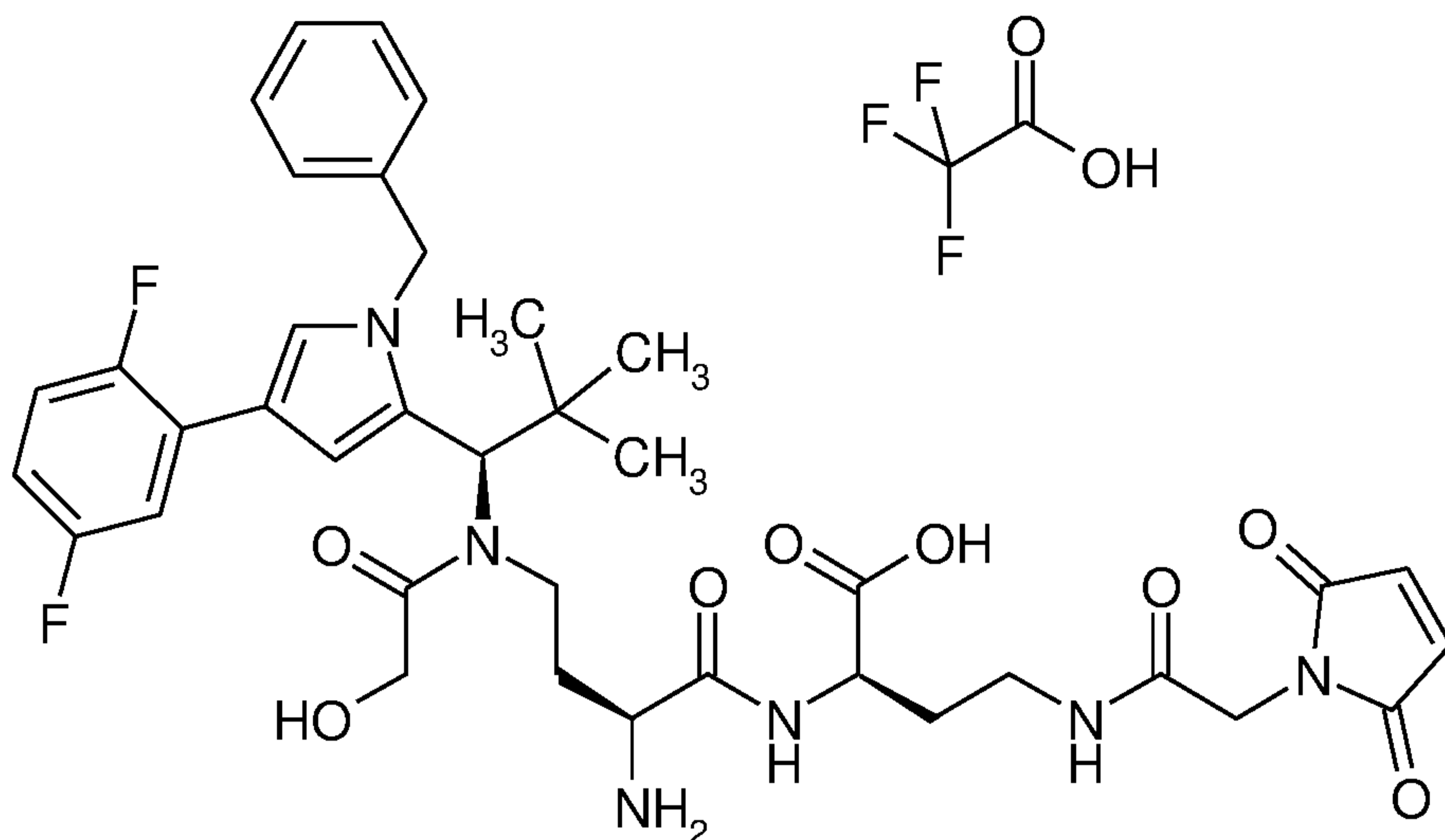
(trimethylsilyl)ethoxy]carbonyl}amino)propyl]carbamoyl)phenyl]  
-L-alaninamide.

11.7 mg (0.01 mmol) of N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-  
5 1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-  
gamma-glutamyl-S-(11-((1R)-1-[1-N-[19-(2,5-dioxo-2,5-dihydro-  
1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-  
oyl]-L-valyl-N-[4-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
1H-pyrrol-2-yl]-2,2-dimethylpropyl)}[3-([2-  
10 (trimethylsilyl)ethoxy]carbonyl}amino)propyl]carbamoyl)phenyl]  
-L-alaninamide were dissolved in 2.0 ml of trifluoroethanol, and  
3.9 mg (0.03 mmol) of zinc dichloride were added. The reaction  
mixture was stirred at 50°C overnight. 8.3 mg (0.03 mmol) of  
ethylenediamine-N,N,N',N'-tetraacetic acid were added, the  
15 reaction mixture was stirred for 10 min and water (0.1% TFA) was  
then added. Purification was carried out directly by preparative  
RP-HPLC (column: Reprosil 125x30; 10µ, flow rate: 50 ml/min,  
MeCN/water, 0.1% TFA). The solvents were evaporated under  
reduced pressure and the residue was dried under high vacuum.  
20 This gave 5.4 mg (47% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.94$  min; MS (ESIpos):  $m/z = 1100$  (M+H)<sup>+</sup>.

### Intermediate F236

25 (2R)-2-((2S)-2-Amino-4-((1R)-1-[1-benzyl-4-(2,5-  
difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
dimethylpropyl)(glycoloyl)amino]butanoyl)amino)-4-[[2,5-  
dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino]butanoic acid /  
30 trifluoroacetic acid (1:1)

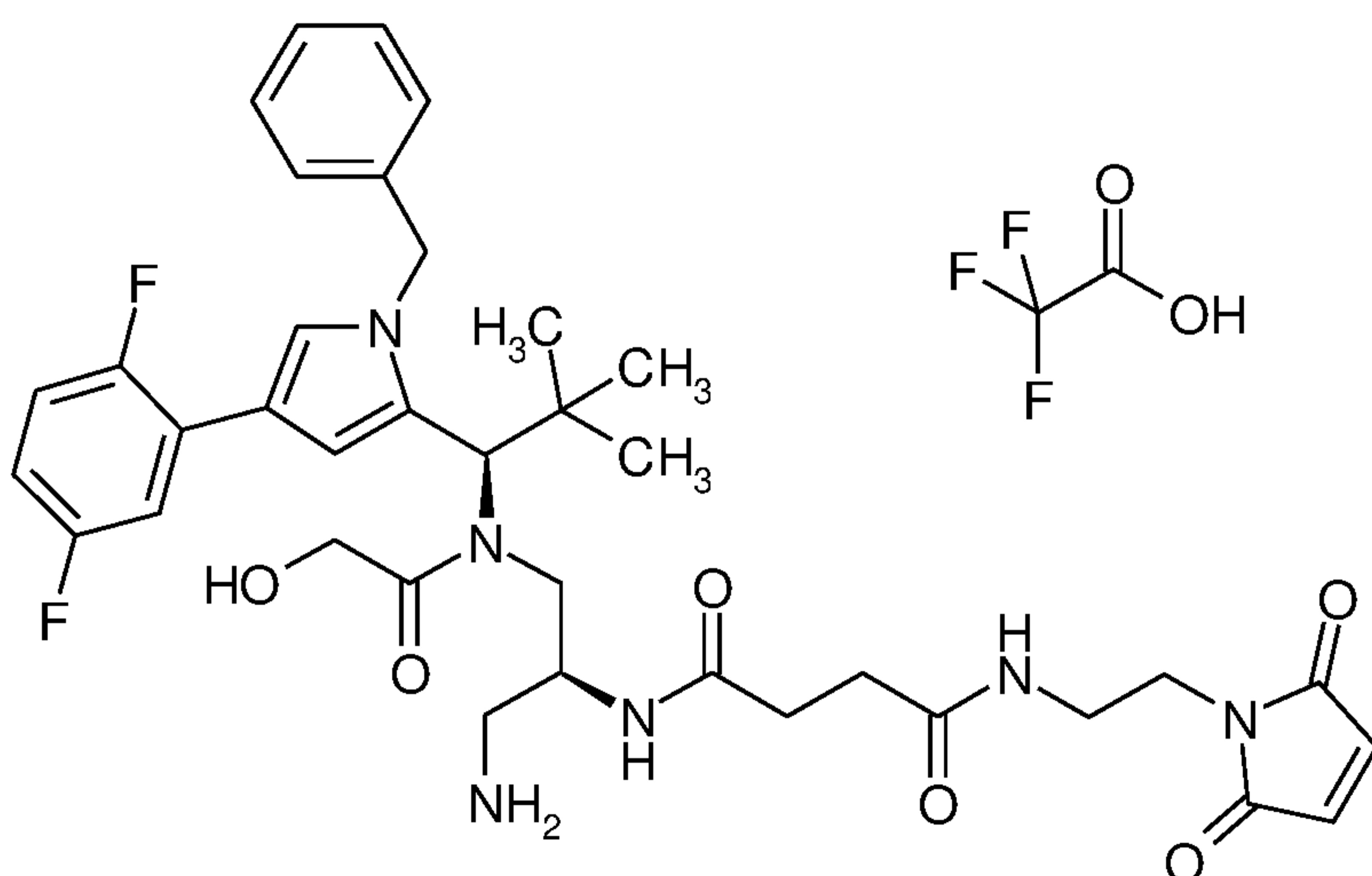


The synthesis of the title compound was carried out analogously to Intermediate F192 from (2R)-4-[[[(benzyloxy)carbonyl]amino]-2-[(tert-butoxycarbonyl)amino]butanoic acid / N-cyclohexylcyclohexanamine (1:1).

LC-MS (Method 4):  $R_t = 1.1$  min; MS (ESIpos):  $m/z = 751$  (M+H)<sup>+</sup>.

#### 10 Intermediate F238

Trifluoroacetic acid / N-{(2S)-1-amino-3-[[[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]propan-2-yl]-N'-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]succinamide (1:1)



18 mg (0.025 mmol) of Intermediate C72 were taken up in 6 ml of

DMF and coupled with 7.5 mg (0.03 mmol) of trifluoroacetic acid / 1-(2-aminoethyl)-1H-pyrrole-2,5-dione (1:1) in the presence of 11.3 mg (0.03 mmol) of HATU and 22  $\mu$ l of *N,N*-diisopropylethylamine. After 1 h of stirring at RT, the reaction was concentrated and the residue was purified by preparative HPLC. The appropriate fractions were concentrated and the residue was lyophilized from acetonitrile/water 1:1. This gave 15 mg (67% of theory) of the intermediate.

10 LC-MS (Method 4):  $R_t = 1.71$  min; MS (EIpos):  $m/z = 873$   $[M+Na]^+$ .

The title compound was then prepared from this intermediate by deprotection with zinc chloride in 4 ml of trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 8.5 mg (63% of theory) of the title compound.

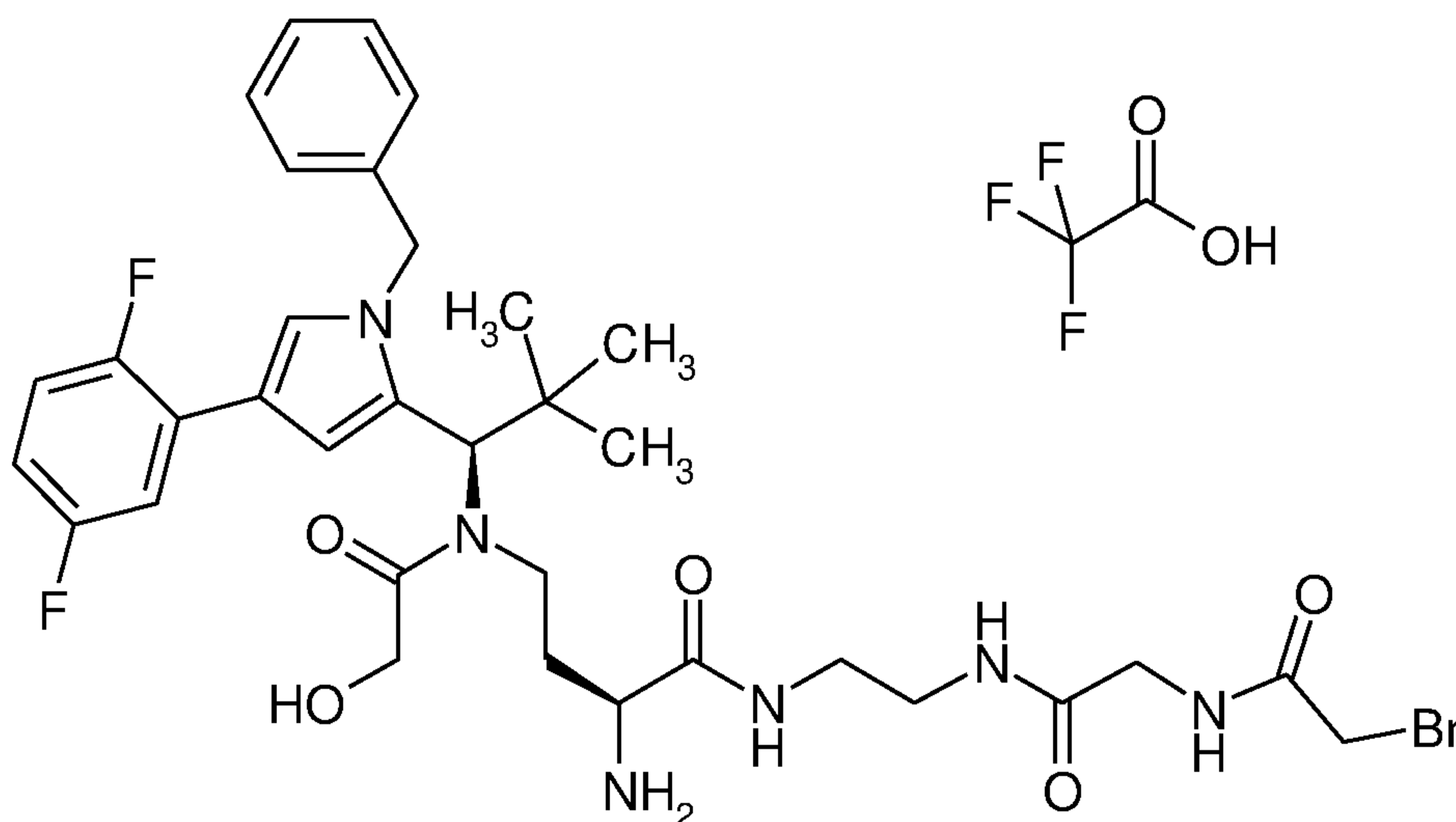
LC-MS (Method 1):  $R_t = 0.86$  min; MS (ESIpos):  $m/z = 707$   $(M+Na)^+$ .

### Intermediate F241

20

Trifluoroacetic acid/ (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-(2-{[N-(bromoacetyl)glycyl]amino}ethyl)butanamide (1:1)

25



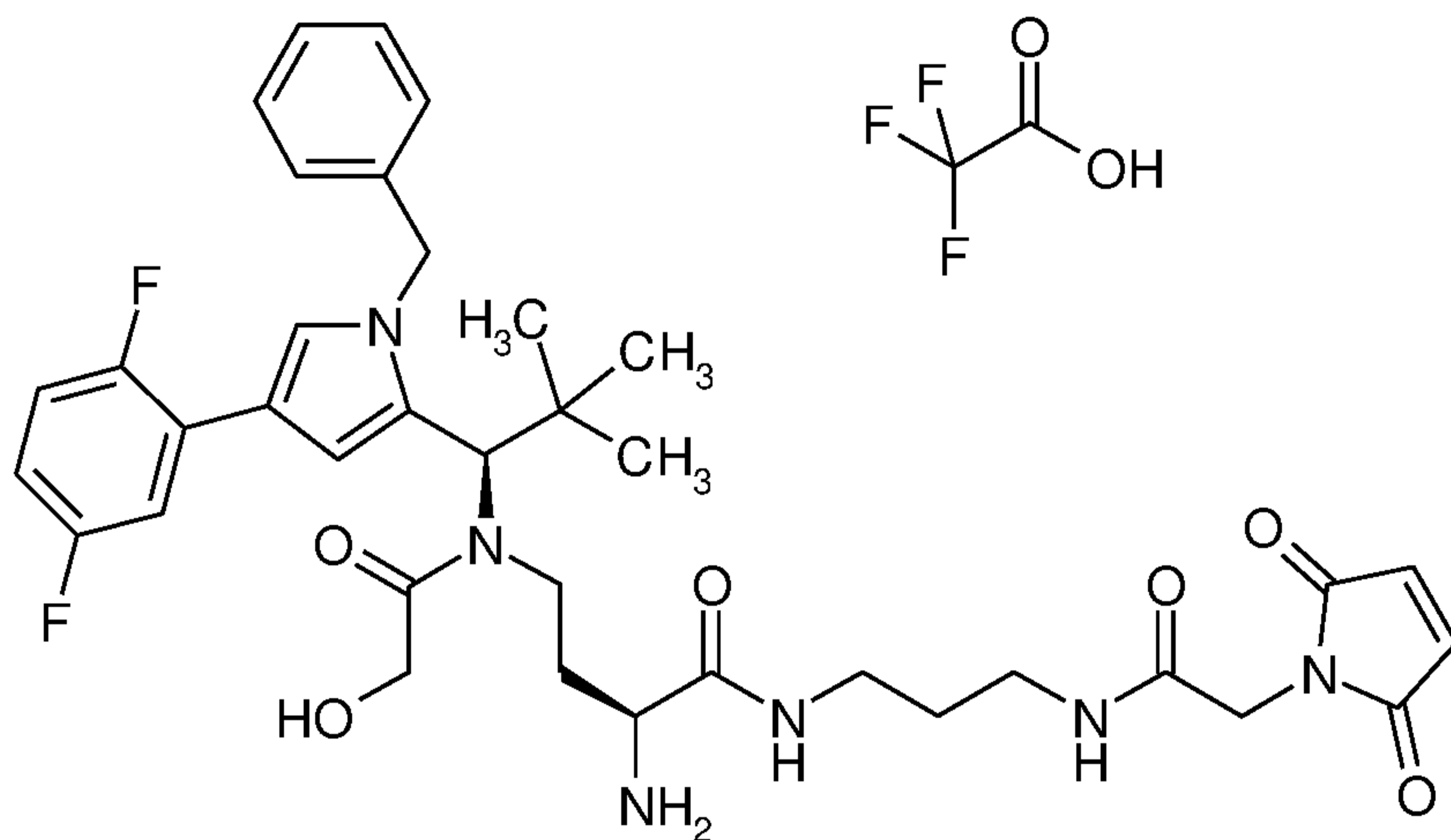
The title compound was prepared analogously from Intermediate

C66 by coupling with commercially available 1-(2-bromoacetoxy)pyrrolidine-2,5-dione and subsequent deblocking with zinc chloride.

- 5 LC-MS (Method 1):  $R_t = 0.84$  min; MS (EIpos):  $m/z = 733$  and  $735$   $[M+H]^+$ .

### Intermediate F242

- 10 Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-(3-{[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino}propyl)butanamide (1:1)



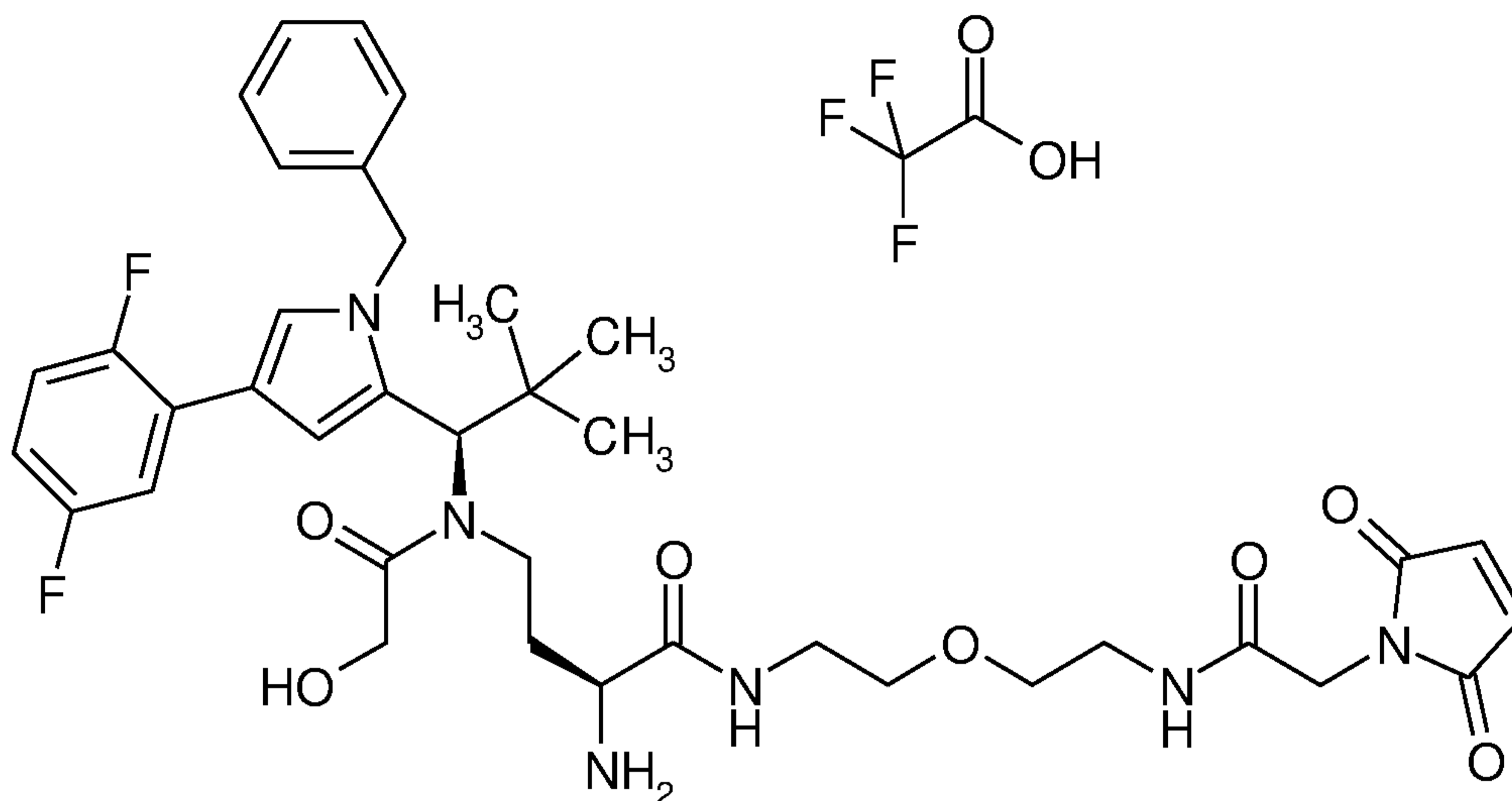
15

The synthesis of the title compound was carried out analogously to Intermediate F104.

- 20 LC-MS (Method 1):  $R_t = 0.84$  min; MS (ESIpos):  $m/z = 707$   $(M+H)^+$ .

### Intermediate F243

- 25 Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-[2-(2-{[(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)acetyl]amino}ethoxy)ethyl]butanamide (1:1)



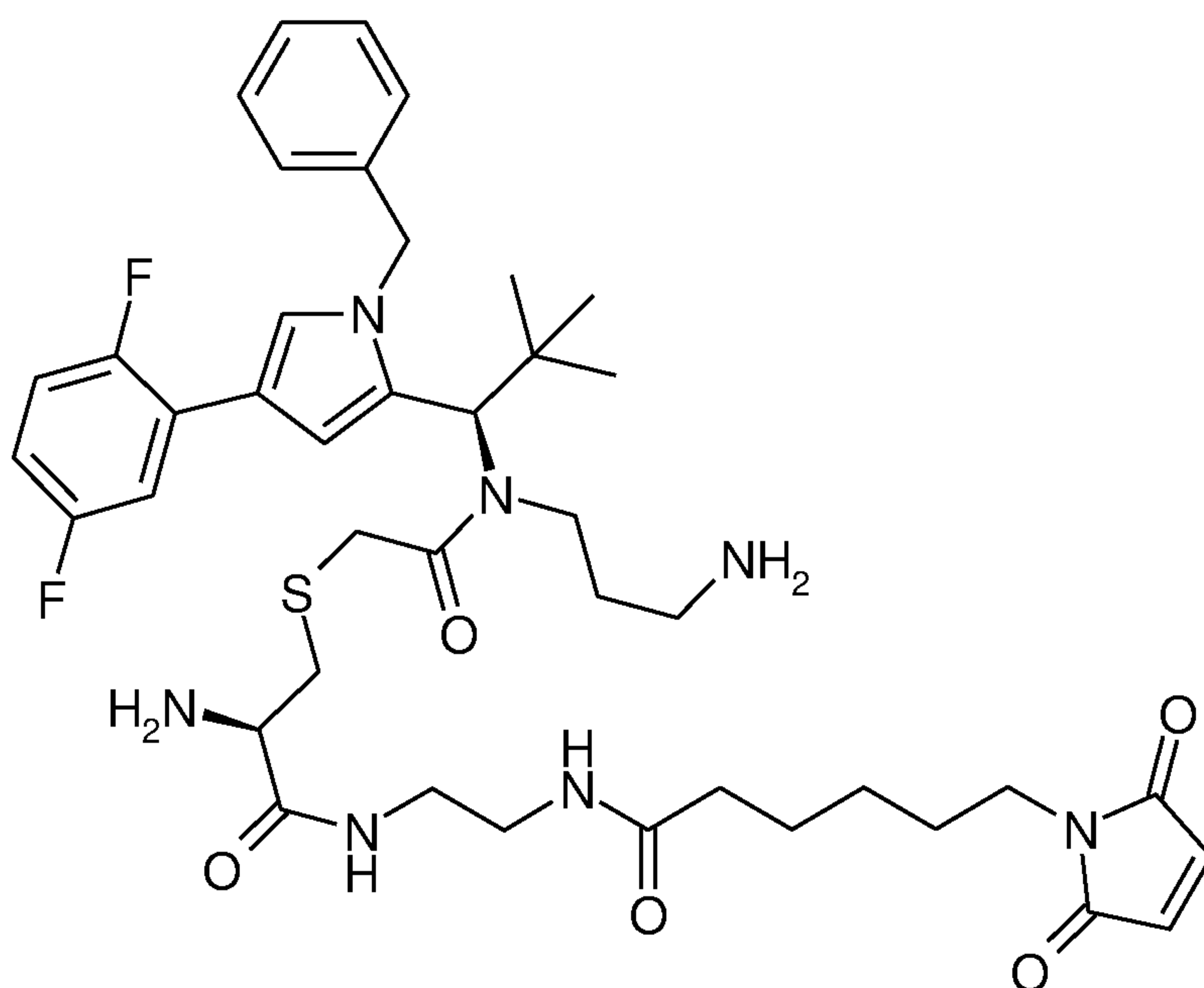
The synthesis of the title compound was carried out analogously to Intermediate F242.

5

LC-MS (Method 1):  $R_t = 0.81$  min; MS (ESIpos):  $m/z = 737$  (M+H)<sup>+</sup>.

#### Intermediate F244

- 10 N-{2-[(S-{2-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-L-cysteinyl)amino]ethyl}-6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanamide



15



100 mg (about 0.101 mmol) of S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-N-[3-(trimethylsilyl)propanoyl]-L-cysteine (Intermediate C 73) were initially charged in 88 ml of dimethylformamide, and with 107 mg (about 0.15 mmol) of N-(2-aminoethyl)-6-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)hexanamide (Intermediate L73 ), 46 mg (0.12 mmol) of HATU and 88  $\mu$ l (0.50 mmol) of were added. The reaction mixture was stirred at RT for 15 minutes. Water/dichloromethane was added to the mixture, and the organic phase was then washed with water and brine, dried over magnesium sulphate, concentrated on a rotary evaporator and dried under high vacuum. The residue was used further without further purification. This gave 92 mg (59%, purity 72%) of the title compound.

LC-MS (Method 1):  $R_t$  = 1.59 min; MS (ESIpos): m/z = 1096 (M+H)<sup>+</sup>.

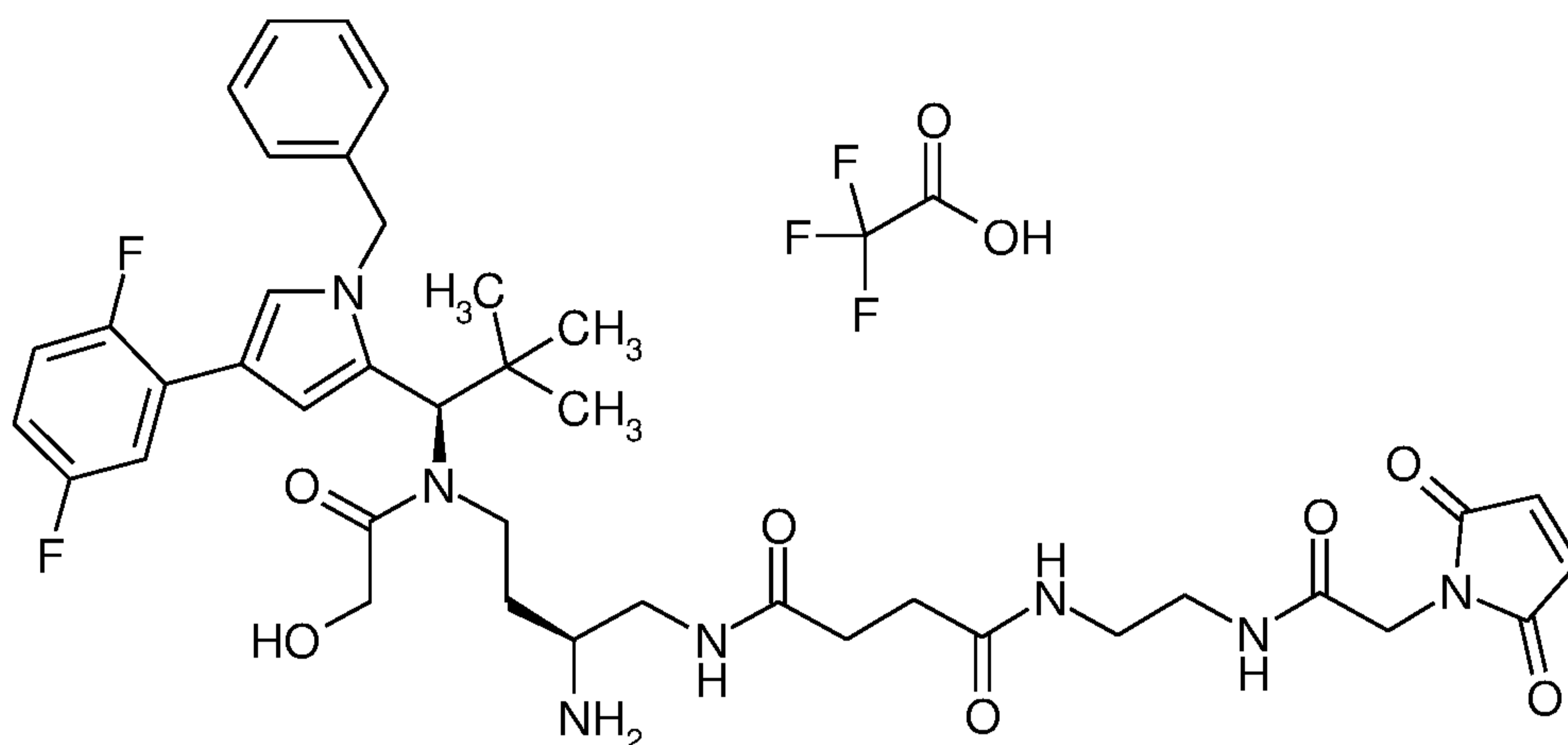
Under argon, 40 mg (0.30 mmol) of zinc chloride were added to a solution of 91 mg (about 0.06 mmol) of 2-(trimethylsilyl)ethyl [(9R)-4-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-20-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-5,10,15-trioxo-9-[[3-(trimethylsilyl)propanoyl]amino]-7-thia-4,11,14-triazaicos-1-yl]carbamate in 1.45 ml of trifluoroethanol. The reaction mixture was stirred at 50°C for 2 h. 30 mg (0.22 mmol) of zinc chloride were then added, and the mixture was stirred at RT for another 1 h. 52 mg (0.18 mmol) of EDTA were added, and after 10 minutes of stirring at RT the mixture was diluted slightly with water/acetonitrile and purified by preparative HPLC (mobile phase: ACN/water +0.1%TFA, gradient). This gave 17 mg (31%) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.80 min; MS (ESIpos): m/z = 808 (M+H)<sup>+</sup>.

### 35 Intermediate F245

Trifluoroacetic acid / N-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-

dimethylpropyl} (glycoloyl) amino]butyl}-N'-(2-{{(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl) acetyl] amino}ethyl) succinamide (1:1)



5

The title compound was prepared by coupling of 10 mg (0.0135 mmol) of Intermediate C65 with 8 mg (0.027 mmol) of Intermediate L1 in 8 ml of DMF in the presence of 15 mg (0.04 mmol) of HATU and 9  $\mu$ l of *N,N*-diisopropylethylamine and subsequent deprotection with zinc chloride in trifluoroethanol as described for Intermediate F119. Purification by preparative HPLC gave 8.8 mg (58% of theory over 2 steps) of the title compound.

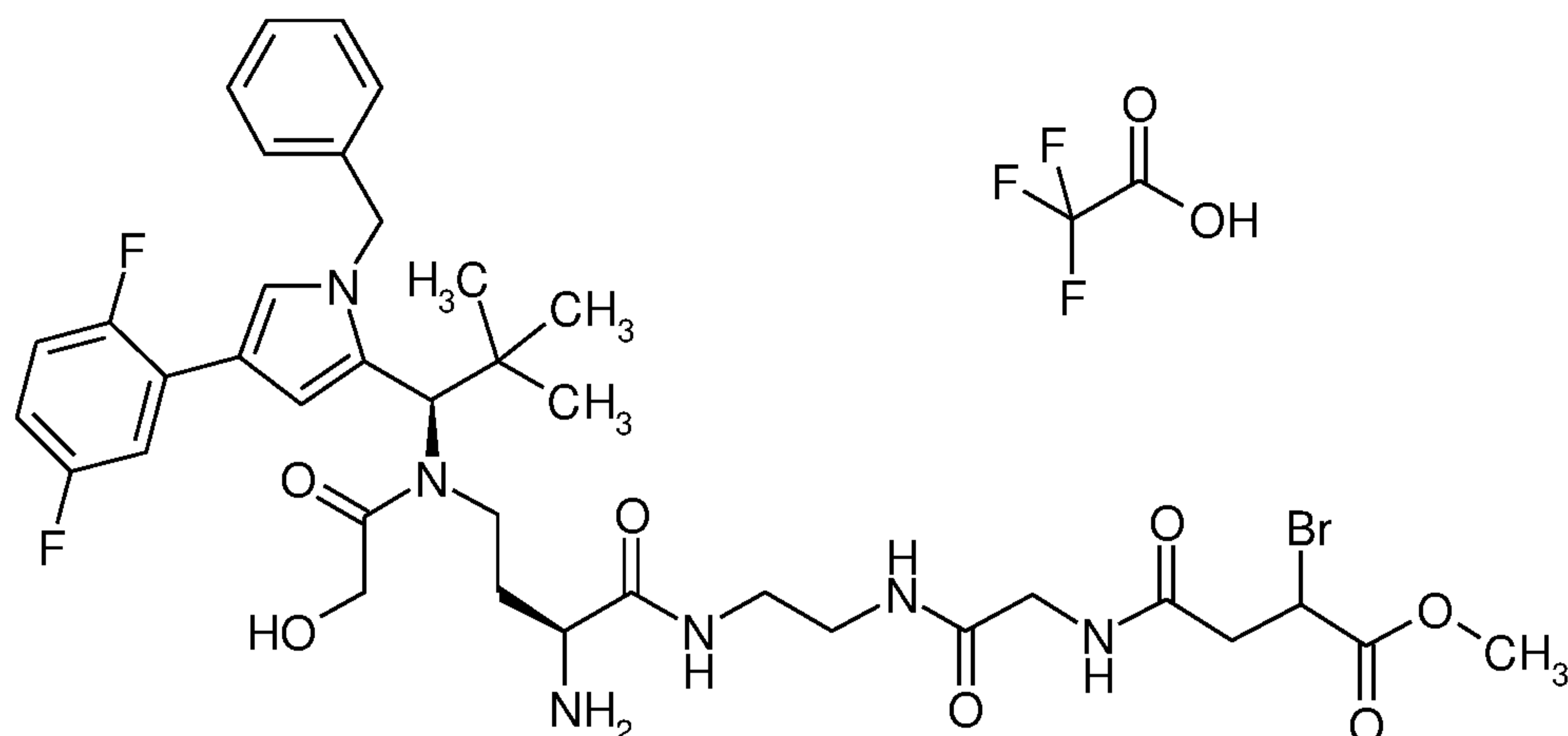
15

LC-MS (Method 1):  $R_t$  = 0.84 min; MS (ESIpos):  $m/z$  = 778 (M+H)<sup>+</sup>.

### Intermediate F247

Trifluoroacetic acid / methyl 4-[(2-{{[2-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl} amino) ethyl] amino}-2-oxoethyl) amino]-2-bromo-4-oxobutanoate (1:1)

20

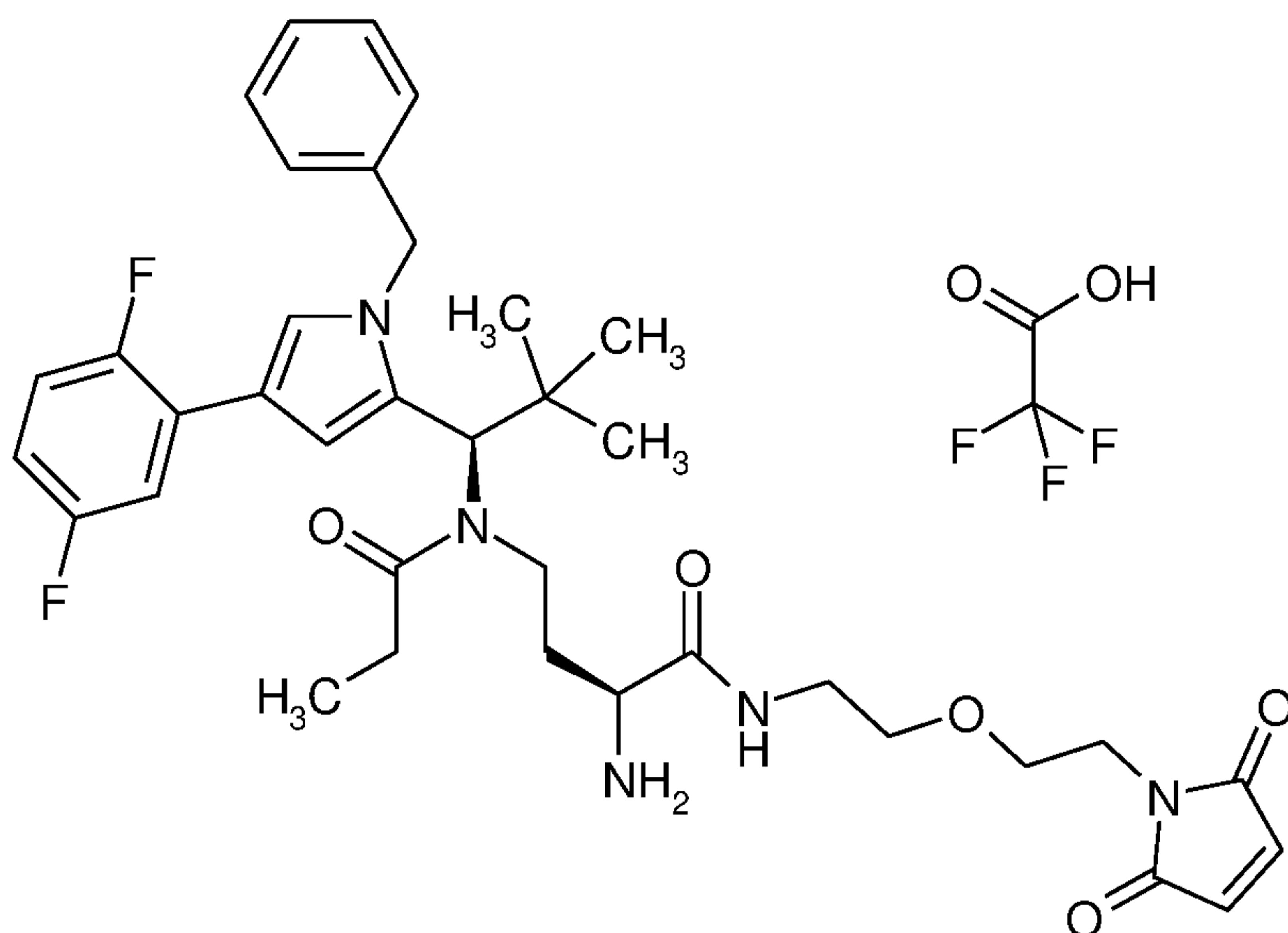


14 mg (0.018 mmol) of Intermediate C66 were dissolved in 14 ml of DCM, and with 10.1 mg (0.037 mmol) of 2-bromo-1-ethylpyridinium tetrafluoroborate (BEP) and, a little at a time, a total of 250  $\mu$ l of pyridine were added, the pH being kept between 5 and 6. The pH was then adjusted to 4 with acetic acid, the reaction was concentrated and the residue was purified by preparative HPLC. Combination of the appropriate fractions, lyophilization and drying gave 4 mg (21% of theory) the protected intermediate, which were then deprotected at the amino function with zinc chloride. HPLC purification and lyophilisation gave 3 mg (72% of theory) of the title compound as a colourless foam.

LC-MS (Method 1):  $R_t$  = 0.88 min; MS (ESIpos):  $m/z$  = 805 and 807 (M+H)<sup>+</sup>.

### Intermediate F248

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]-N-{2-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethoxy]ethyl}butanamide (1:1)



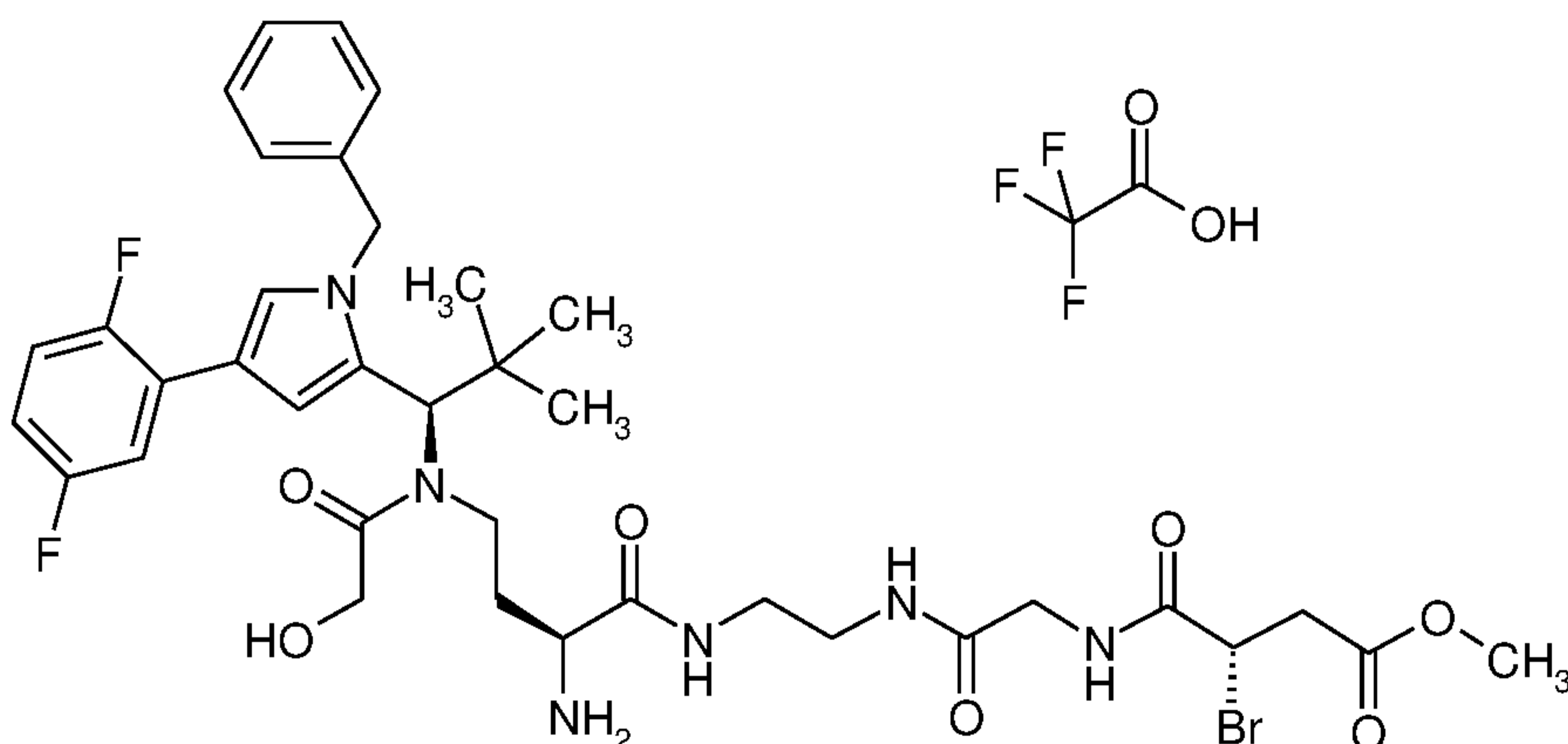
The title compound was prepared by coupling of 10 mg (0.015 mmol) of Intermediate C58 with 5 mg (0.017 mmol) of Intermediate L12 in the presence of HATU and subsequent deprotection with zinc chloride. This gave 6.5 mg (52% of theory over 2 steps) of the title compound.

LC-MS (Method 1):  $R_t = 0.91$  min; MS (ESIpos):  $m/z = 680$  (M+H)<sup>+</sup>.

10

### Intermediate F254

Trifluoroacetic acid / methyl (3S)-4-[(2-[[2-((2S)-2-amino-4-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl] amino}-2-oxoethyl)amino]-3-bromo-4-oxobutanoate (1:1)



20 The title compound was prepared analogously to Intermediate 247

by coupling of 15 mg (0.02 mmol) of Intermediate C66 with 21 mg  
 (0.099 mmol) of (2S)-2-bromo-4-methoxy-4-oxobutanoic acid which  
 had been synthesized as described in (J.Org.Chem. 200, 65, 517-  
 522) from (2S)-2-amino-4-methoxy-4-oxobutanoic acid  
 5 hydrochloride (1:1).

LC-MS (Method 1):  $R_t = 0.89$  min; MS (ESIpos):  $m/z = 805$  and  $807$   
 (M+H)<sup>+</sup>.

10 **B: Preparation of antibody/active compound conjugates (ADC)**

B-1. General process for generating anti-TWEAKR antibodies

The anti-TWEAKR antibodies were generated, for example, by  
 15 screening of a phage display library for recombinant human  
 TWEAKR SEQ ID NO: 138 and murine TWEAKR SEQ ID NO: 137. The  
 antibodies obtained in this manner were reformatted into the  
 human IgG1 format and used for the working examples described  
 here. In addition, antibodies which bind to TWEAKR are known to  
 20 the person skilled in the art, see, for example,  
 WO2009/020933(A2) or WO2009140177 (A2).

SEQ ID NO:138 (polypeptide):

25 EQAPGTAPCSRGSWSADLDKCMDCASCRARPHSDFCLGCAAAPPAPFRLWPRSDKTHT

CPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVH

NAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPRE

30

PQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFF

LYSKLTVDKSRWQQGNVFCSSVMHEALHNHYTQKSLSLSPGK

35 SEQ ID NO:137 (polypeptide):

EQAPGTSPCSSGSSWSADLDKCMDCASCPCARPHSDFCLGCAAAPPAHFRLWPRSDKTHT

CPPCPAPELLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVH

NAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPRE

5 PQVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFF

LYSKLTVDKSRWQQGNVFNCSVMHEALHNHYTQKSLSLSPGK

10 B-2. General process for expressing anti-TWEAKR antibodies  
in mammalian cells

The antibodies, for example TPP-2090, were produced in transient mammalian cell cultures as described by Tom et al., Chapter 12 in Methods Express: Expression Systems edited by Micheal R. Dyson and Yves Durocher, Scion Publishing Ltd, 2007 (see AK-Example 1).

20 B-3. General process for purifying antibodies from cell supernatants

The antibodies, for example TPP-2090, were obtained from the cell culture supernatants. The cell supernatants were clarified by centrifugation of cells. The cell supernatant was then purified by affinity chromatography on a MabSelect Sure (GE Healthcare) chromatography column. To this end, the column was equilibrated in DPBS pH 7.4 (Sigma/Aldrich), the cell supernatant was applied and the column was washed with about 10 column volumes of DPBS pH 7.4 + 500 mM sodium chloride. The antibodies were eluted in 50 mM sodium acetate pH 3.5 + 500 mM sodium chloride and then purified further by gel filtration chromatography on a Superdex 200 column (GE Healthcare) in DPBS pH 7.4.

35 The commercially available antibody cetuximab (trade name Erbitux) was purified from the commercial product by standard chromatographic methods (protein A, preparative SEC).

The commercially available antibody trastuzumab (trade name

Herceptin) was purified from the commercial product by standard chromatographic methods (protein A, preparative SEC).

From the commercial product (trade name CIMAher), the antibody  
5 nimotuzumab was purified from the commercial product by standard chromatographic methods (protein A, preparative SEC).

From the commercial product (trade name Vectibix), the antibody  
10 panitumumab was purified from the commercial product by standard chromatographic methods (protein A, preparative SEC).

#### B-4. General process for coupling to cysteine side chains

The following antibodies were used for the coupling reactions:

15

cetuximab (anti EGFR AK)

anti-TWEAKR AK 1 (TPP-2090)

20

trastuzumab (anti-Her2 AK)

nimotuzumab (anti-EGFR AK)

panitumumab (anti-EGFR AK)

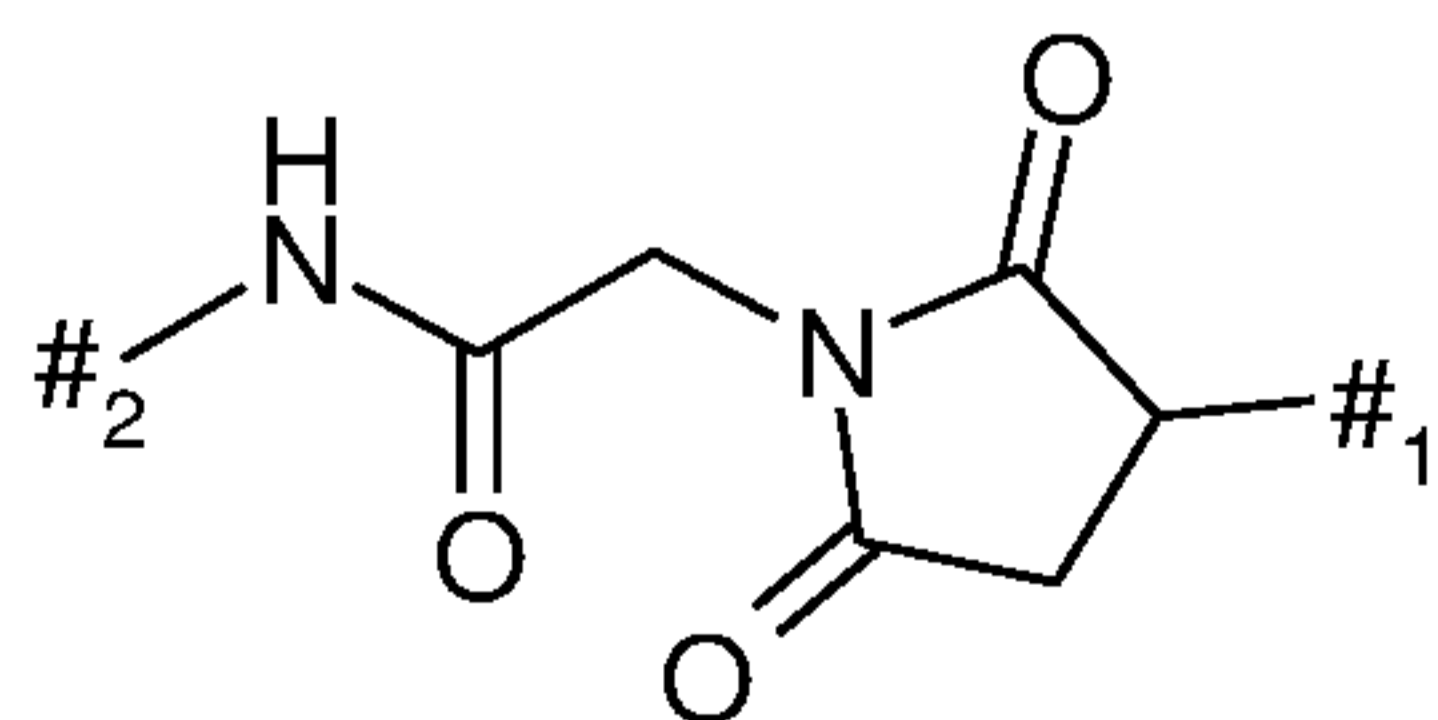
25

Between 2 and 5 equivalents of tris(2-carboxyethyl)phosphine hydrochloride (TCEP), dissolved in PBS buffer, were added to a solution of the appropriate antibody in PBS buffer in the concentration range between 1 mg/ml and 20 mg/ml, preferably in  
30 the range of about 10 mg/ml to 15 mg/ml, and the mixture was stirred at RT for 1h. For this purpose, the solution of the respective antibody used can be employed at the concentrations stated in the working examples, or it may optionally also be diluted with PBS buffer to about half of the stated starting  
35 concentrations in order to get into the preferred concentration range. Subsequently, depending on the intended loading from 2 to 12 equivalents, preferably about 5-10 equivalents of the maleinimide precursor compound or halide precursor compound to

be coupled can be added as a solution in DMSO. Here, the amount of DMSO should not exceed 10% of the total volume. The reaction was stirred in the case of maleinimide precursors for 60-240 min at RT and in the case of halide precursors between 8 and 24 h at RT and then applied to PBS-equilibrated PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) and eluted with PBS buffer. Generally, unless indicated otherwise, 5 mg of the antibody in question in PBS buffer were used for the reduction and the subsequent coupling. Purification on the PD10 column thus in each case afforded solutions of the respective ADCs in 3.5 ml PBS buffer. The sample was then concentrated by ultracentrifugation and optionally rediluted with PBS buffer. If required, for better removal of low-molecular weight components, concentration by ultrafiltration was repeated after redilution with PBS buffer. For biological tests, if required, the concentrations of the final ADC samples were optionally adjusted to the range of 0.5-15 mg/ml by redilution. The respective protein concentrations, stated in the working examples, of the ADC solutions were determined. Furthermore, antibody loading (drug/mAb ratio) was determined using the methods described under B-7.

Unless indicated otherwise, the immunoconjugates shown in the examples were prepared by this process. Depending on the linker, the ADCs shown in the examples may also be present to a lesser or higher degree in the form of the hydrolysed open-chain succinamides attached to the antibodies.

In particular the KSP-I-ADCs attached through the linker substructure



to thiol groups of the antibodies may optionally also be prepared in a targeted manner by rebuffering after the coupling and



stirring at pH 8 for about 20 h according to Scheme 26 via the ADCs attached via open-chain succinamides.

#1 represents the sulphur bridge to the antibody, and #2 the  
5 point of attachment to the modified KSP inhibitor

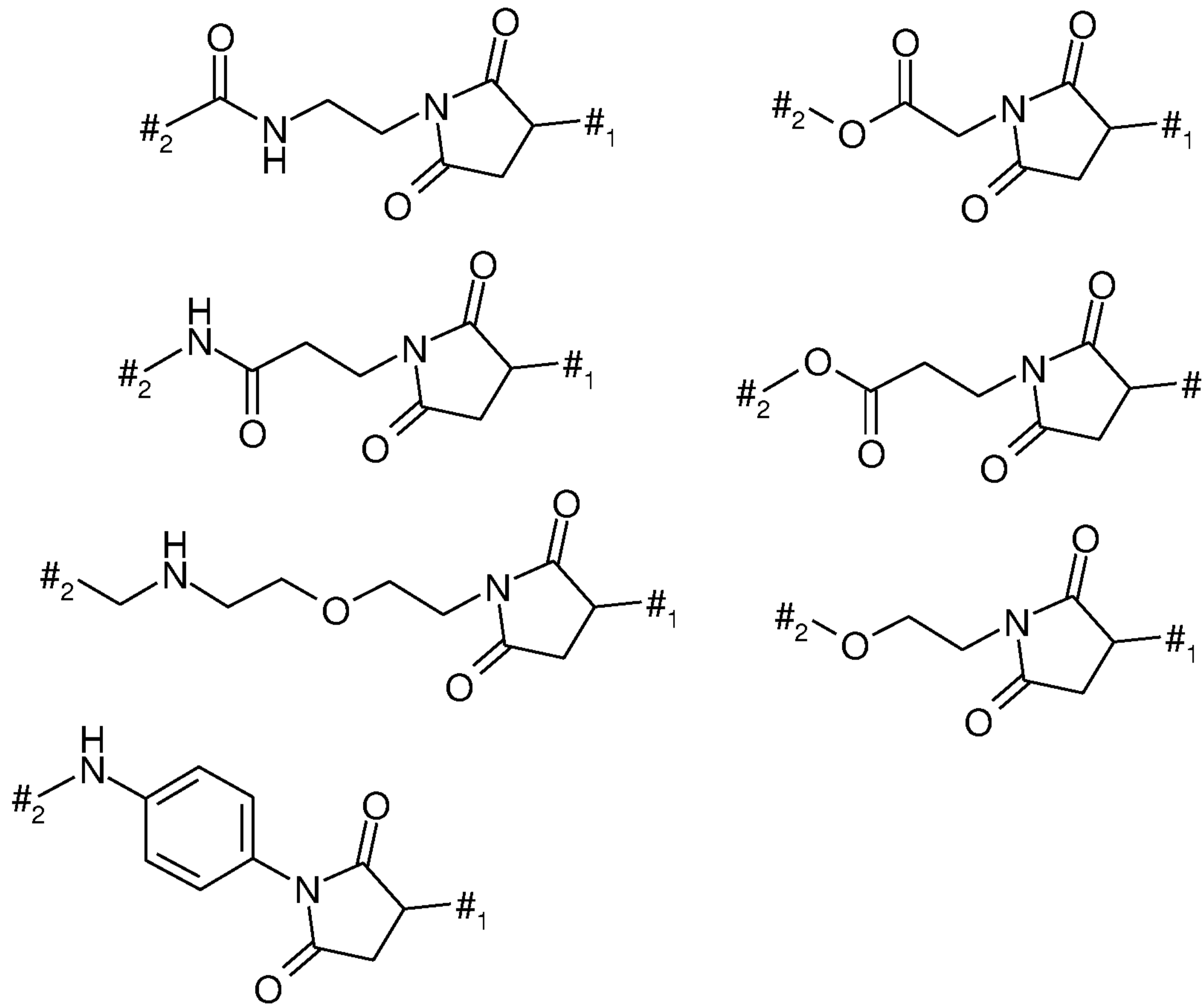
Such ADCs where the linker is attached to the antibodies through hydrolysed open-chain succinamides may optionally also be prepared in a targeted manner by an exemplary procedure as  
10 follows:

Under argon, a solution of 0.344 mg TCEP in 100 µl of PBS buffer was added to 60 mg of the antibody in question in 5 ml of PBS buffer (c~12 mg/ml). The reaction was stirred at RT for 30 min,  
15 and 0.003 mmol of a maleinimide precursor compound dissolved in 600 µl of DMSO was then added. After a further 1.5 h - 2 h of stirring at RT, the reaction was diluted with 1075 µl of PBS buffer which had been adjusted to pH 8 beforehand.

20 This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was diluted with PBS buffer pH 8 to a total volume of 14 ml. This solution was stirred at RT under argon overnight. If required, the  
25 solution was then rebuffered to pH 7.2. The ADC solution was concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and then optionally concentrated again to a concentration of about 10 mg/ml.

30 Other potentially hydrolysis-sensitive thianylsuccinimide bridges to the antibody in the working examples contain the following linker substructures, where #1 represents the thioether linkage to the antibody and #2 the point of attachment to the modified KSP inhibitor:

35



These linker substructures represent the linking unit to the antibody and have (in addition to the linker composition) a significant effect on the structure and the profile of the metabolites formed in the tumour cells.

In the structural formulae shown, AK<sub>1A</sub>, AK<sub>1B</sub>, AK<sub>1E</sub>, AK<sub>1I</sub>, AK<sub>1H</sub>, AK<sub>1K</sub> have the following meanings:

10

AK<sub>1A</sub> = cetuximab (partially reduced) - S<sup>1</sup>

AK<sub>1B</sub> = anti-TWEAKR AK-1 (partially reduced) - S<sup>1</sup>

15

AK<sub>1E</sub> = trastuzumab (partially reduced) - S<sup>1</sup>

AK<sub>1I</sub> = nimotuzumab (partially reduced) - S<sup>1</sup>

AK<sub>1H</sub> = panitumumab (partially reduced) - S<sup>1</sup>

20

where

§<sup>1</sup> represents the linkage to the succinimide group or to any isomeric hydrolysed open-chain succinamides or the alkylene radical resulting therefrom,

5

and

S represents the sulphur atom of a cysteine residue of the partially reduced antibody.

10

#### B-5. General process for coupling to lysine side chains

The following antibodies were used for the coupling reactions:

15 cetuximab (anti EGFR AK)

anti-TWEAKR AK 1 (TPP-2090)

trastuzumab (anti-Her2 AK)

20

nimotuzumab (anti-EGFR AK)

panitumumab (anti-EGFR AK)

25 From 2 to 8 equivalents of the precursor compound to be coupled were added as a solution in DMSO to a solution of the antibody in question in PBS buffer in a concentration range between 1 mg/ml and 20 mg/ml, preferably about 10 mg/ml, depending on the intended loading. After 30 min to 6 h of stirring at RT, the  
30 same amount of precursor compound in DMSO was added again. Here, the amount of DMSO should not exceed 10% of the total volume. After a further 30 min to 6 h of stirring at RT, the reaction was applied to PD 10 columns (Sephadex® G-25, GE Healthcare) equilibrated with PBS and eluted with PBS buffer. Generally,  
35 unless indicated otherwise, 5 mg of the antibody in question in PBS buffer were used for the reduction and the subsequent coupling. Purification on the PD10 column thus in each case afforded solutions of the respective ADCs in 3.5 ml PBS buffer.

The sample was then concentrated by ultracentrifugation and optionally rediluted with PBS buffer. If required, for better removal of low-molecular weight components, concentration by ultrafiltration was repeated after redilution with PBS buffer.

5 For biological tests, if required, the concentrations of the final ADC samples were optionally adjusted to the range of 0.5-15 mg/ml by redilution.

The respective protein concentrations, stated in the working examples, of the ADC solutions were determined. Furthermore, antibody loading (drug/mAb ratio) was determined using the methods described under B-7.

10

In the structural formulae shown, AK<sub>2A</sub>, AK<sub>2B</sub>, AK<sub>2G</sub>, AK<sub>2E</sub>, AK<sub>2I</sub>, AK<sub>2H</sub>, AK<sub>2K</sub> have the following meanings:

15

AK<sub>2A</sub> = cetuximab (anti-EGFR AK)-NH§<sup>2</sup>

AK<sub>2B</sub> = anti-TWEAKR AK-1- NH§<sup>2</sup>

20

AK<sub>2E</sub> = trastuzumab- NH§<sup>2</sup>

AK<sub>2I</sub> = nimotuzumab-NH§<sup>2</sup>

25 AK<sub>2H</sub> = panitumumab-NH§<sup>2</sup>

where

§<sup>2</sup> represents the linkage to the carbonyl group

30

and

NH represents the side-chain amino group of a lysine residue of the antibody.

35

B-6a. General process for preparing closed succinimide-cysteine adducts:

In an exemplary embodiment, 10  $\mu\text{mol}$  of the maleinimide precursor compounds described above were taken up in 3-5 ml of DMF, and 2.1 mg (20  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 2 h to 24 h, then dried under reduced pressure and then purified by preparative HPLC.

B-6aa. General process for preparing isomeric open succinamide-cysteine adducts:

In an exemplary embodiment, 68  $\mu\text{mol}$  of the maleinimide precursor compounds described above were taken up in 15 ml of DMF, and with 36 mg (136  $\mu\text{mol}$ ) of N- $\{[2-(\text{trimethylsilyl})\text{ethoxy}]\text{carbonyl}\}$ -L-cysteine were added. The reaction mixture was stirred at RT for ~20 h, then dried under reduced pressure and then purified by preparative HPLC. The appropriate fractions were combined and the solvents were evaporated under reduced pressure, and the residue was then dissolved in 15 ml of THF/water 1:1. 131  $\mu\text{l}$  of a 2M aqueous lithium hydroxide solution were added, and the reaction was stirred at RT for 1 h. The reaction was then neutralized with a 1M hydrochloric acid, the solvent was evaporated under reduced pressure and the residue was purified by preparative HPLC. This gave ~50% of theory of the regioisomeric protected intermediates as a colourless foam.

In the last step, 0.023 mmol of these regioisomeric hydrolysis products were dissolved in 3 ml of 2,2,2-trifluoroethanol. 12.5 mg (0.092 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 4 h. 27 mg (0.092 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were then added, and the solvent was evaporated under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave the hydrolysed open sulphanylsuccinamides as a regioisomer mixture.

B-6b. General process for preparing lysine adducts:

In an exemplary embodiment, 10  $\mu\text{mol}$  of the activated ester

precursor compounds described above were taken up in 3-5 ml of DMF, and  $\alpha$ -amino-protected L-lysine was added in the presence of 30  $\mu$ mol of *N,N*-diisopropylethylamine. The reaction mixture was stirred at RT for 2 h to 24 h, then dried under reduced pressure and then purified by preparative HPLC. The protective group was then removed by known methods.

Further purification and characterization of the conjugates according to the invention

10

After the reaction, in some instances the reaction mixture was concentrated, for example by ultrafiltration, and then desalted and purified by chromatography, for example using a Sephadex® G-25 column. Elution was carried out, for example, with phosphate-buffered saline (PBS). The solution was then sterile filtered and frozen. Alternatively, the conjugate can be lyophilized.

15

B-7. Determination of the antibody, the toxophor loading and the proportion of open cysteine adducts

20

For protein identification in addition to molecular weight determination after deglycosylation and/or denaturing, a tryptic digestion was carried out which, after denaturing, reduction and derivatization, confirms the identity of the protein via the tryptic peptides found.

25

The toxophor loading of the PBS buffer solutions obtained of the conjugates described in the working example was determined as follows:

30

Determination of toxophor loading of lysine-linked ADCs was carried out by mass spectrometric determination of the molecular weights of the individual conjugate species. Here, the antibody conjugates were first deglycosylated with PNGaseF, and the sample was acidified and, after HPLC separation/desalting, analysed by mass spectrometry using ESI-MicroTofQ (Bruker Daltonik). All spectra over the signal in the TIC (Total Ion

35

Chromatogram) were added and the molecular weight of the different conjugate species was calculated based on MaxEnt deconvolution. The DAR (= drug/antibody ratio) was then calculated after signal integration of the different species.

5

The toxophor loading of cysteine-linked conjugates was determined by reversed-phase chromatography of the reduced and denatured ADCs. Guanidinium hydrochloride (GuHCl) (28.6 mg) and a solution of DL-dithiothreitol (DTT) (500 mM, 3  $\mu$ l) were added  
10 to the ADC solution (1 mg/ml, 50  $\mu$ l). The mixture was incubated at 55°C for one hour and analysed by HPLC.

HPLC analysis was carried out on an Agilent 1260 HPLC system with detection at 220 nm. A Polymer Laboratories PLRP-S  
15 polymeric reversed-phase column (catalogue number PL1912-3802) (2.1 x150 mm, 8  $\mu$ m particle size, 1000 Å) was used at a flow rate of 1 ml/min with the following gradient: 0 min, 25%B; 3 min, 25%B; 28 min, 50%B. Mobile phase A consisted of 0.05% trifluoroacetic acid (TFA) in water, mobile phase B of 0.05%  
20 trifluoroacetic acid in acetonitrile.

The detected peaks were assigned by retention time comparison with the light chain (L0) and the heavy chain (H0) of the non-conjugated antibody. Peaks detected exclusively in the  
25 conjugated sample were assigned to the light chain with one toxophor (L1) and the heavy chains with one, two and three toxophors (H1, H2, H3).

Average loading of the antibody with toxophors was calculated  
30 from the peak areas determined by integration as double the sum of the toxophor number weighed integration results of all peaks divided by the sum of the singly weighed integration results of all peaks. In individual cases, it may be possible that, owing to co-elution of some peaks, it is not possible to determine  
35 toxophor loading accurately.

In the cases where light and heavy chains could not be separated sufficiently by HPLC, determination of toxophor loading of

cysteine-linked conjugates was carried out by mass spectrometric determination of the molecular weights of the individual conjugate species at light and heavy chain.

5 Guanidinium hydrochloride (GuHCl) (28.6 mg) and a solution of DL-dithiothreitol (DTT) (500 mM, 3  $\mu$ l) were added to the ADC solution (1 mg/ml, 50  $\mu$ l). The mixture was incubated for one hour at 55°C and analysed by mass spectrometry after online desalting using ESI-MicroTof<sub>0</sub> (Bruker Daltonik).

10

For the DAR determination, all spectra were added over the signal in the TIC (Total Ion Chromatogram), and the molecular weight of the different conjugate species at light and heavy chain was calculated based on MaxEnt deconvolution. The average loading of the antibody with toxophores was calculated by integration of certain molecular weight areas as double the sum of the toxophor number weighed integration results of all peaks divided by the sum of the singly weighed integration results of all peaks.

20

To determine the proportion of the open cysteine adduct, the molecular weight area ratio of closed to open cysteine adduct (molecular weight delta 18 Dalton) of all singly conjugated light and heavy chains was determined. The mean of all variants yielded the proportion of the open cysteine adduct.

25

#### B-8. Checking the antigen-binding of the ADCs

The capability of the binder of binding to the target molecule was checked after coupling had taken place. The person skilled in the art is familiar with multifarious methods which can be used for this purpose; for example, the affinity of the conjugate can be checked using ELISA technology or surface plasmon resonance analysis (BIAcore™ measurement). The conjugate concentration can be measured by the person skilled in the art using customary methods, for example for antibody conjugates by protein determination. (see also Doronina et al.; Nature Biotechnol. 2003; 21:778-784 and Polson et al., Blood 2007;

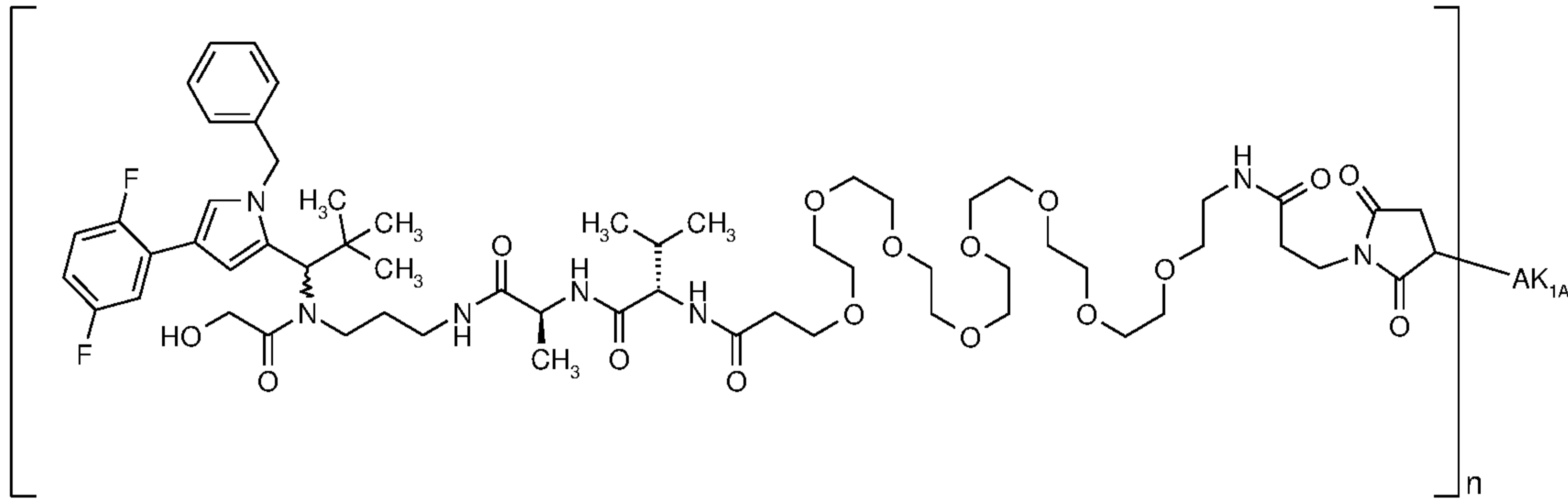
35



1102:616-623).

### Working Examples ADCs

#### 5 Example 82A



10 Here, 5.0 mg of cetuximab in PBS (c=5.90 mg/ml) were used for coupling with Intermediate F82, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 1.59 mg/ml

Drug/mAb ratio: 2.3

#### 20 Example 82B

25 Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=10.10 mg/ml) were used for coupling with Intermediate F82, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

30

Protein concentration: 1.59 mg/ml

Drug/mAb ratio: 1.8

**Example 82E**

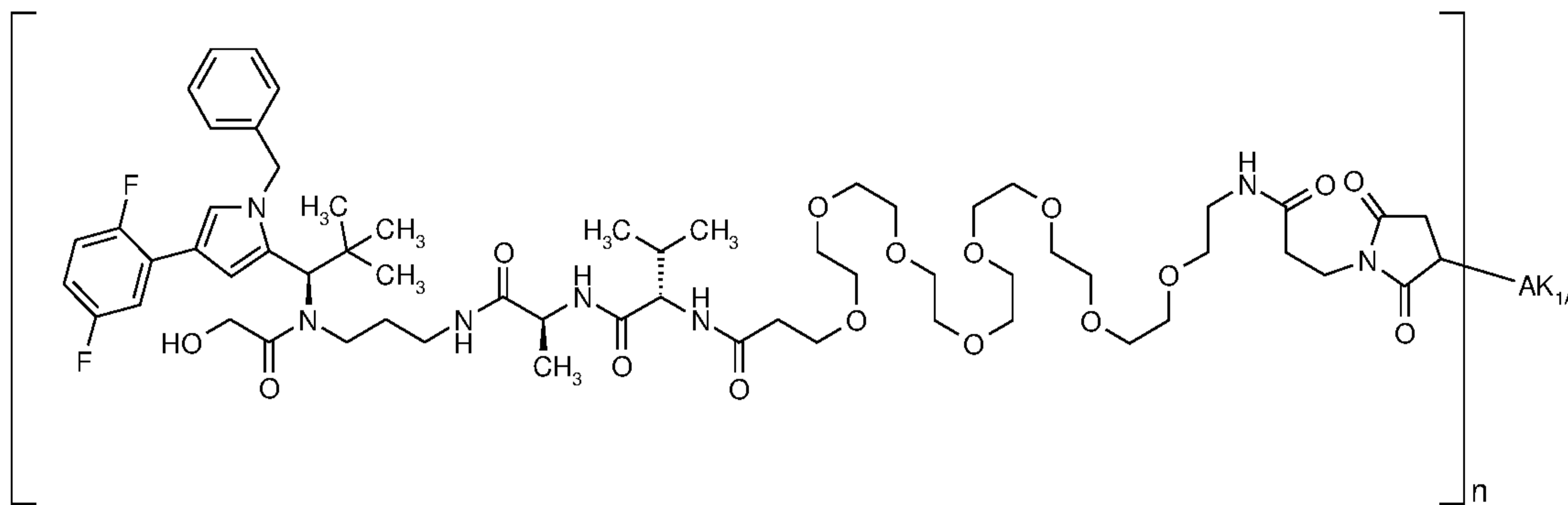
Here, 5.0 mg of trastuzumab antibody in PBS (c=11.50 mg/ml) were used for coupling with Intermediate F82, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 2.15 mg/ml

Drug/mAb ratio: 2.7

**Example 83A**

15



Here, 5.0 mg of cetuximab in PBS (c=5.90 mg/ml) were used for coupling with Intermediate F83, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

25 Protein concentration: 1.77 mg/ml

Drug/mAb ratio: 2.0

**Example 83B**

30

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=12.87 mg/ml)

were used for coupling with Intermediate F83, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides  
5 attached to the antibody.

Protein concentration: 1.29 mg/ml

Drug/mAb ratio: 1.3

10

#### Example 83E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F83, and the reaction was,  
15 after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

20 Protein concentration: 1.91 mg/ml

Drug/mAb ratio: 2.1

#### Example 83H

25

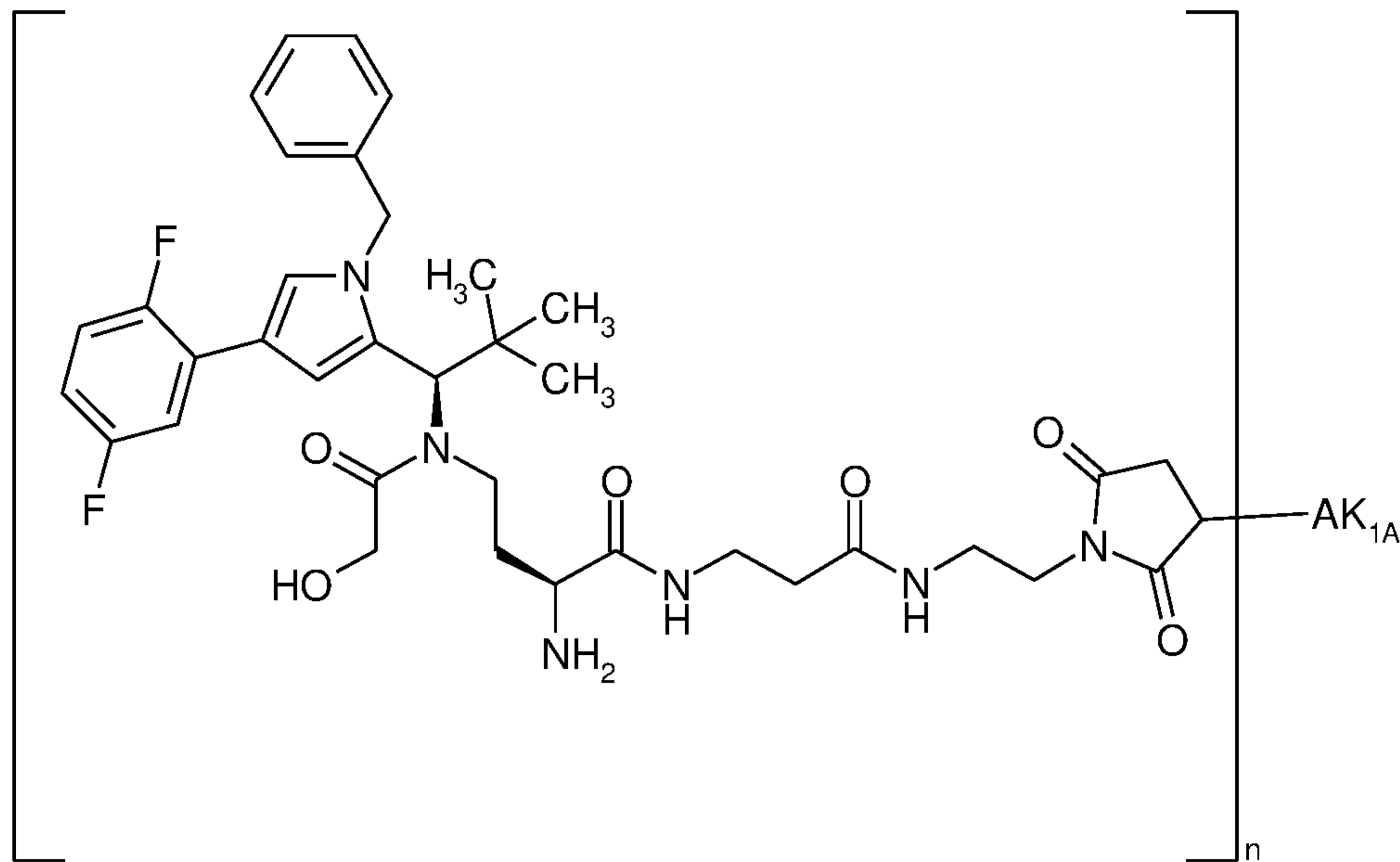
Here, 5.0 mg of panitumumab in PBS (c = 10 mg/ml) were used for coupling with Intermediate F83. The time for the reduction with TCEP was increased to 4 h and stirring time for the ADC coupling was increased to 20 h. The reaction was then, after Sephadex  
30 purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.57 mg/ml

35

Drug/mAb ratio: 0.9

#### Example 84A



Here, 5 mg of cetuximab in PBS (c=11.02 mg/ml) were used for  
 5 coupling with Intermediate F84, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS. Some of the ADC may also be present in the  
 form of the hydrolysed open-chain succinamides attached to the  
 antibody.

10

Protein concentration: 1.94 mg/ml

Drug/mAb ratio: 2.9

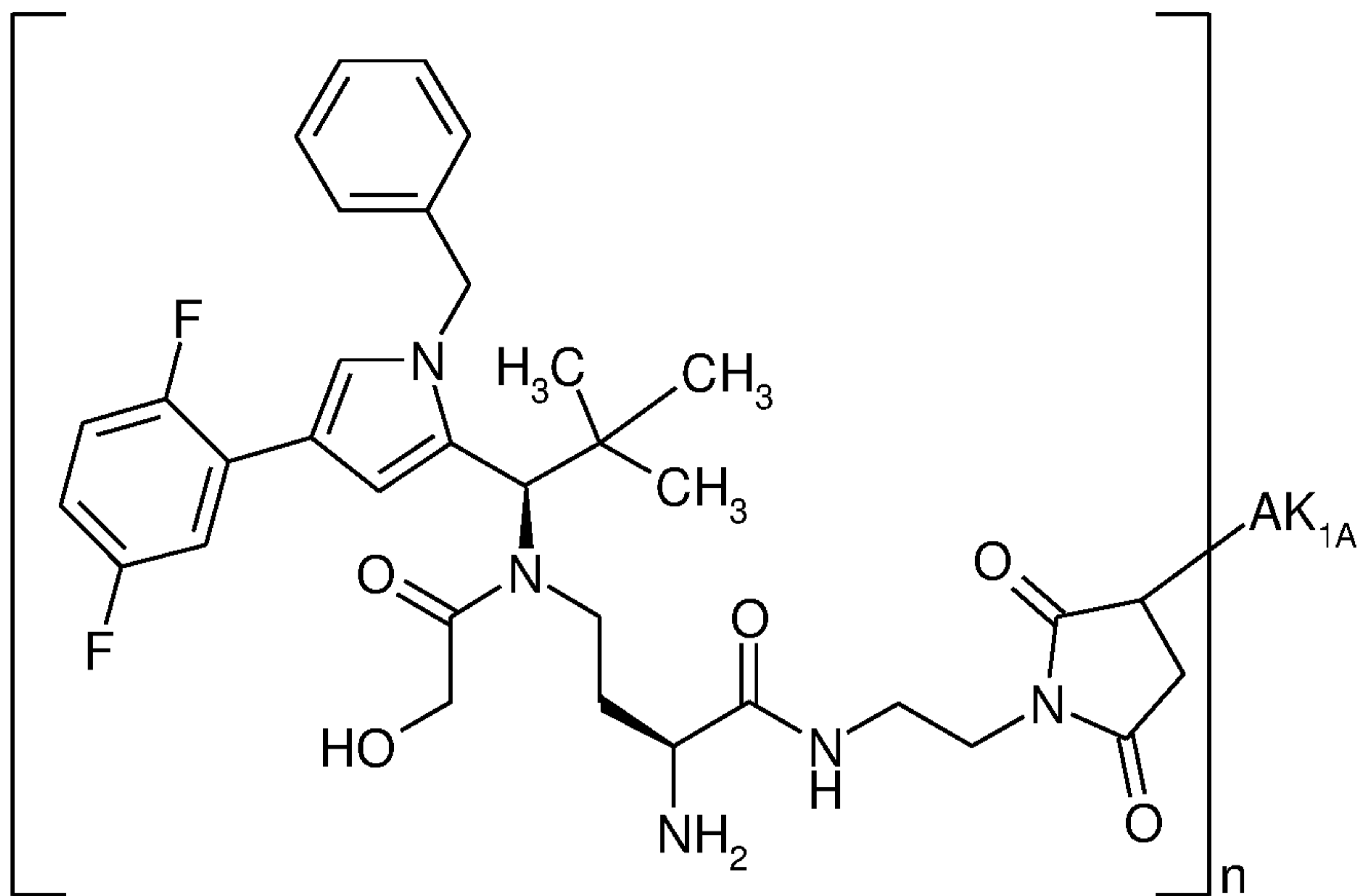
### 15 Example 84B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used  
 for coupling with Intermediate F84, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 20 rediluted with PBS. Some of the ADC may also be present in the  
 form of the hydrolysed open-chain succinamides attached to the  
 antibody.

25

Protein concentration: 1.77 mg/ml

Drug/mAb ratio: 3.0

Example 85A

5

Here, 5 mg of cetuximab in PBS (c=11.02 mg/ml) were used for coupling with Intermediate F85, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10

Protein concentration: 2.13 mg/ml

15 Drug/mAb ratio: 3.4

Example 85B

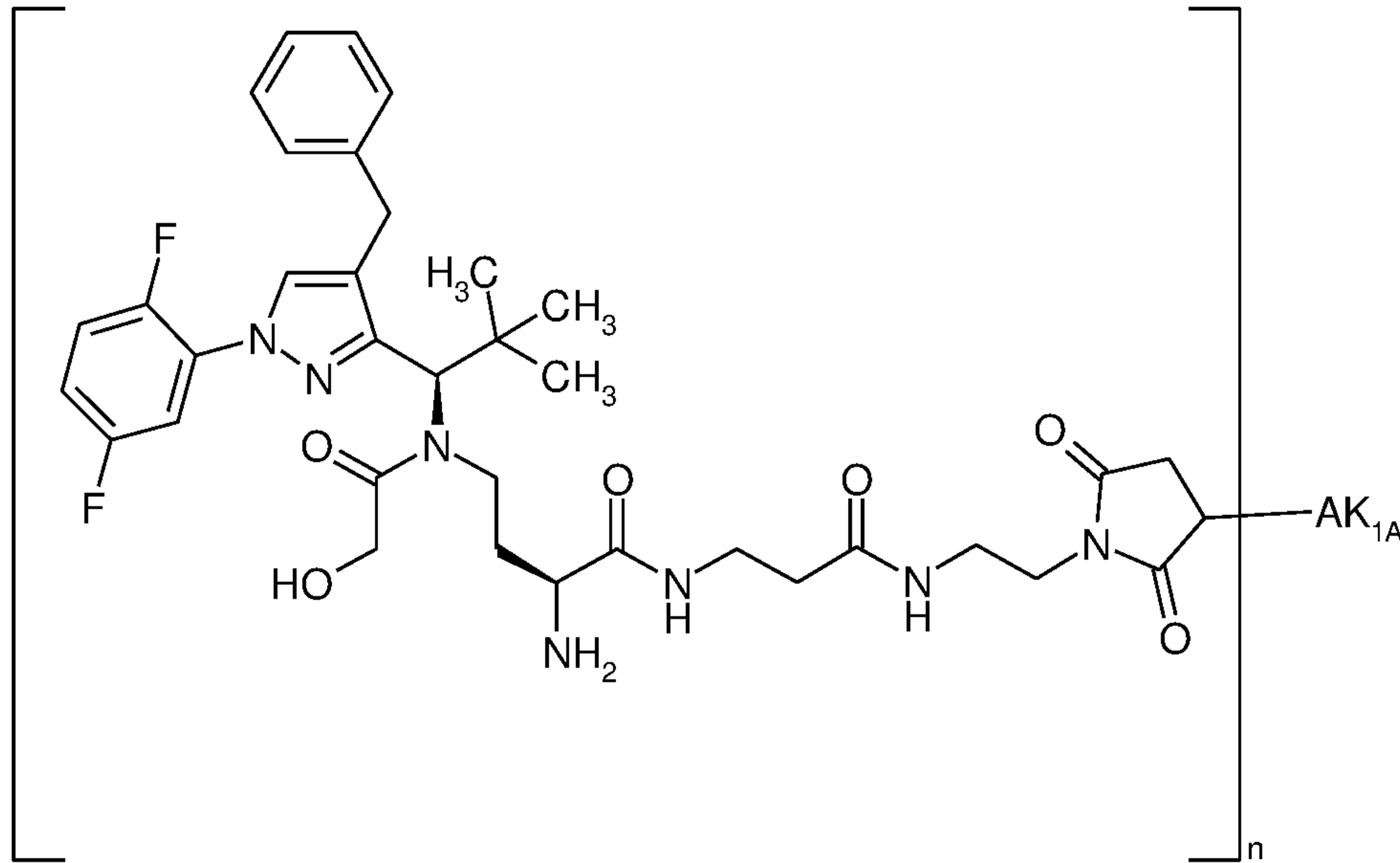
Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F85, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

25

Protein concentration: 1.63 mg/ml

Drug/mAb ratio: 3.2

**Example 86A**



5

Here, 5 mg of cetuximab in PBS (c=11.59 mg/ml) were used for coupling with Intermediate F86, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.96 mg/ml

15

Drug/mAb ratio: 3.2

**Example 86B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F86, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

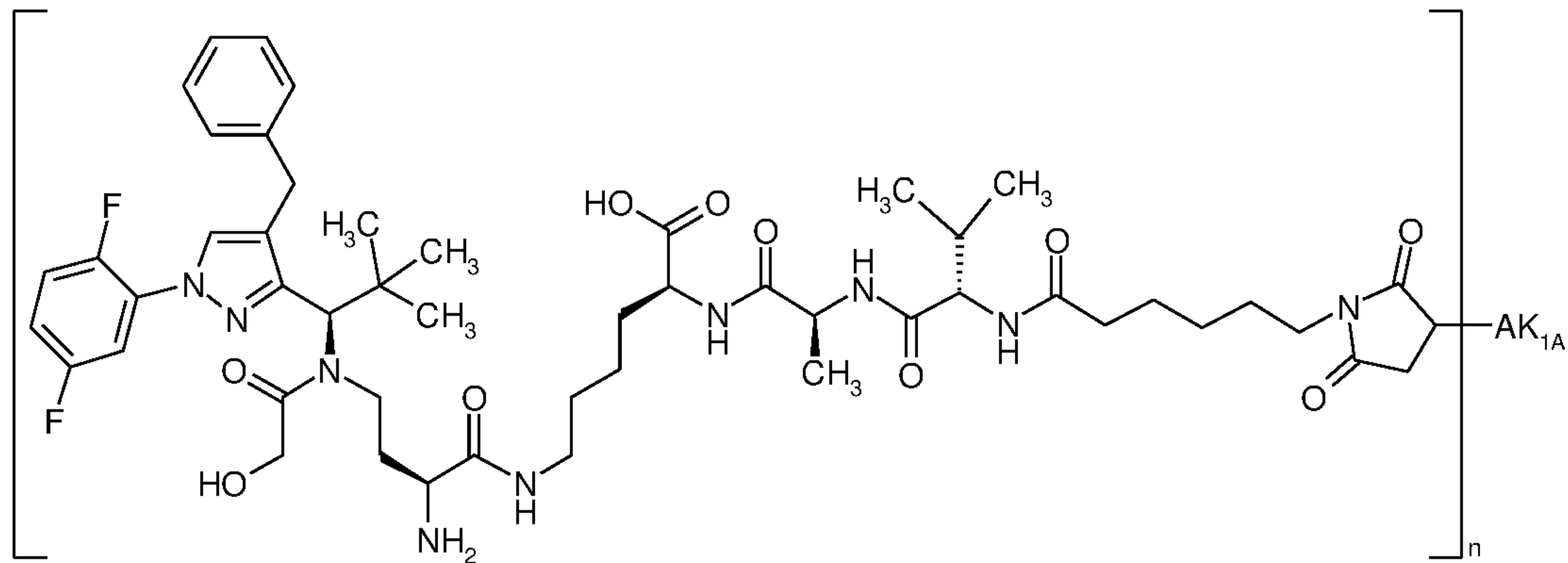
25



Drug/mAb ratio: 2.2

**Example 88A**

5



Here, 5 mg of cetuximab in PBS (c=11.02 mg/ml) were used for coupling with Intermediate F88, and the reaction was, after  
 10 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.03 mg/ml

15 Drug/mAb ratio: 3.1

**Example 88B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used  
 20 for coupling with Intermediate F88, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

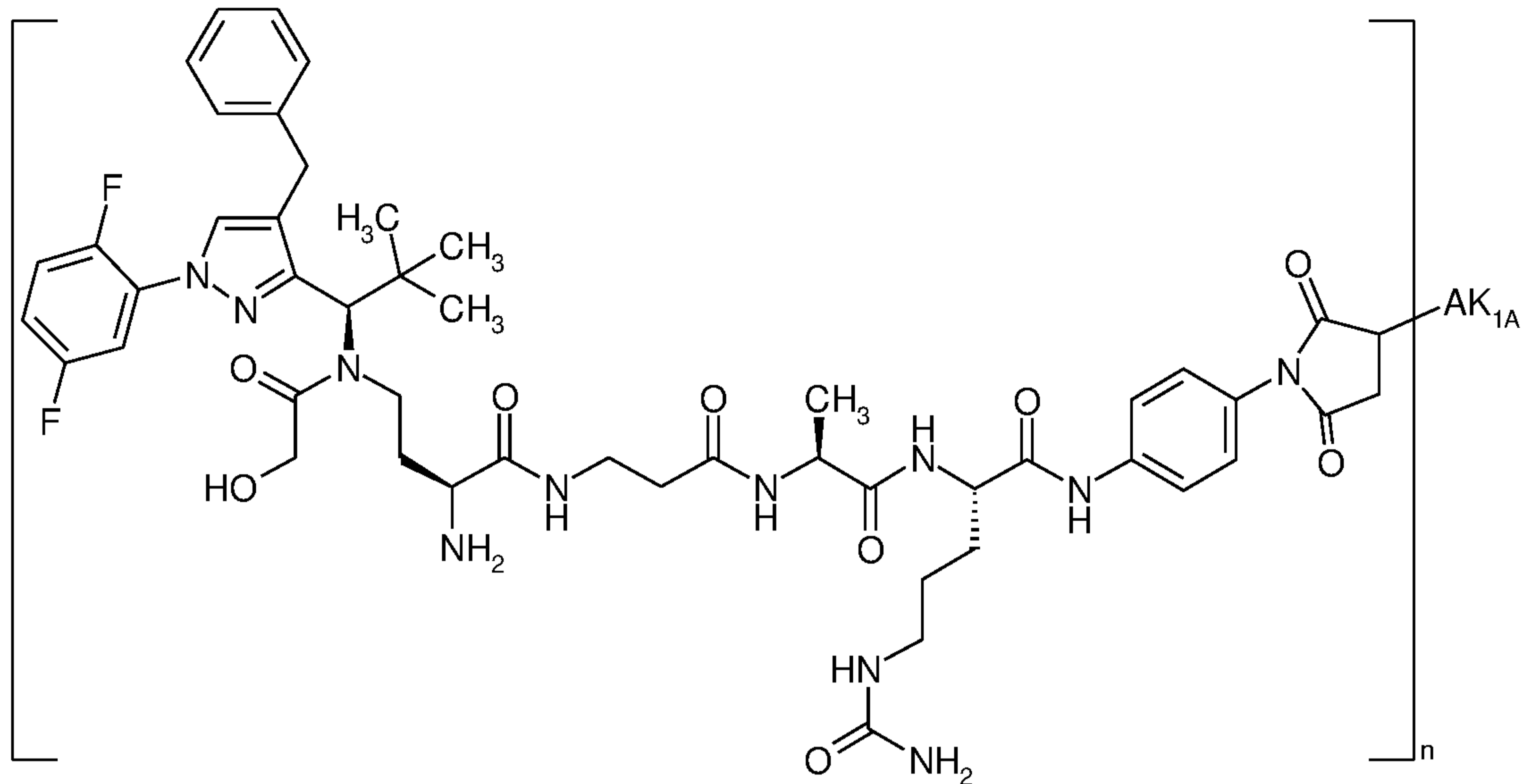
Protein concentration: 1.9 mg/ml

25

Drug/mAb ratio: 3.3

**Example 89A**





Here, 5 mg of cetuximab in PBS (c=11.02 mg/ml) were used for coupling with Intermediate F89, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 2.2 mg/ml

Drug/mAb ratio: 3.3

#### Example 89B

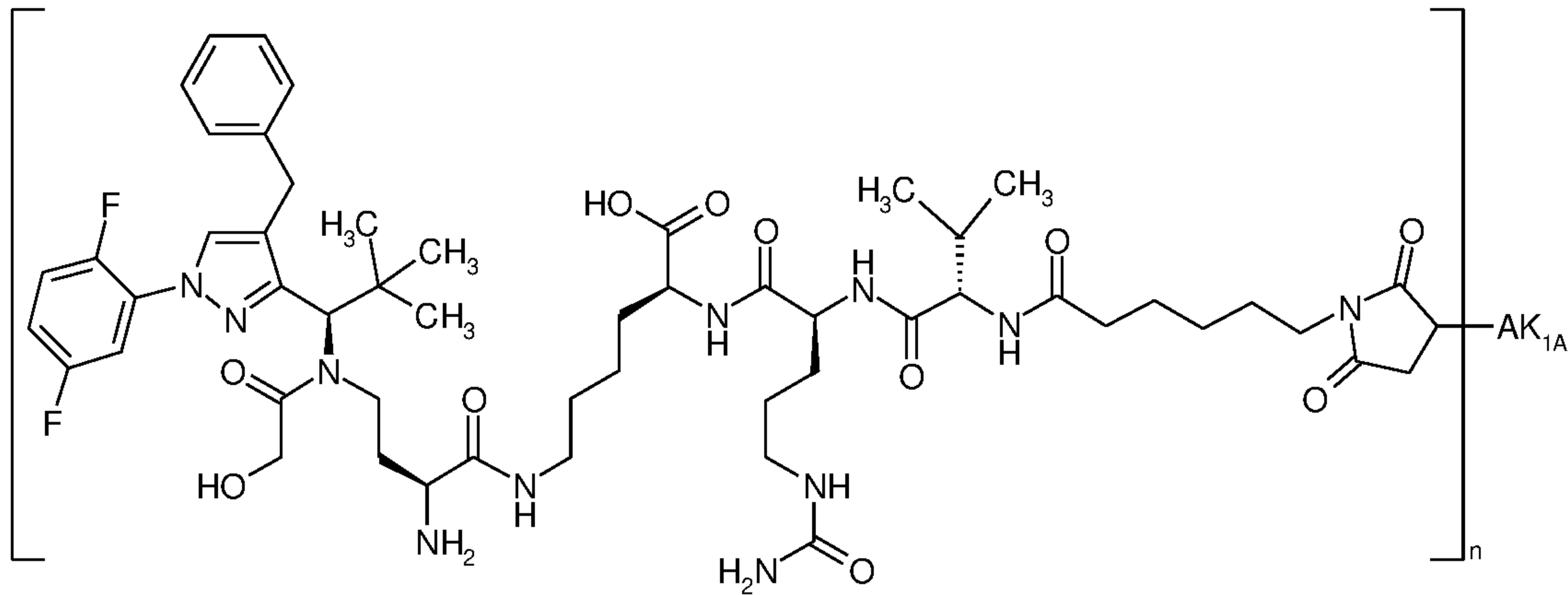
15

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F89, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the  
 20 form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 2.03 mg/ml

25 Drug/mAb ratio: 3.4

#### Example 90A



Here, 5 mg of cetuximab in PBS (c=13.33 mg/ml) were used for  
 5 coupling with Intermediate F90, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

Protein concentration: 2.19 mg/ml

10

Drug/mAb ratio: 3.0

#### Example 90B

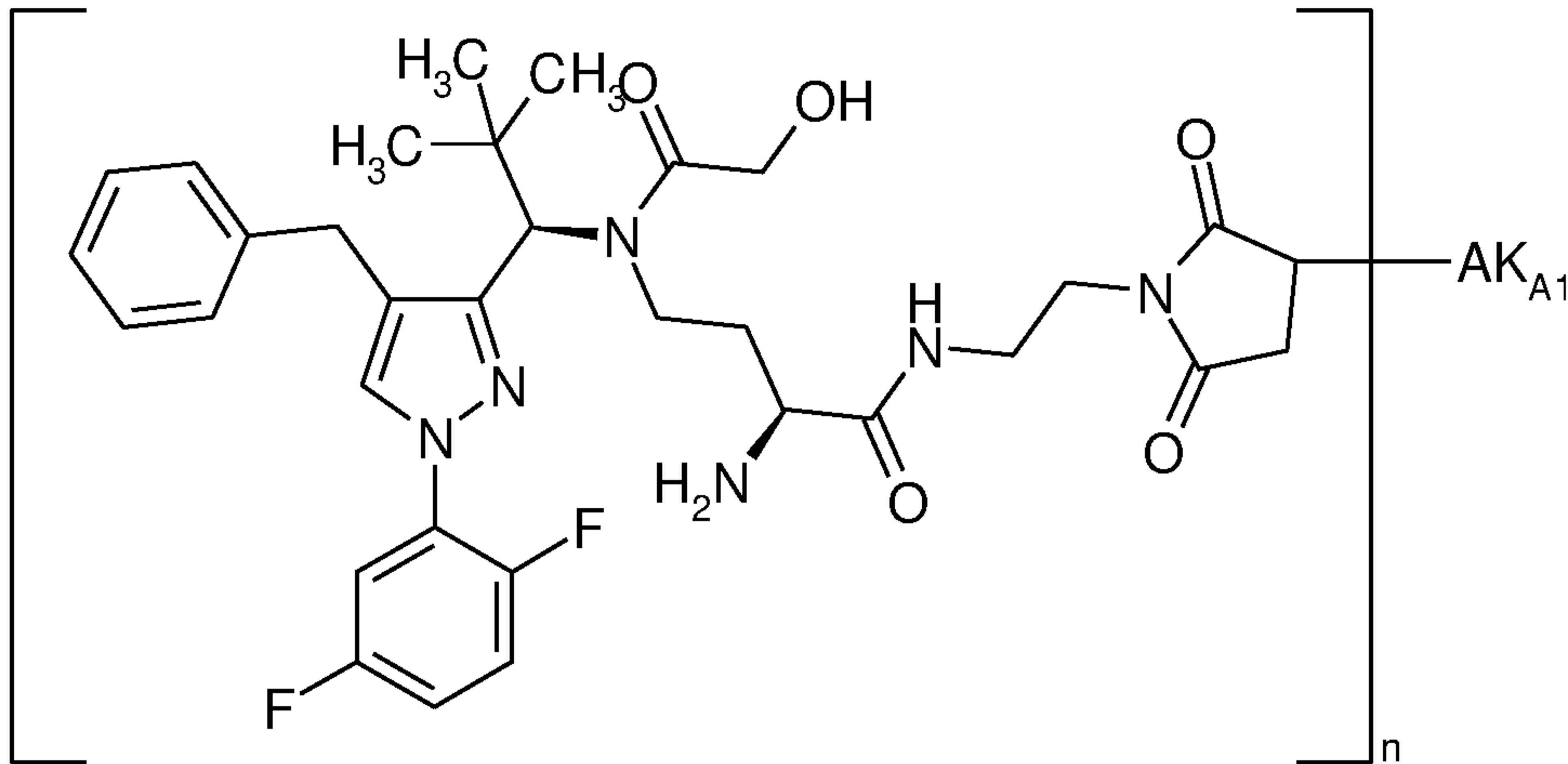
15 Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used  
 for coupling with Intermediate F90, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

20 Protein concentration: 1.97 mg/ml

Drug/mAb ratio: 2.9

#### Example 91A

25



Here, 80 mg of cetuximab in PBS (c=5.9 mg/ml) were used for coupling with Intermediate F91, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 12.75 mg/ml

Drug/mAb ratio: 3.7

### Example 91B

15

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F91, and the reaction was, after  
 20 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 5.71 mg/ml

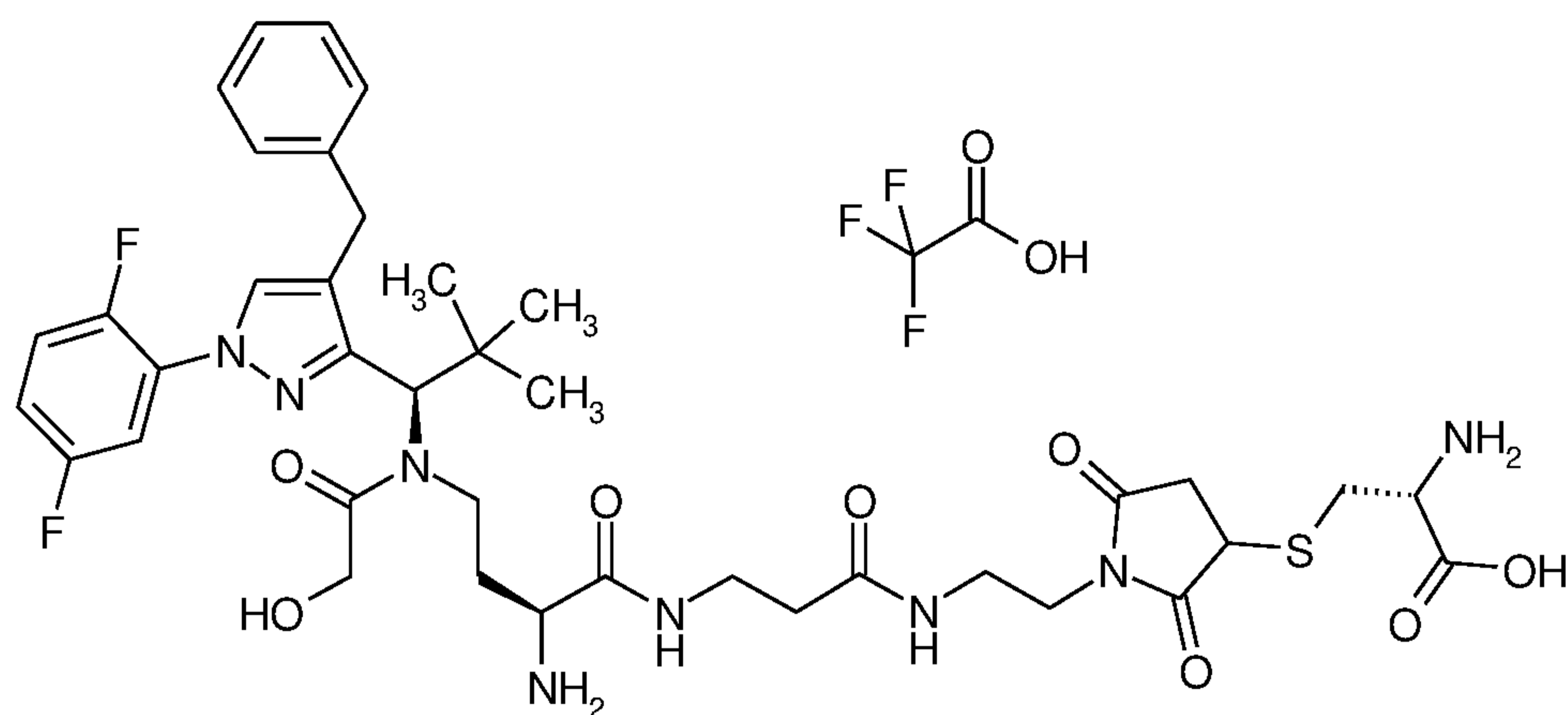
25 Drug/mAb ratio: 4.0

### Example 92

S-(1-{2-[ (N-{ (2S)-2-Amino-4- [ { (1R)-1-[4-benzyl-1-(2,5-

difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)amino]ethyl}-2,5-dioxopyrrolidin-3-yl)-L-cysteine trifluoroacetic acid (1:1)

5



3 mg (4  $\mu\text{mol}$ ) of Intermediate F86 were taken up in 3 ml of DCM/water 10:1, and 1.3 mg (11  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 10 min, then concentrated under reduced pressure and then purified by preparative HPLC.

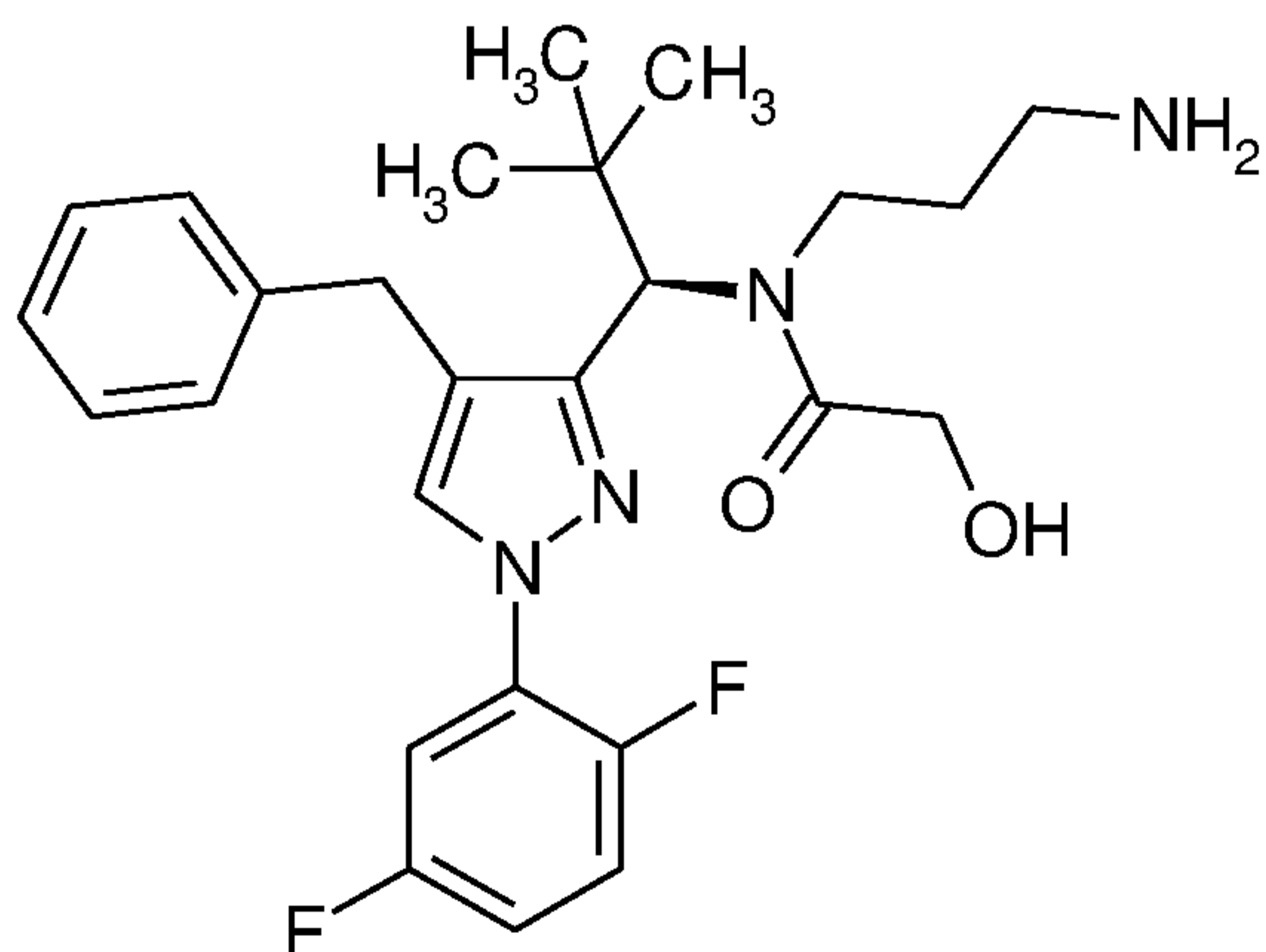
LC-MS (Method 1):  $R_t = 0.77$  min; MS (EIpos):  $m/z = 829$   $[\text{M}+\text{H}]^+$ .

15

### Example 96

N-(3-Aminopropyl)-N-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)-2-hydroxyacetamide

20

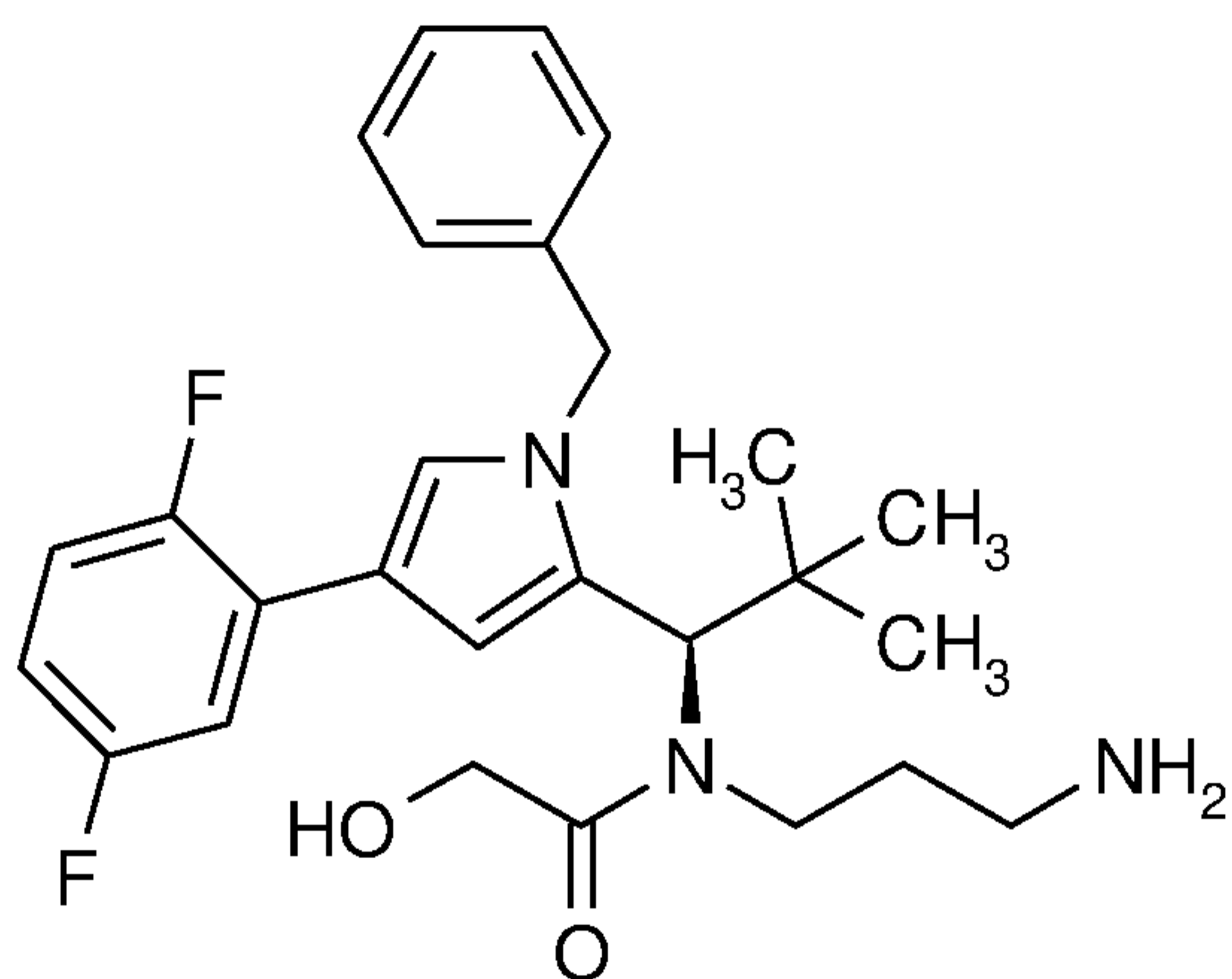


101 mg (0.16 mmol) of 2-({(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}[3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propyl]amino)-2-oxoethyl acetate were initially charged in 2 ml of absolute ethanol, and  
5 244 mg (3.14 mmol, 225  $\mu$ l) of a 40% strength solution of methylamine in water were added. The mixture was stirred at 50°C for 1 h, another 244 mg (3.14 mmol, 225  $\mu$ l) of a 40% strength solution of methylamine in water were then added and after a total of 3.5 h the mixture was purified directly by preparative  
10 HPLC (mobile phase: ACN/water + 1.0% NEt<sub>3</sub>, gradient). This gave 52 mg (70% of theory) of the target compound.

LC-MS (Method 3): R<sub>t</sub> = 2.56 min; MS (EIpos): m/z = 471 [M+H]<sup>+</sup>.

### 15 Example 98

N-(3-Aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide



150.0 mg (0.42 mmol) of (1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropan-1-amine (Intermediate C52) were initially charged in 2.0 ml of dichloromethane, and 29.2 mg (0.49 mmol) of HOAc and 125.6 mg (0.59 mmol) of sodium triacetoxyborohydride were added and the  
25 mixture was stirred at RT for 5 min. 98.9 mg (0.49 mmol) of 3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propanal were added. The reaction mixture was stirred at RT overnight. The reaction  
30 mixture was then diluted with ethyl acetate and the organic

phase was washed twice with saturated sodium carbonate solution and once with saturated NaCl solution. After drying over magnesium sulphate, the solvent was evaporated under reduced pressure and the residue was purified using silica gel (mobile  
5 phase: dichloromethane/methanol 100:1). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 188.6 mg (74%) of the compound 2-[3-({(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)propyl]-1H-isoindole-1,3(2H)-dione.

10

LC-MS (Method 1):  $R_t = 1.00$  min; MS (ESIpos):  $m/z = 541$  [M+H]<sup>+</sup>.

171.2 mg (0.32 mmol) of 2-[3-({(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)propyl]-1H-isoindole-1,3(2H)-dione were  
15 initially charged in 5.0 ml of dichloromethane, and 73.6 mg (0.73 mmol) of triethylamine were added. At 0°C, 94.9 mg (0.70 mmol) of acetoxyacetyl chloride were added, and the reaction mixture was stirred at RT overnight. The reaction mixture was  
20 diluted with ethyl acetate and the organic phase was washed twice with saturated sodium bicarbonate solution and once with sat. NaCl solution. After drying over magnesium sulphate, the solvent was evaporated under reduced pressure and the residue was purified using Biotage Isolera (silica gel, column 10 g  
25 SNAP, flow rate 12 ml/min, ethyl acetate/cyclohexane 1:3). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 159.0 mg (77%) of the compound 2-({(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propyl]amino)-2-oxoethyl acetate.  
30

LC-MS (Method 1):  $R_t = 1.35$  min; MS (ESIpos):  $m/z = 642$  [M+H]<sup>+</sup>.

147.2 mg (0.23 mmol) of 2-({(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[3-(1,3-dioxo-1,3-dihydro-2H-isoindol-2-yl)propyl]amino)-2-oxoethyl acetate were initially charged in 4.0 ml of ethanol, and 356.2 mg (4.59 mmol) of methanamine (40% in water) were added. The  
35

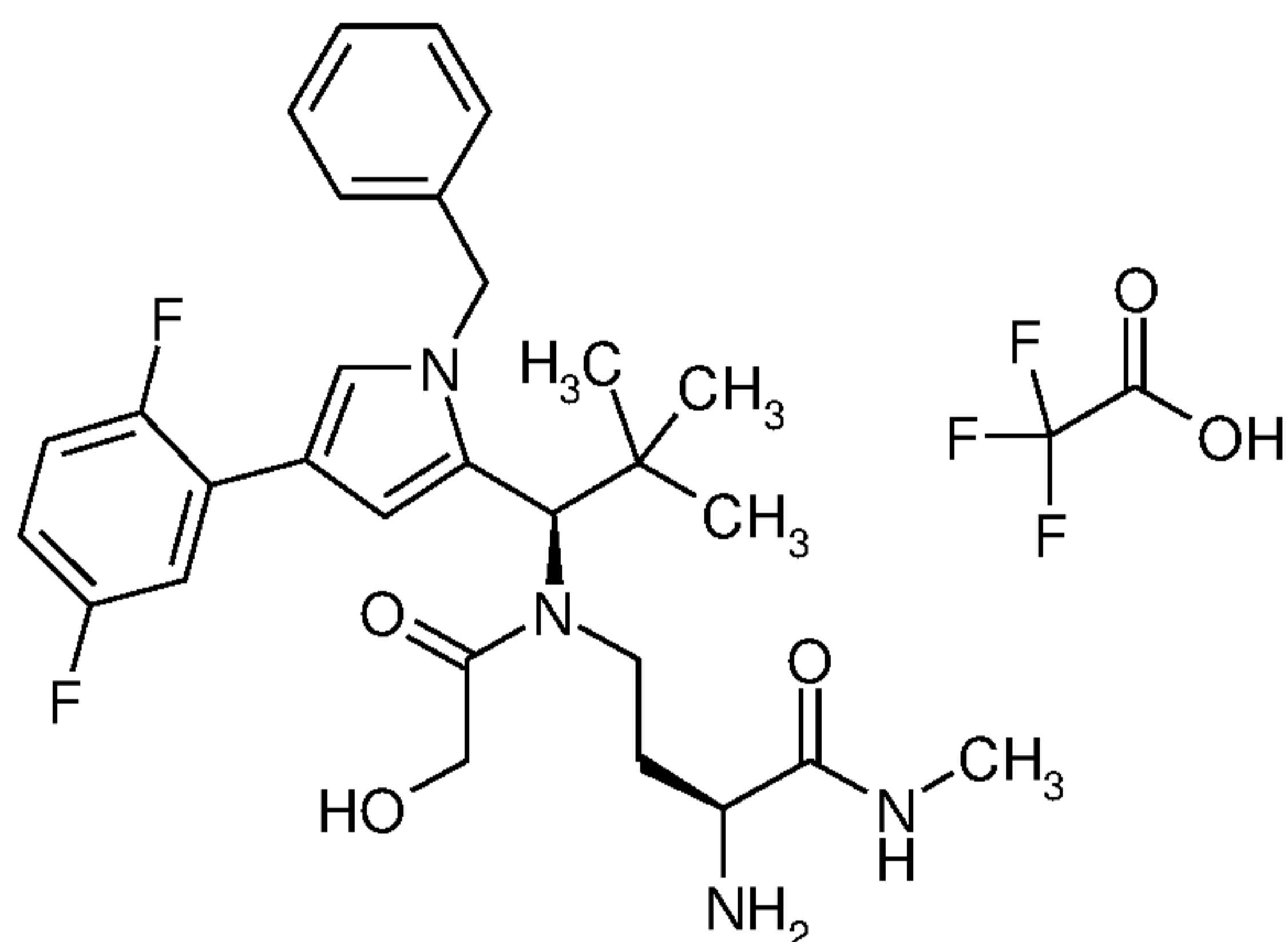
reaction mixture was stirred at 50°C overnight. The solvent was evaporated under reduced pressure and the residue was co-distilled with toluene three times. The residue was purified using silica gel (mobile phase: dichloromethane/methanol 10:1).  
 5 The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 67.4 mg (63%) of the title compound.

LC-MS (Method 1):  $R_t = 0.91$  min; MS (ESIpos):  $m/z = 470$   $[M+H]^+$ .

10

### Example 99

Trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-methylbutanamide (1:1)  
 15



First, Intermediate C52 was reductively alkylated with benzyl  
 20 (2S)-2-[[ (benzyloxy) carbonyl] amino]-4-oxobutanoate analogously to C2. The secondary amino group was then acylated with 2-chloro-2-oxoethyl acetate as described in Intermediate C27.

190 mg (0.244 mmol) of this intermediate were taken up in 7.5  
 25 ml of ethanol, and 0.35 ml of a 40% strength solution of methanamine in water was added. The reaction was stirred at 50°C for 3 h, and the same amount of methanamine was then added again. After a further 5 h of stirring at 50°C, the reaction was concentrated and the residue was purified by preparative HPLC.  
 30 This gave 78 mg (48% of theory) of this title compound.

LC-MS (Method 1):  $R_t = 1.32$  min; MS (EIpos):  $m/z = 661$   $[M+H]^+$ .

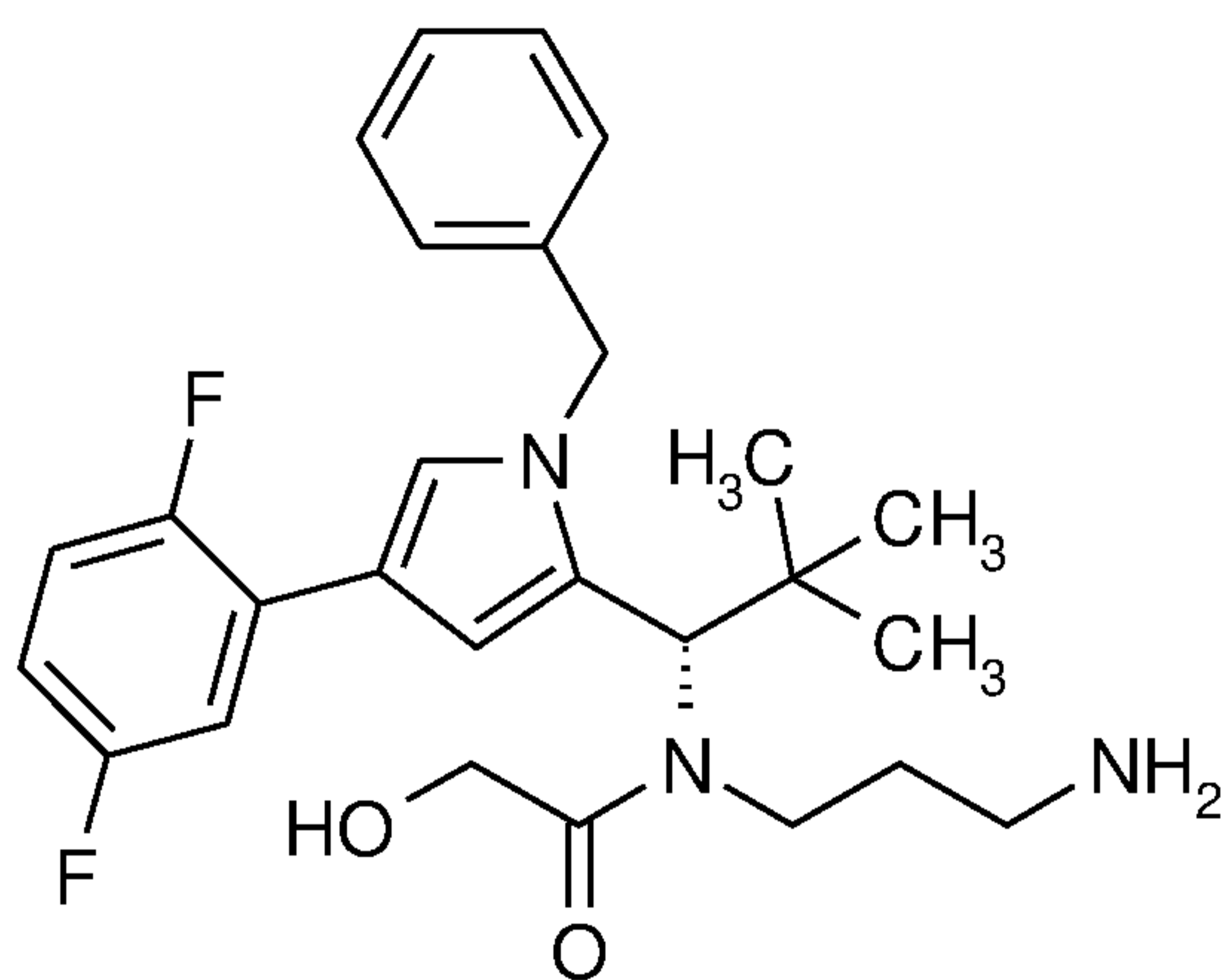
78 mg (0.118 mmol) of this intermediate were dissolved in 8 ml of ethanol and, after addition of 15 mg of 10% palladium on activated carbon, hydrogenated under standard hydrogen pressure at RT for 3 min. The catalyst was then filtered off, the solvent was removed under reduced pressure and the product was purified by preparative HPLC. After lyophilization from acetonitrile/water, 33 mg (44% of theory) of the title compound were obtained.

LC-MS (Method 1):  $R_t = 0.88$  min; MS (ESIpos):  $m/z = 527$   $(M+H)^+$ .

$^1H$  NMR (500 MHz, DMSO- $d_6$ ):  $\delta = 8.1$  (m, 1H), 8.0 (m, 3H), 7.9 (m, 1H), 7.65 (m, 1H), 7.5 (s, 1H), 7.15-7.35 (m, 5H) 7.0 (m, 1H), 6.85 (m, 1H), 5.6 (s, 1H), 4.9 and 5.2 (2d, 2H), 4.02 and 4.22 (2d, 2H), 3.2-3.5 (m, 6H), 0.7 and 1.46 (2m, 2H), 0.8 (s, 9H).

### Example 102

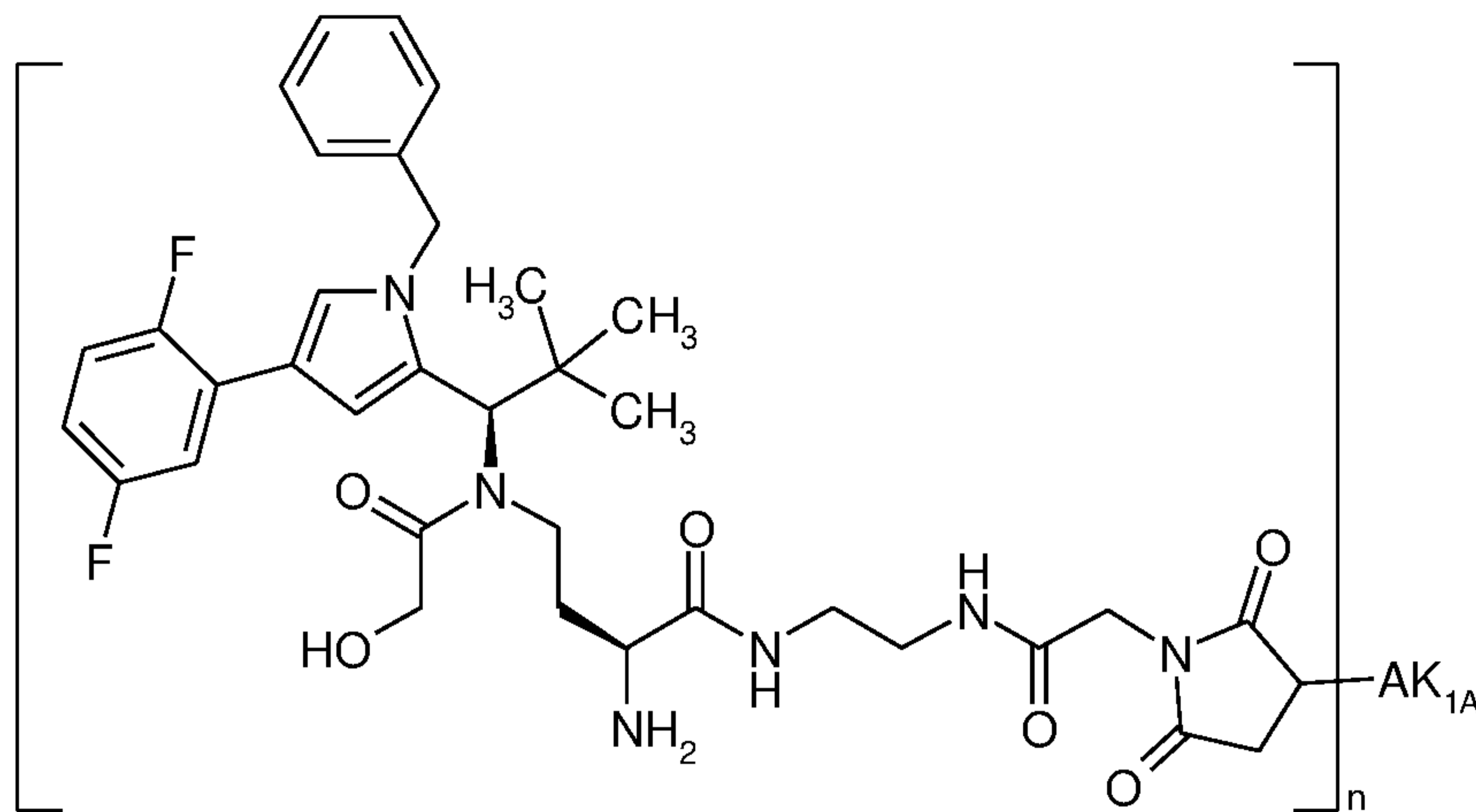
N-(3-Aminopropyl)-N-{(1S)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxyacetamide



The synthesis was carried out analogously to the synthesis of Example 98 using the corresponding S-isomer intermediate.

LC-MS (Method 1):  $R_t = 0.92$  min; MS (ESIpos):  $m/z = 470$   $[M+H]^+$ .



**Example 104A**

5 Here, 5 mg of cetuximab in PBS (c=15.33 mg/ml) were used for  
 coupling with Intermediate F104, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS. Some of the ADC may also be present in the  
 form of the hydrolysed open-chain succinamides attached to the  
 10 antibody.

Protein concentration: 1.95 mg/ml

Drug/mAb ratio: 3.7

15

**Example 104B**

Here, 35 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used  
 for coupling with Intermediate F104, and the reaction was, after  
 20 Sephadex purification, concentrated by ultracentrifugation,  
 rediluted with PBS and concentrated again. Some of the ADC may  
 also be present in the form of the hydrolysed open-chain  
 succinamides attached to the antibody.

25 Protein concentration: 10.93 mg/ml

Drug/mAb ratio: 3.2

**Example 104E**

Here, 5.0 mg of trastuzumab antibody in PBS (c=8.23 mg/ml) were used for coupling with Intermediate F104, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 2.11 mg/ml

Drug/mAb ratio: 2.8

#### Example 104I

Here, 5.0 mg of nimotuzumab in PBS (c=13.1 mg/ml) were used for coupling with Intermediate F104, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.89 mg/ml

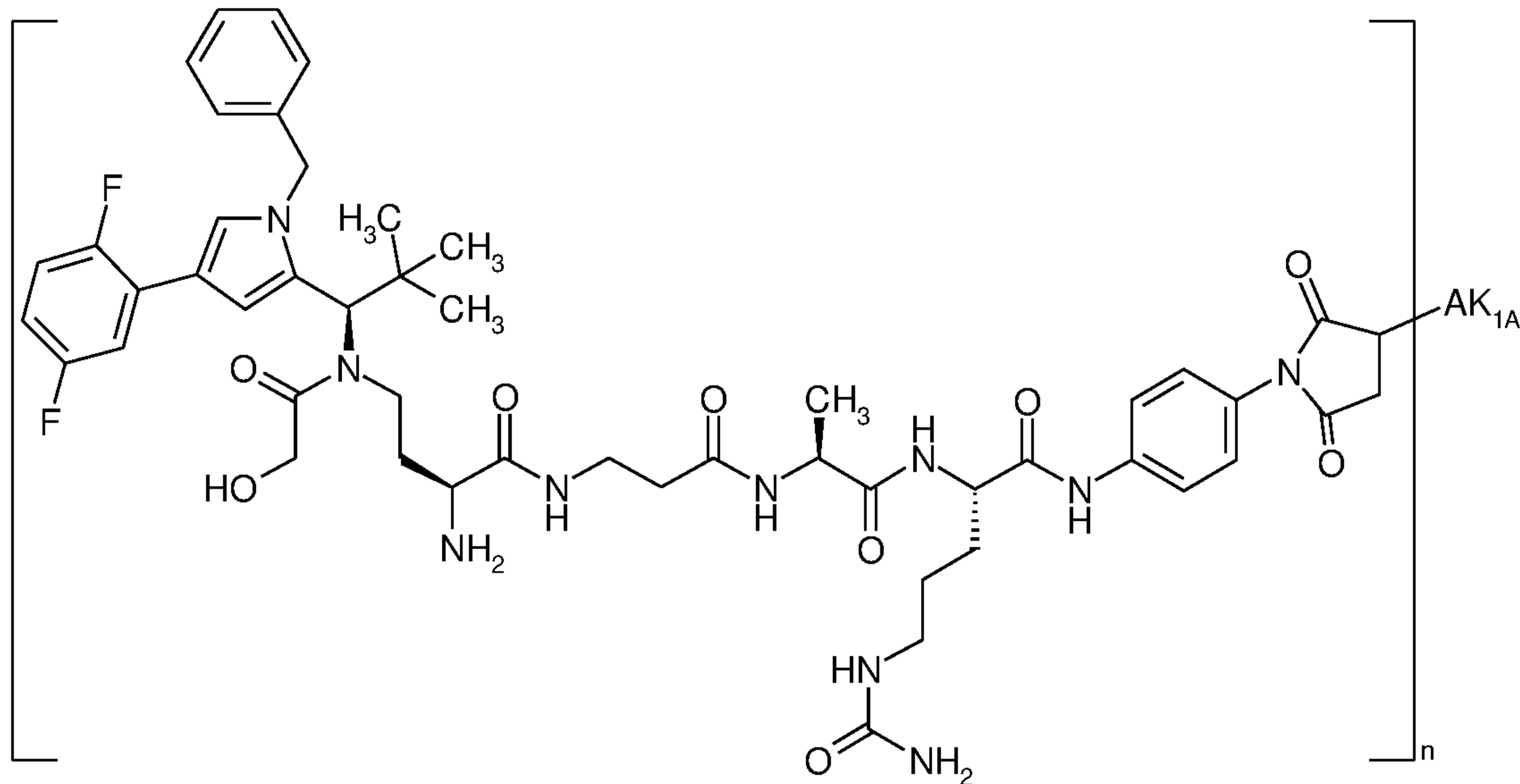
Drug/mAb ratio: 3.5

#### Example 104H

Here, 5.0 mg of panitumumab in PBS (c = 12 mg/ml) were used for coupling with Intermediate F104. The time for the reduction with TCEP was increased to 4 h and stirring time for the ADC coupling was increased to 20 h. The reaction was then, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.79 mg/ml

Drug/mAb ratio: 2.2

**Example 106A**

5

Here, 5 mg of cetuximab in PBS ( $c=15.33$  mg/ml) were used for coupling with Intermediate F106, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.91 mg/ml

15 Drug/mAb ratio: 3.3

**Example 106B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS ( $c=12.87$  mg/ml) were used for coupling with Intermediate F106, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

25

Protein concentration: 1.76 mg/ml

Drug/mAb ratio: 3.0

**Example 106E**

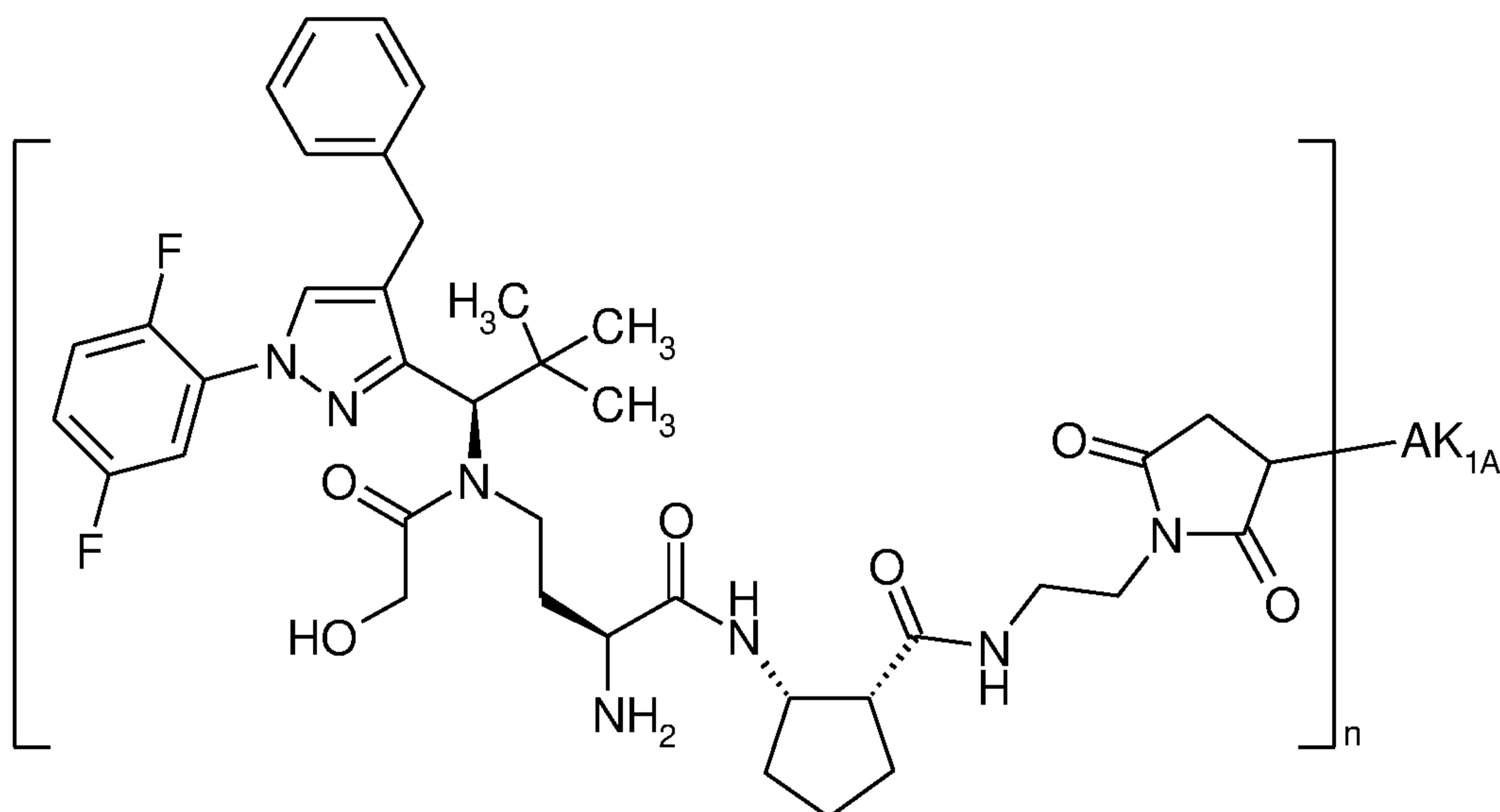
5 Here, 5 mg of trastuzumab in PBS (c=8.23 mg/ml) were used for coupling with Intermediate F106, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the  
10 antibody.

Protein concentration: 2.5 mg/ml

Drug/mAb ratio: 2.4

15

**Example 107A**



20 Here, 5 mg of cetuximab in PBS (c=12.3 mg/ml) were used for coupling with Intermediate F107, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the  
25 antibody.

Protein concentration: 2.16 mg/ml

Drug/mAb ratio: 3.3

**Example 107B**

5

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F107, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the  
10 form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.9 mg/ml

15 Drug/mAb ratio: 3.1

**Example 107E**

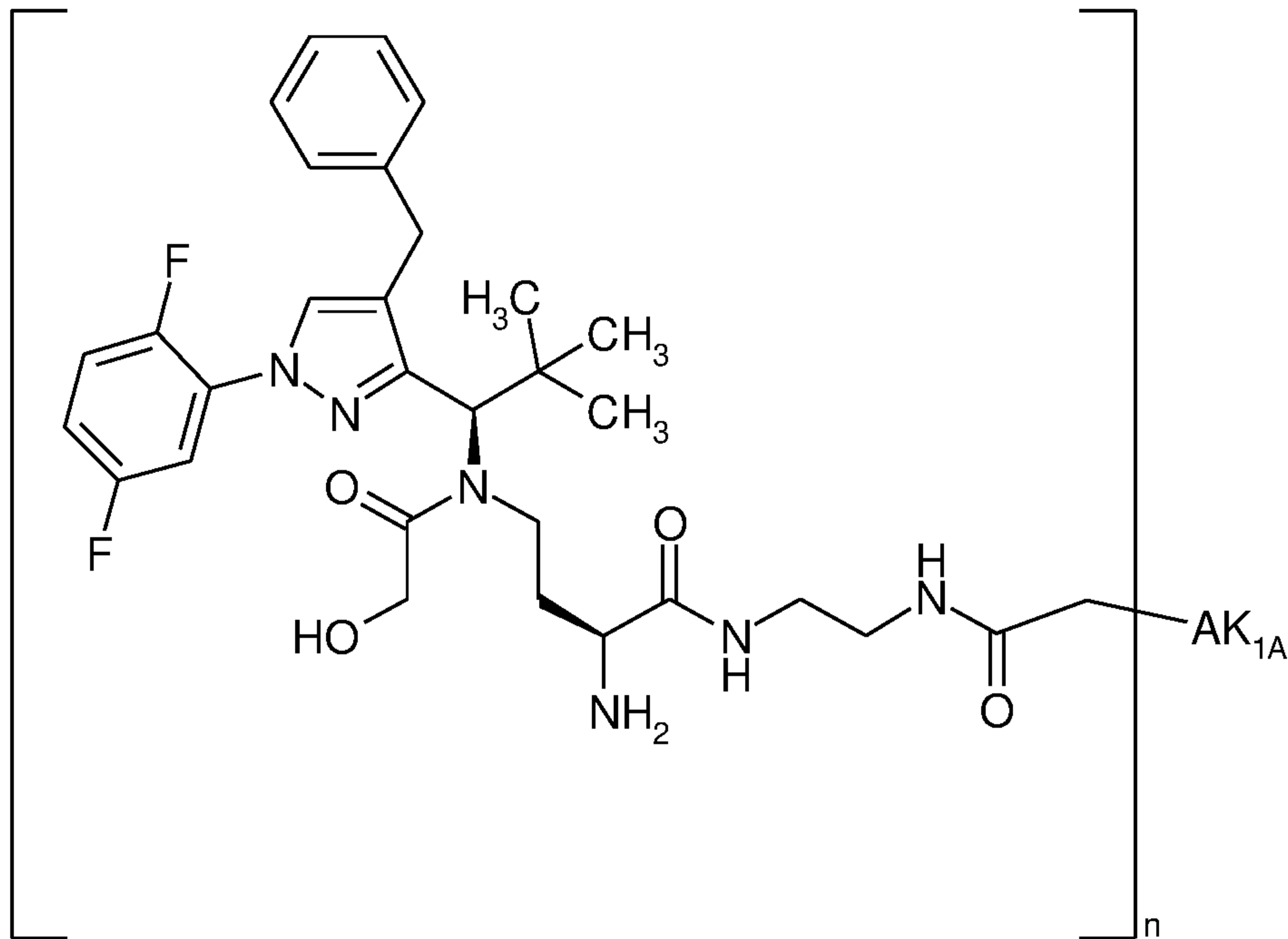
Here, 5 mg of trastuzumab in PBS (c=8.23 mg/ml) were used for  
20 coupling with Intermediate F107, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

25

Protein concentration: 2.48 mg/ml

Drug/mAb ratio: 2.9

30 **Example 109A**



Here, 5 mg of cetuximab in PBS ( $c=12.33$  mg/ml) were used for coupling with Intermediate F109. After TCEP reduction, coupling with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.1 mg/ml

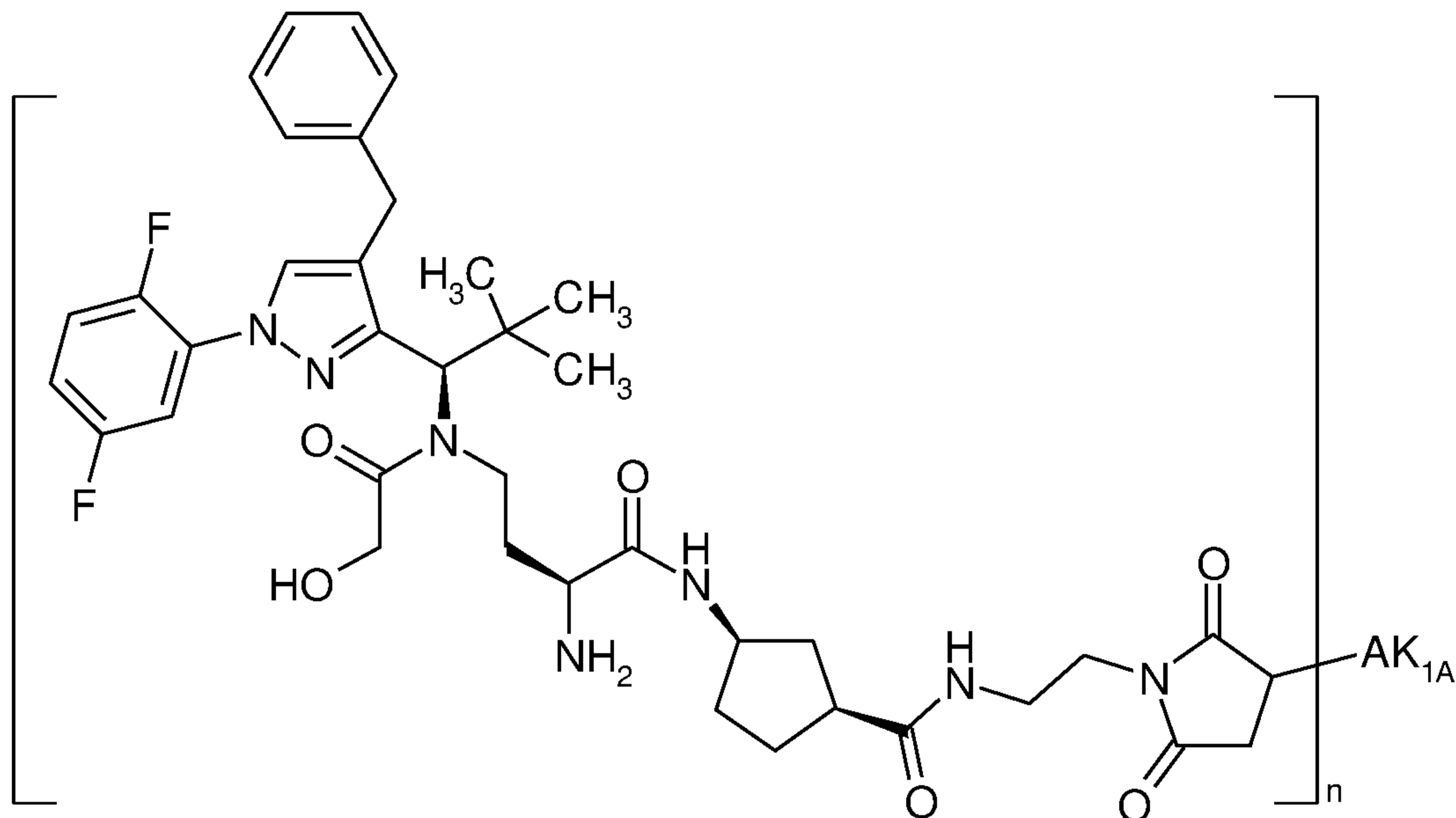
Drug/mAb ratio: 3.1

#### Example 109B

Here, 5 mg of anti-TWEAKR AK-1 in PBS ( $c=34.4$  mg/ml) were used for coupling with Intermediate F109. After TCEP reduction, coupling with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.63 mg/ml

Drug/mAb ratio: 2.7

Example 112A

5

Here, 5 mg of cetuximab in PBS (c=12.33 mg/ml) were used for coupling with Intermediate F122, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10

Protein concentration: 2.2 mg/ml

15 Drug/mAb ratio: 3

Example 112B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=34.4 mg/ml) were used for coupling with Intermediate F112, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

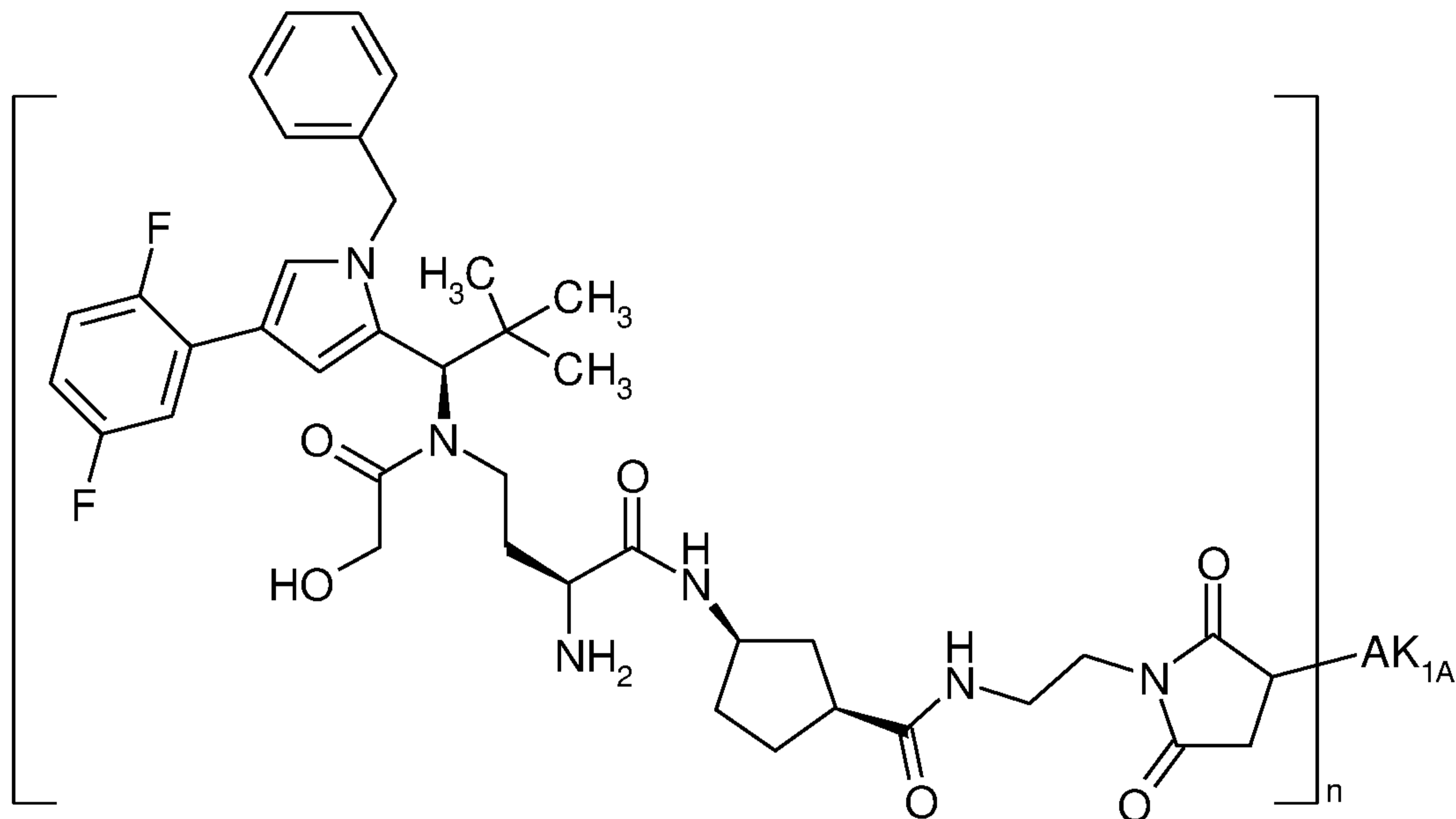
25

Protein concentration: 2.39 mg/ml

Drug/mAb ratio: 2.1

**Example 113A**

5



Here, 5 mg of cetuximab in PBS ( $c=12.33$  mg/ml) were used for coupling with Intermediate F113, and the reaction was, after  
 10 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15 Protein concentration: 1.95 mg/ml

Drug/mAb ratio: 2.2

**Example 113B**

20

Here, 5 mg of anti-TWEAKR AK-1 in PBS ( $c=34.4$  mg/ml) were used for coupling with Intermediate F113, and the reaction was, after  
 25 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

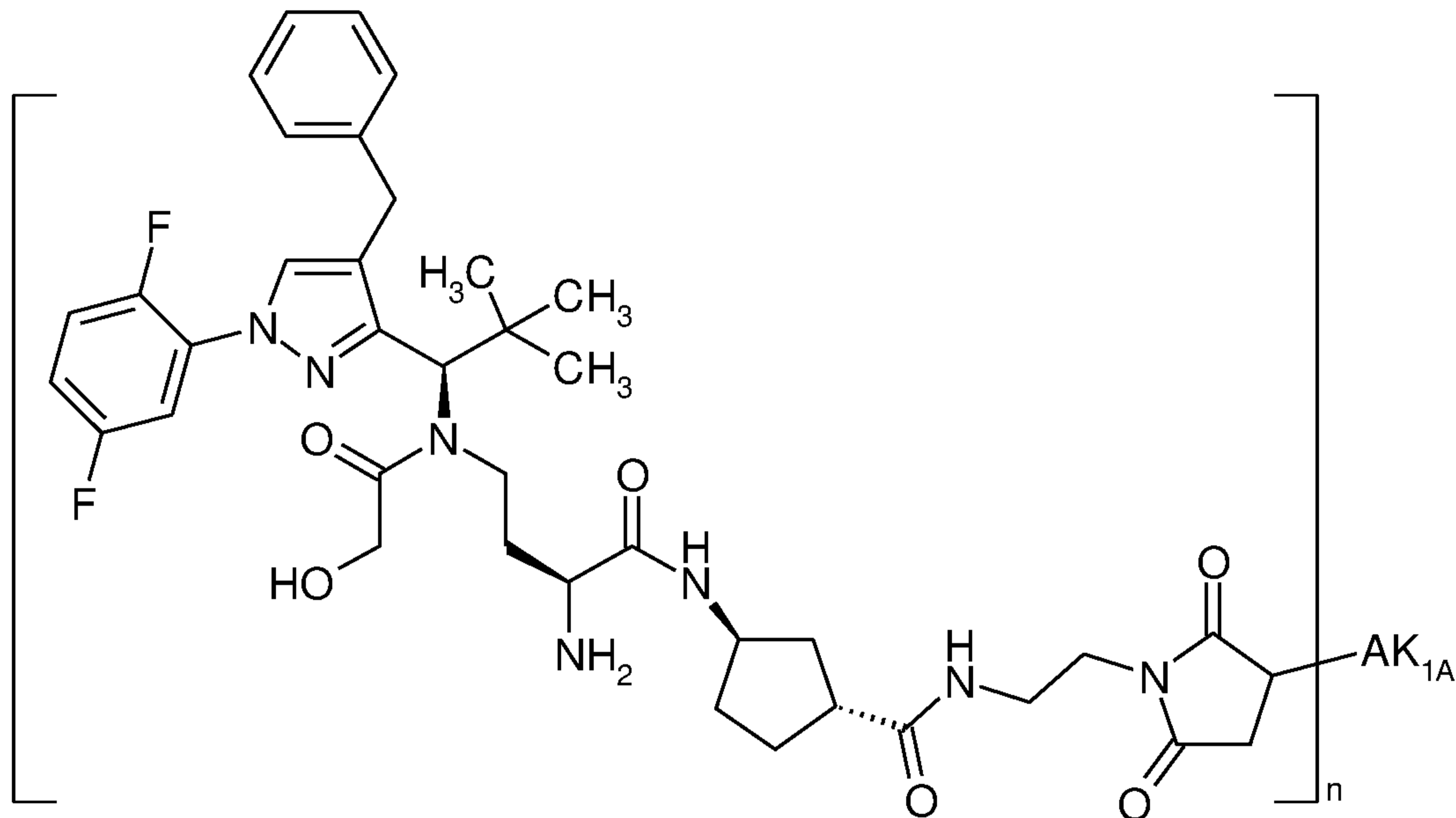


Protein concentration: 1.8 mg/ml

Drug/mAb ratio: 2.2

5

**Example 115A**



10 Here, 5 mg of cetuximab in PBS (c=12.33 mg/ml) were used for  
coupling with Intermediate F115, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS. Some of the ADC may also be present in the  
15 form of the hydrolysed open-chain succinamides attached to the  
antibody.

Protein concentration: 2.18 mg/ml

Drug/mAb ratio: 3.7

20

**Example 115B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=34.4 mg/ml) were used  
for coupling with Intermediate F115, and the reaction was, after  
25 Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS. Some of the ADC may also be present in the

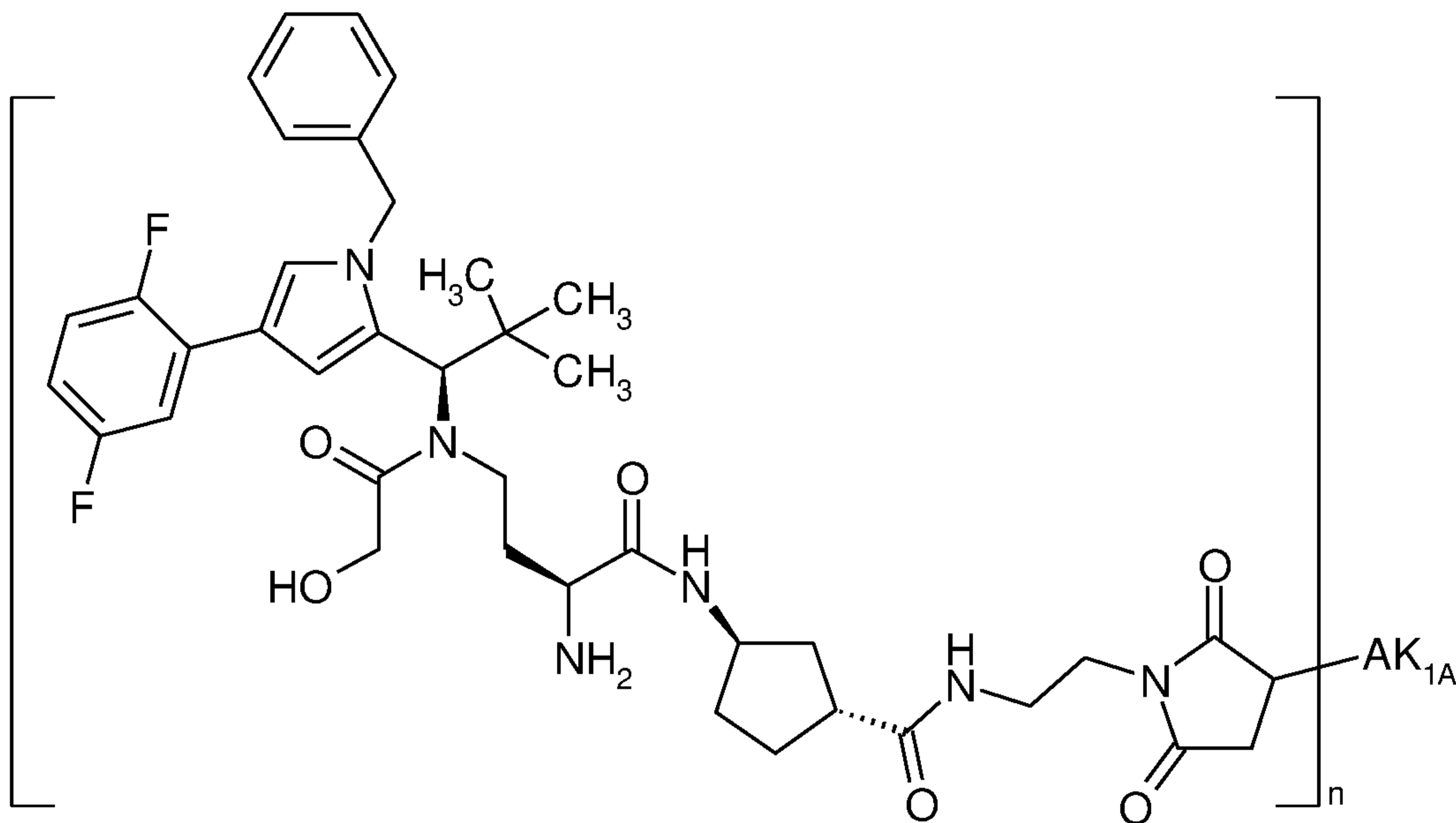
form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 2.0 mg/ml

5

Drug/mAb ratio: 3.0

### Example 116A



10

Here, 5 mg of cetuximab in PBS (c=12.33 mg/ml) were used for coupling with Intermediate F116, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 2.03 mg/ml

20

Drug/mAb ratio: 4.4

### Example 116B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=34.4 mg/ml) were used for coupling with Intermediate F116, and the reaction was, after

25

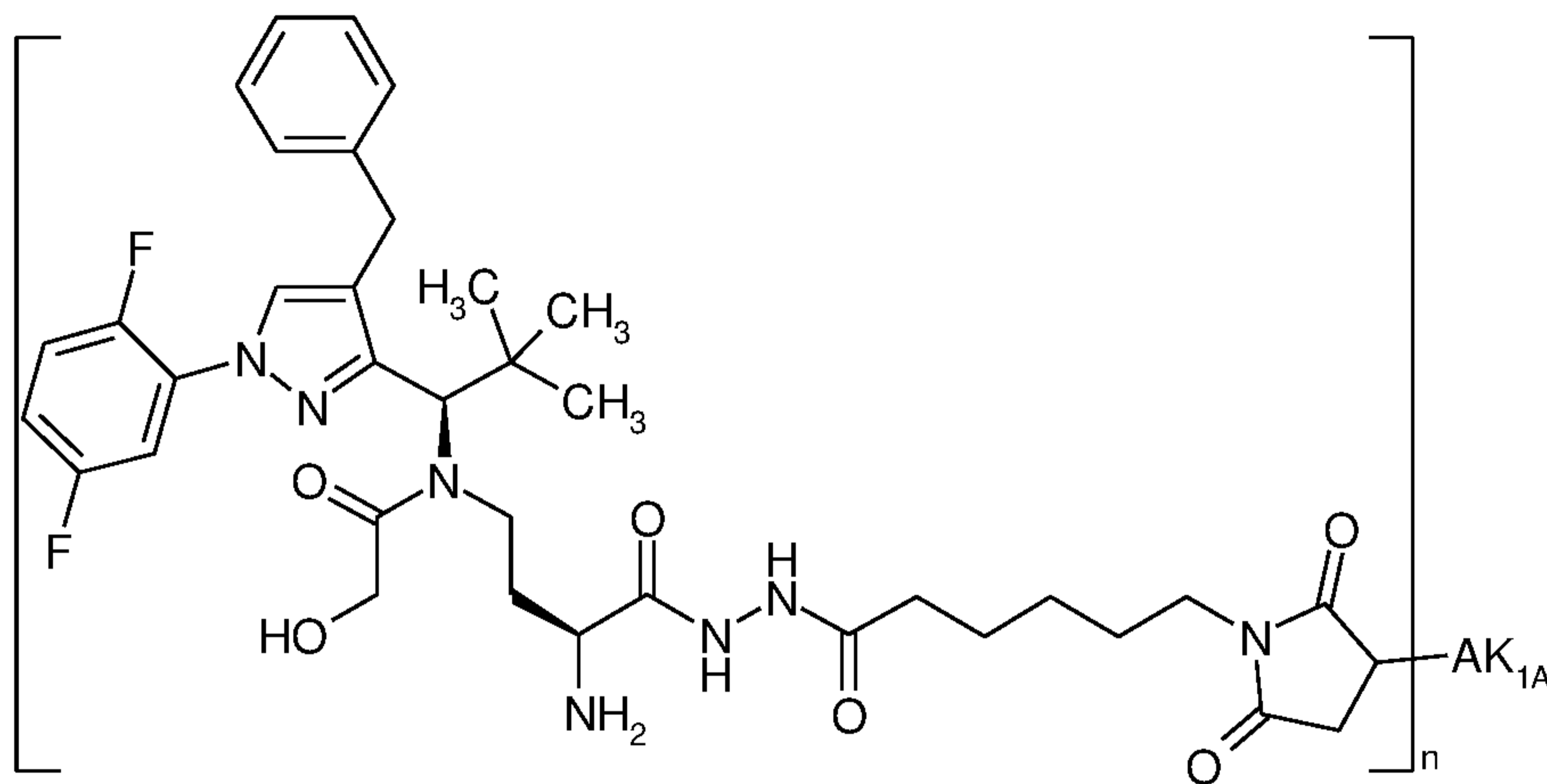
Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 1.96 mg/ml

Drug/mAb ratio: 2.9

### 10 Example 117A



Here, 5 mg of cetuximab in PBS (c=12.33 mg/ml) were used for  
 15 coupling with Intermediate F117, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

Protein concentration: 2.02 mg/ml

20

Drug/mAb ratio: 2.7

### Example 117B

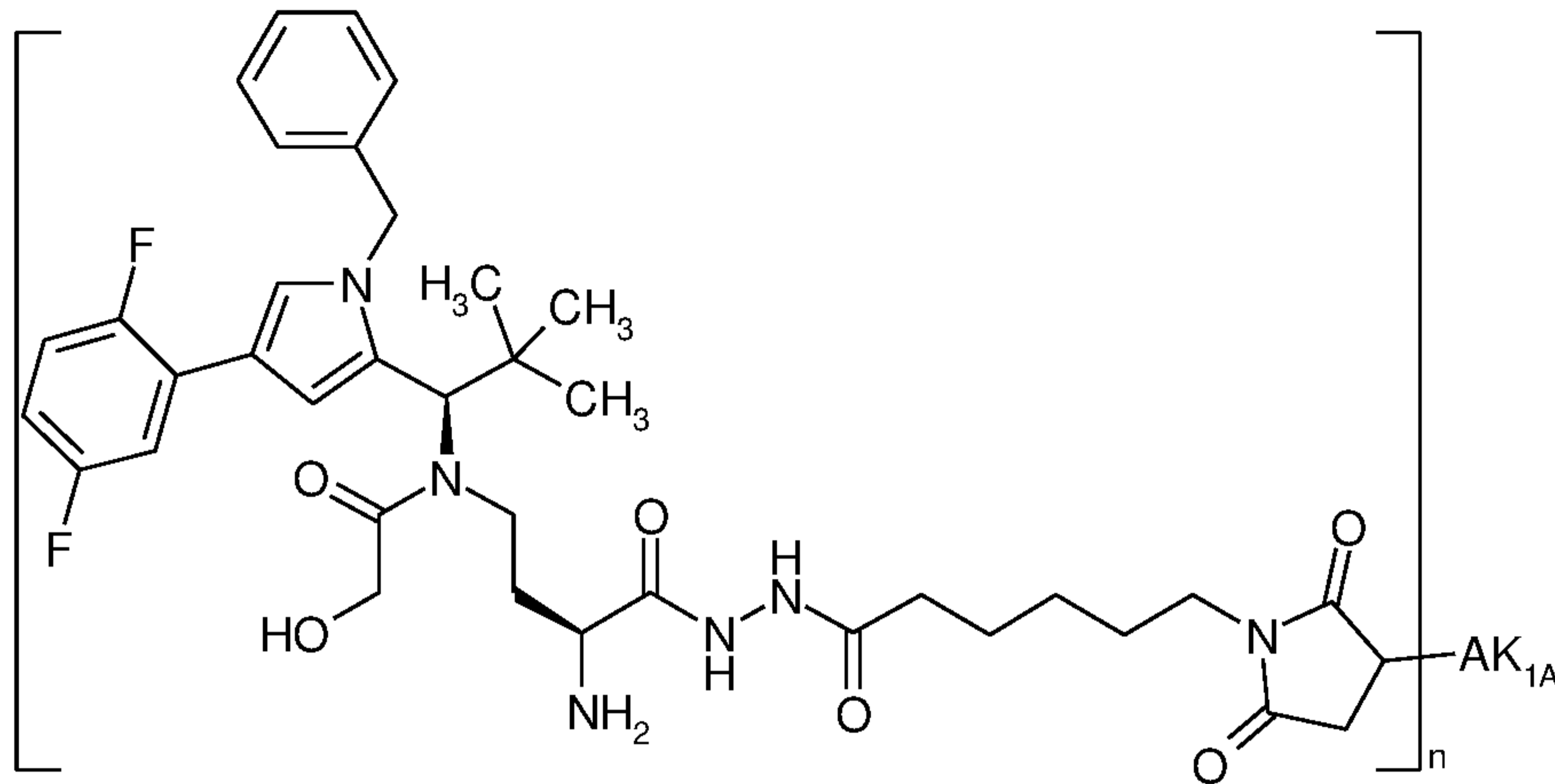
25 Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=34.4 mg/ml) were used  
 for coupling with Intermediate F117, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

30 Protein concentration: 1.77 mg/ml

Drug/mAb ratio: 2.7

**Example 118A**

5



Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F118, and the reaction was, after  
 10 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.38 mg/ml

15 Drug/mAb ratio: 3.6

**Example 118B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.87 mg/ml) were used  
 20 for coupling with Intermediate F118, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.14 mg/ml

25

Drug/mAb ratio: 2.9

**Example 118E**

30 Here, 5 mg of trastuzumab in PBS (c=8.23 mg/ml) were used for

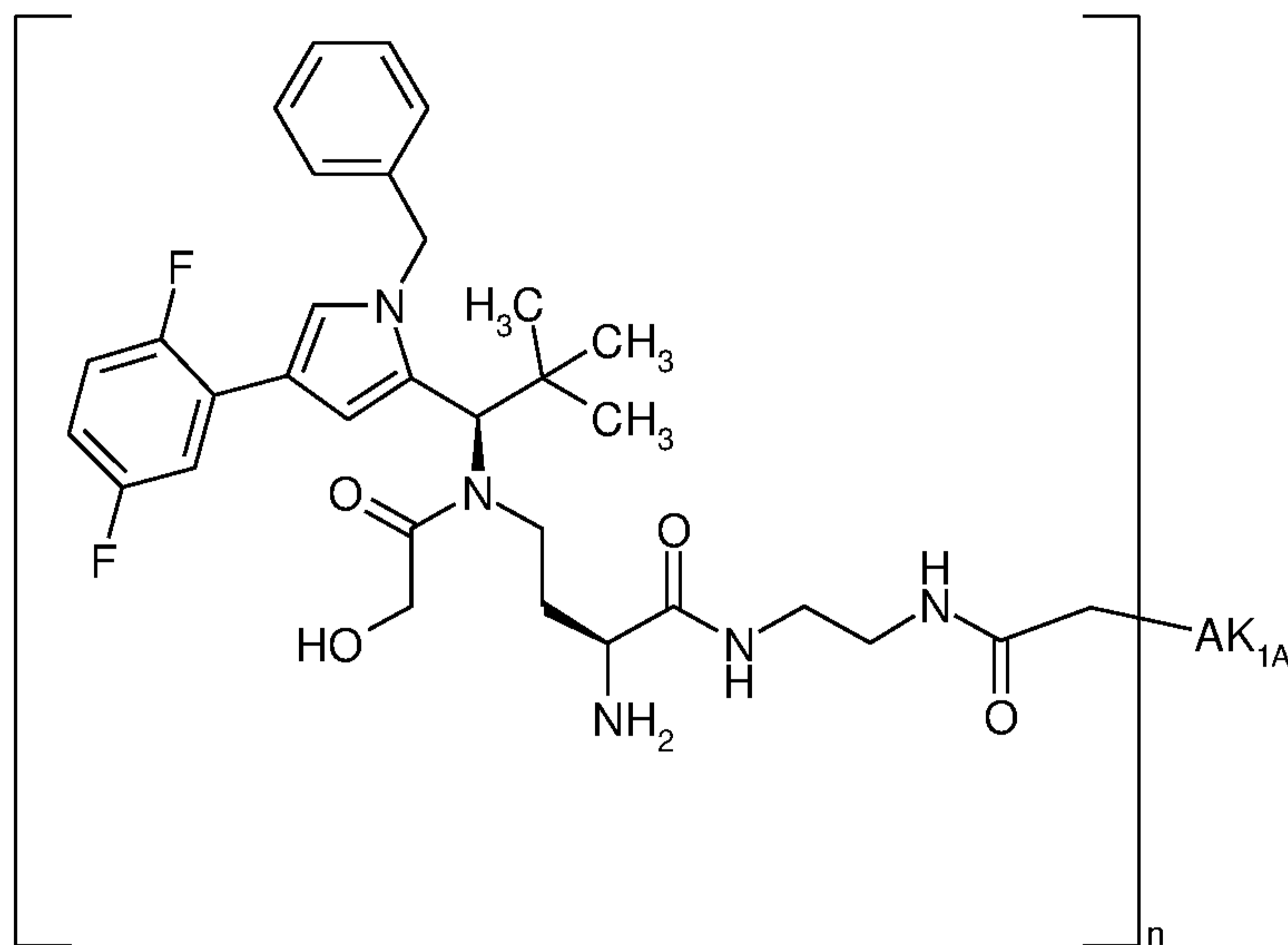
coupling with Intermediate F118, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

5 Protein concentration: 2.27 mg/ml

Drug/mAb ratio: 3

**Example 119A**

10



Here, 5 mg of cetuximab in PBS (c=26.8 mg/ml) were used for coupling with Intermediate F119. After TCEP reduction, coupling  
 15 with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

20 Protein concentration: 2.14 mg/ml

Drug/mAb ratio: 3.9

**Example 119B**

25

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.87 mg/ml) were used for coupling with Intermediate F119. After TCEP reduction,

coupling with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

5

Protein concentration: 0.91 mg/ml

Drug/mAb ratio: 4.1

### 10 Example 119E

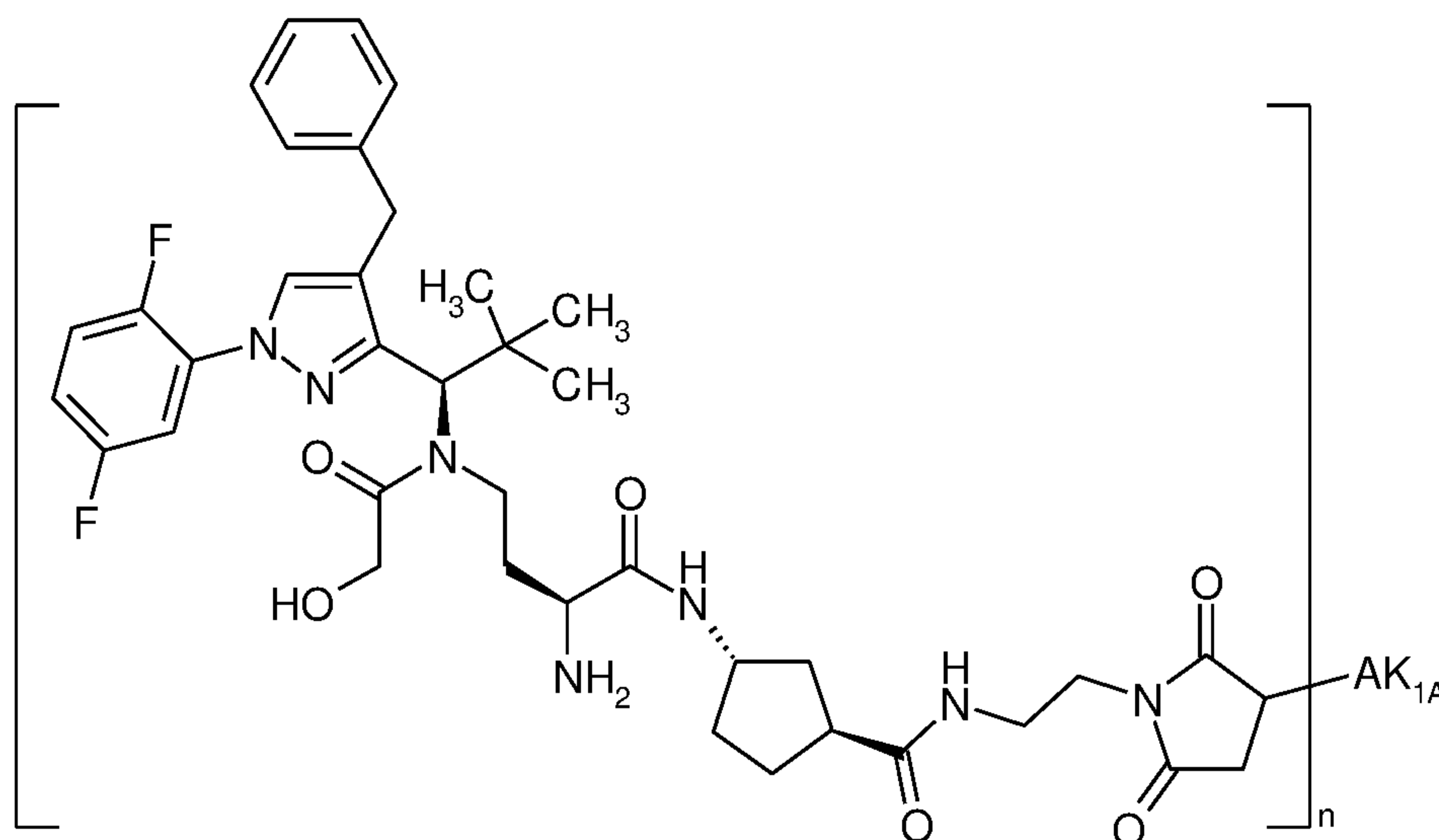
Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F119. After TCEP reduction, coupling with the antibody was carried out with stirring overnight,  
 15 followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.69 mg/ml

20

Drug/mAb ratio: 4.4

### Example 121A



25

Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F121, and the reaction was, after

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 2.1 mg/ml

Drug/mAb ratio: 3.2

### 10 Example 121B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.87 mg/ml) were used for coupling with Intermediate F121, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

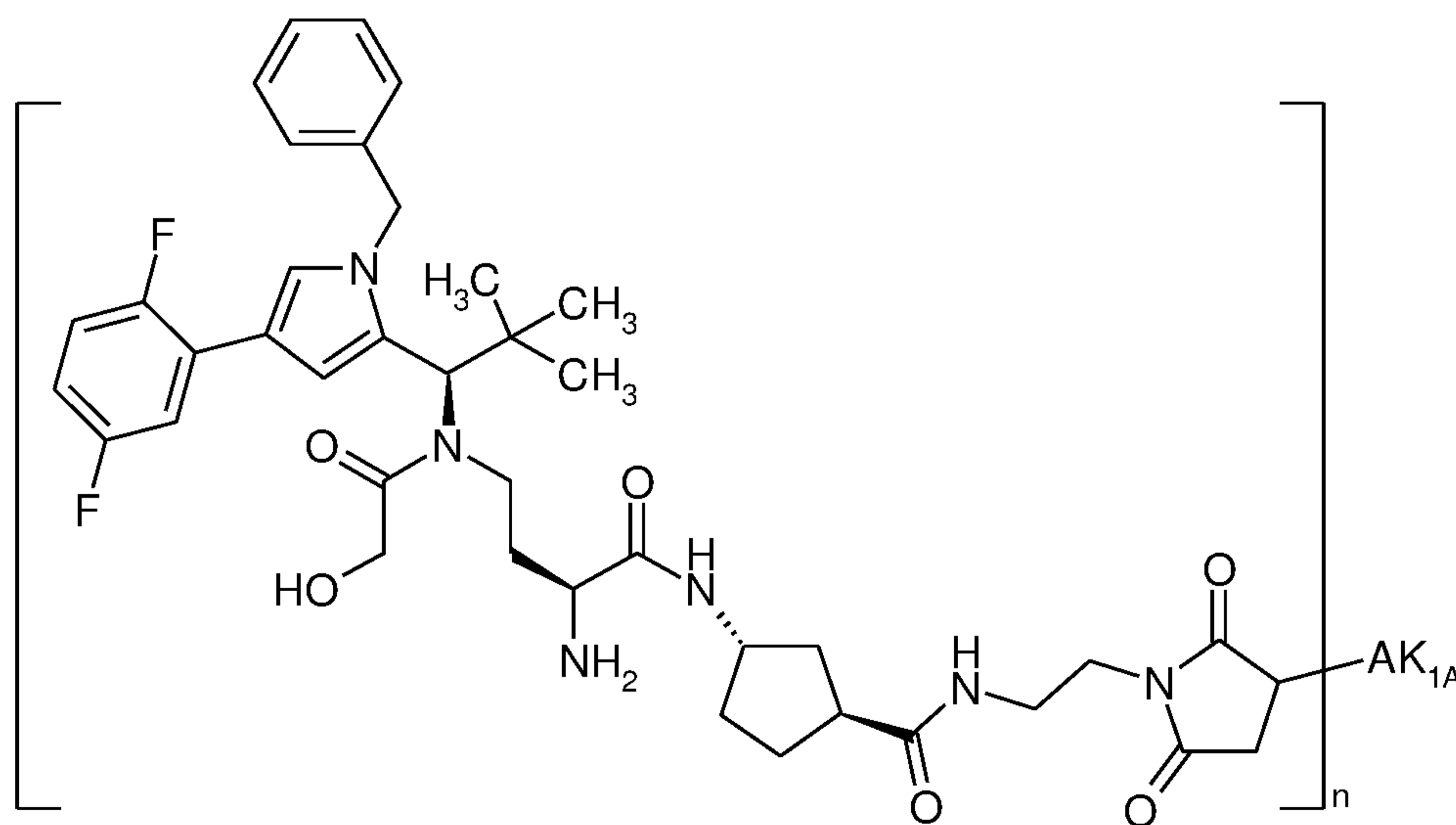
15

Protein concentration: 1.88 mg/ml

20

Drug/mAb ratio: 3.4

### Example 122A



25

Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F122, and the reaction was, after

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 1.78 mg/ml

Drug/mAb ratio: 3.2

### 10 Example 122B

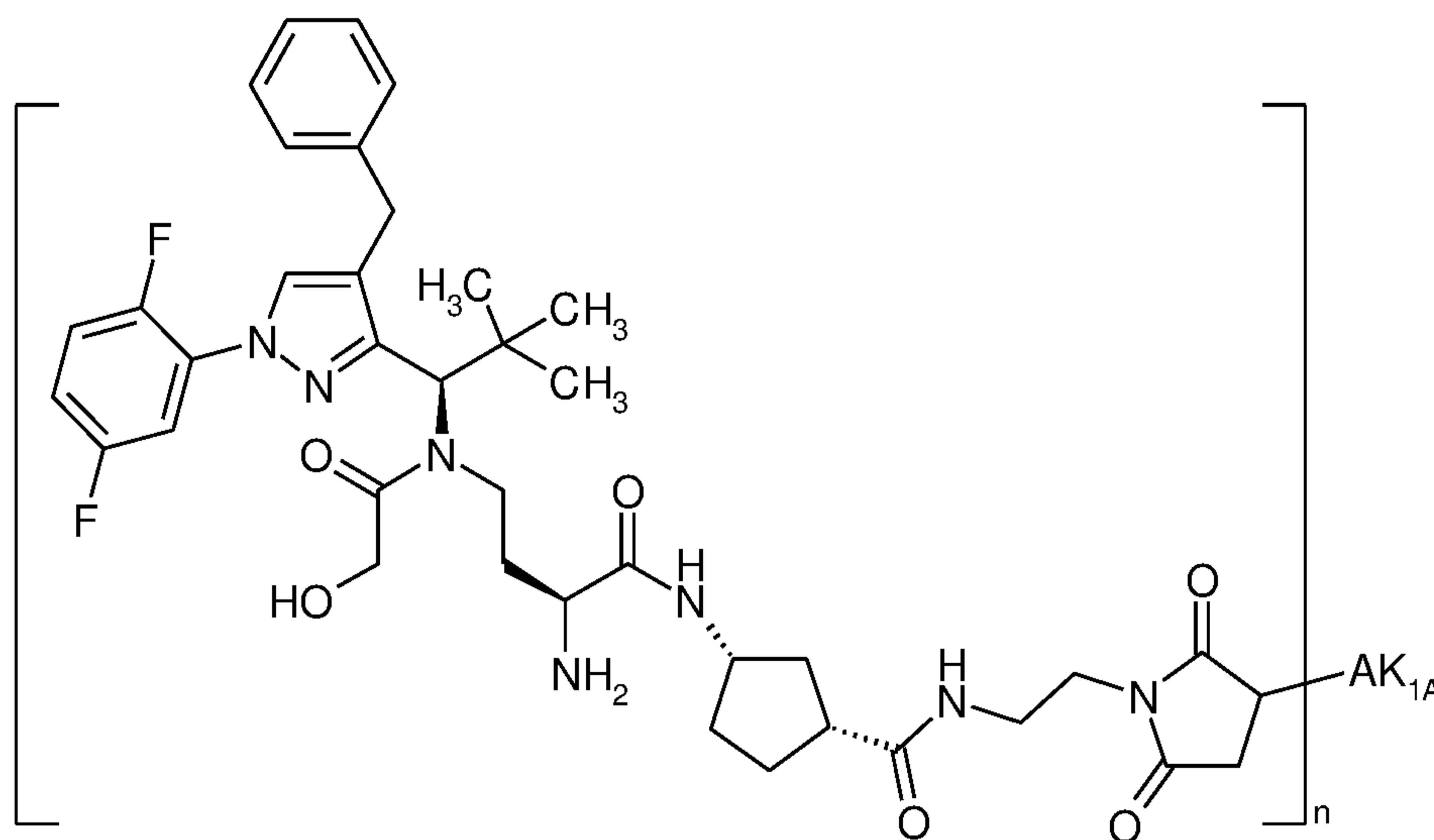
Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.87 mg/ml) were used for coupling with Intermediate F122, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 1.64 mg/ml

Drug/mAb ratio: 3.4

### 20 Example 124A



25

Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F124, and the reaction was, after



Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 1.93 mg/ml

Drug/mAb ratio: 2.8

### 10 Example 124B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.87 mg/ml) were used for coupling with Intermediate F124, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

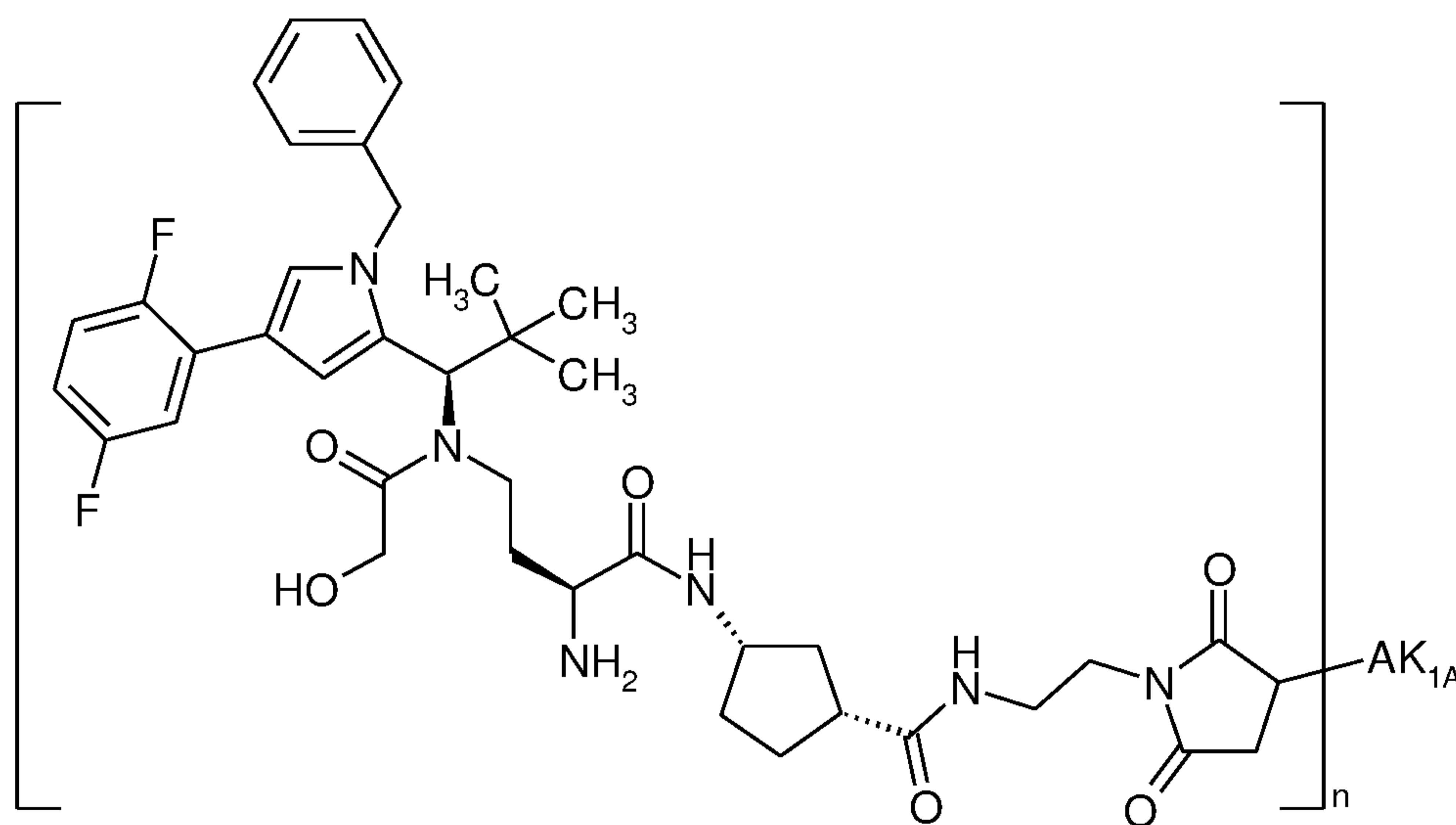
15

Protein concentration: 1.84 mg/ml

20

Drug/mAb ratio: 3.0

### Example 125A



25

Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F125, and the reaction was, after

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 2.14 mg/ml

Drug/mAb ratio: 2.9

### 10 Example 125B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.87 mg/ml) were used for coupling with Intermediate F125, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

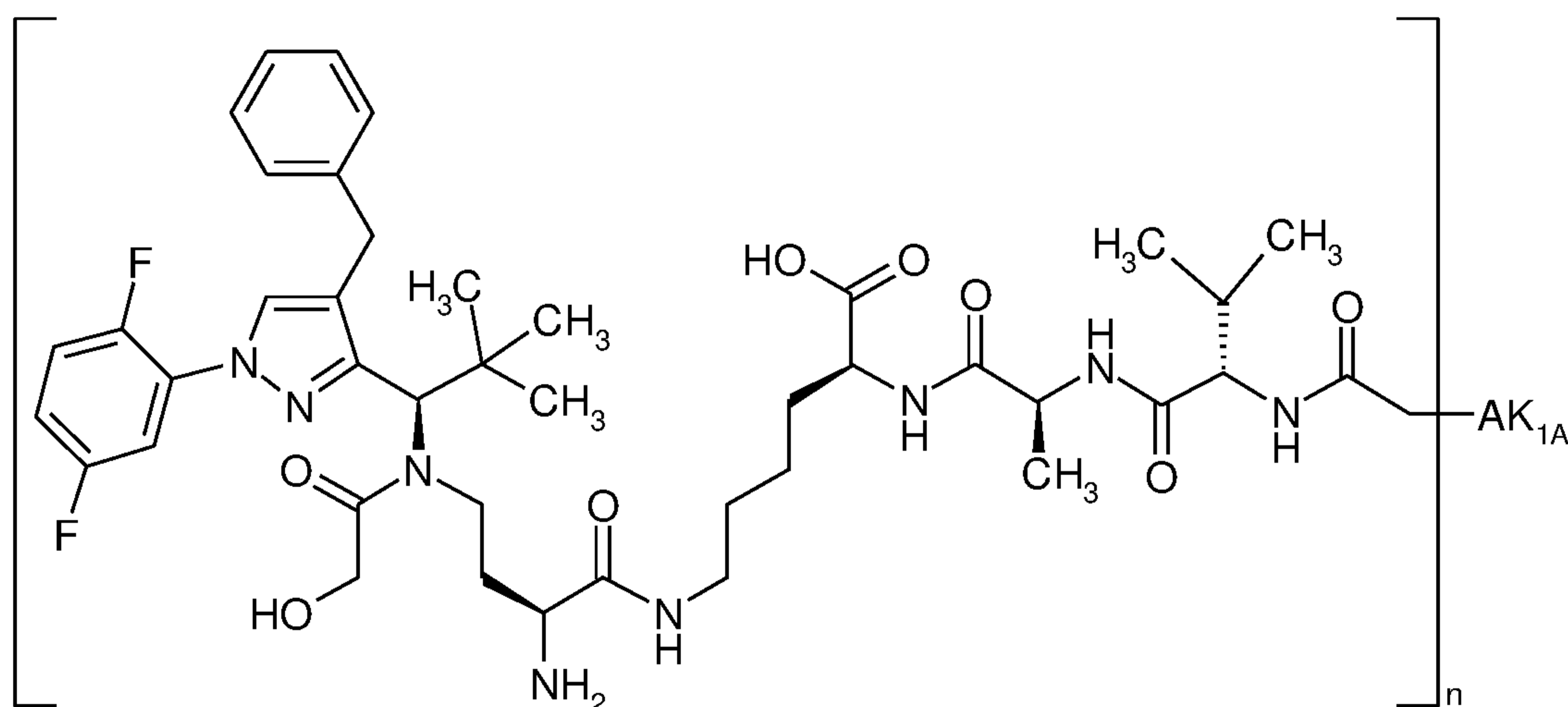
15

Protein concentration: 1.83 mg/ml

20

Drug/mAb ratio: 2.3

### Example 126A



25

Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F126. After TCEP reduction, coupling

with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

5

Protein concentration: 1.89 mg/ml

Drug/mAb ratio: 2.5

#### 10 Example 126B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F126. After TCEP reduction, coupling with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

15

Protein concentration: 1.62 mg/ml

20

Drug/mAb ratio: 2.8

#### Example 126E

Here, 5 mg of trastuzumab in PBS (c=8.23 mg/ml) were used for coupling with Intermediate F126. After TCEP reduction, coupling with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

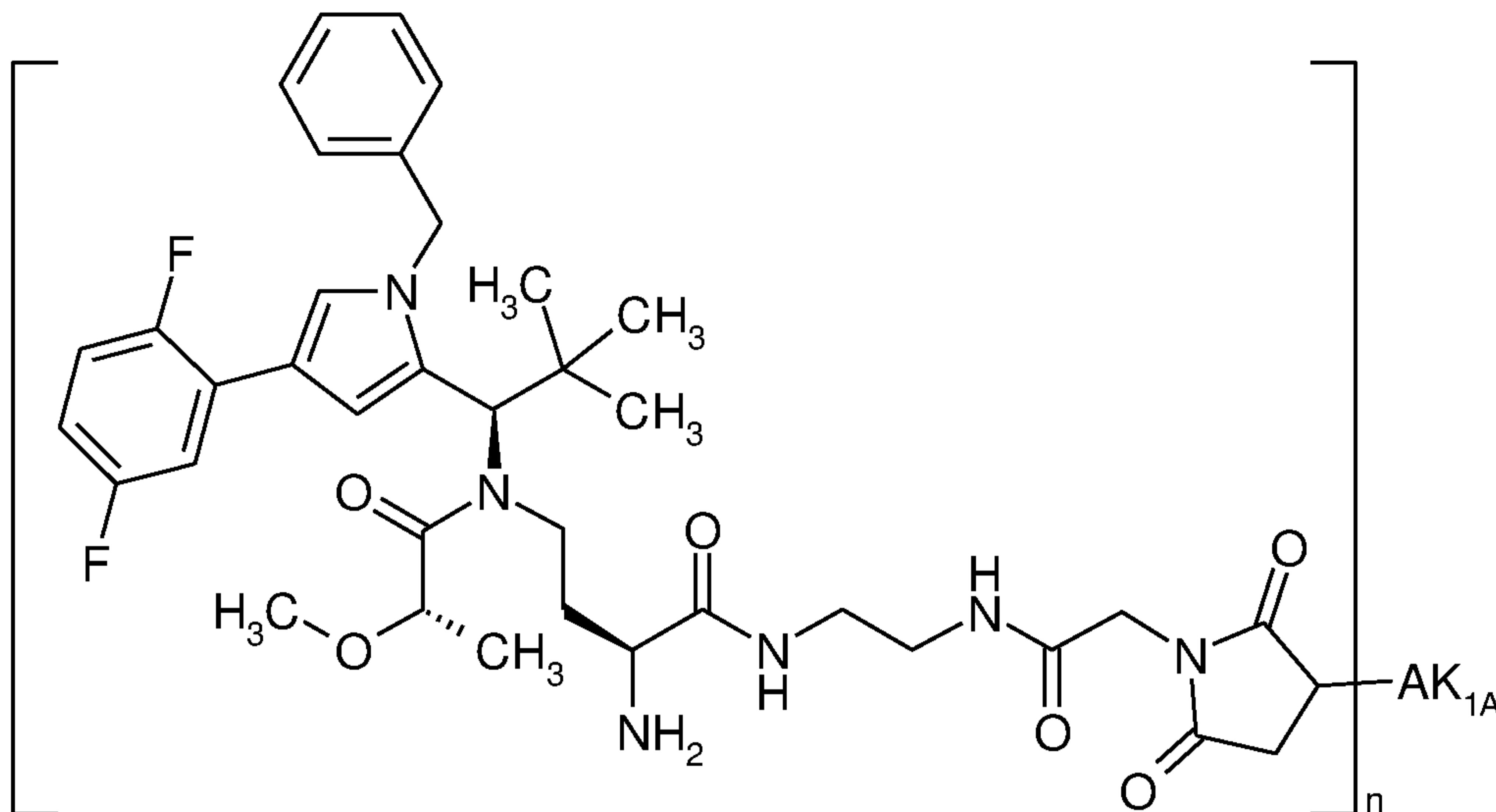
30

Protein concentration: 1.93 mg/ml

Drug/mAb ratio: 1.9

35

#### Example 127A



Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F127, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.54 mg/ml

Drug/mAb ratio: 3.3

### Example 127B

15

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F127, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the  
 20 form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.62 mg/ml

25 Drug/mAb ratio: 3.3

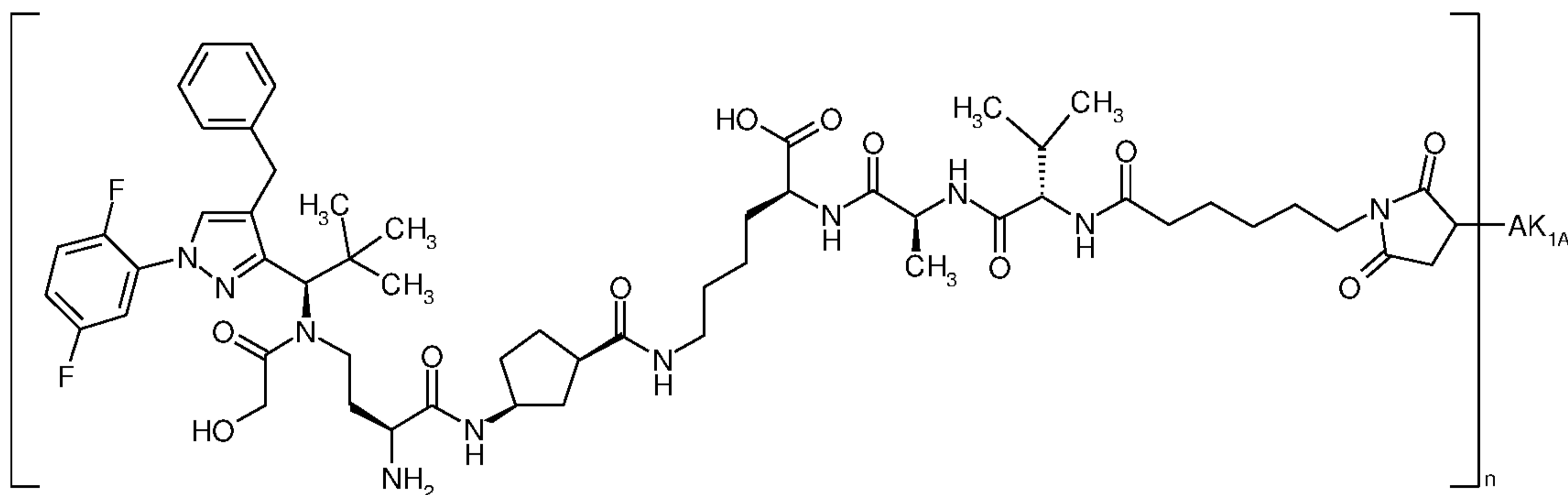
### Example 127E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F127, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 2.07 mg/ml

10 Drug/mAb ratio: 3.6

### Example 129A



15

Here, 5 mg of cetuximab in PBS (c=26.84 mg/ml) were used for coupling with Intermediate F129, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

20

Protein concentration: 2.28 mg/ml

Drug/mAb ratio: 2.9

### Example 129B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F129, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

30

Protein concentration: 2.06 mg/ml

Drug/mAb ratio: 3.2

**Example 129E**

5

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F129, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

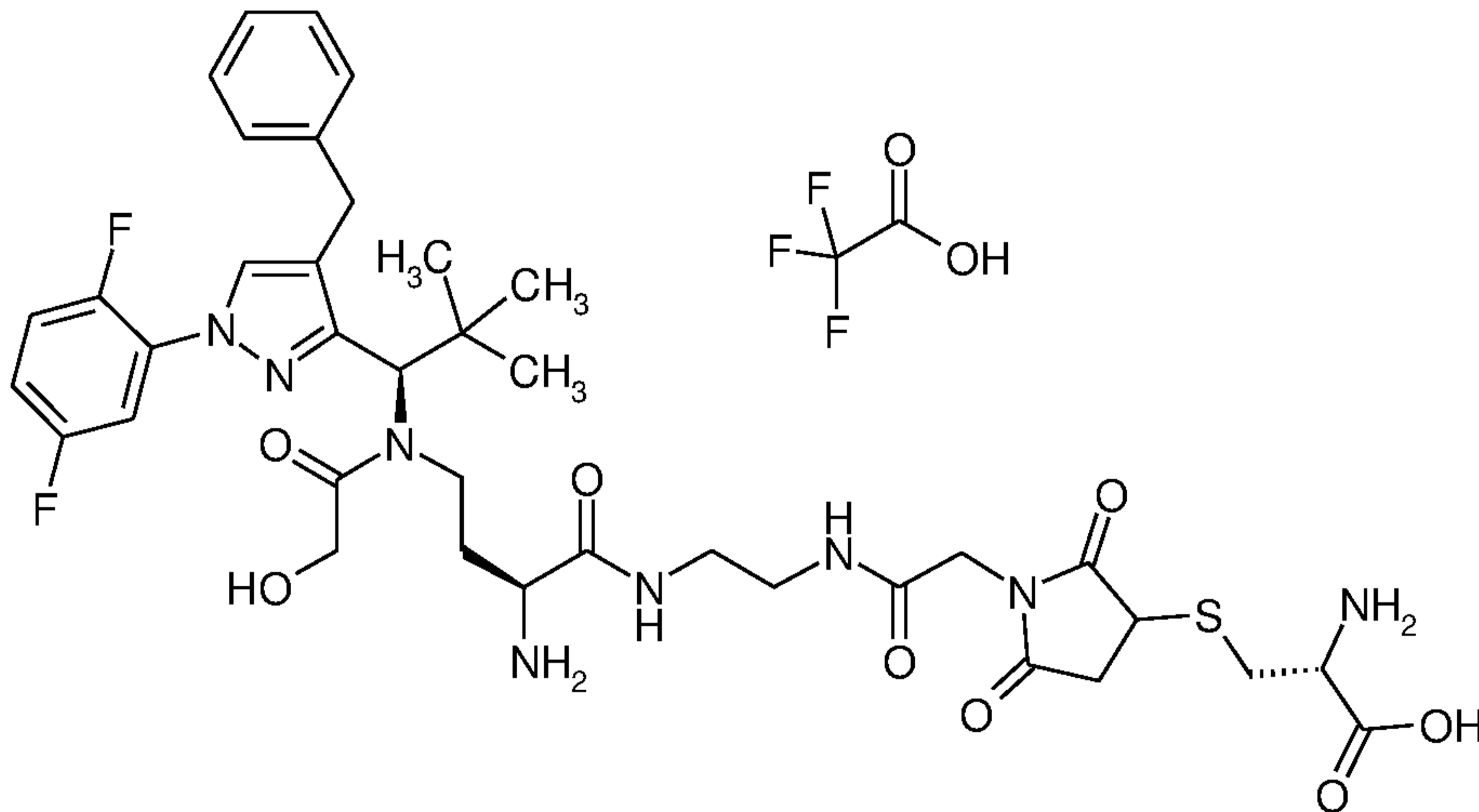
10

Protein concentration: 2.0 mg/ml

Drug/mAb ratio: 3.4

15 **Example 131**

S-[1-(2-{{[2-((2S)-2-Amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-oxoethyl)-2,5-dioxopyrrolidin-3-yl]-L-cysteine /  
20 trifluoroacetic acid (1:1)

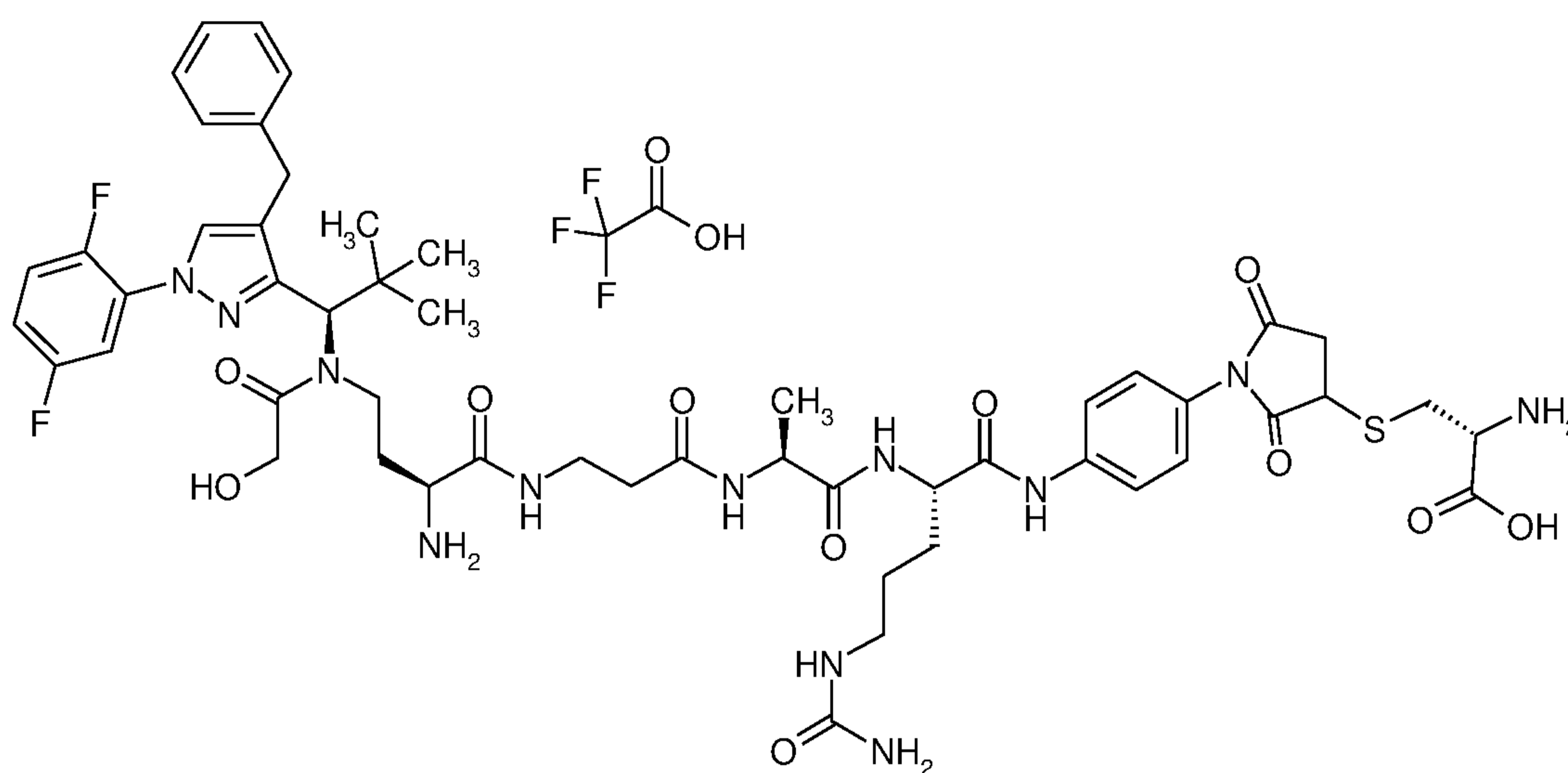


25 5 mg (6  $\mu$ mol) of Intermediate F87 were taken up in 1 ml of DMF, and 7.5 mg (62  $\mu$ mol) of L-cysteine were added. The reaction mixture was stirred at RT for 20 h, then concentrated under reduced pressure and then purified by preparative HPLC.

LC-MS (Method 1):  $R_t = 0.81$  min; MS (EIpos):  $m/z = 815$   $[M+H]^+$ .

### Example 132

5 N-{(2S)-2-Amino-4-[[{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
1H-pyrazol-3-yl]-2,2-  
dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl-L-  
alanyl-N-[4-(3-[[{(2R)-2-amino-2-carboxyethyl]sulphonyl}-2,5-  
dioxopyrrolidin-1-yl]phenyl]-N<sup>5</sup>-carbamoyl-L-  
10 ornithinamide/trifluoroacetic acid (1:1)



5 mg (5  $\mu$ mol) of Intermediate F89 were taken up in 2 ml of  
15 DMF/water 10:1, and 1.7 mg (14  $\mu$ mol) of L-cysteine were added.  
The reaction mixture was stirred at RT for 1 h and then  
concentrated under reduced pressure. The residue was taken up  
in acetonitrile/water, and the mixture was adjusted to pH 2  
using TFA and then concentrated under reduced pressure and then  
20 purified by preparative HPLC. The appropriate fractions were  
concentrated, giving, after lyophilization of the residue from  
acetonitrile/water, 3 mg of the title compound as a white foam.

LC-MS (Method 4):  $R_t = 0.93$  min; MS (EIpos):  $m/z = 1105$   $[M+H]^+$ .

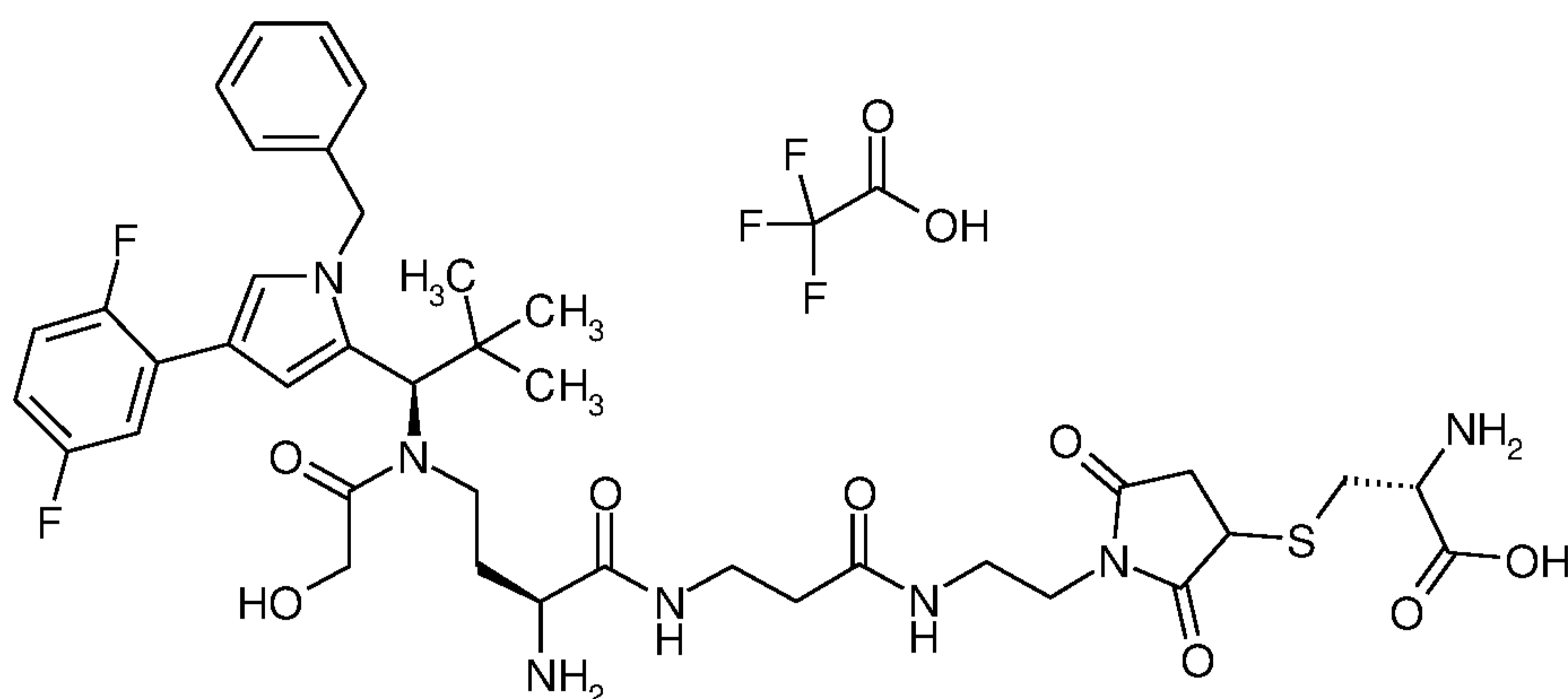
25

### Example 133

S-(1-{2-[(N-{(2S)-2-Amino-4-[[{(1R)-1-[1-benzyl-4-(2,5-

difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)amino]ethyl}-2,5-dioxopyrrolidin-3-yl)-L-cysteine / trifluoroacetic acid (1:1)

5



1.6 mg (2  $\mu$ mol) of Intermediate F84 were taken up in 1.5 ml of DMF/water 10:1, and 0.74 mg (6  $\mu$ mol) of L-cysteine were added. The reaction mixture was stirred at RT for 10 min and then concentrated under reduced pressure. The residue was taken up in acetonitrile/water, and the mixture was adjusted to pH 2 using TFA and then concentrated under reduced pressure and then purified by preparative HPLC. The appropriate fractions were concentrated, giving, after lyophilization of the residue from acetonitrile/water, 1.9 mg (89% of theory) of the title compound as a white foam.

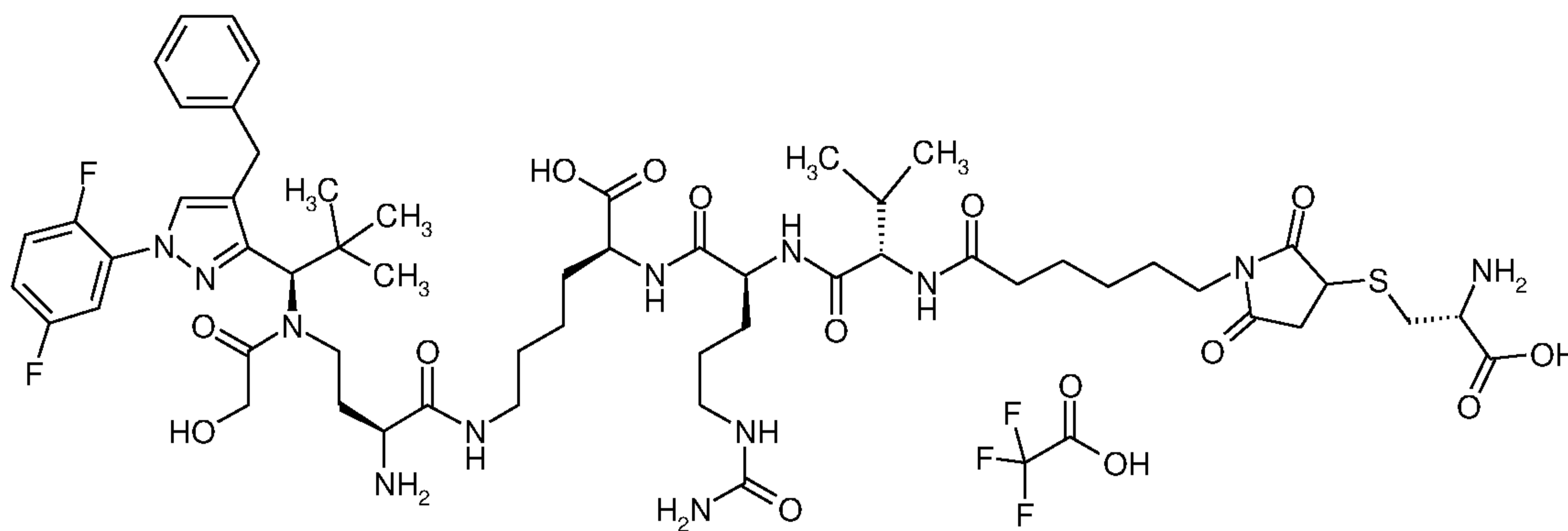
LC-MS (Method 1):  $R_t$  = 0.8 min; MS (EIpos):  $m/z$  = 828  $[M+H]^+$ .

20

### Example 134

N-[6-(3-{(2R)-2-Amino-2-carboxyethyl}sulphanyl)-2,5-dioxopyrrolidin-1-yl)hexanoyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithyl-N<sup>6</sup>-{(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-lysine / trifluoroacetic acid (1:1)



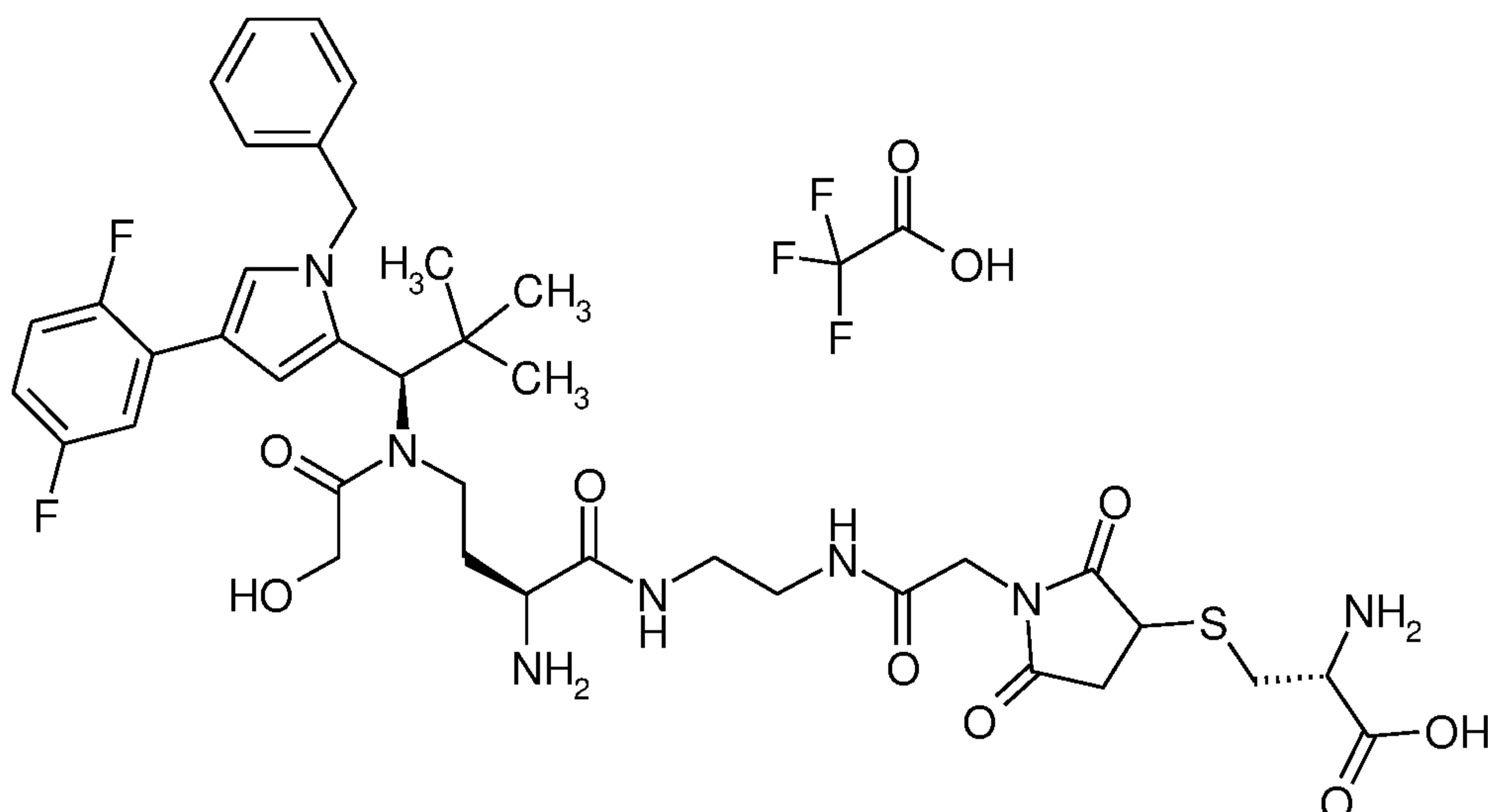


3.8 mg (3  $\mu$ mol) of Intermediate F90 were taken up in 1.5 ml of DMF/water 10:1, and 1.2 mg (9  $\mu$ mol) of L-cysteine were added.  
 5 The reaction mixture was stirred at RT for 15 min and then concentrated under reduced pressure. The residue was taken up in acetonitrile/water 1:1, concentrated again and then purified by preparative HPLC. The appropriate fractions were concentrated, giving, after lyophilization of the residue from  
 10 acetonitrile/water, 2.3 mg (56% of theory) of the title compound as a white foam.

LC-MS (Method 4):  $R_t$  = 1.0 min; MS (EIpos):  $m/z$  = 1213  $[M+H]^+$ .

15 **Example 135**

S-[1-(2-{{2-((2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-oxoethyl)-2,5-dioxopyrrolidin-3-yl]-L-cysteine /  
 20 trifluoroacetic acid (1:1)

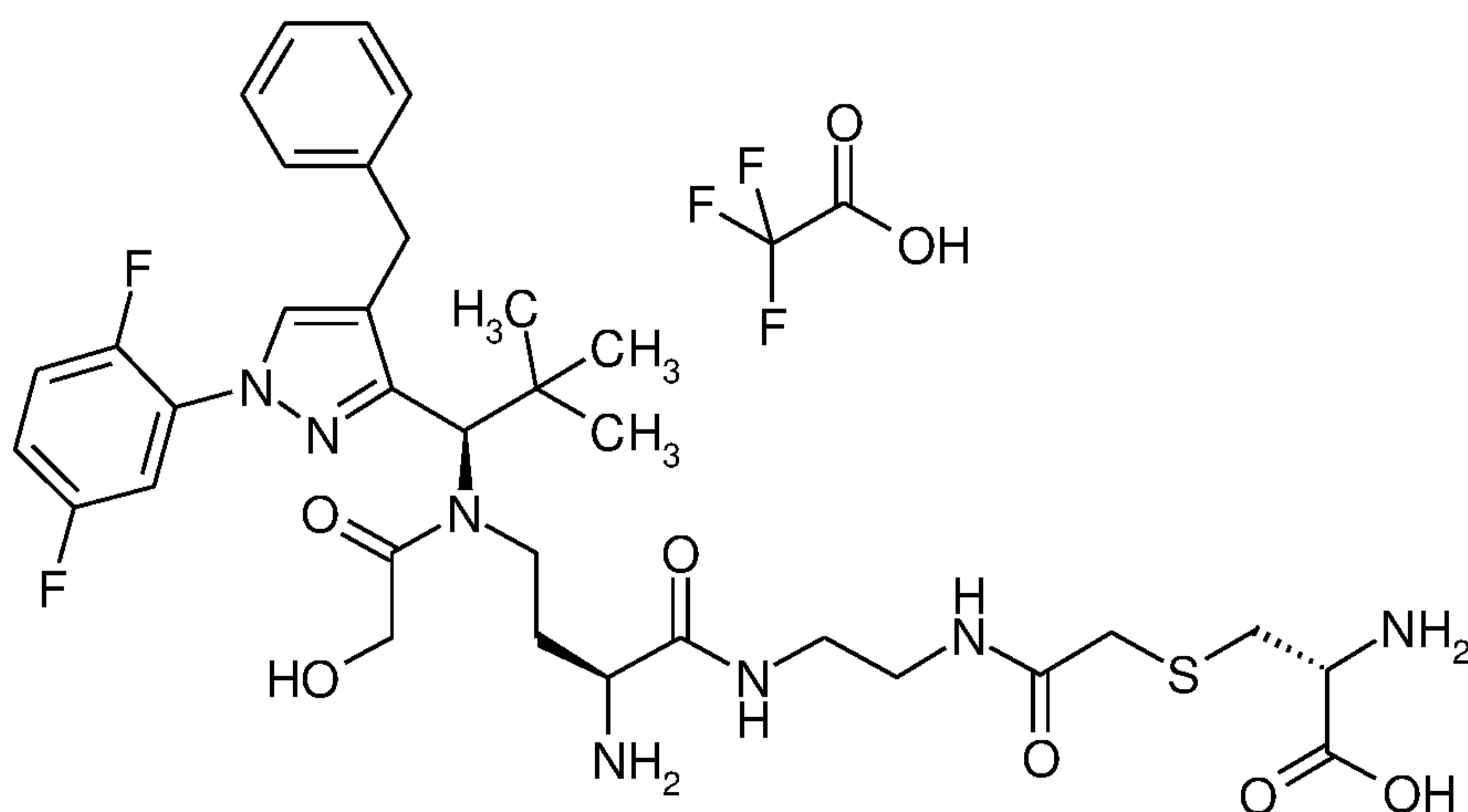


1.8 mg (2  $\mu\text{mol}$ ) of Intermediate F104 were taken up in 1 ml of DMF, and 2.7 mg (22  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 20 h, then concentrated under reduced pressure and then purified by preparative HPLC. 0.6 mg (26% of theory) of the title compound remained as a colourless foam.

10 LC-MS (Method 1):  $R_t = 0.80$  min; MS (EIpos):  $m/z = 814$   $[\text{M}+\text{H}]^+$ .

### Example 136

S-(2-([2-((2S)-2-Amino-4-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)(glycoloyl)amino]butanoyl)amino)ethyl]amino)-2-oxoethyl)-L-cysteine/trifluoroacetic acid (1:1)

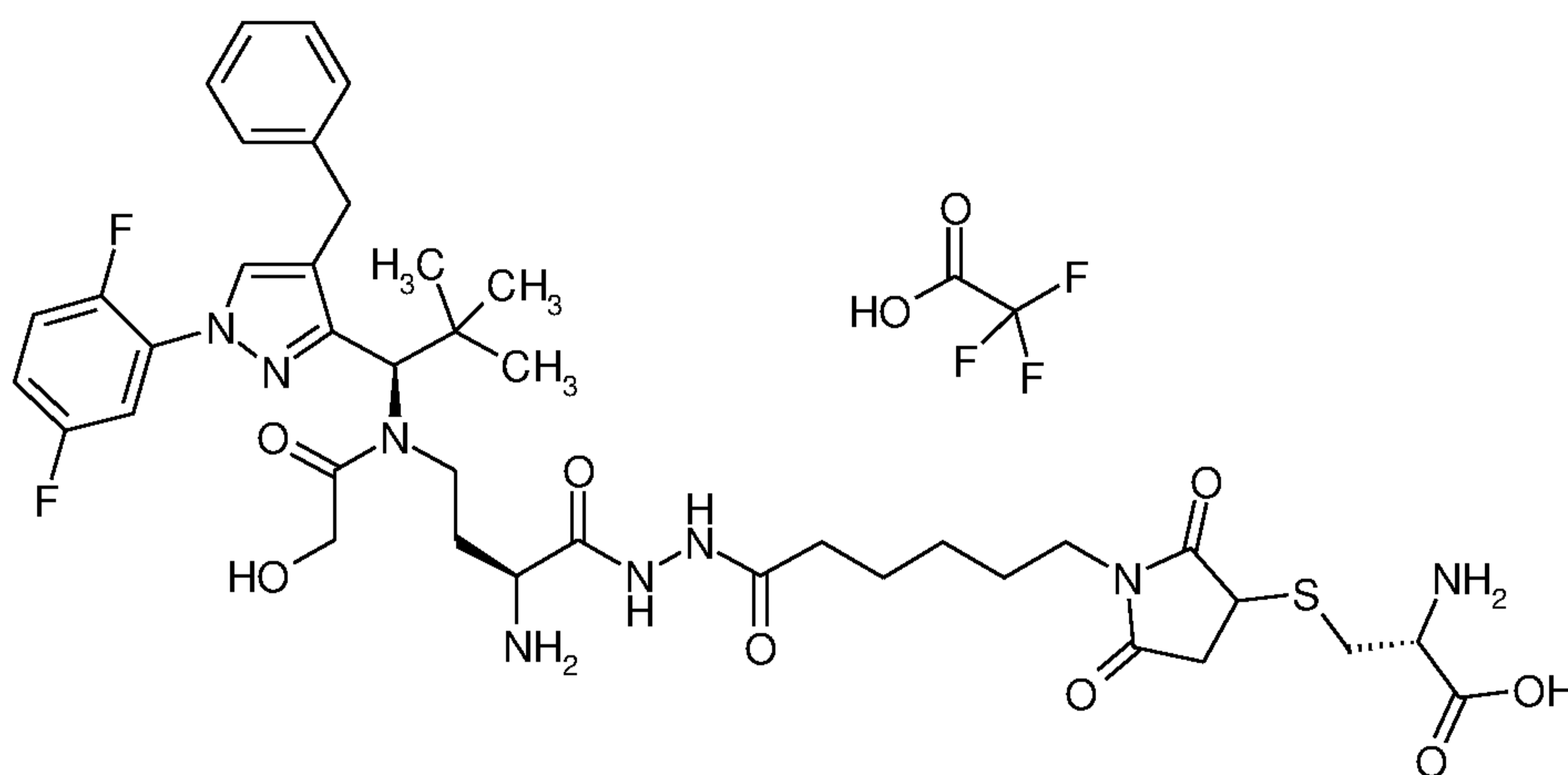


3.3 mg (4  $\mu\text{mol}$ ) of Intermediate F109 were taken up in 2 ml of DMF/water 10:1, and 1.5 mg (13  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 30 min and then concentrated under reduced pressure. The residue was taken up in acetonitrile/water 1:1, concentrated again and then purified by preparative HPLC. The appropriate fractions were concentrated, giving, after lyophilization of the residue from acetonitrile/water, 1.9 mg (55% of theory) of the title compound as a white foam.

LC-MS (Method 1):  $R_t = 0.75$  min; MS (EIpos):  $m/z = 718$   $[\text{M}+\text{H}]^+$ .

### Example 137

S-{1-[6-(2-((2S)-2-Amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}hydrazino)-6-oxohexyl]-2,5-dioxopyrrolidin-3-yl}-L-cysteine / trifluoroacetic acid (1:1)



3.2 mg (4  $\mu\text{mol}$ ) of Intermediate F117 were taken up in 2 ml of DMF/water 10:1, and 1.6 mg (13  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 30 min and then concentrated under reduced pressure. The residue was taken up in acetonitrile/water 1:1, concentrated again and then purified by preparative HPLC. The appropriate fractions were concentrated, giving, after lyophilization of the residue from

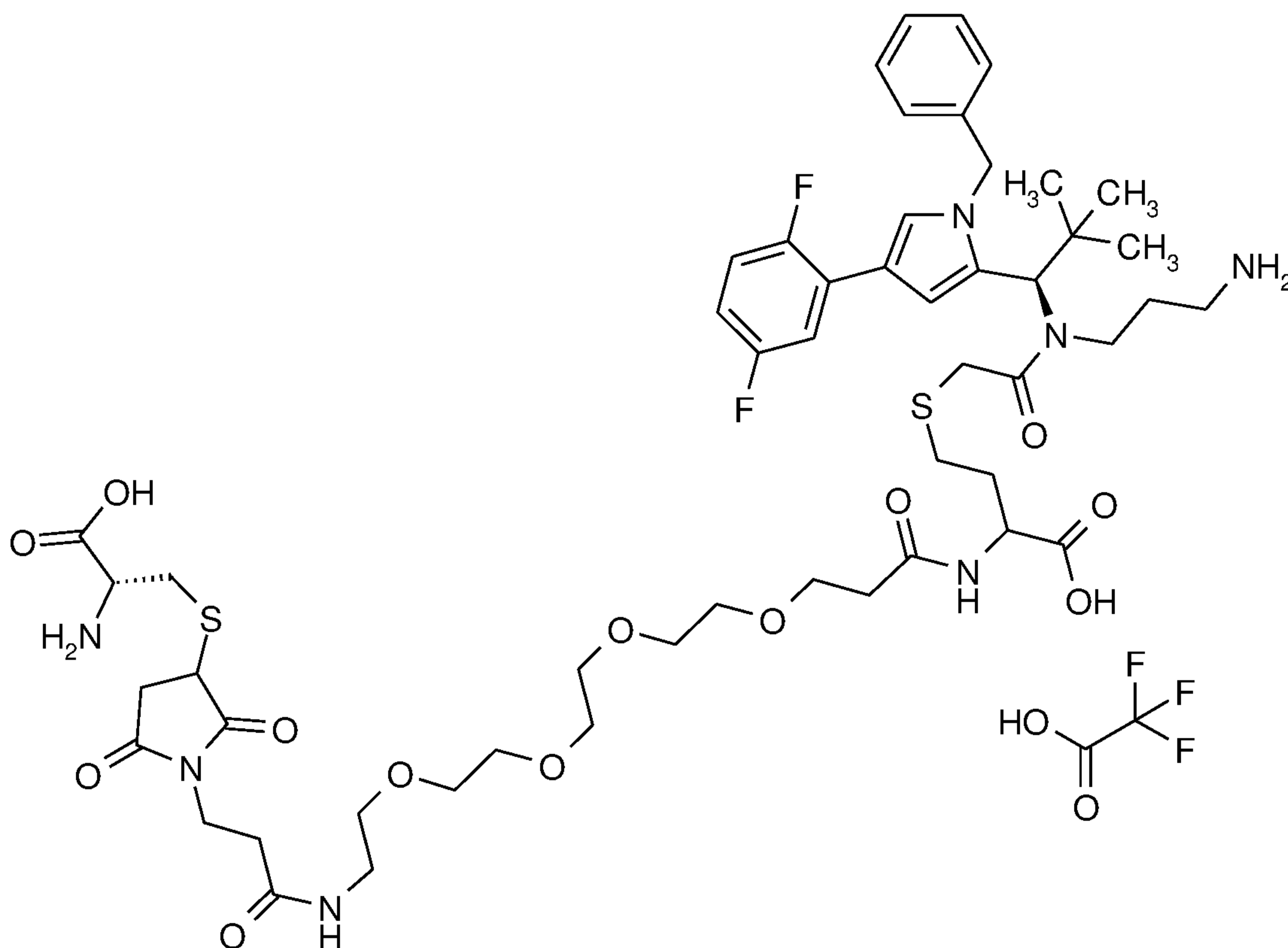
acetonitrile/water, 2 mg (47% of theory) of the title compound as a white foam.

LC-MS (Method 1):  $R_t = 0.76$  min; MS (EIpos):  $m/z = 843$   $[M+H]^+$ .

5

### Example 138

N-[19-(3(R/S)-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-2,5-dioxopyrrolidin-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-  
 10 azanonadecan-1-oyl]-R/S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}homocysteine / trifluoroacetic acid (1:1)



15

6.0 mg (0.01 mmol) of R/S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-  
 20 azanonadecan-1-oyl]-homocysteine / trifluoroacetic acid (1:1) (Intermediate F146) were initially charged in 2.2 ml of

DMF/water (10:1), 2.0 mg (0.02 mmol) of L-cysteine were added and the mixture was stirred at RT for 10 min. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 5.5 mg (76% of theory) of the title compound.

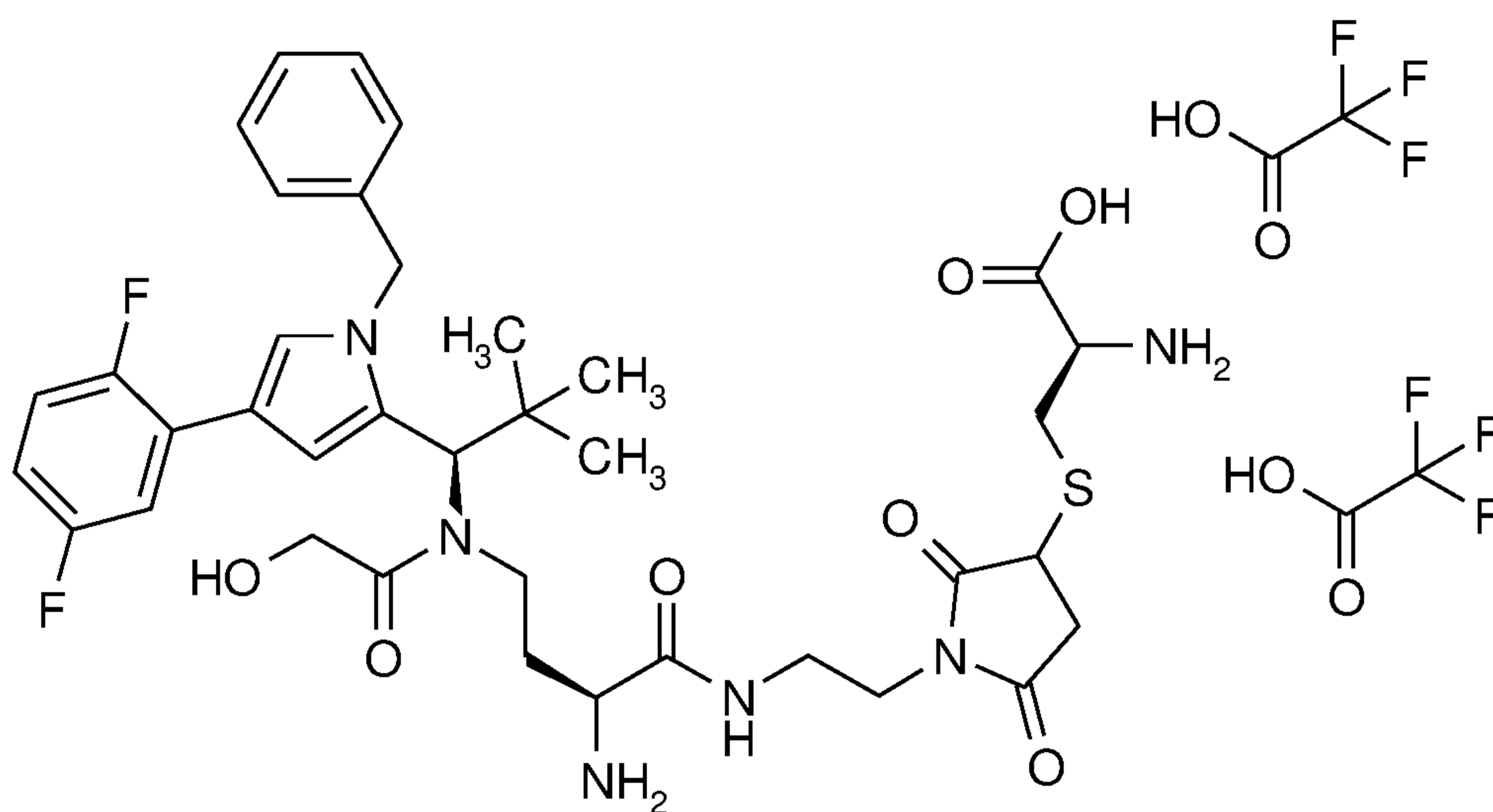
LC-MS (Method 4):  $R_t$  = 1.07 min; MS (ESIpos):  $m/z$  = 1106 (M+H)<sup>+</sup>.

10

### Example 139

S-{(3R/S)-1-[2-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl)amino)ethyl]-2,5-dioxopyrrolidin-3-yl}-L-cysteine/trifluoroacetic acid (1:2)

15



20 The synthesis was carried out analogously to the synthesis of compound Example 138.

6.0 mg (0.01 mmol) of trifluoroacetic acid / (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]butanamide (1:1).

25

2.6 mg (0.02 mmol) of L-cysteine.

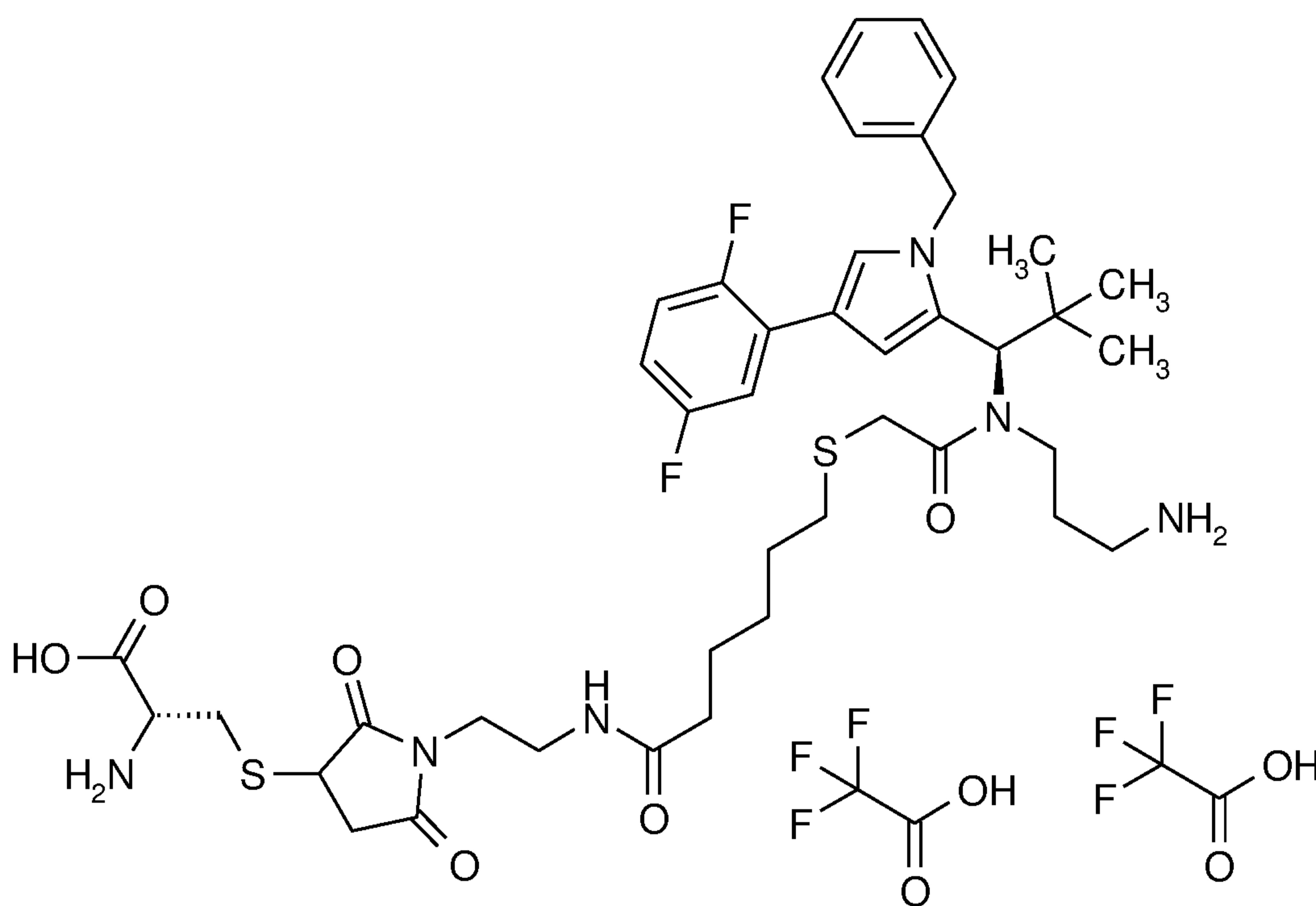
This gave 3.4 mg (43% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.82$  min; MS (ESIpos):  $m/z = 757$  (M+H)<sup>+</sup>.

5

### Example 141

S-[(3R/S)-1-(2-{[6-({2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulphonyl)hexanoyl]amino}ethyl)-2,5-dioxopyrrolidin-3-yl]-L-cysteine / trifluoroacetic acid (1:2)



15

The synthesis was carried out analogously to the synthesis of compound Example 138.

10.07 mg (0.01 mmol) of trifluoroacetic acid / 6-({2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulphonyl)-N-[2-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)ethyl]hexanamide (1:1) (Intermediate F143).

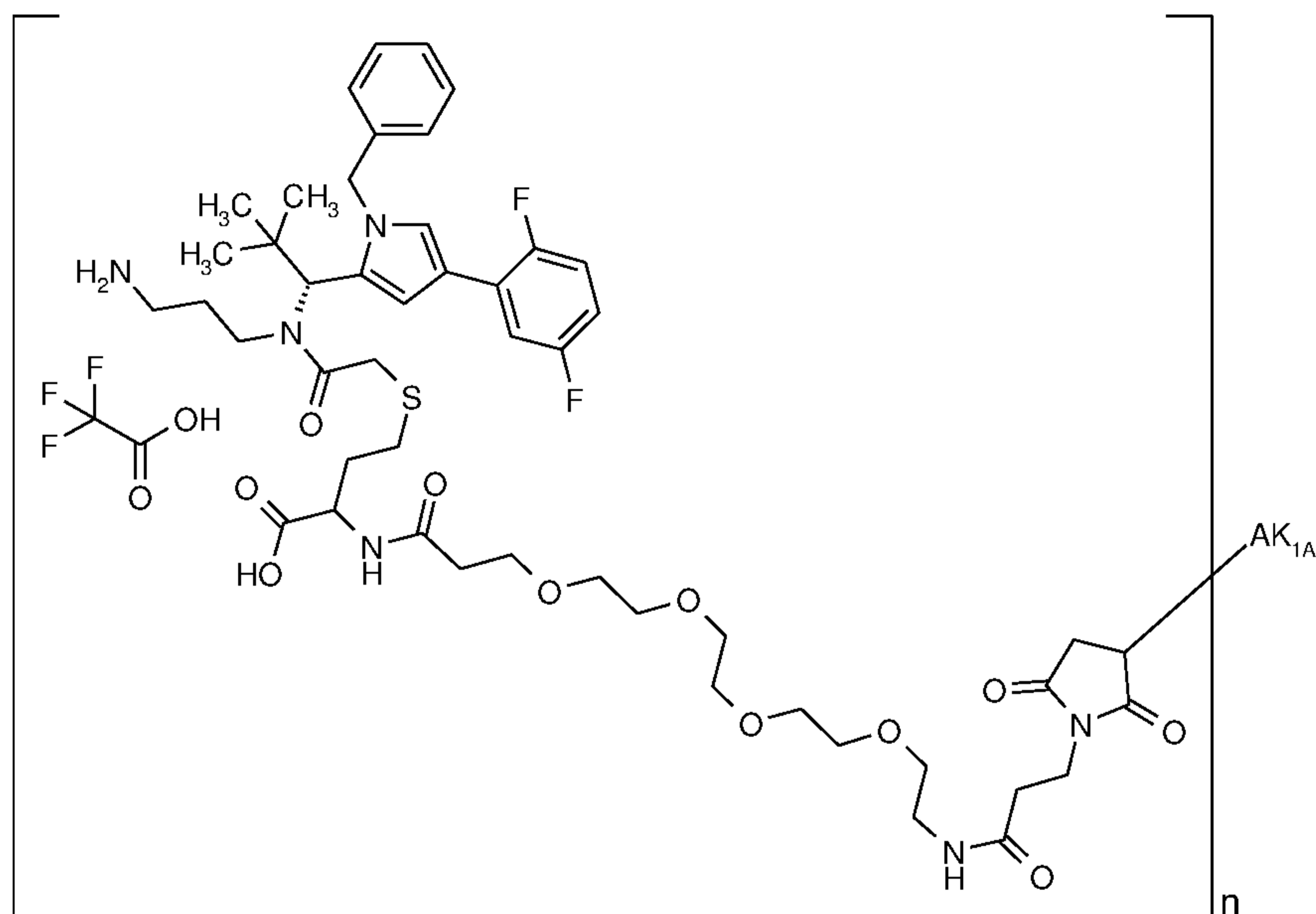
20

9.3 mg (0.08 mmol) of L-cysteine.

This gave 9.2 mg (67% of theory) of the title compound.

5 LC-MS (Method 4):  $R_t = 1.05$  min; MS (ESIpos):  $m/z = 843$  (M+H)<sup>+</sup>.

### Example 142A



10

Here, 5.0 mg of cetuximab in PBS ( $c=12.33$  mg/ml) were used for coupling with Intermediate F142, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 2.08 mg/ml

20 Drug/mAb ratio: 2.1

### Example 142B

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS ( $c=34.42$  mg/ml) were used for coupling with Intermediate F142, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be

25

present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.84 mg/ml

5

Drug/mAb ratio: 2.0

#### Example 142E

10 Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F142, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain  
15 succinamides attached to the antibody.

Protein concentration: 1.92 mg/ml

Drug/mAb ratio: 2.3

20

#### Example 142I

Here, 5.0 mg of nimotuzumab in PBS (c=13.8 mg/ml) were used for coupling with Intermediate F142, and the reaction was, after  
25 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

30 Protein concentration: 1.92 mg/ml

Drug/mAb ratio: 2.5

#### Example 142H

35

Here, 5.0 mg of panitumumab in PBS (c = 2.1 mg/ml) were used for coupling with Intermediate F142. The reduction time with TCEP was 1 h and the stirring time for the ADC coupling was 1.5 h.



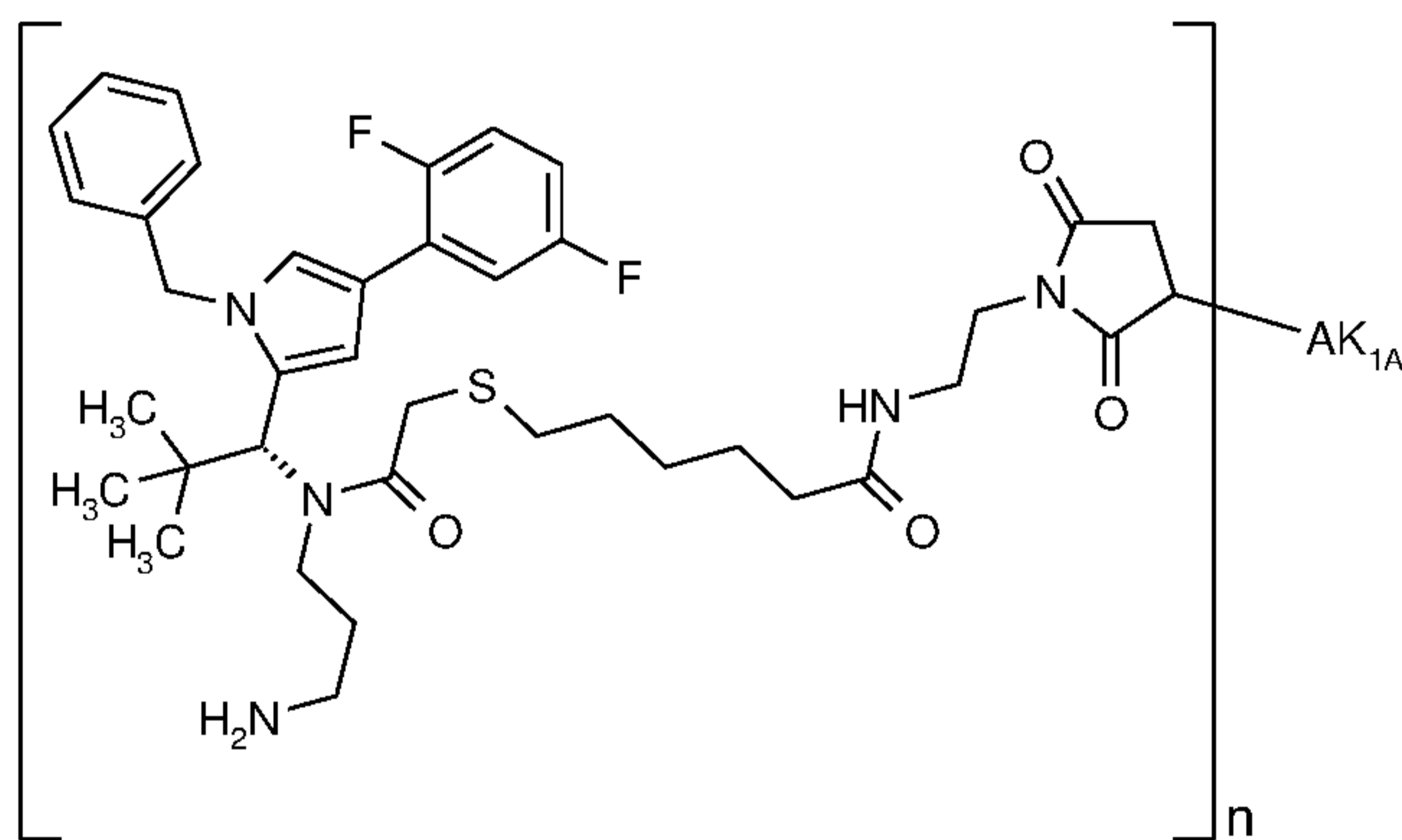
After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 1.7 mg/ml

Drug/mAb ratio: 2.2

### 10 Example 143A



Here, 5.0 mg of cetuximab in PBS ( $c=12.33$  mg/ml) were used for coupling with Intermediate F143, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

20

Protein concentration: 1.84 mg/ml

Drug/mAb ratio: 2.5

### 25 Example 143B

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS ( $c=34.42$  mg/ml) were used for coupling with Intermediate F143, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides

30

attached to the antibody.

Protein concentration: 1.05 mg/ml

5 Drug/mAb ratio: 1.6

### Example 143E

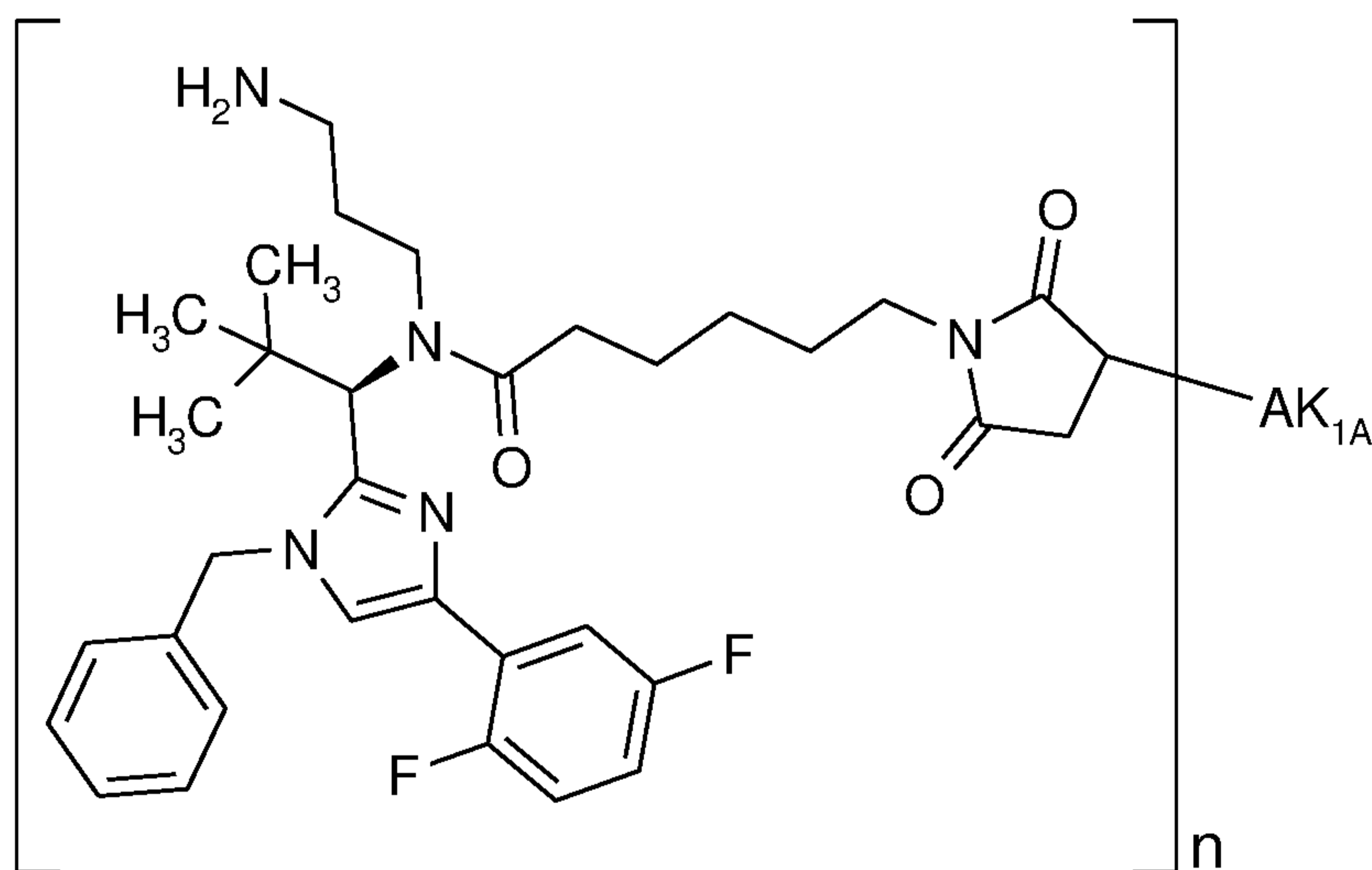
10 Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F143, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 1.95 mg/ml

Drug/mAb ratio: 2.3

### 20 Example 144B

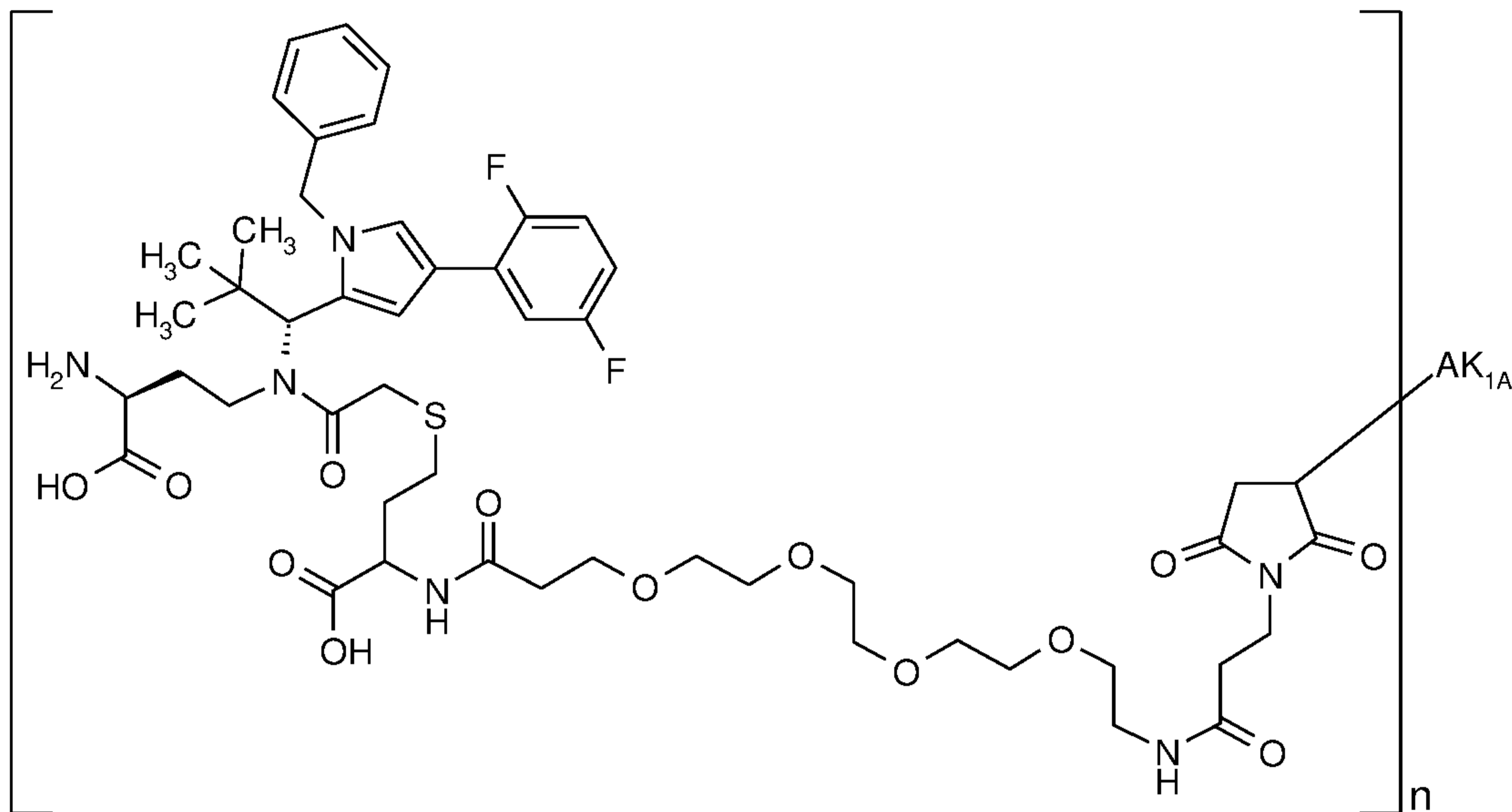


25 Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=12.87 mg/ml) were used for coupling with Intermediate F144, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted.

Protein concentration: 1.53 mg/ml

Drug/mAb ratio: 2.0

5 **Example 146A**



10 Here, 5.0 mg of cetuximab in PBS (c=12.33 mg/ml) were used for coupling with Intermediate F146, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 2.02 mg/ml

Drug/mAb ratio: 2.4

20 **Example 146B**

25 Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=34.42 mg/ml) were used for coupling with Intermediate F146, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.87 mg/ml

Drug/mAb ratio: 2.4

5

**Example 146E**

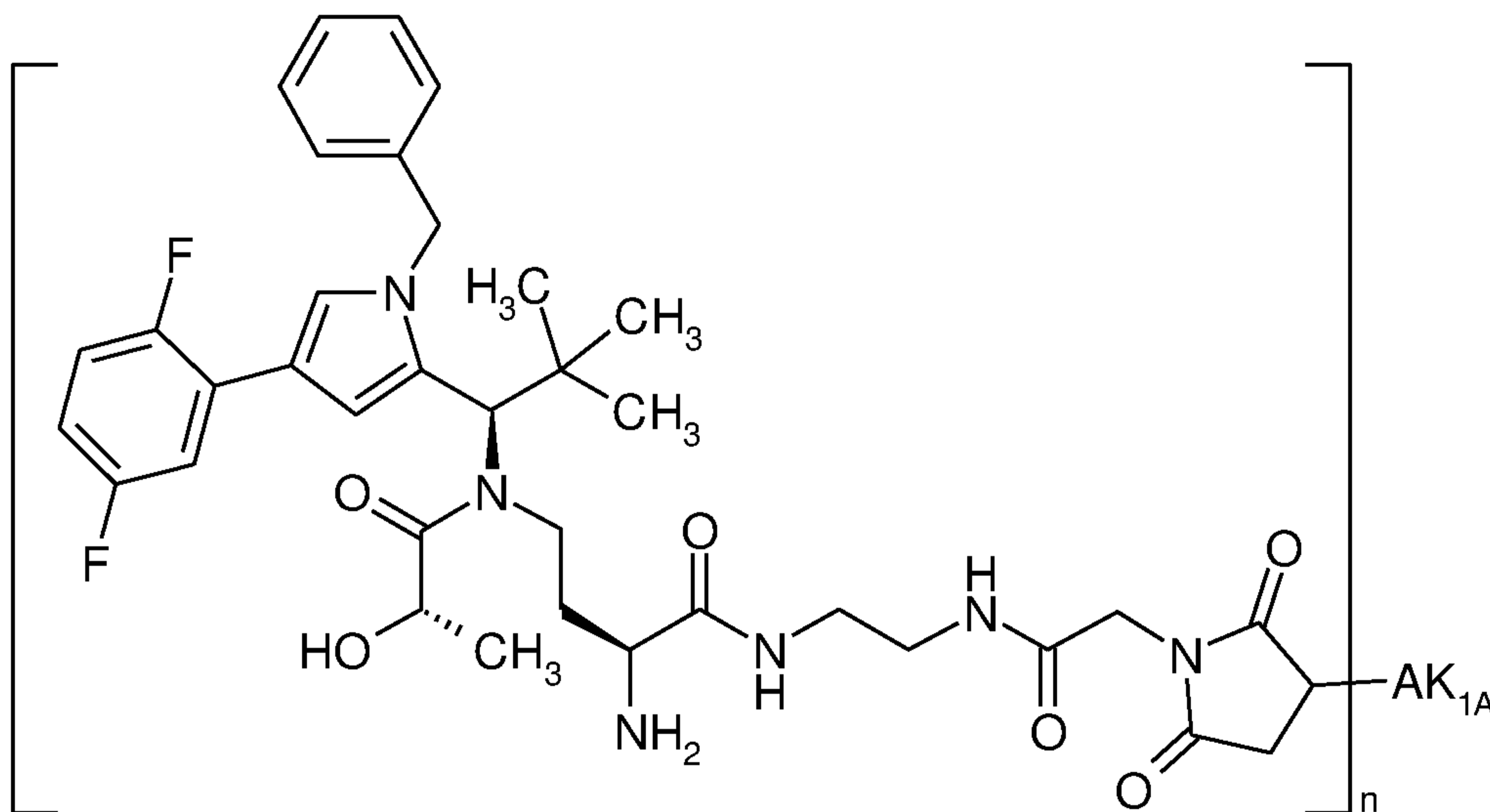
Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F146, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15 Protein concentration: 1.92 mg/ml

Drug/mAb ratio: 2.5

**Example 153A**

20



Here, 5 mg of cetuximab in PBS (c=21.32 mg/ml) were used for coupling with Intermediate F153, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.93 mg/ml

Drug/mAb ratio: 3.2

5

**Example 153B**

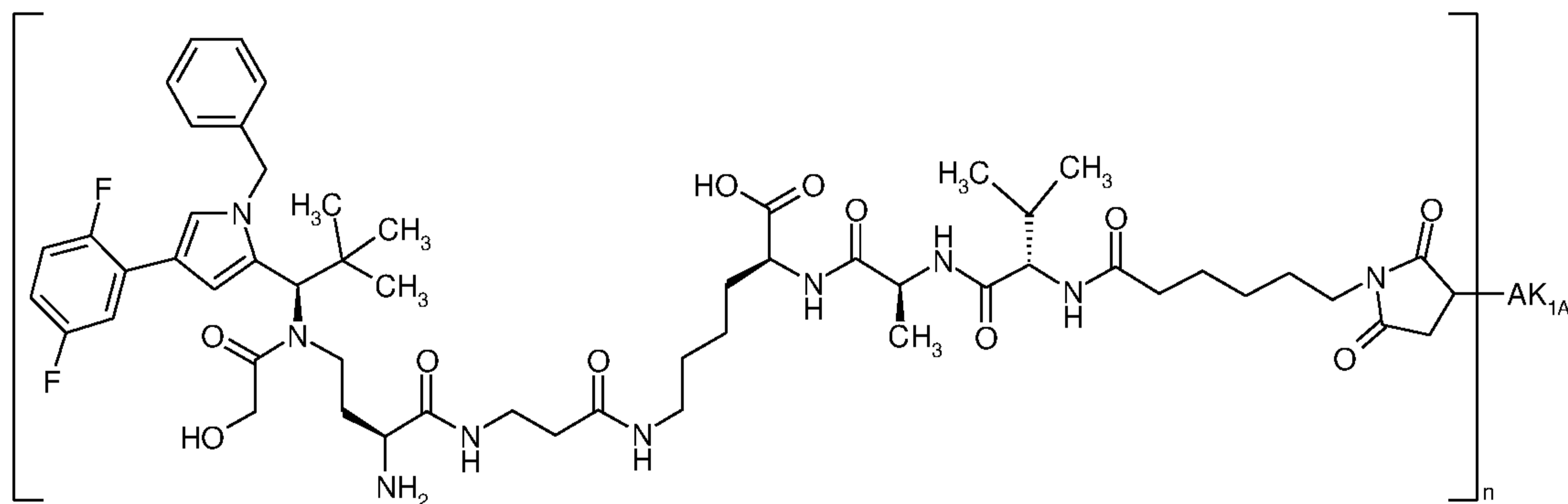
Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F153, and the reaction was, after  
 10 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15 Protein concentration: 1.71 mg/ml

Drug/mAb ratio: 2.8

**Example 155A**

20



Here, 50 mg of cetuximab in PBS (c=8.51 mg/ml) were used for coupling with Intermediate F155, and the reaction was, after  
 25 Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again.

Protein concentration: 14.85 mg/ml

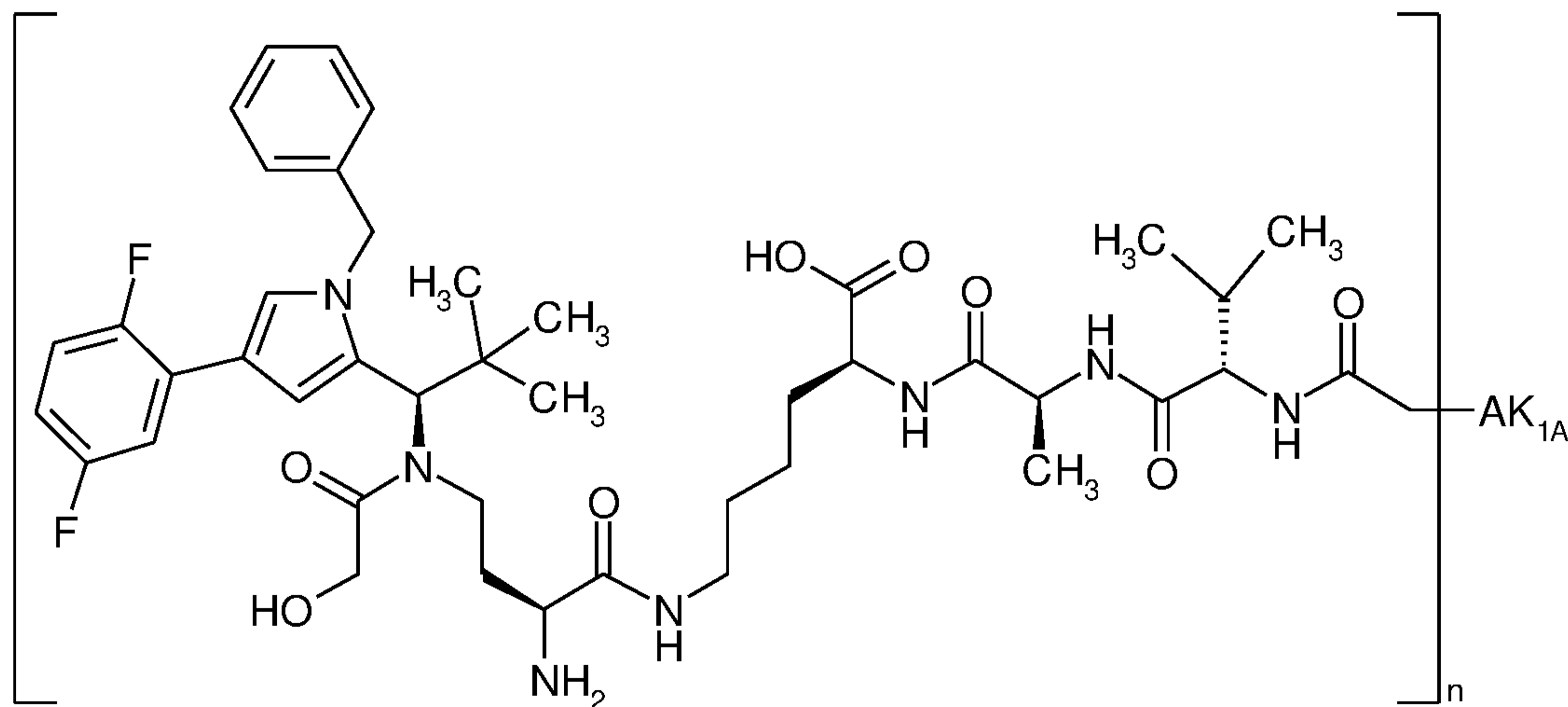
30 Drug/mAb ratio: 2.5

**Example 155B**

Here, 40 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F155, and the reaction was, after  
5 Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again.

Protein concentration: 11.25 mg/ml

10 Drug/mAb ratio: 3.1

**Example 156A**

15

Here, 5 mg of cetuximab in PBS (c=21.3 mg/ml) were used for coupling with Intermediate F156. After TCEP reduction, coupling with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After  
20 Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.83 mg/ml

25 Drug/mAb ratio: 3.6

**Example 156B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F156, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

5

Protein concentration: 1.79 mg/ml

Drug/mAb ratio: 3.9

#### 10 Example 156E

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F156, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

15

Protein concentration: 1.91 mg/ml

Drug/mAb ratio: 4.2

20

#### Example 157

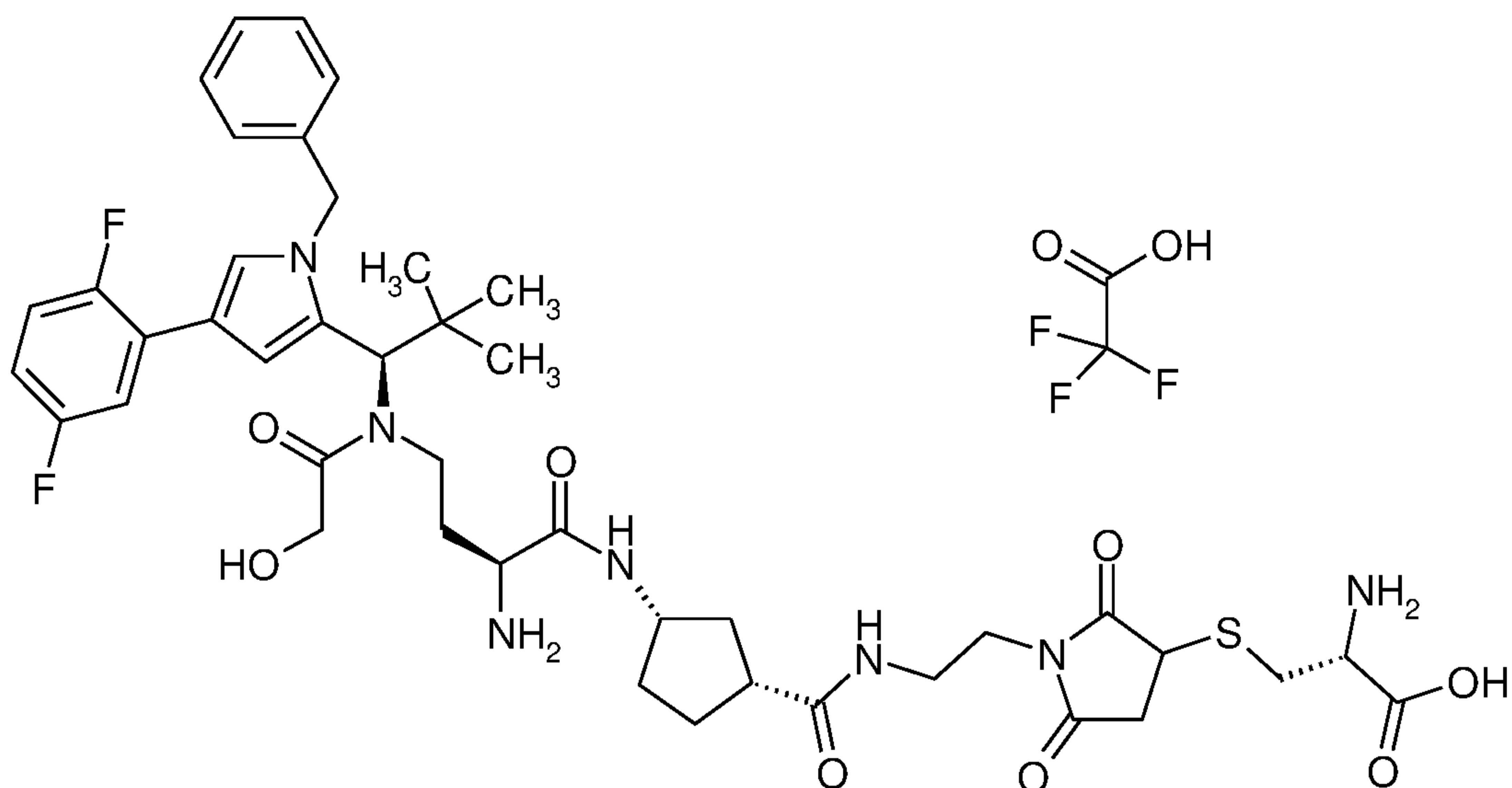
S-{1-[2-({[(1R,3S)-3-({(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-

(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-

25 dimethylpropyl}(glycoloyl)amino]butanoyl}amino)cyclopentyl]car

bonyl}amino)ethyl]-2,5-dioxopyrrolidin-3-yl}-L-cysteine /

trifluoroacetic acid (1:1)



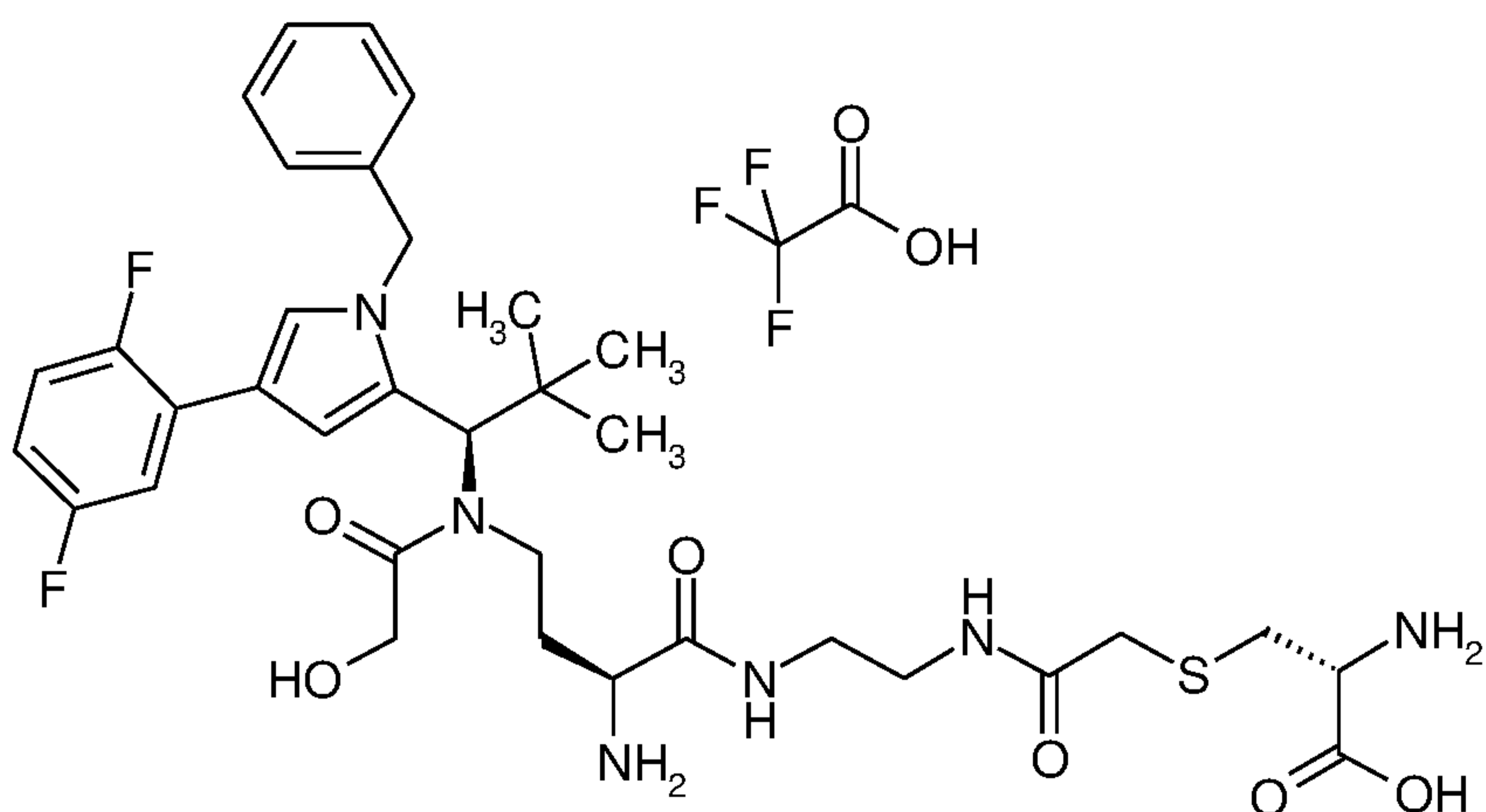
2 mg (2  $\mu$ mol) of Intermediate F125 were taken up in 2 ml of DMF/water 10:1, and 0.8 mg (6  $\mu$ mol) of L-cysteine were added.  
 5 The reaction mixture was stirred at RT for 20 h, then concentrated under reduced pressure and then purified by preparative HPLC.

LC-MS (Method 1):  $R_t$  = 0.81 min; MS (EIpos):  $m/z$  = 868  $[M+H]^+$ .

10

### Example 158

S-(2-([2-((2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino)-2-oxoethyl)-L-cysteine/trifluoroacetic acid (1:1)





6 mg (8  $\mu\text{mol}$ ) of Intermediate F119 were taken up in 3 ml of DMF, and 1.8 mg (15  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 6 h and then allowed to stand at RT for 3 days. The reaction was then concentrated under reduced pressure, and the product was purified by preparative HPLC.

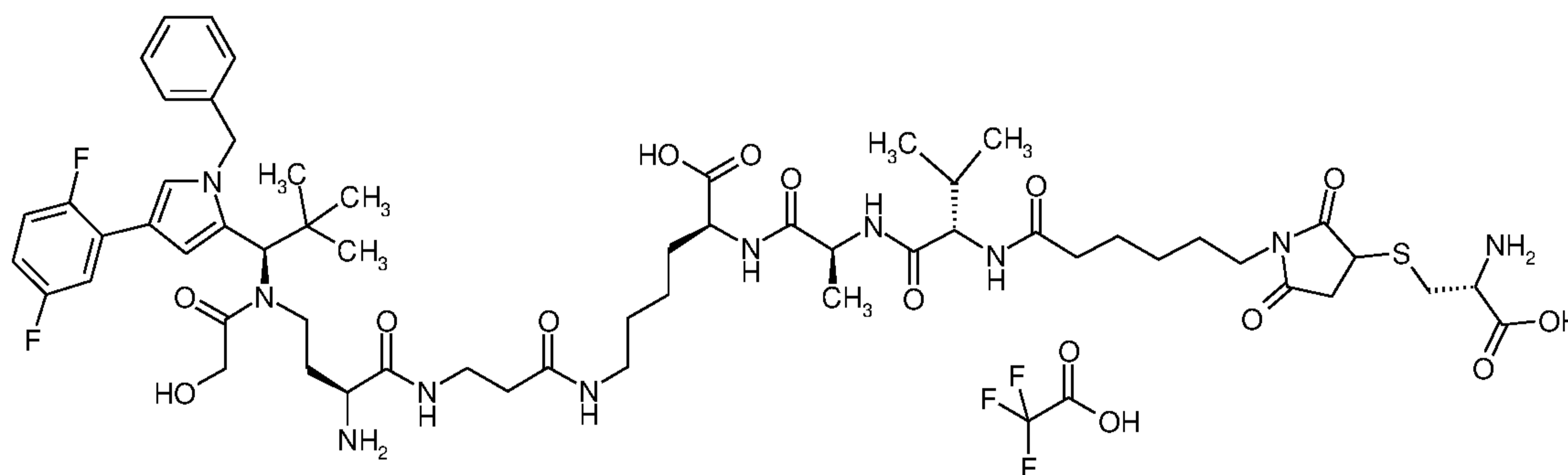
LC-MS (Method 1):  $R_t = 0.81$  min; MS (ESIpos):  $m/z = 717$  (M+H)<sup>+</sup>.

### Example 160

10

N<sup>6</sup>- (N- { (2S)-2-Amino-4- [ { (1R)-1- [1-benzyl-4- (2,5-difluorophenyl)-1H-pyrrol-2-yl] -2,2-dimethylpropyl } (glycoloyl) amino] butanoyl } -beta-alanyl) -N<sup>2</sup>- { N- [6- (3- { [(2R)-2-amino-2-carboxyethyl] sulfanyl } -2,5-dioxopyrrolidin-1-yl) hexanoyl] -L-valyl-L-alanyl } -L-lysine / trifluoroacetic acid (1:1)

15



20 4 mg (3  $\mu\text{mol}$ ) of Intermediate F155 were taken up in 2.5 ml of DMF/water 10:1, and 1.2 mg (10  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 30 min, then concentrated under reduced pressure, taken up in acetonitrile/water 1:1 and then purified by preparative HPLC.

25

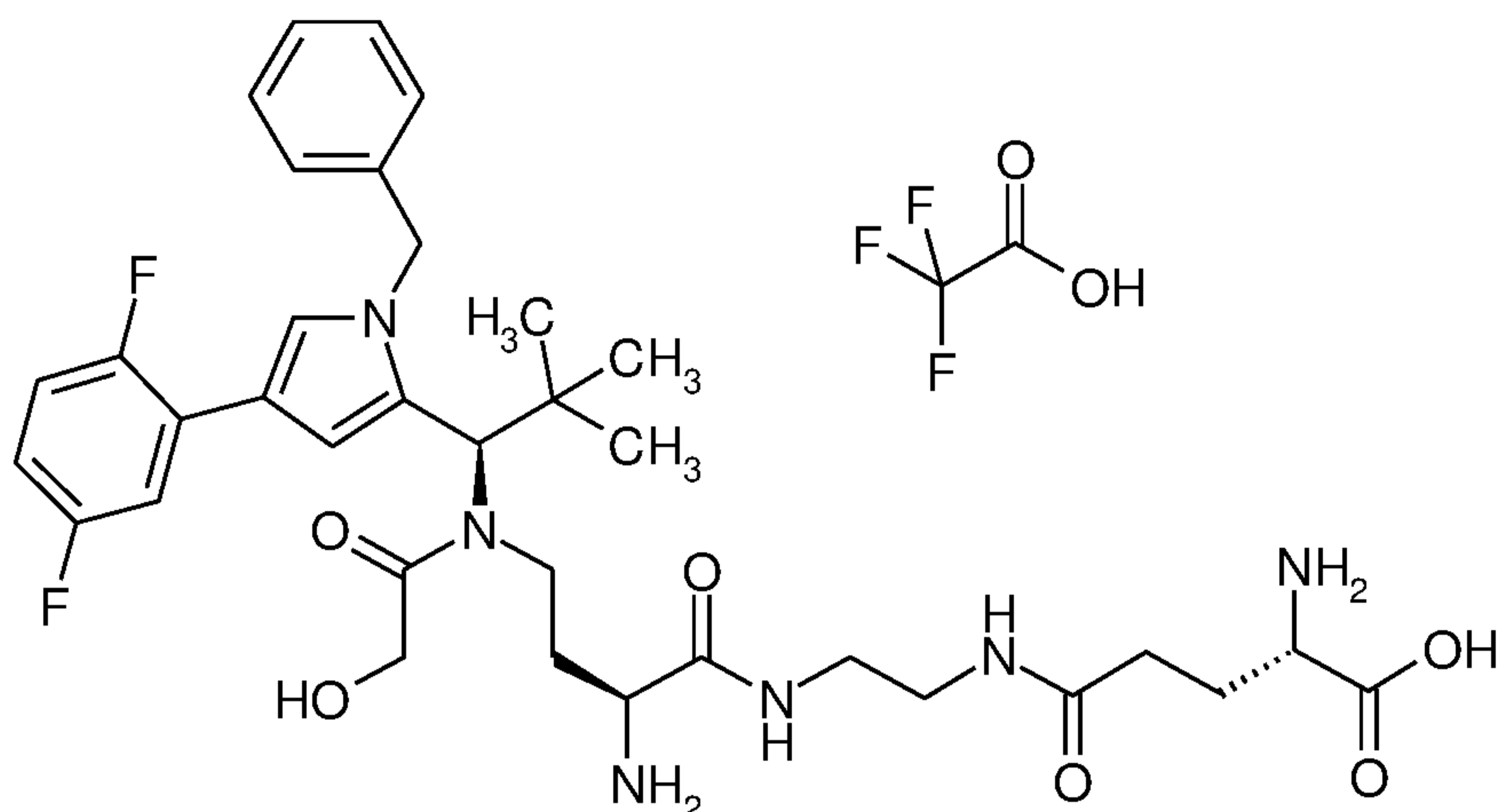
LC-MS (Method 1):  $R_t = 0.81$  min; MS (EIpos):  $m/z = 1197$  [M+H]<sup>+</sup>.

### Example 161

30

N- [2- ( { (2S)-2-Amino-4- [ { (1R)-1- [1-benzyl-4- (2,5-difluorophenyl)-1H-pyrrol-2-yl] -2,2-dimethylpropyl } (glycoloyl) amino] butanoyl } amino) ethyl] -L-

glutamine / trifluoroacetic acid (1:1)



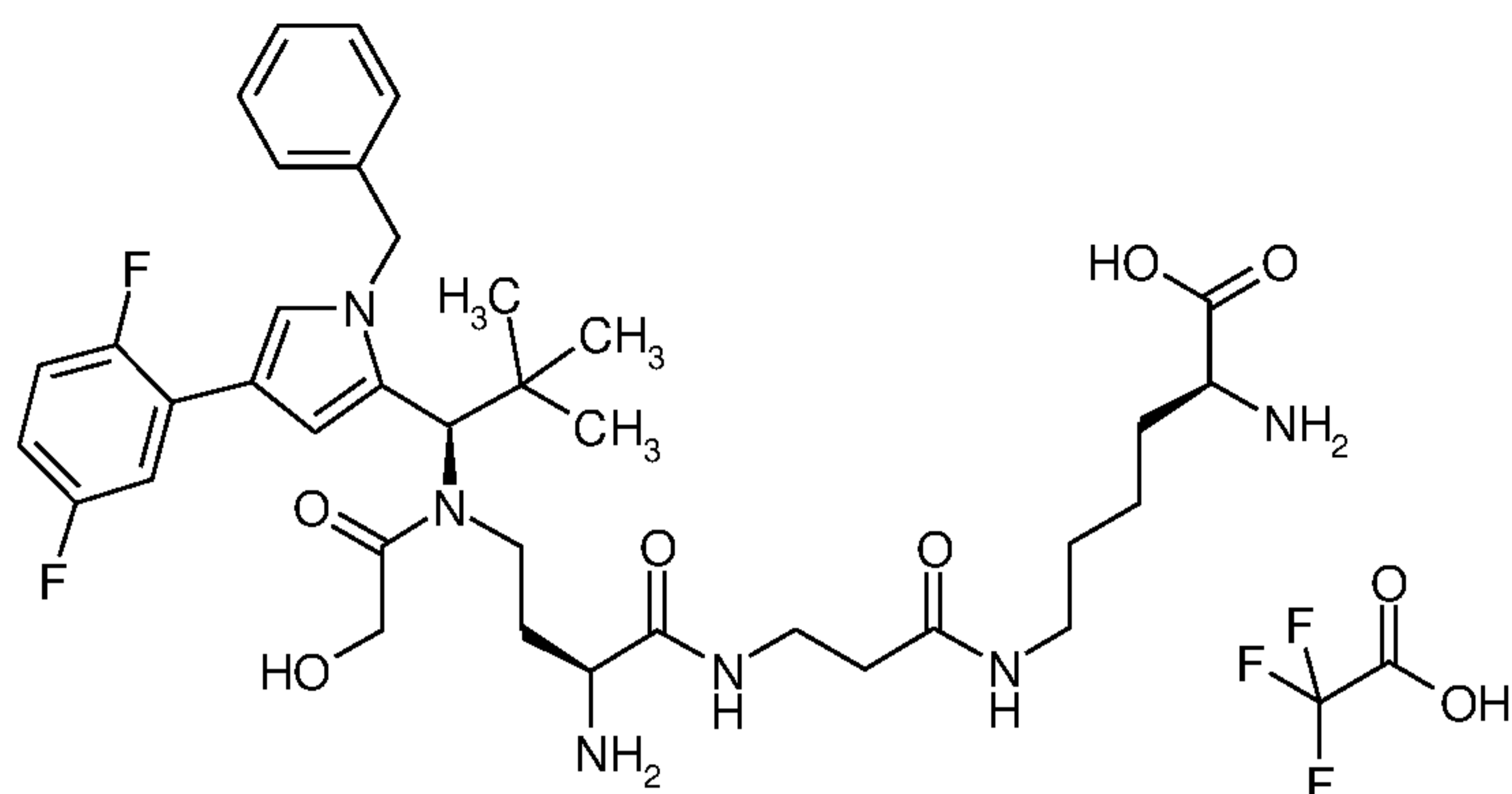
5 First, trifluoroacetic acid / benzyl N-(2-aminoethyl)-N<sup>2</sup>-  
 [(benzyloxy)carbonyl]-L-glutamate (1:1) was prepared using  
 classical methods of peptide chemistry. In the presence of HATU,  
 this intermediate was then coupled with Intermediate C58.  
 Subsequently, first the benzyloxycarbonyl protective group and  
 10 the benzyl ester were removed by hydrogenolytic cleavage, and  
 then the 2-(trimethylsilyl)ethoxycarbonyl protective group was  
 removed using zinc chloride.

LC-MS (Method 6):  $R_t = 1.91$  min; MS (EIpos):  $m/z = 685$  [M+H]<sup>+</sup>.

15

### Example 162

N<sup>6</sup>-(N-{(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-  
 difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 20 dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanyl)-L-  
 lysine / trifluoroacetic acid (1:1)



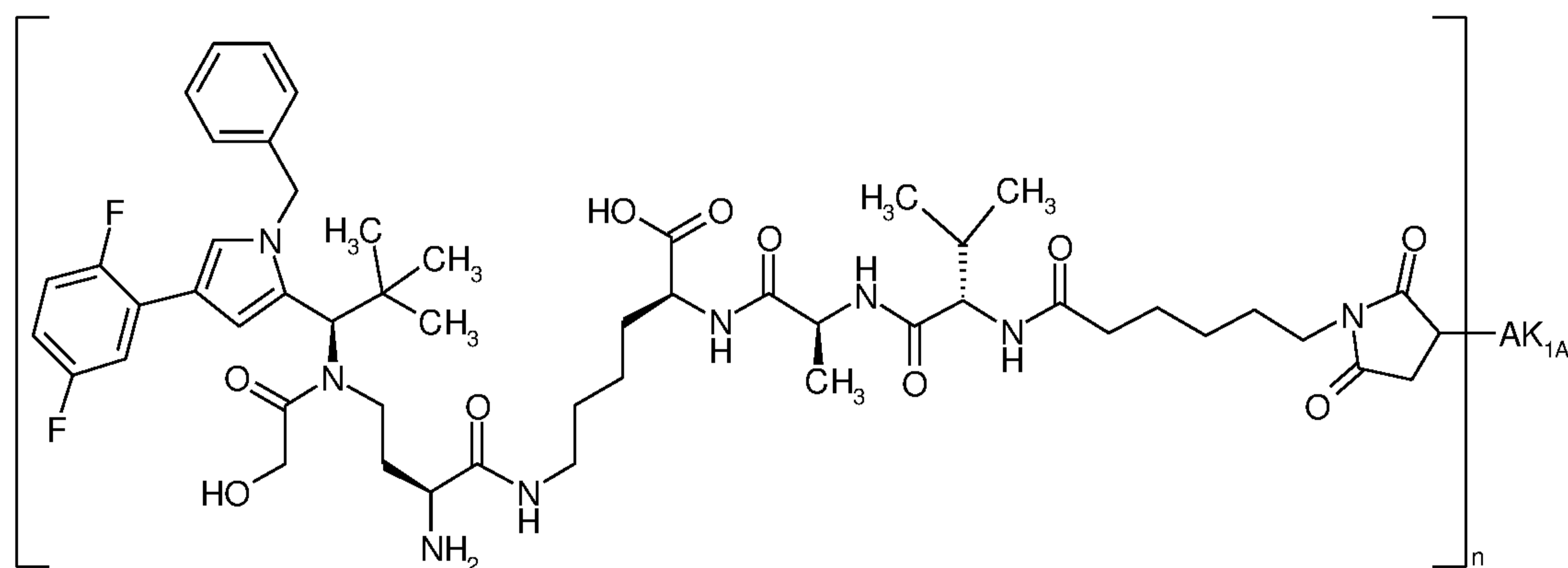
Initially, trifluoroacetic acid / 2-(trimethylsilyl)ethyl-N2-  
 [(benzyloxy)carbonyl]-L-lysinate (1:1) was prepared using  
 5 classical protective group operations known in peptide  
 chemistry. In the presence of HATU, this intermediate was then  
 coupled with Intermediate C61. Subsequently, first the 2-  
 (trimethylsilyl)ethoxycarbonyl protective group and the 2-  
 (trimethylsilyl)ethyl ester were cleaved using zinc chloride.  
 10 Finally, the title compound was obtained by hydrogenolytical  
 cleavage of the benzyloxycarbonyl protective group and  
 purification by preparative HPLC.

HPLC (Method 11):  $R_t = 1.65$  min;

15

LC-MS (Method 1):  $R_t = 0.76$  min; MS (EIpos):  $m/z = 713$   $[M+H]^+$ .

### Example 163A



20

Here, 5 mg of cetuximab in PBS ( $c=11.3$  mg/ml) were used for  
 coupling with Intermediate F163, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and

rediluted with PBS.

Protein concentration: 2.02 mg/ml

5 Drug/mAb ratio: 3.3

**Example 163B**

10 Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F163, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

15 Protein concentration: 1.76 mg/ml

Drug/mAb ratio: 2.5

**Example 163H**

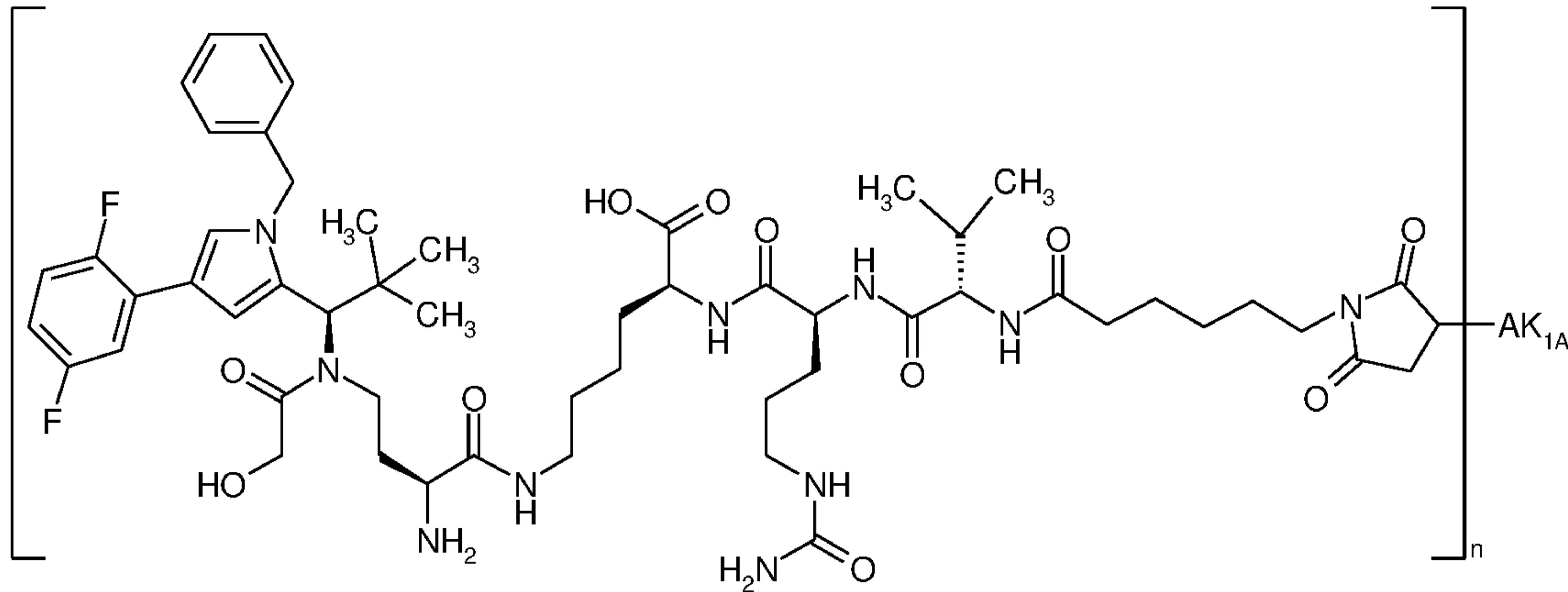
20 Here, 5.0 mg of panitumumab in PBS (c = 10 mg/ml) were used for coupling with Intermediate F163. The reduction time with TCEP was 30 min and the stirring time for the ADC coupling was 2 h. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

25

Protein concentration: 1.61 mg/ml

Drug/mAb ratio: 2.6

30 **Example 164A**



Here, 5 mg of cetuximab in PBS (c=16.9 mg/ml) were used for coupling with Intermediate F164, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.15 mg/ml

10 Drug/mAb ratio: 3.5

#### Example 164B

Here, 30 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used  
 15 for coupling with Intermediate F164, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again.

Protein concentration: 14.8 mg/ml

20

Drug/mAb ratio: 2.8

#### Example 164E

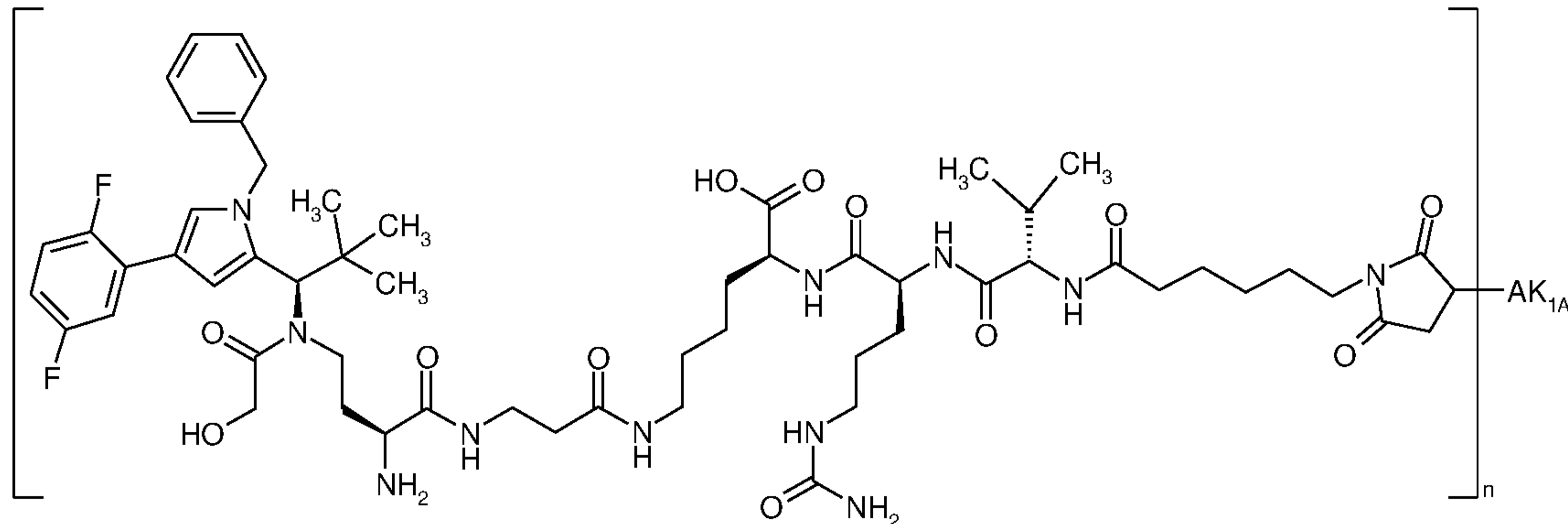
25 Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F164, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

30 Protein concentration: 2.11 mg/ml

Drug/mAb ratio: 3.8

**Example 165A**

5



Here, 5 mg of cetuximab in PBS (c=16.9 mg/ml) were used for coupling with Intermediate F165, and the reaction was, after  
 10 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.93 mg/ml

15 Drug/mAb ratio: 3.4

**Example 165B**

Here, 40 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used  
 20 for coupling with Intermediate F165, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again.

Protein concentration: 12.02 mg/ml

25

Drug/mAb ratio: 3.3

**Example 165E**

30 Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F165, and the reaction was, after

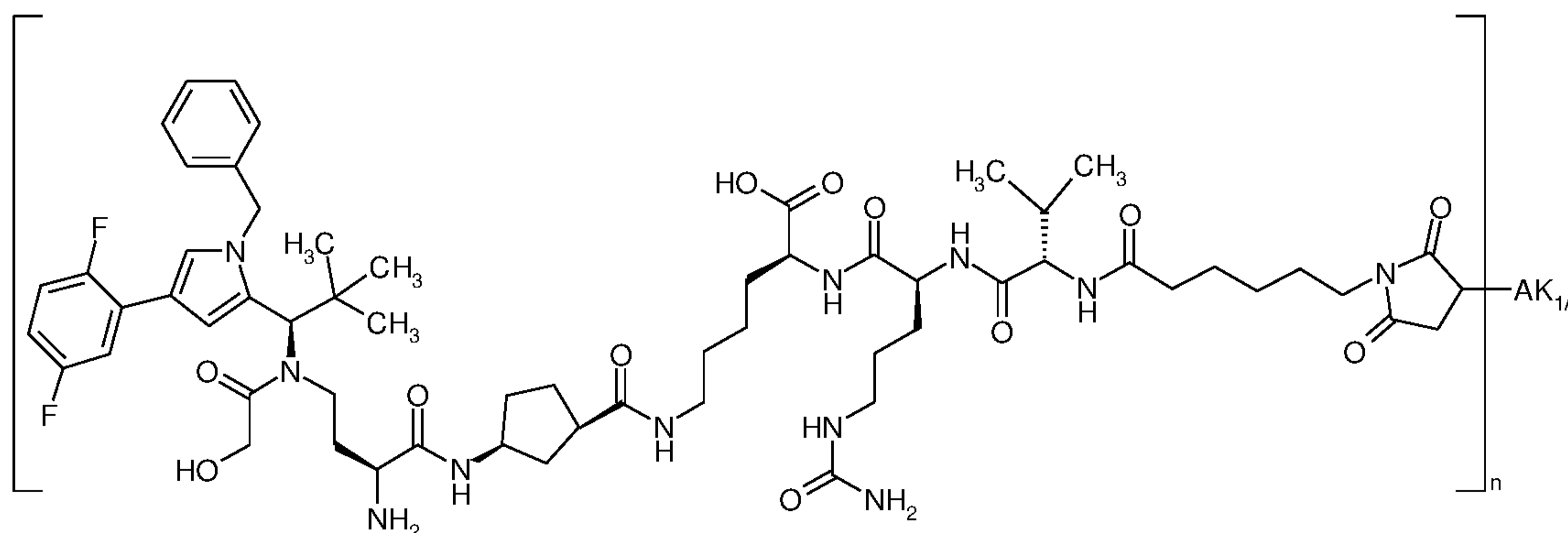
Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.94 mg/ml

5

Drug/mAb ratio: 3.5

### Example 166A



10

Here, 5 mg of cetuximab in PBS ( $c=21.3$  mg/ml) were used for coupling with Intermediate F166, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

15

Protein concentration: 2.0 mg/ml

Drug/mAb ratio: 3.0

20

### Example 166B

Here, 5 mg of anti-TWEAKR AK-1 in PBS ( $c=18.6$  mg/ml) were used for coupling with Intermediate F166, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

25

Protein concentration: 1.76 mg/ml

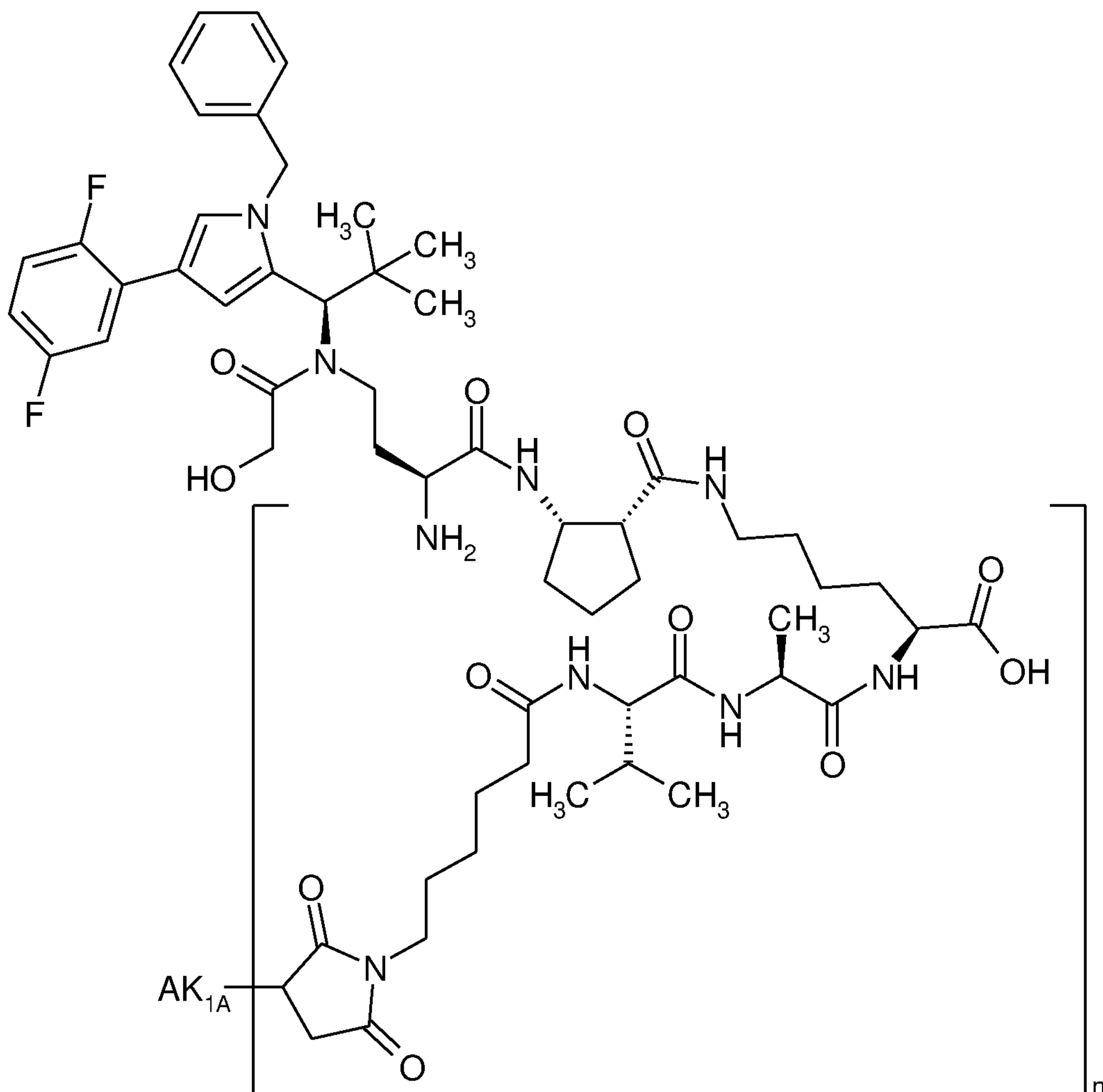
30 Drug/mAb ratio: 3.4

Example 166E

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F166, and the reaction was, after 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.01 mg/ml

10 Drug/mAb ratio: 3.6

Example 168A

15

Here, 5 mg of cetuximab in PBS (c=11.3 mg/ml) were used for coupling with Intermediate F168, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and



rediluted with PBS.

Protein concentration: 1.94 mg/ml

5 Drug/mAb ratio: 3.0

#### **Example 168B**

10 Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=12.9 mg/ml) were used for coupling with Intermediate F168, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.33 mg/ml

15

Drug/mAb ratio: 2.8

#### **Example 168E**

20 Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F168, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

25 Protein concentration: 1.89 mg/ml

Drug/mAb ratio: 3.0

#### **Example 168H**

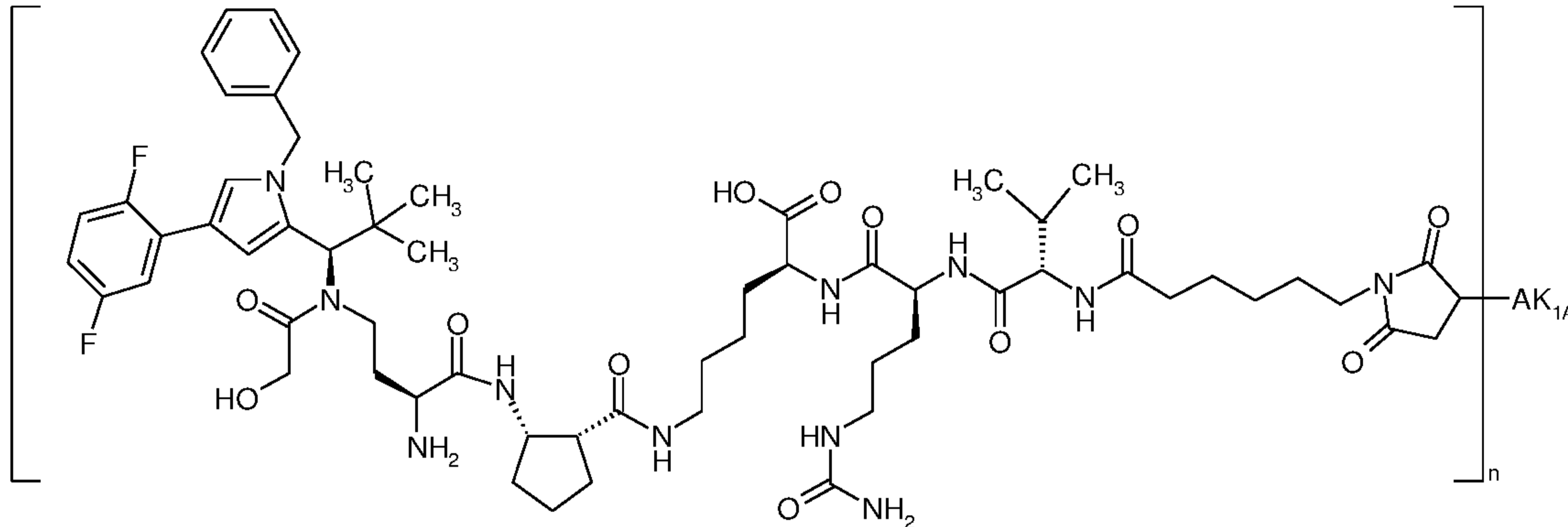
30

Here, 5.0 mg of panitumumab in PBS (c = 10 mg/ml) were used for coupling with Intermediate F168. The reduction time with TCEP was 4 h and the stirring time for the ADC coupling was 20 h. After Sephadex purification, the reaction was concentrated by  
35 ultracentrifugation and rediluted with PBS.

Protein concentration: 1.76 mg/ml

Drug/mAb ratio: 2.8

**Example 169A**



5

Here, 5 mg of cetuximab in PBS (c=16.9 mg/ml) were used for coupling with Intermediate F169, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

10

Protein concentration: 1.98 mg/ml

Drug/mAb ratio: 3.4

15

**Example 169B**

Here, 40 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F169, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again.

20

Protein concentration: 11.2 mg/ml

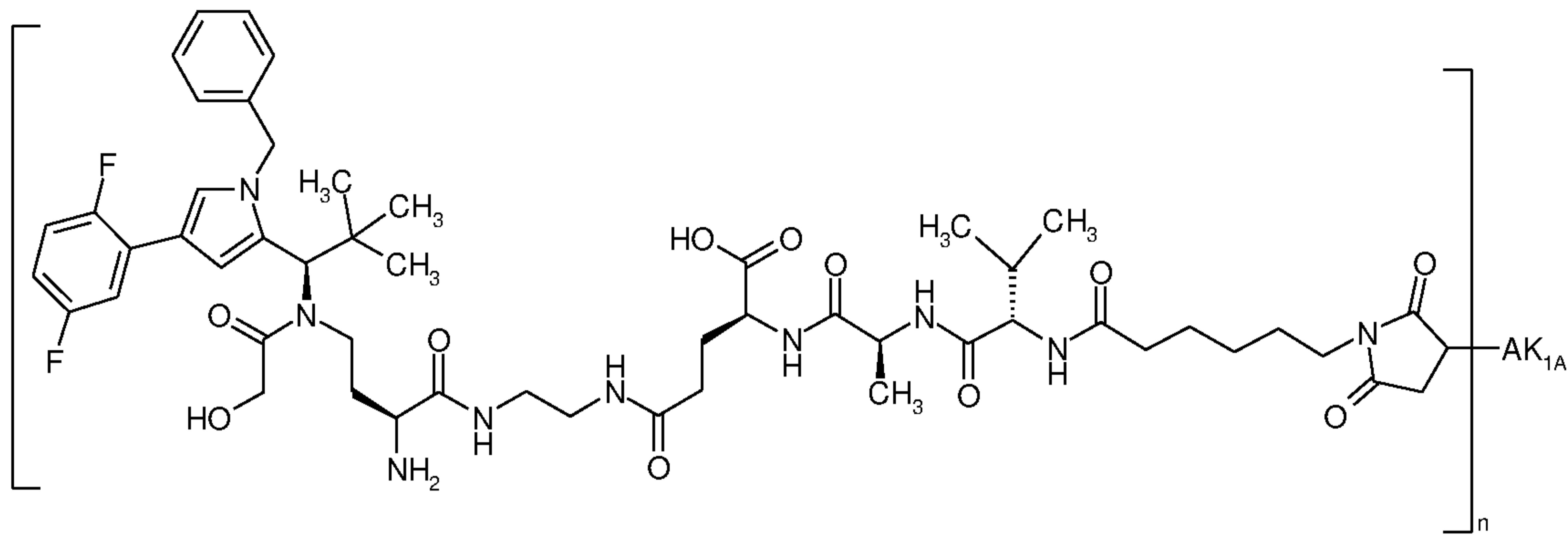
25 Drug/mAb ratio: 2.4

**Example 169E**

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F169, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and

30





Here, 5 mg of cetuximab in PBS (c=11.3 mg/ml) were used for  
 5 coupling with Intermediate F173, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

Protein concentration: 2.1 mg/ml

10

Drug/mAb ratio: 3.6

#### Example 173B

15 Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used  
 for coupling with Intermediate F173, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation,  
 rediluted with PBS and concentrated again.

20 Protein concentration: 12.26 mg/ml

Drug/mAb ratio: 3.4

#### Example 173E

25

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for  
 coupling with Intermediate F173, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

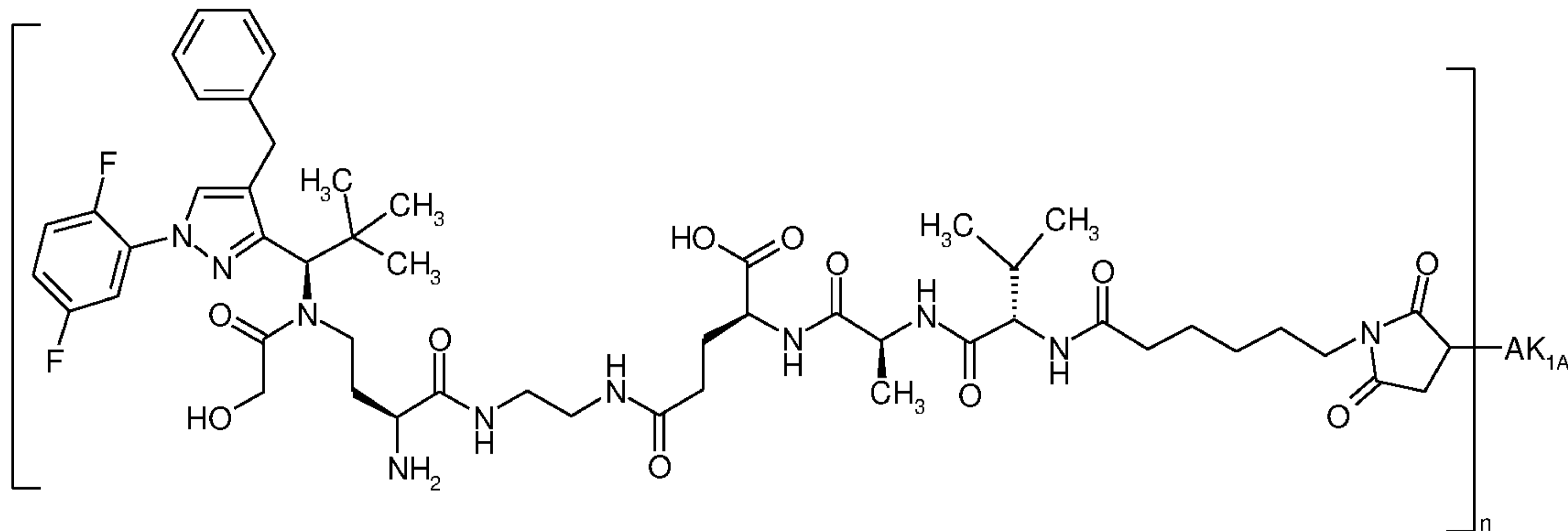
30

Protein concentration: 2.33 mg/ml

Drug/mAb ratio: 3.9

**Example 174A**

5



Here, 5 mg of cetuximab in PBS (c=11.3 mg/ml) were used for coupling with Intermediate F174, and the reaction was, after  
 10 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.18 mg/ml

15 Drug/mAb ratio: 3.1

**Example 174B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used  
 20 for coupling with Intermediate F174, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 1.99 mg/ml

25

Drug/mAb ratio: 3.1

**Example 174E**

30 Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F174, and the reaction was, after

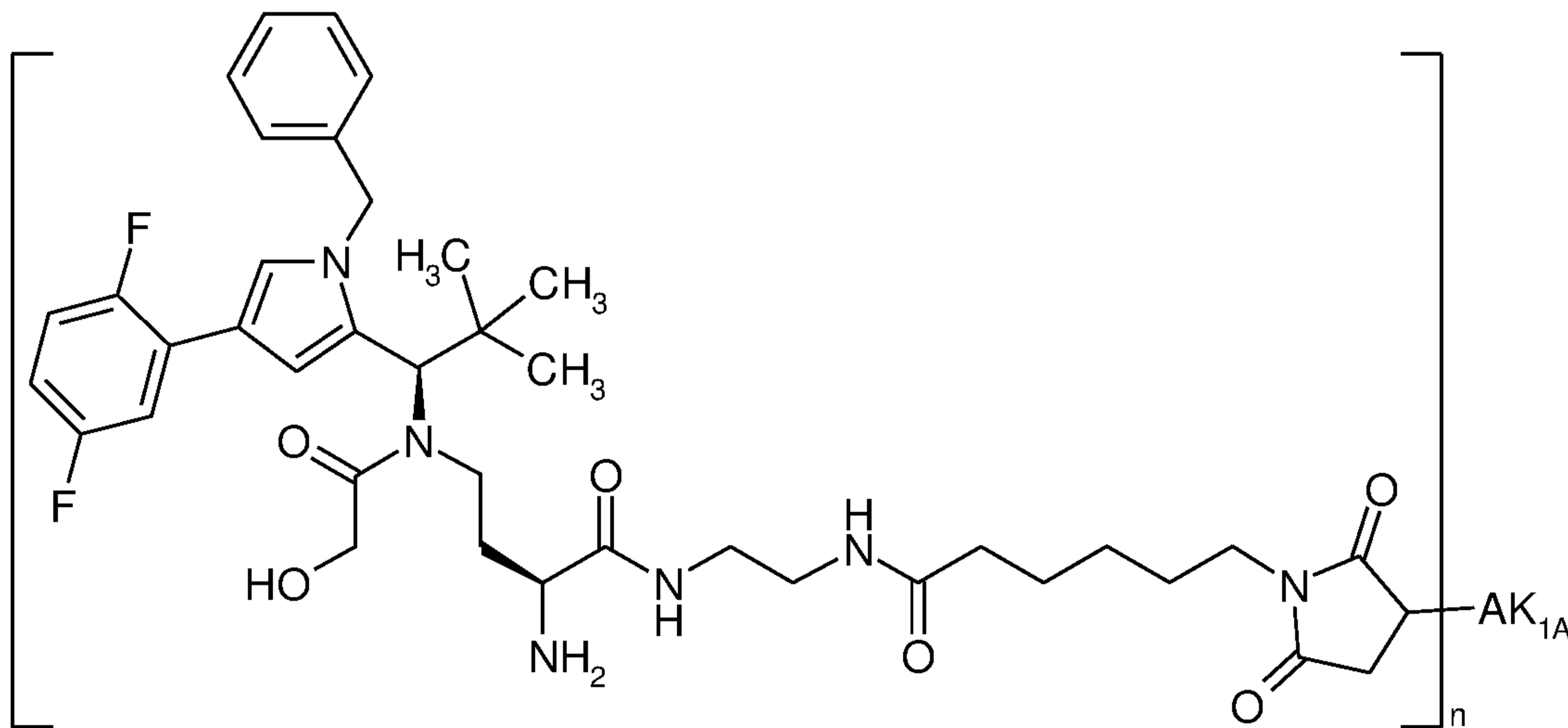
Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.03 mg/ml

5

Drug/mAb ratio: 3.5

**Example 175A**



10

Here, 5 mg of cetuximab in PBS (c=16.9 mg/ml) were used for coupling with Intermediate F175, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

15

Protein concentration: 1.98 mg/ml

Drug/mAb ratio: 3.8

20

**Example 175B**

Here, 40 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F175, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again.

25

Protein concentration: 9.8 mg/ml

Drug/mAb ratio: 2.8

**Example 175E**

5

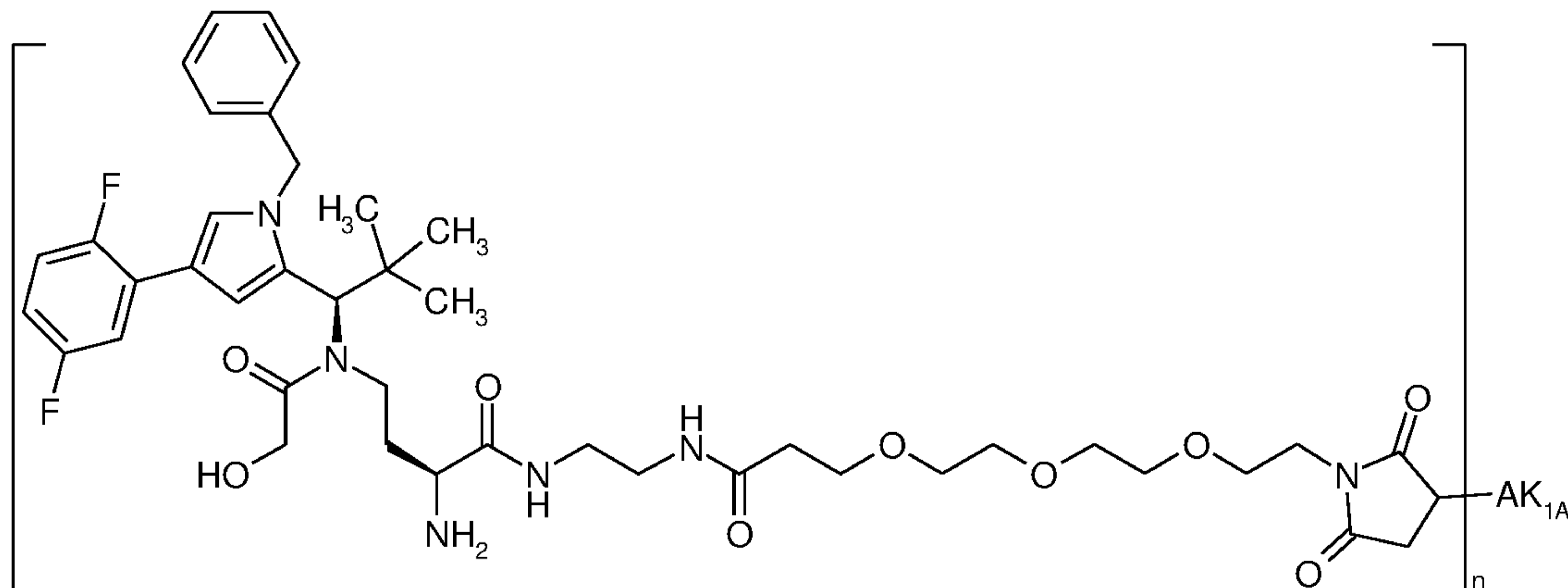
Here, 5.0 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F175, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

10

Protein concentration: 1.97 mg/ml

Drug/mAb ratio: 4.2

15 **Example 176A**



20 Here, 5 mg of cetuximab in PBS (c=21.3 mg/ml) were used for coupling with Intermediate F176, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

25

Protein concentration: 1.98 mg/ml

Drug/mAb ratio: 2.6

30 **Example 176B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F176, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.93 mg/ml

10

Drug/mAb ratio: 2.8

#### Example 176E

15 Here, 5.0 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F176, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

20

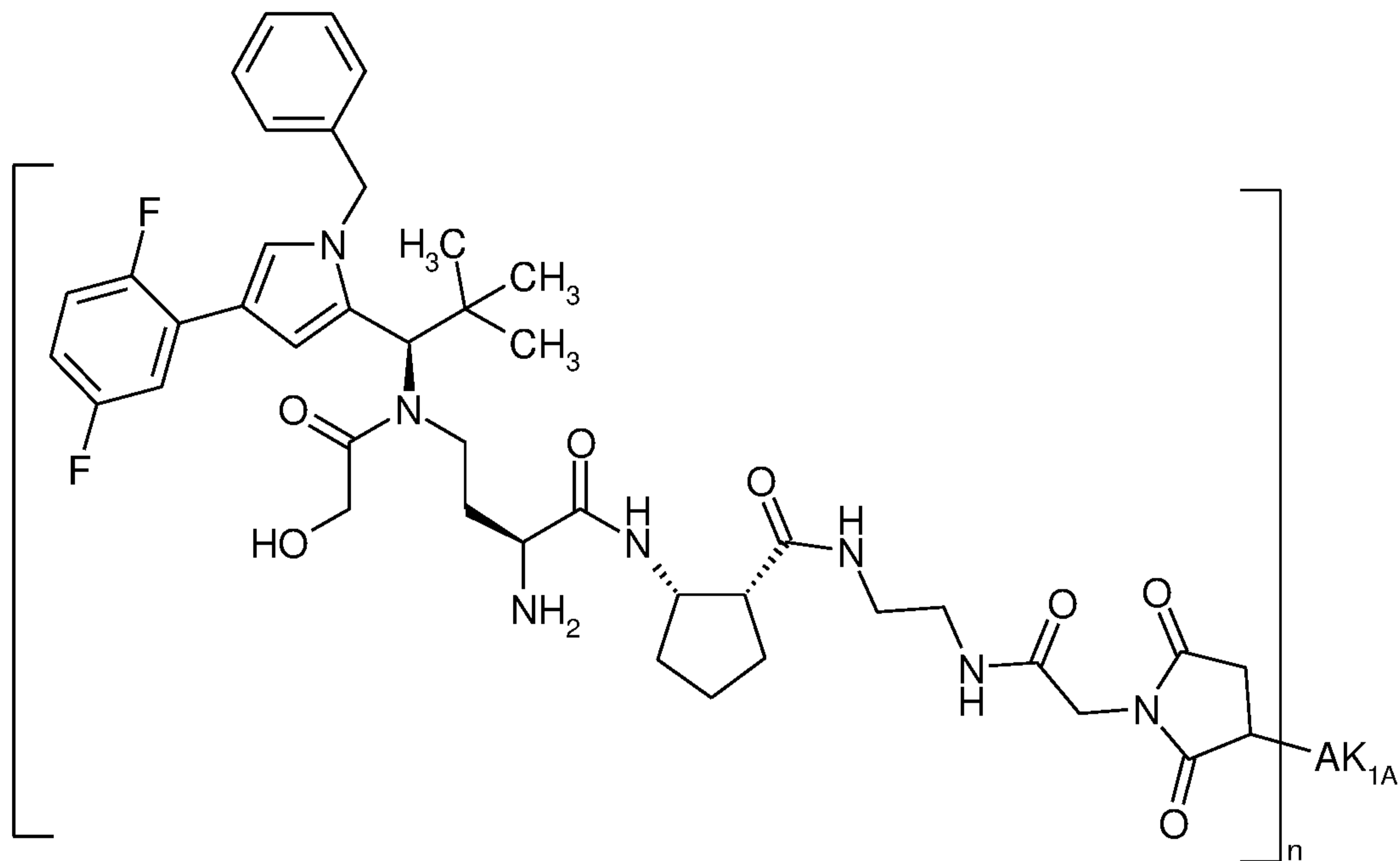
Protein concentration: 1.85 mg/ml

Drug/mAb ratio: 3.3

25

#### Example 177A





Here, 5 mg of cetuximab in PBS (c=21.3 mg/ml) were used for coupling with Intermediate F177, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.96 mg/ml

Drug/mAb ratio: 2.8

### Example 177B

15

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F177, and the reaction was, after  
 20 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.2 mg/ml

25 Drug/mAb ratio: 2.3

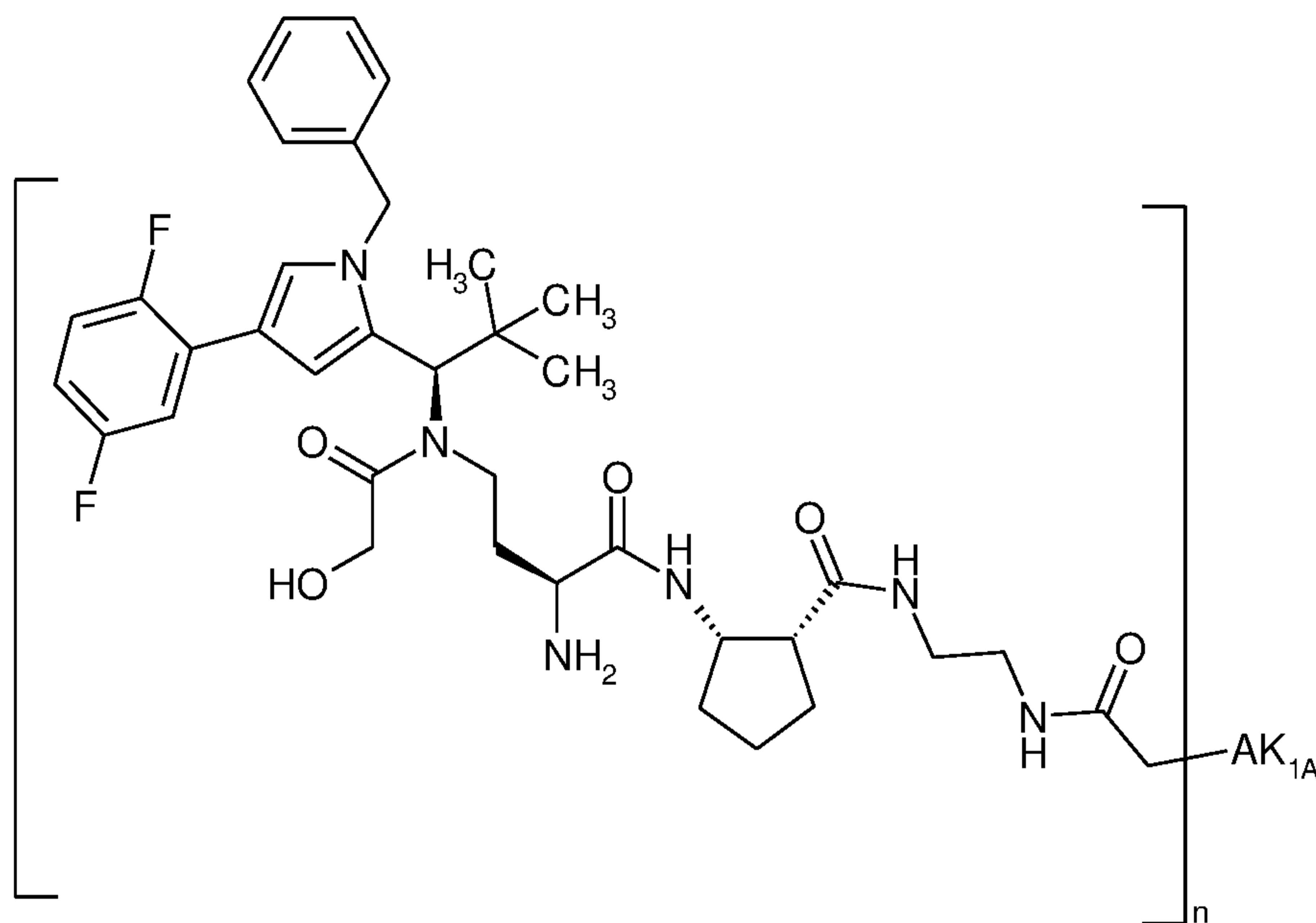
**Example 177E**

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for  
5 coupling with Intermediate F177, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS. Some of the ADC may also be present in the  
form of the hydrolysed open-chain succinamides attached to the  
antibody.

10

Protein concentration: 1.83 mg/ml

Drug/mAb ratio: 3.1

**Example 178A**

Here, 5 mg of cetuximab in PBS (c=21.3 mg/ml) were used for  
20 coupling with Intermediate F178, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS.

Protein concentration: 1.8 mg/ml

25

Drug/mAb ratio: 2.1

**Example 178B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used  
 5 for coupling with Intermediate F178, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

Protein concentration: 1.45 mg/ml

10

Drug/mAb ratio: 2.4

**Example 178E**

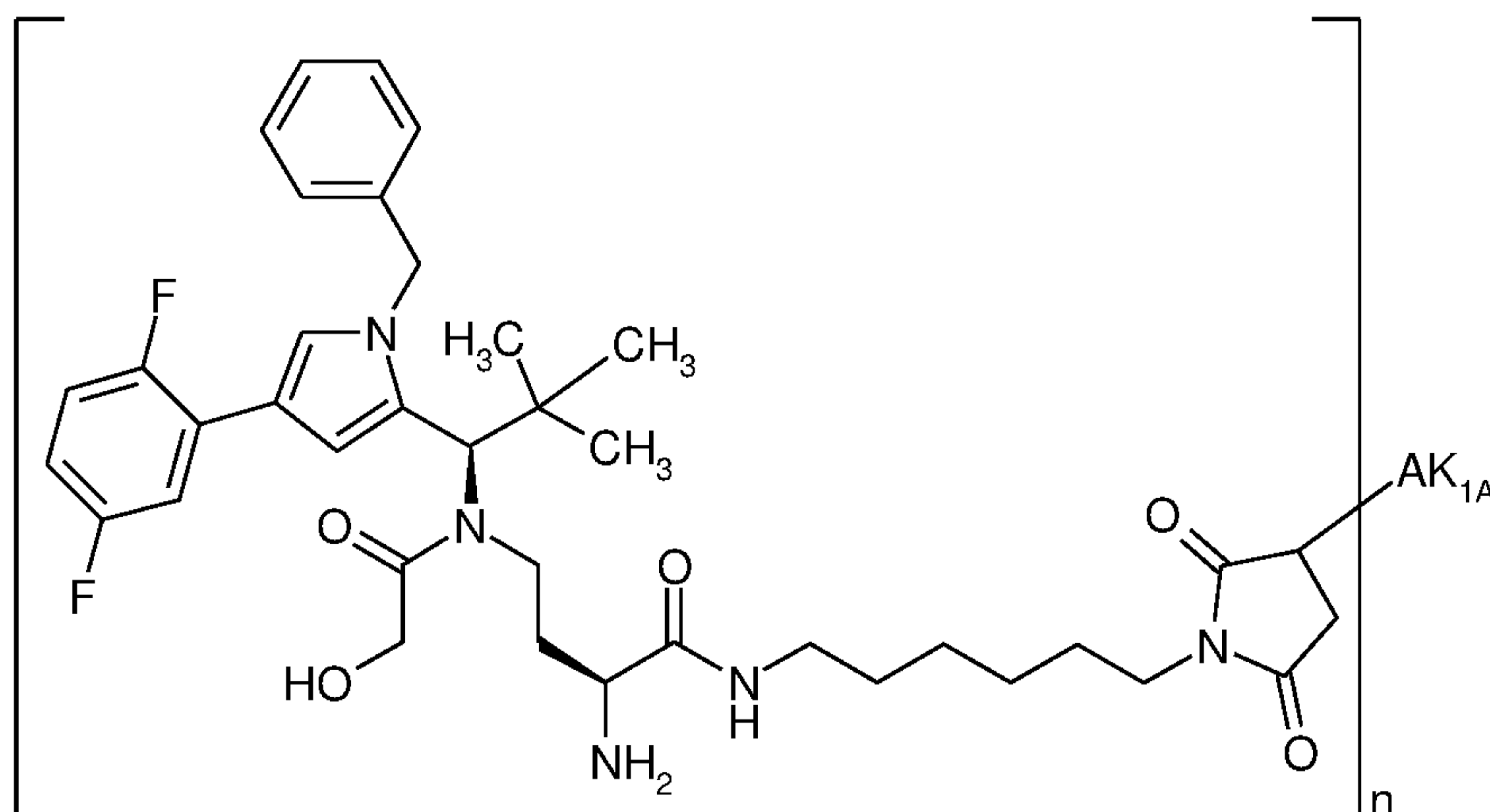
15 Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for  
 coupling with Intermediate F178, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation and  
 rediluted with PBS.

20 Protein concentration: 1.8 mg/ml

Drug/mAb ratio: 2.6

**Example 179A**

25



Here, 5 mg of cetuximab in PBS (c=11.3 mg/ml) were used for  
 coupling with Intermediate F179, and the reaction was, after

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

Protein concentration: 2.04 mg/ml

5

Drug/mAb ratio: 3.1

**Example 179B**

10 Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F179, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

15 Protein concentration: 1.65 mg/ml

Drug/mAb ratio: 3.1

**Example 179E**

20

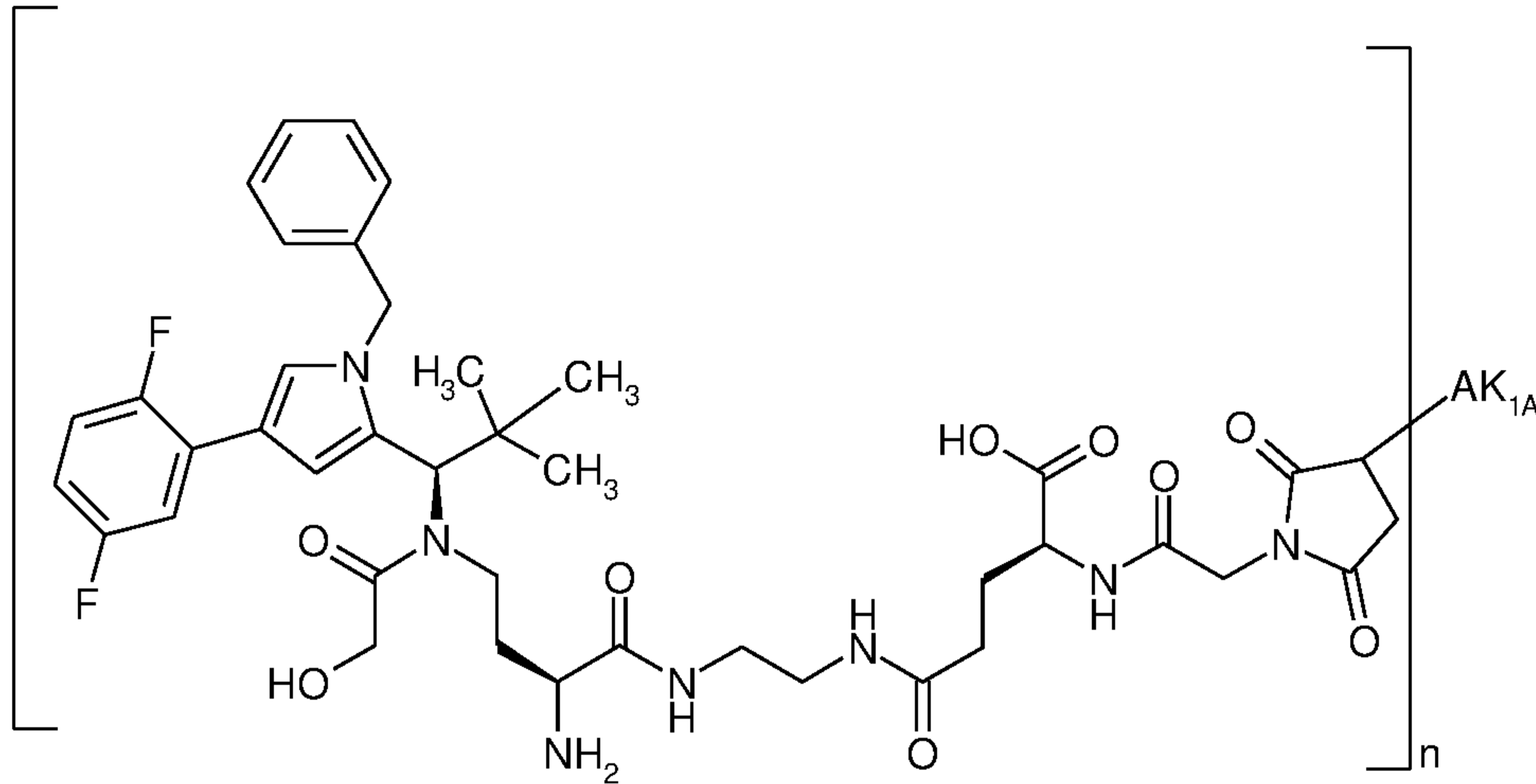
Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F179, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

25

Protein concentration: 1.89 mg/ml

Drug/mAb ratio: 3.3

30 **Example 180A**



Here, 5 mg of cetuximab in PBS (c=8.51 mg/ml) were used for coupling with Intermediate F180, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.72 mg/ml

Drug/mAb ratio: 3.5

#### Example 180B

15

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F180, and the reaction was, after  
 20 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.82 mg/ml

25 Drug/mAb ratio: 3.4

#### Example 180E

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for

coupling with Intermediate F180, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS. Some of the ADC may also be present in the  
form of the hydrolysed open-chain succinamides attached to the  
5 antibody.

Protein concentration: 2.01 mg/ml

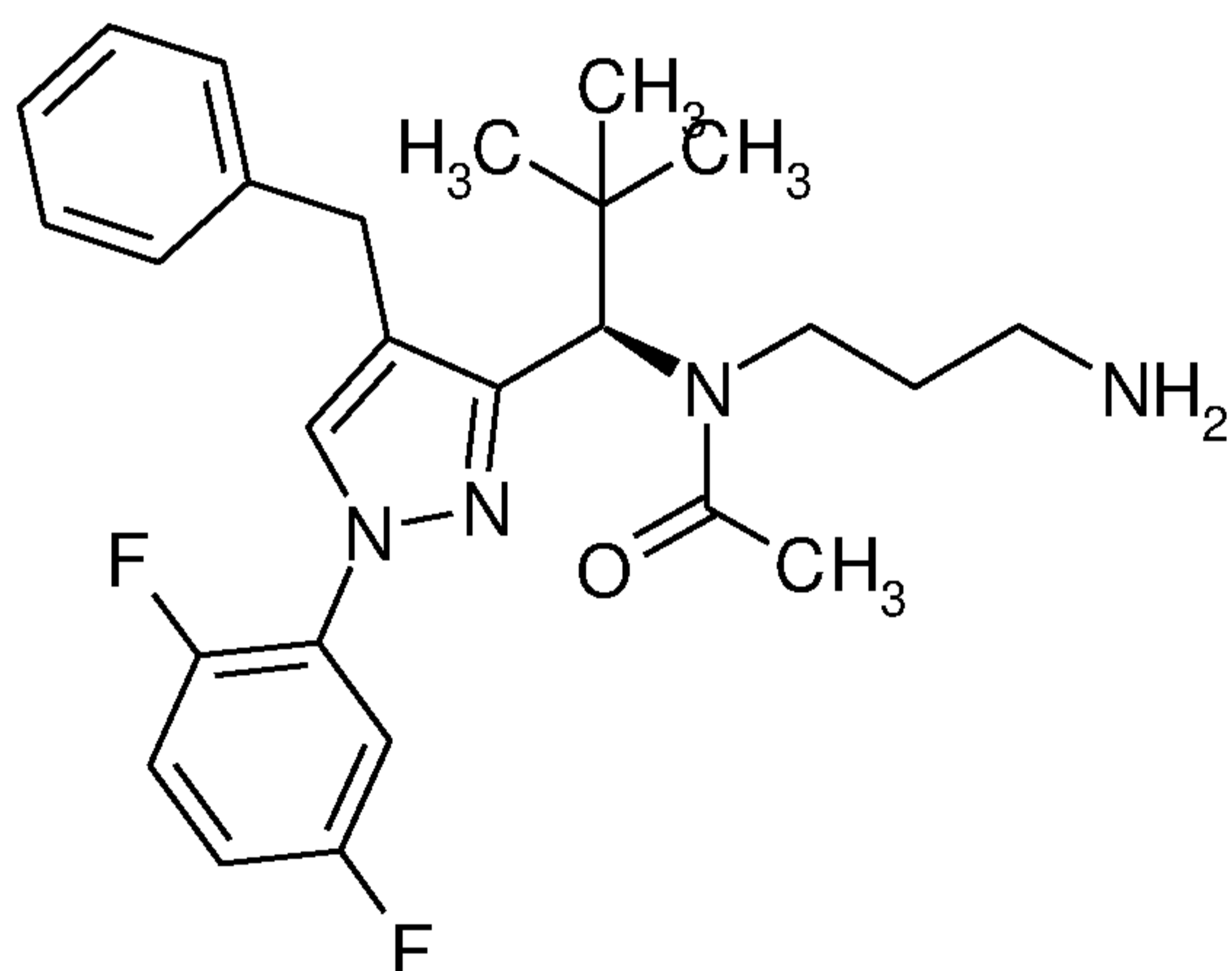
Drug/mAb ratio: 4.7

10

**Example 181**

N-(3-Aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
1H-pyrazol-3-yl]-2,2-dimethylpropyl}acetamide

15



1.01 g (2.84 mmol) of (1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
1H-pyrazol-3-yl]-2,2-dimethylpropan-1-amine were initially  
20 charged in 20 ml of 1,2-dichloroethane, and 0.84 g (3.98 mmol)  
of sodium triacetoxyborohydride and 2.56 g (42.65 mmol) of  
acetic acid were added, and the mixture was stirred at RT for 5  
min. A solution of 0.54 g (3.13 mmol) of tert-butyl (3-  
oxopropyl)carbamate in 5 ml of 1,2-dichloroethane was then  
25 added, and the reaction mixture was stirred overnight. The  
mixture was then evaporated to dryness, the residue was taken  
up in ethyl acetate and filtered and the product-containing  
filtrate was evaporated to dryness.

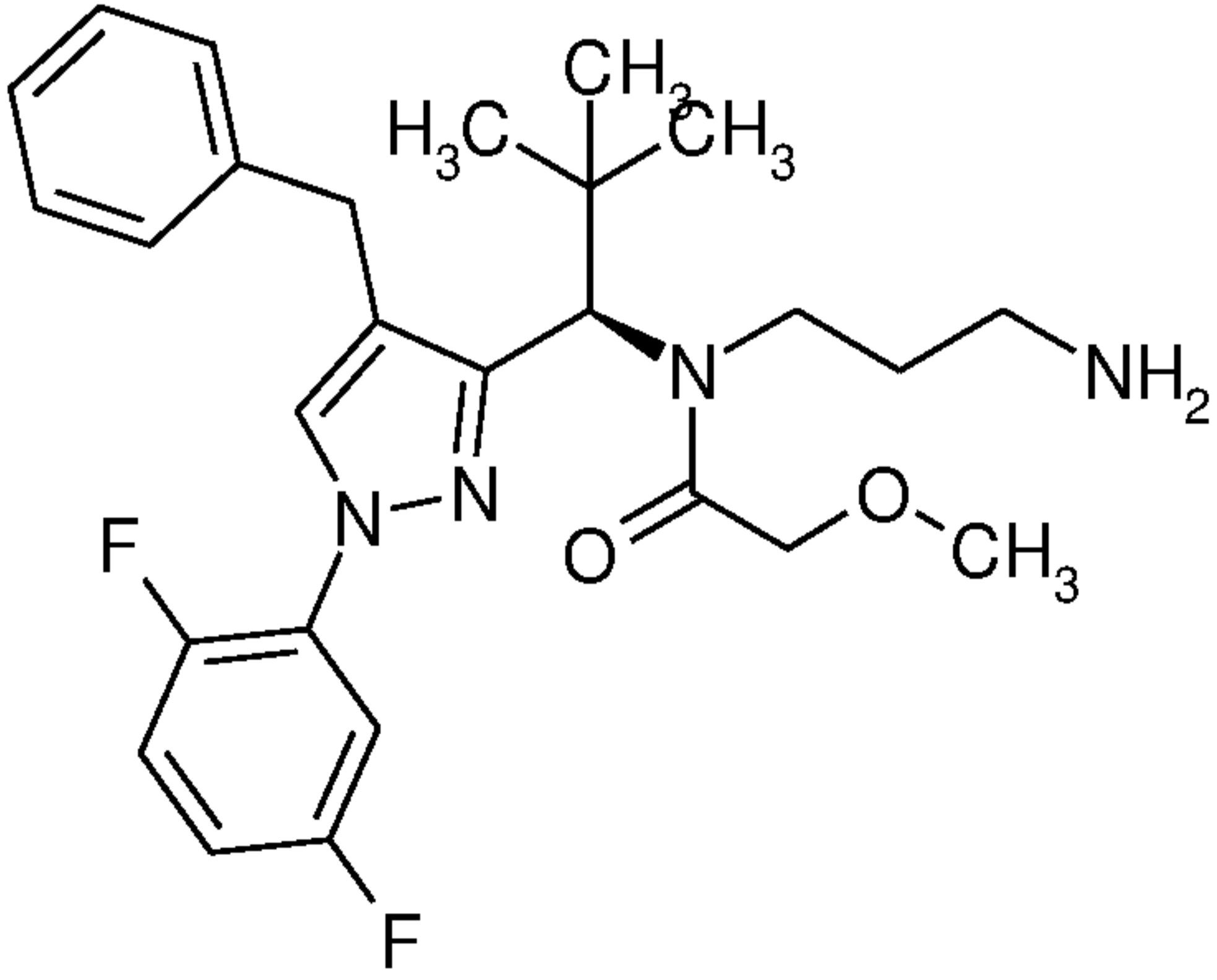
30 51.26 mg of the residue were dissolved in 0.8 ml of 1,2-

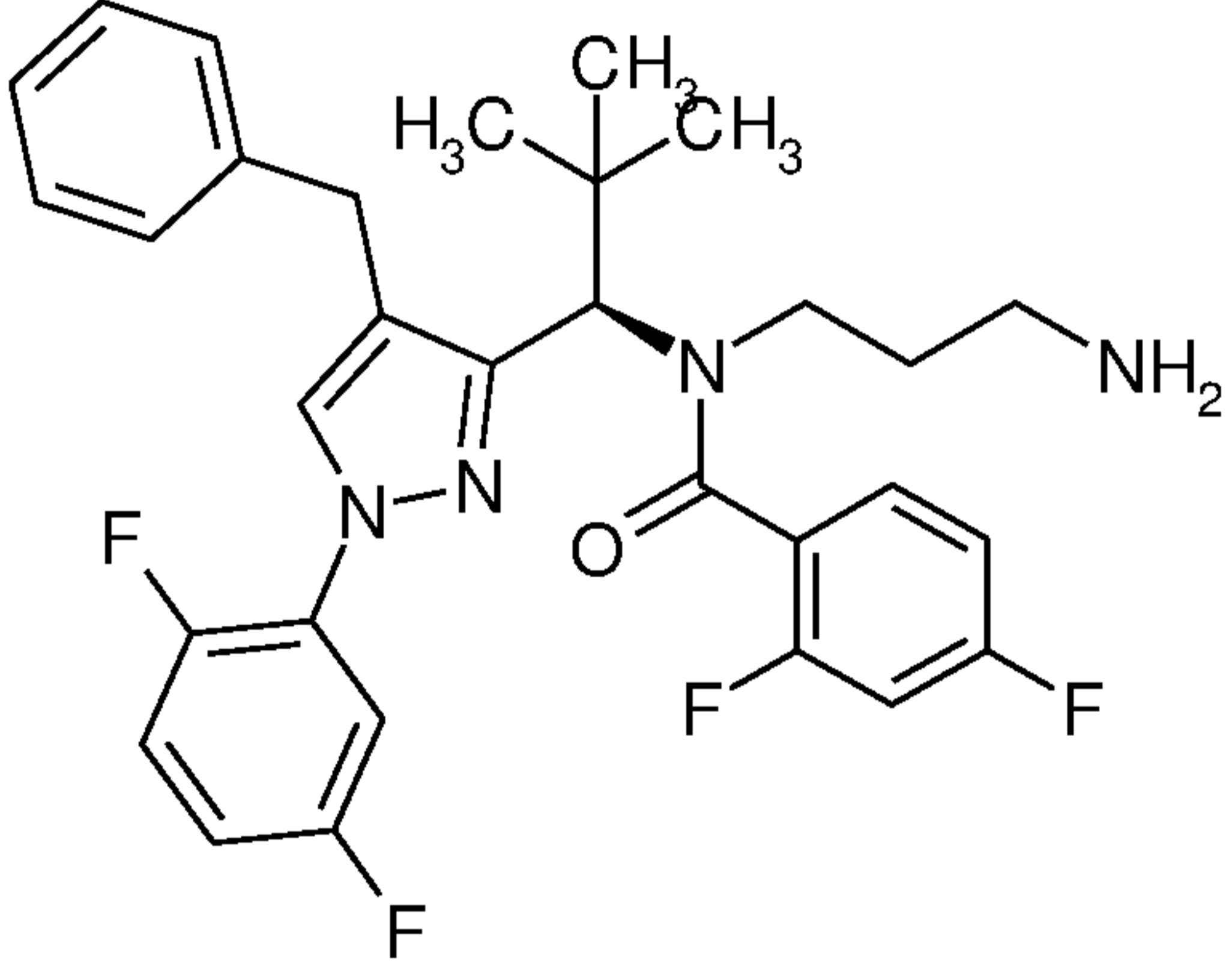
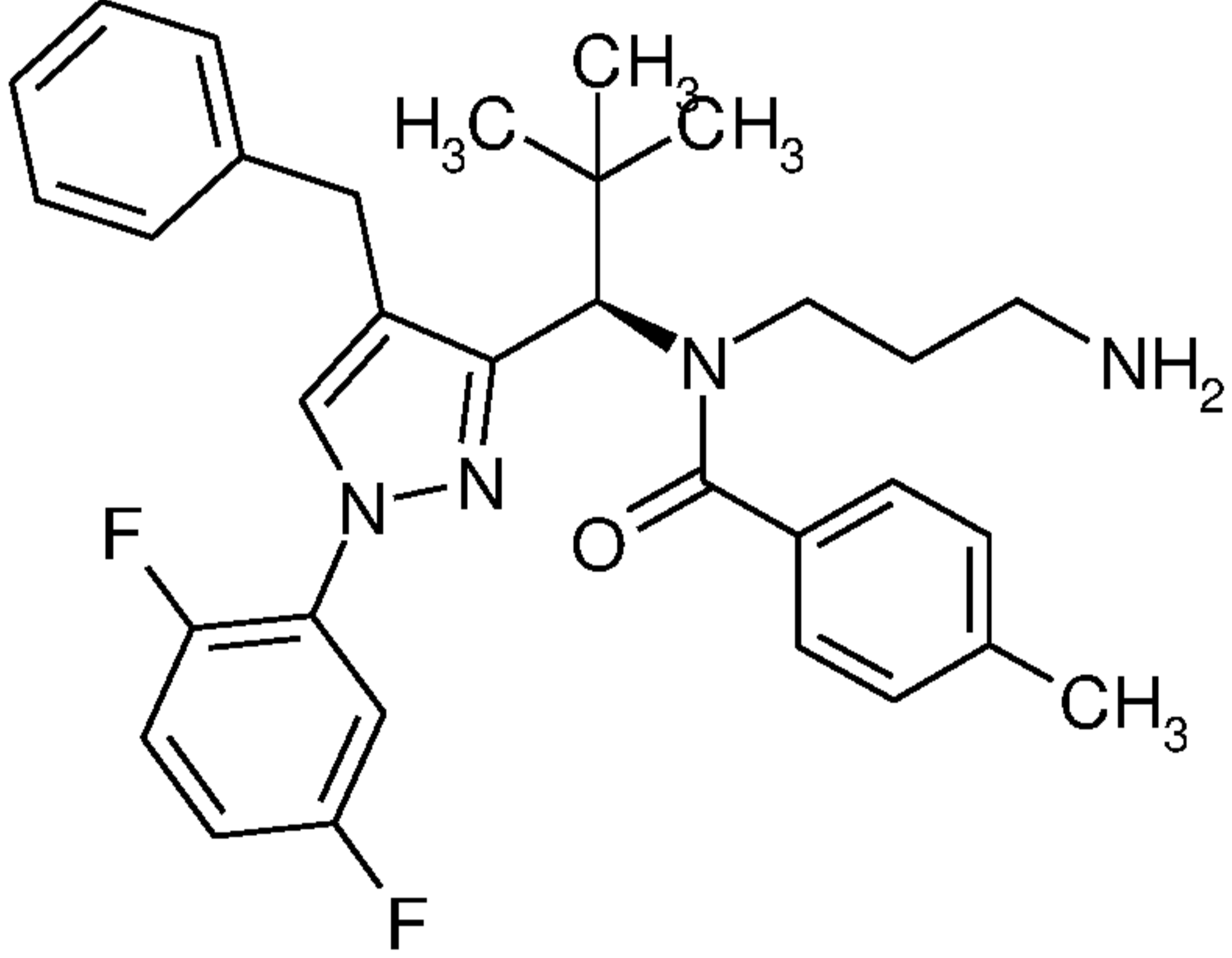
dichloromethane and added to 7.85 mg (0.1 mmol) of acetyl chloride on a deep 96-well multitre plate 25.8 mg (0.2 mmol) of *N,N*-diisopropylethylamine were then added and the mixture was shaken at RT overnight. The solvent was then removed completely using a centrifugal drier, 0.4 ml of 1,2-dichlorethane and 0.4 ml of trifluoroacetic acid were added and the mixture was shaken overnight. The solvent was then removed completely using a centrifugal drier, and 0.8 ml of DMF were added to the residue. The mixture was then filtered and the target compound was isolated from the filtrate by preparative LC-MS (Method 9). The product-containing fractions were concentrated under reduced pressure using a centrifugal dryer. The residue of each product fraction was dissolved in 0.6 ml of DMSO. These were combined and finally freed of the solvent in a centrifugal dryer. This gave 12.2 mg (27% of theory; purity 100%) of the title compound.

LC-MS (Method 10):  $R_t = 0.95$  min; MS (ESIpos):  $m/z = 455$   $[M+H]^+$

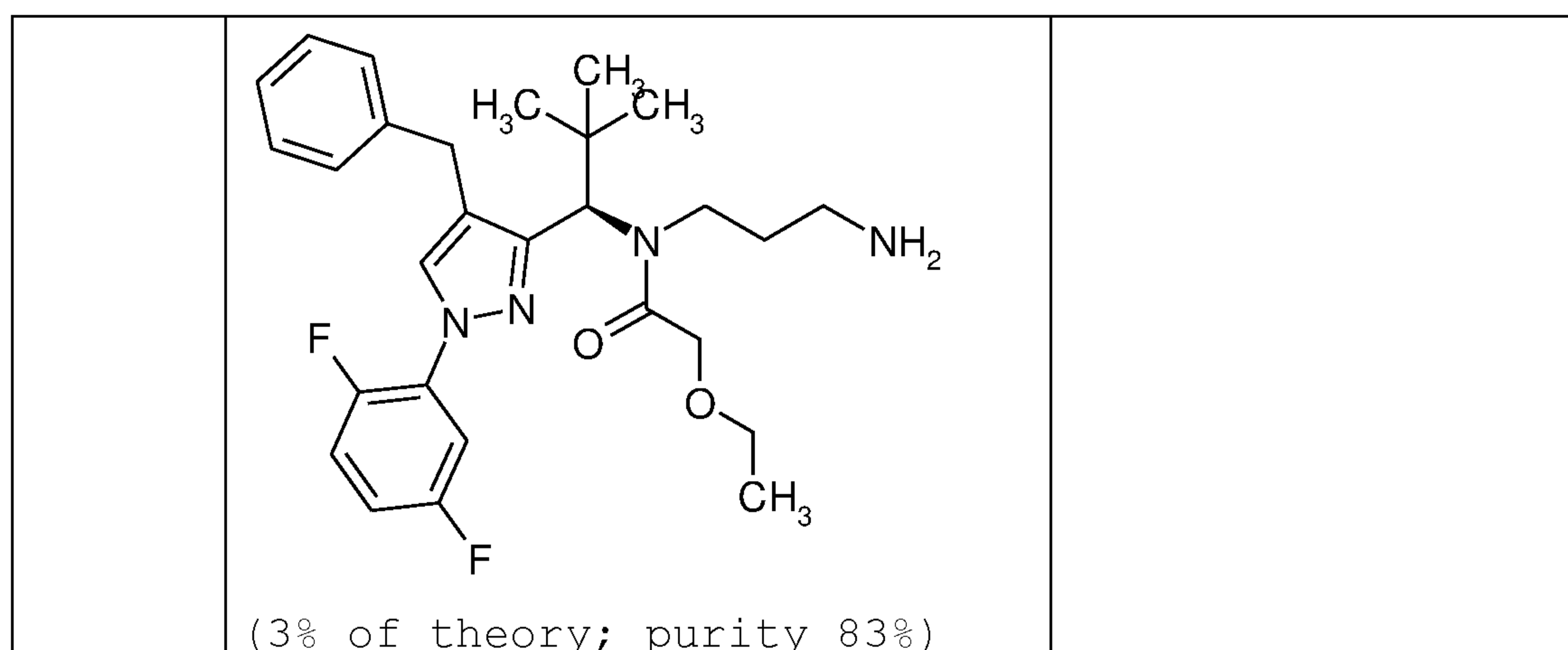
The exemplary compounds shown in Table XA were prepared analogously to Example 181:

**Table XA with Examples 182-185**

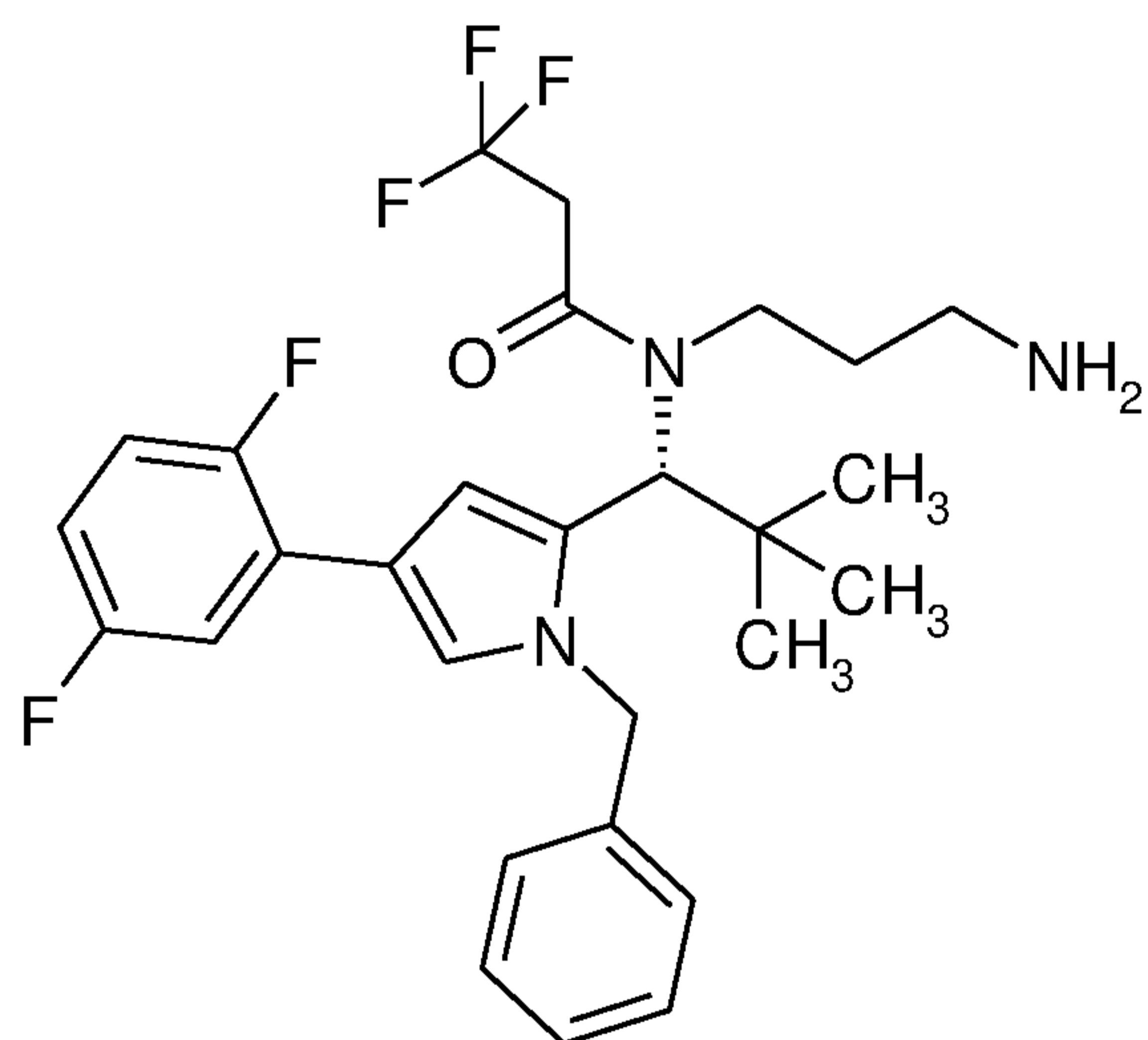
Ex-ample	IUPAC name / structure (Yield)	Analytical data
182	N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-methoxyacetamide 	LC-MS (Method 10): $R_t = 0.95$ min MS (ESpos): $m/z = 485$ $(M+H)^+$

	(14% of theory; purity 98%)	
<b>183</b>	<p>N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2,4-difluorobenzamide</p>  <p>(20% of theory; purity 100%)</p>	<p>LC-MS (Method 10):  <math>R_t = 1.02</math> min  MS (ESpos): <math>m/z = 553</math> (M+H)<sup>+</sup></p>
<b>184</b>	<p>N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-4-methylbenzamide</p>  <p>(14% of theory; purity 100%)</p>	<p>LC-MS (Method 10):  <math>R_t = 1.04</math> min  MS (ESpos): <math>m/z = 531</math> (M+H)<sup>+</sup></p>
<b>185</b>	<p>N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamide</p>	<p>LC-MS (Method 10):  <math>R_t = 0.98</math> min  MS (ESpos): <math>m/z = 499</math> (M+H)<sup>+</sup></p>



**Example 186**

N-(3-Aminopropyl)-N-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
5 1H-pyrrol-2-yl]-2,2-dimethylpropyl)-3,3,3-trifluoropropanamide



1.0 g (2.82 mmol) of (1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
10 1H-pyrrol-2-yl]-2,2-dimethylpropan-1-amine were initially  
charged in 20 ml of 1,2-dichloroethane, and 1.4 g (3.95 mmol)  
of sodium triacetoxyborohydride and 2.54 g (42.32 mmol) of  
acetic acid were added, and the mixture was stirred at RT for 5  
15 min. A solution of 0.54 g (3.10 mmol) of tert-butyl (3-  
oxopropyl)carbamate in 5 ml of 1,2-dichloroethane was then  
added, and the reaction mixture was stirred overnight. The  
mixture was then evaporated to dryness, the residue was taken  
up in ethyl acetate and filtered and the product-containing  
filtrate was evaporated to dryness.

51.16 mg of the residue were dissolved in 0.8 ml of 1,2-dichloromethane and added to 14.65 mg (0.1 mmol) of 3,3,3-trifluoropropanoyl chloride on a deep 96-well multtitre plate  
 5 25.8 mg (0.2 mmol) of *N,N*-diisopropylethylamine were then added and the mixture was shaken at RT overnight. The solvent was then removed completely using a centrifugal drier, 0.4 ml of 1,2-dichloroethane and 0.4 ml of trifluoroacetic acid were added and the mixture was shaken overnight. The solvent was then removed  
 10 completely using a centrifugal drier, and 0.8 ml of DMF were added to the residue. The mixture was then filtered and the target compound was isolated from the filtrate by preparative LC-MS (Method 9). The product-containing fractions were concentrated under reduced pressure using a centrifugal dryer.  
 15 The residue of each product fraction was dissolved in 0.6 ml of DMSO. These were combined and finally freed of the solvent in a centrifugal dryer. This gave 1.0 mg (2% of theory; purity 82%) of the title compound.

LC-MS (Method 10):  $R_t = 1.01$  min MS (ESIpos):  $m/z = 522$  [M+H]<sup>+</sup>

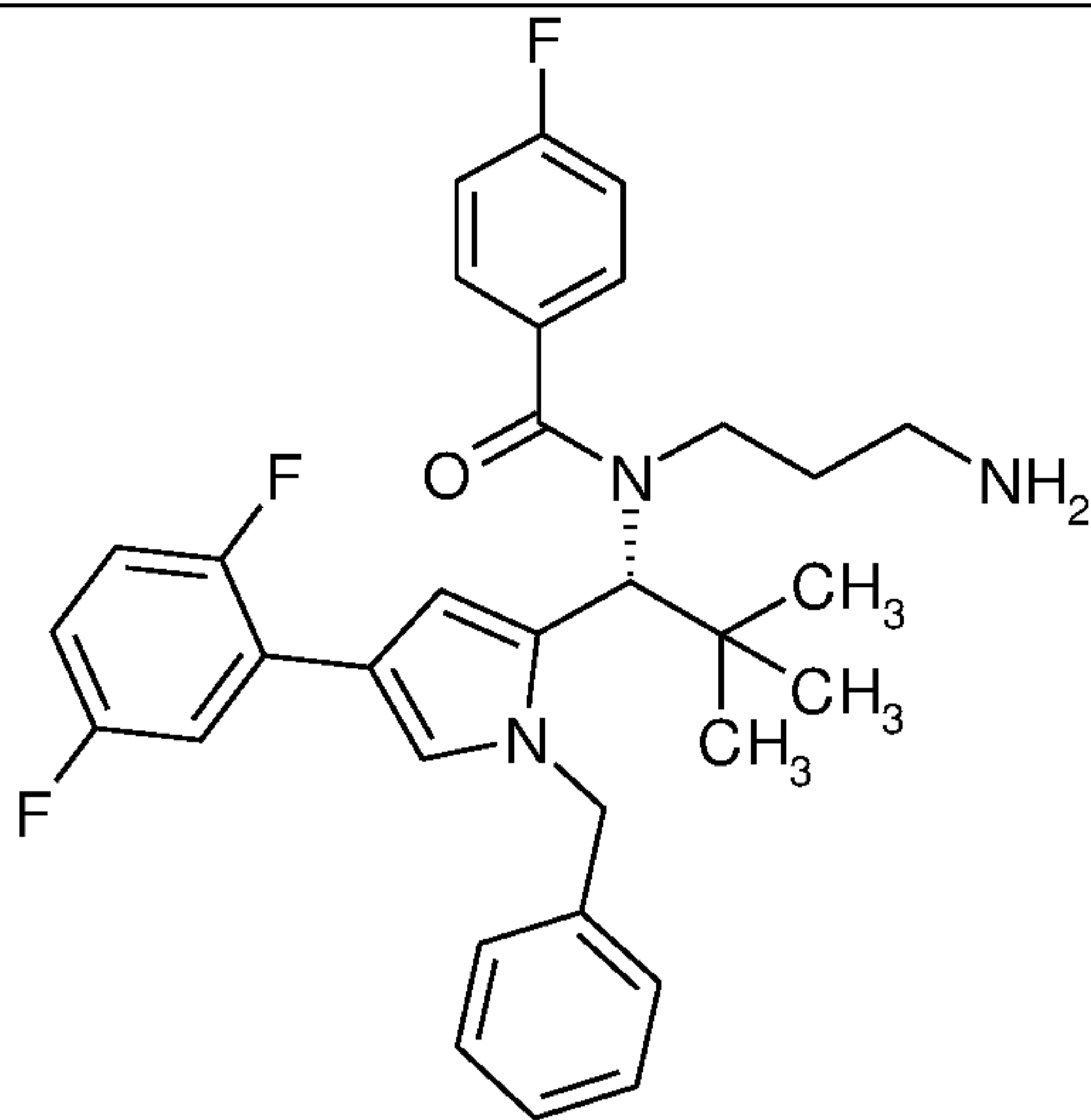
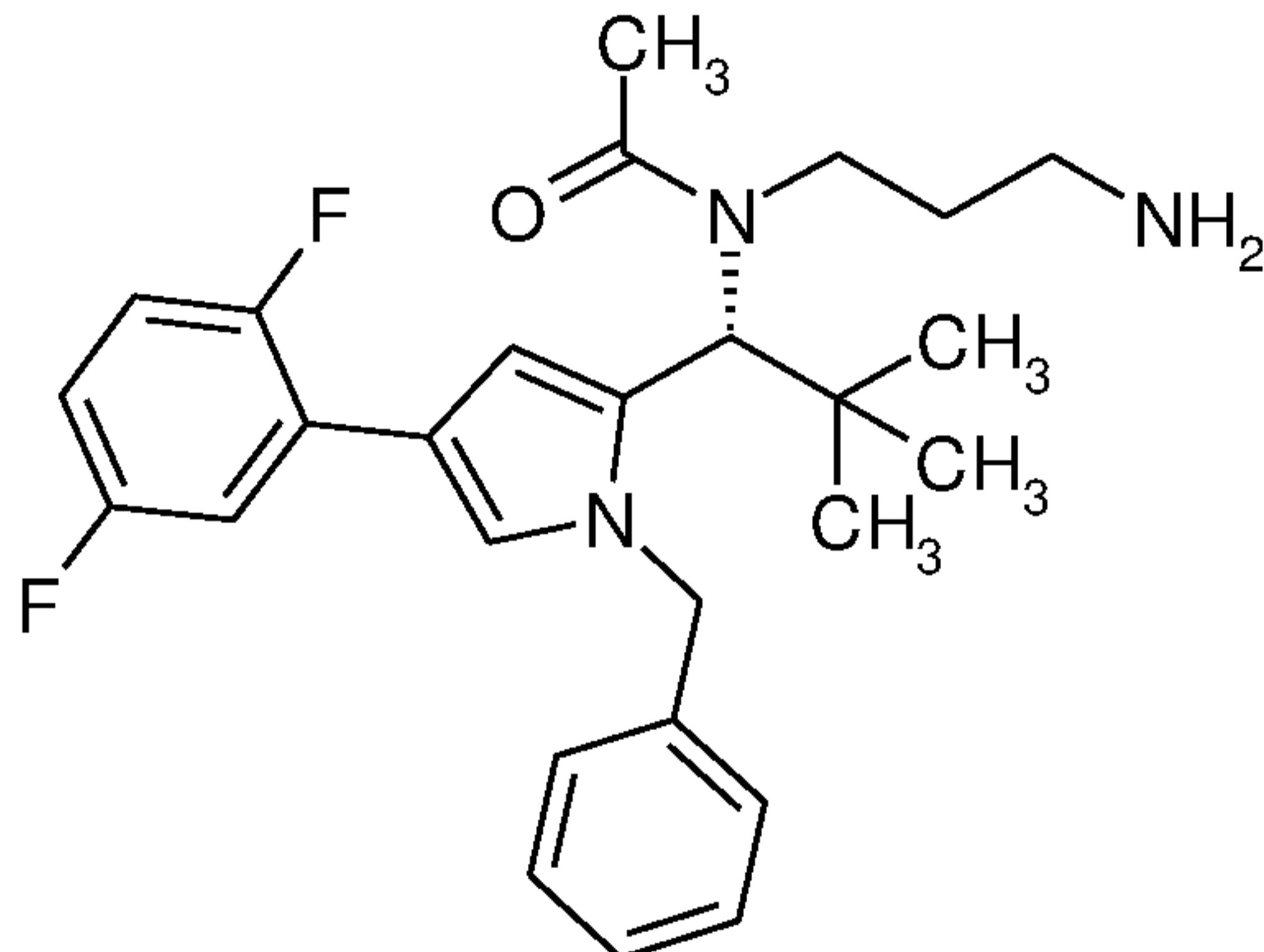
20

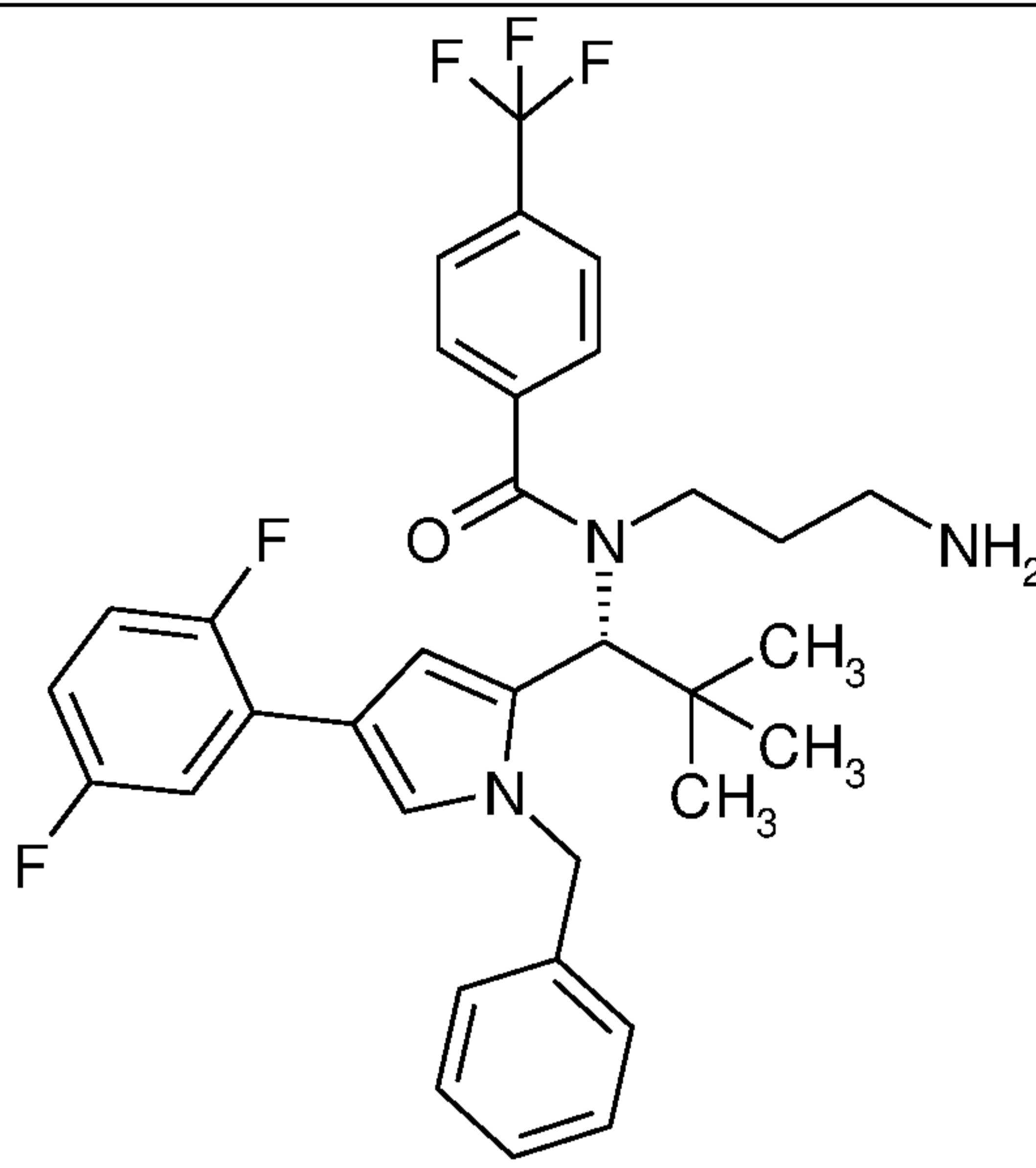
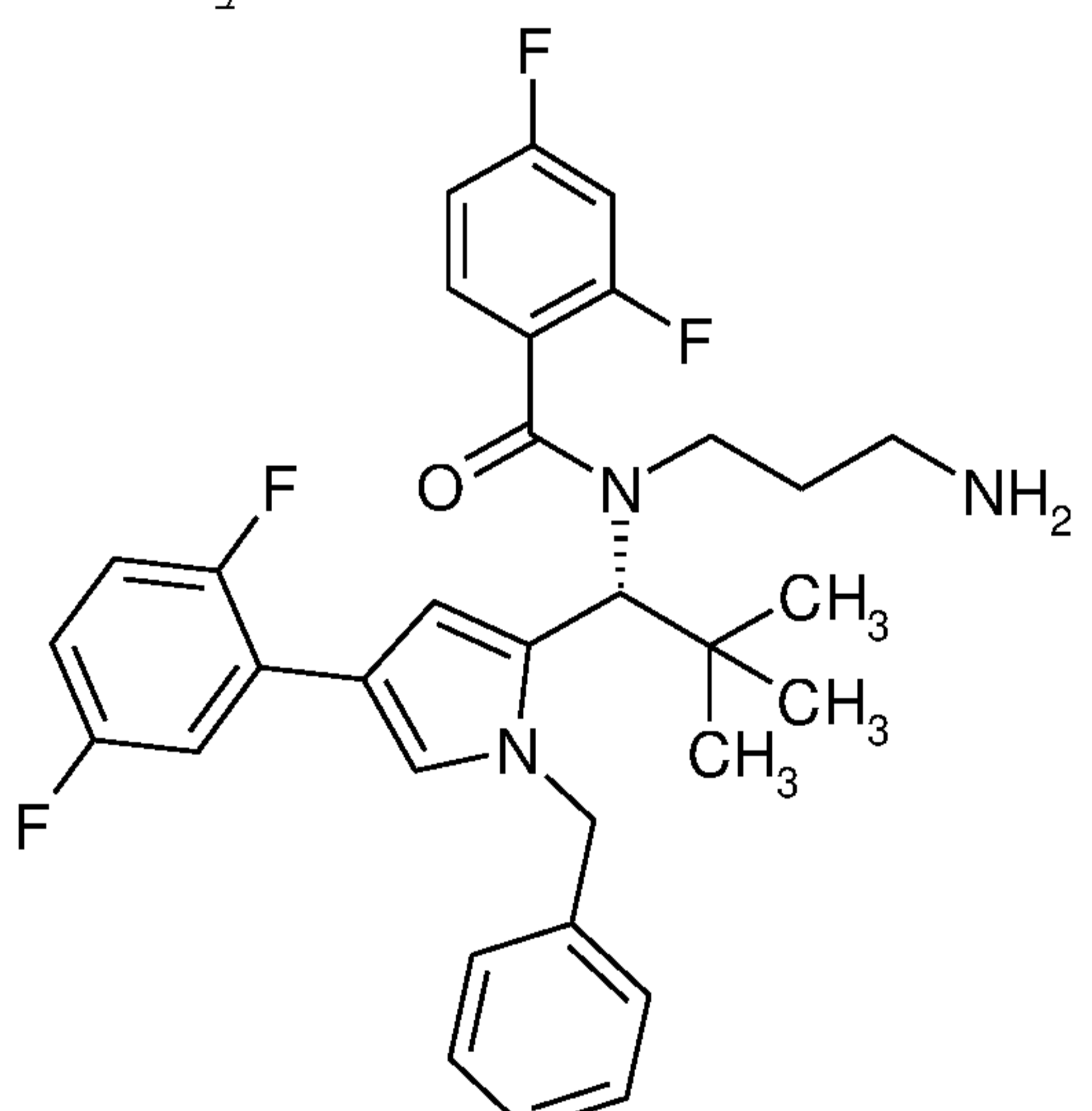
The exemplary compounds shown in Table XA1 were prepared analogously to Example 186:

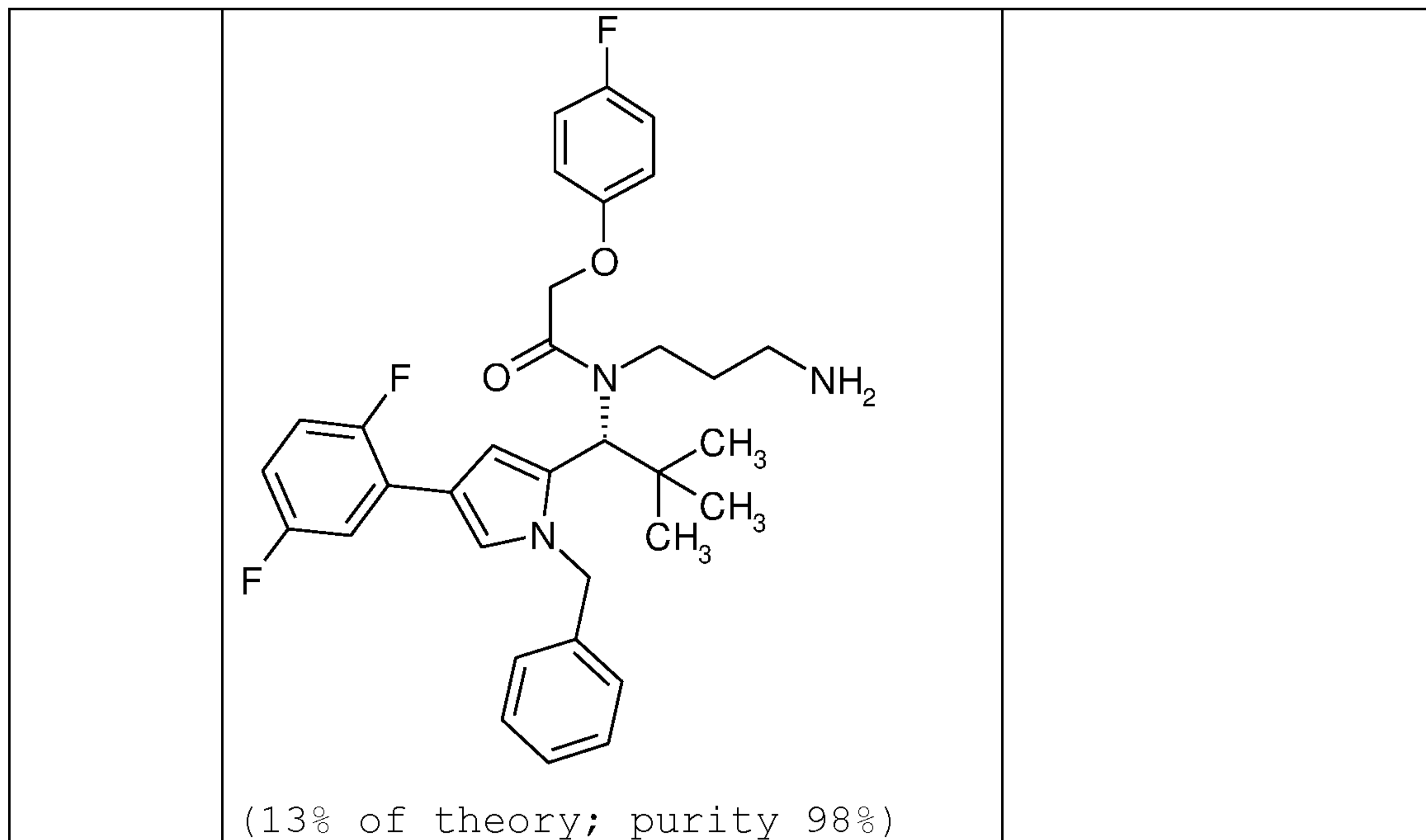
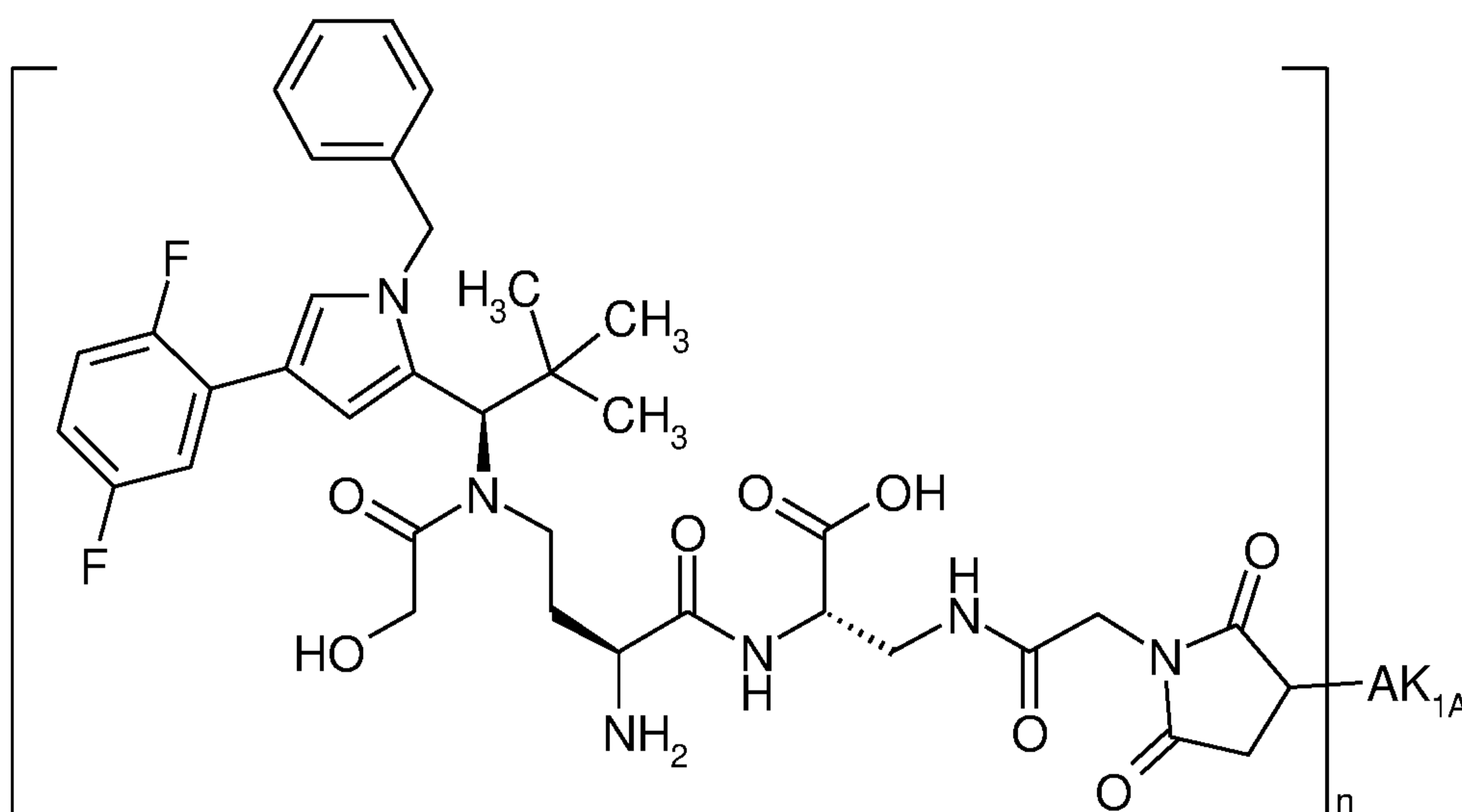
**Table XAI with Examples 187-191**

25

Ex-ample	IUPAC name / structure (Yield)	Analytical data
187	N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-fluorobenzamide	LC-MS (Method 10): $R_t = 1.01$ min MS (ESpos): $m/z = 534$ (M+H) <sup>+</sup>

	 <p>(3% of theory; purity 83%)</p>	
188	<p>N-(3-aminopropyl)-N-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)acetamide</p>  <p>(2% of theory; purity 87%)</p>	<p>LC-MS (Method 10): <math>R_t = 0.94</math> min  MS (ESpos): <math>m/z = 454</math> (M+H)<sup>+</sup></p>
189	<p>N-(3-aminopropyl)-N-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-4-(trifluoromethyl)benzamide</p>	<p>LC-MS (Method 10): <math>R_t = 1.05</math> min  MS (ESpos): <math>m/z = 584</math> (M+H)<sup>+</sup></p>

	 <p>(6% of theory; purity 90%)</p>	
190	<p>N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamide</p>  <p>(7% of theory; purity 83%)</p>	<p>LC-MS (Method 10): <math>R_t = 0.98</math> min  MS (ESpos): <math>m/z = 552</math> (M+H)<sup>+</sup></p>
191	<p>N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamide</p>	<p>LC-MS (Method 10): <math>R_t = 1.02</math> min  MS (ESpos): <math>m/z = 564</math> (M+H)<sup>+</sup></p>

**Example 192A**

5

Here, 5 mg of cetuximab in PBS (c=21.3 mg/ml) were used for coupling with Intermediate F192, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10

Protein concentration: 1,968 mg/ml

Drug/mAb ratio: 2.9

**Example 192B**

5 Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used  
for coupling with Intermediate F192, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS. Some of the ADC may also be present in the  
10 form of the hydrolysed open-chain succinamides attached to the  
antibody.

Protein concentration: 1.92 mg/ml

Drug/mAb ratio: 2.6

15

**Example 192E**

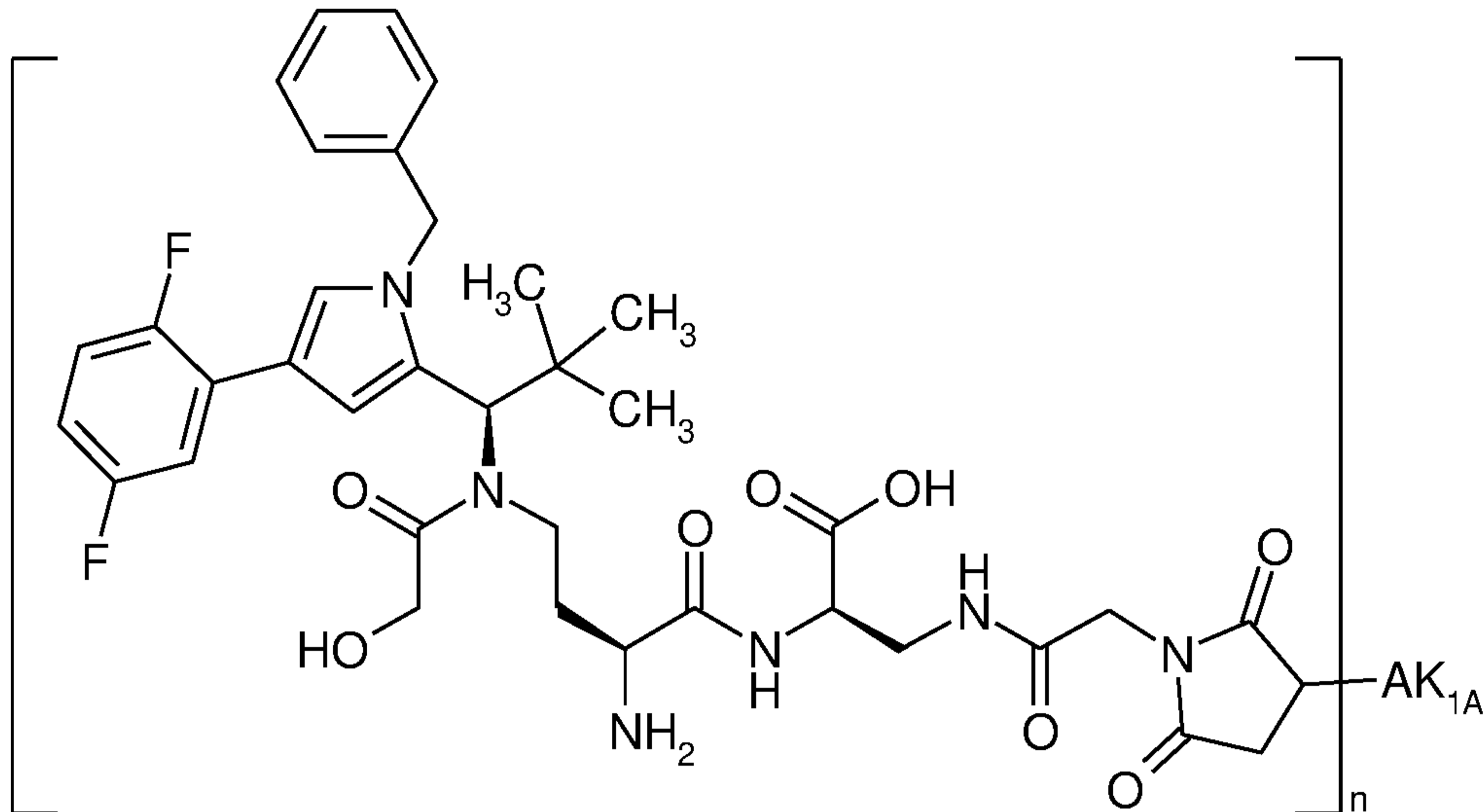
Here, 5 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were  
used for coupling with Intermediate F192, and the reaction was,  
20 after Sephadex purification, concentrated by  
ultracentrifugation and rediluted with PBS. Some of the ADC may  
also be present in the form of the hydrolysed open-chain  
succinamides attached to the antibody.

25 Protein concentration: 2.04 mg/ml

Drug/mAb ratio: 3.3

**Example 193A**

30



Here, 5 mg of cetuximab in PBS (c=23.1 mg/ml) were used for coupling with Intermediate F193, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.98 mg/ml

Drug/mAb ratio: 2.9

### Example 193B

15

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F193, and the reaction was, after  
 20 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 2.03 mg/ml

25 Drug/mAb ratio: 3.2

### Example 193E

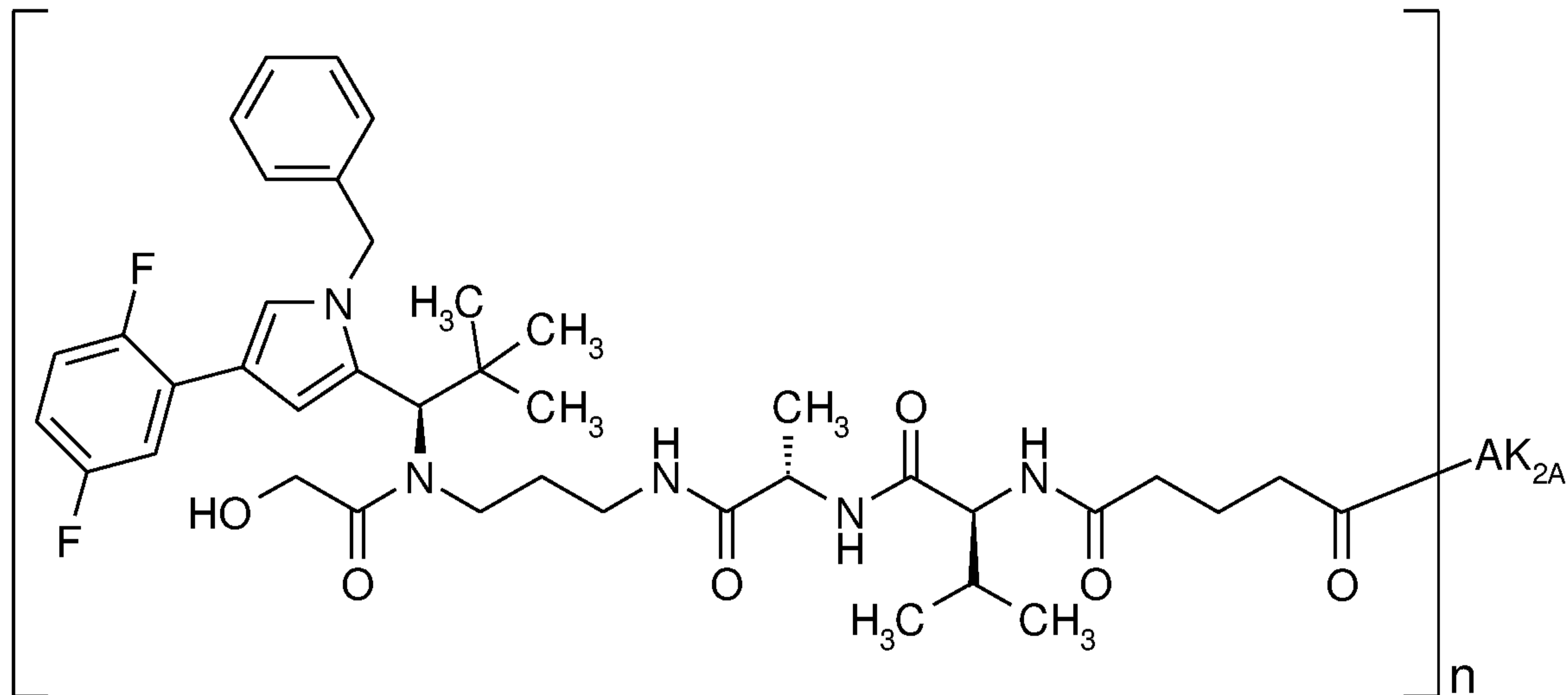
Here, 5 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F193, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.61 mg/ml

10

Drug/mAb ratio: 3.3

### Example 194A



15

Here, 5 mg of cetuximab in PBS (c=23.1 mg/ml) were used for coupling with Intermediate F194, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

20

Protein concentration: 1.67 mg/ml

Drug/mAb ratio: 1.9

25

### Example 194B

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml)



were used for coupling with Intermediate F194, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

5 Protein concentration: 0.99 mg/ml

Drug/mAb ratio: 3.8

**Example 194E**

10

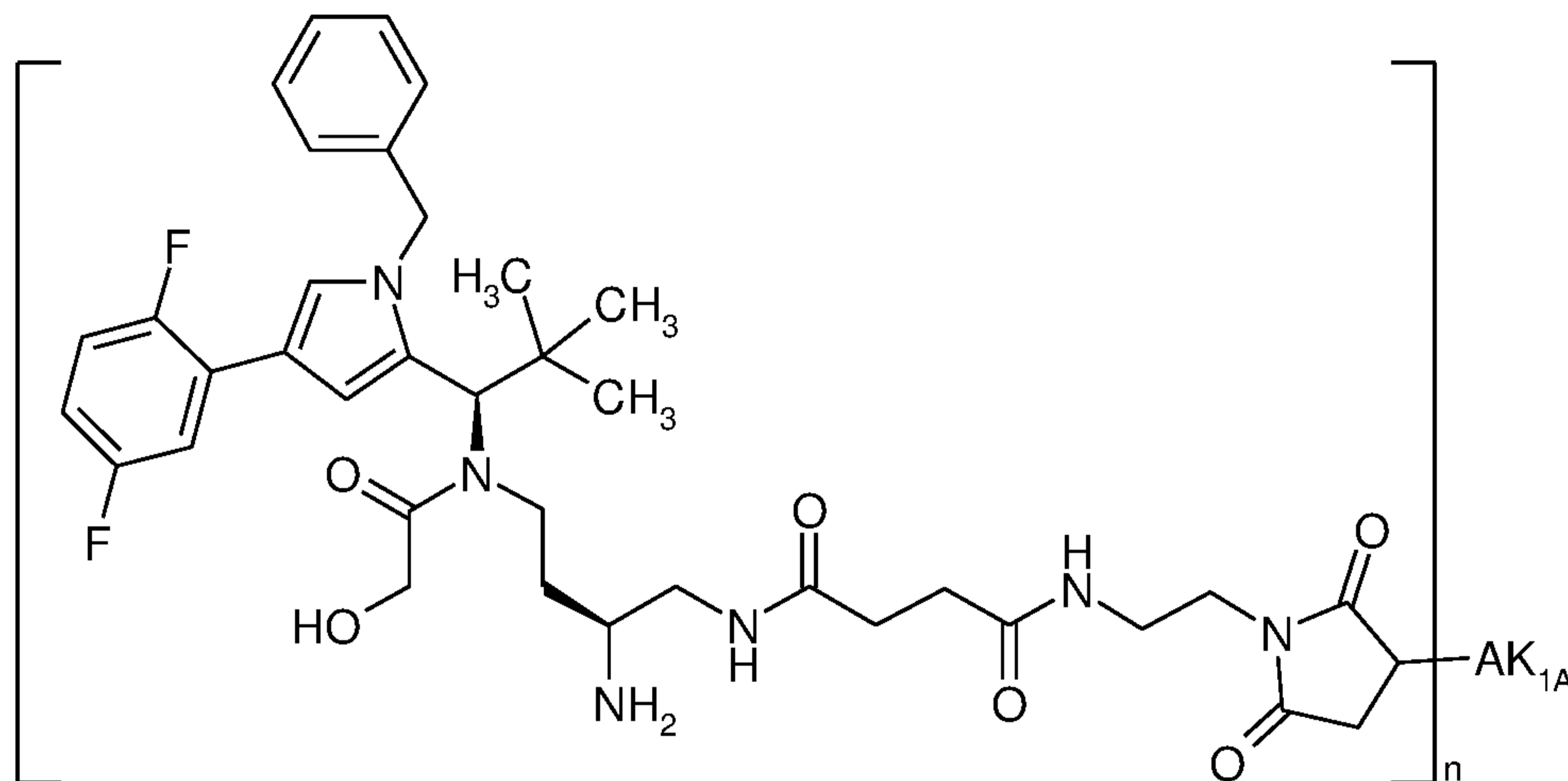
Here, 5.0 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F194, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

15

Protein concentration: 1.39 mg/ml

Drug/mAb ratio: 2.4

20 **Example 195A**



25 Here, 5 mg of cetuximab in PBS (c=23.1 mg/ml) were used for coupling with Intermediate F195, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.79 mg/ml

Drug/mAb ratio: 3.2

5

**Example 195B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F195, and the reaction was, after  
10 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15 Protein concentration: 1.53 mg/ml

Drug/mAb ratio: 3.3

**Example 195E**

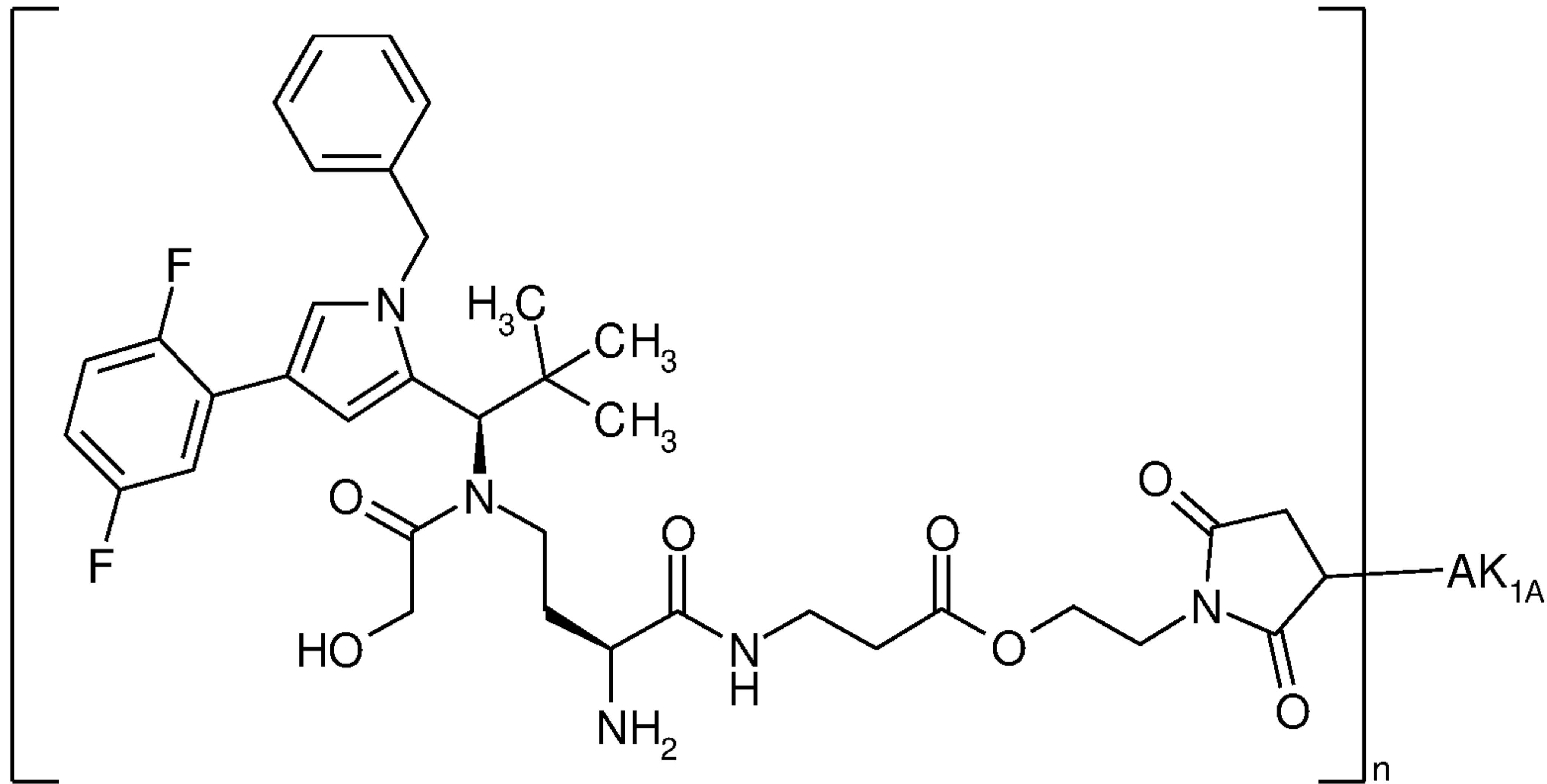
20

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F195, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may  
25 also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.91 mg/ml

30 Drug/mAb ratio: 3.4

**Example 196A**



Here, 5 mg of cetuximab in PBS (c=23.1 mg/ml) were used for coupling with Intermediate F196, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.85 mg/ml

Drug/mAb ratio: 3.0

#### Example 196B

15

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F196, and the reaction was, after  
 20 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.73 mg/ml

25 Drug/mAb ratio: 3.1

#### Example 196E

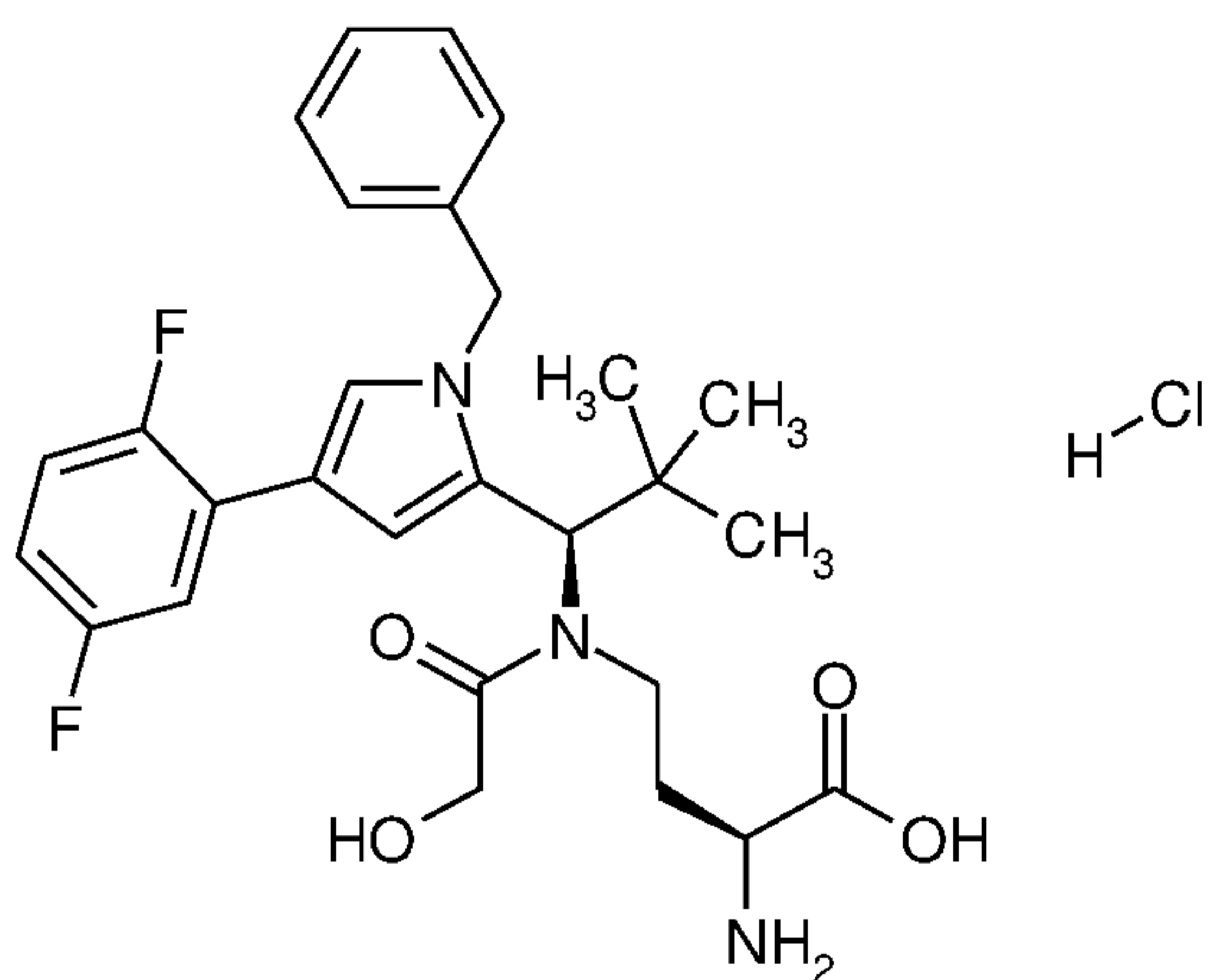
Here, 5.0 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F196, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.86 mg/ml

10 Drug/mAb ratio: 3.4

### Example 197

(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoic acid hydrochloride (1:1)



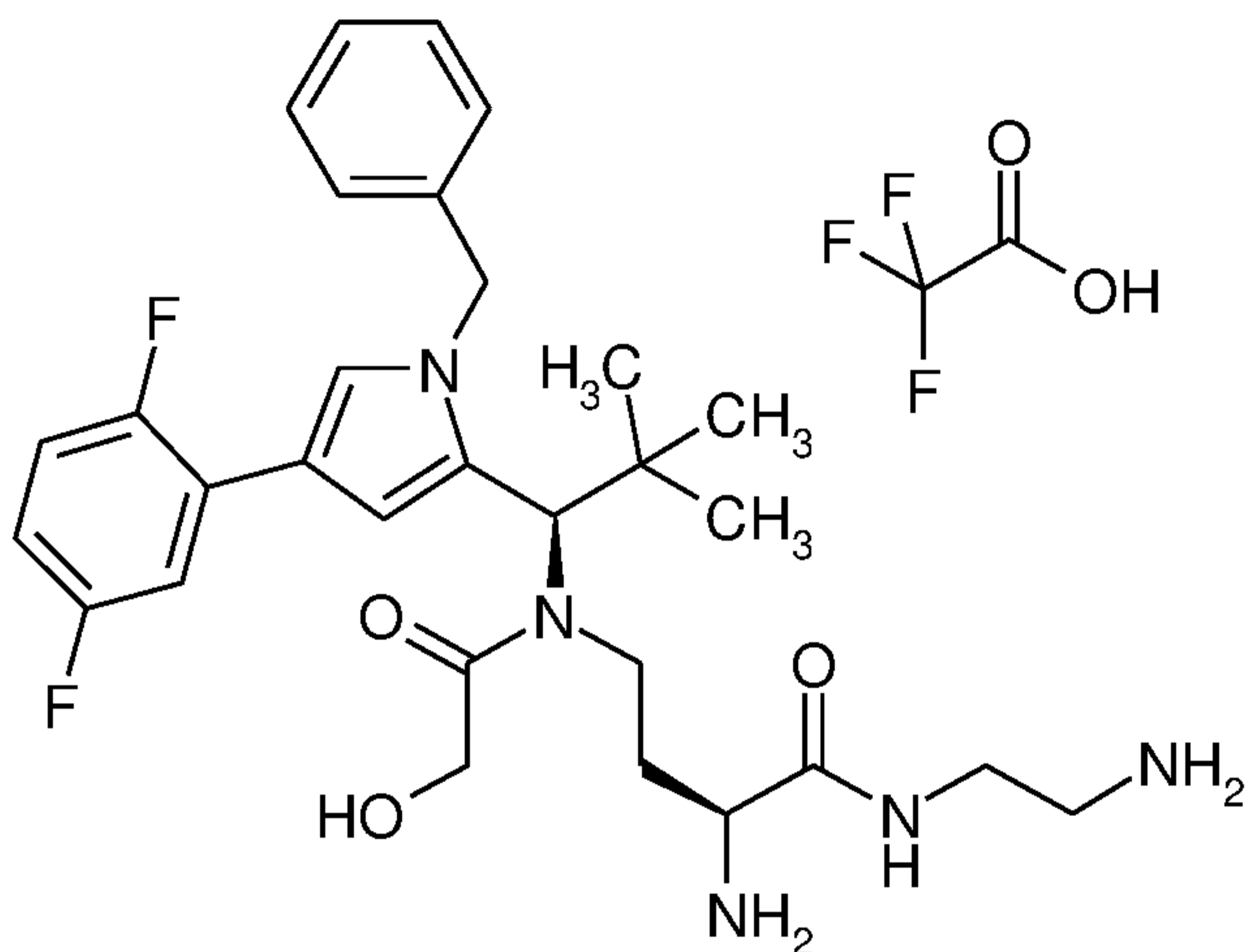
20 150 mg (0.2 mmol) of Intermediate C53 were dissolved in 15 ml of DMF, and 2.29 g (20.39 mmol) of DABCO. The reaction was treated in an ultrasonic bath for 30 min. By addition of 1.17 ml of acetic acid, the reaction was then adjusted to pH 3-4, and the mixture was concentrated under reduced pressure. The residue was purified by preparative HPLC and the appropriate fractions were concentrated at RT under reduced pressure. The residue was taken up in acetonitrile/water 1:1, 5 ml of a 4N hydrochloric acid were added and the mixture was then lyophilized. This gave 81 mg (68% of theory) of the title compound.

30

LC-MS (Method 5):  $R_t = 2.69$  min; MS (EIpos):  $m/z = 514$   $[M+H]^+$ .

### Example 198

5 Trifluoroacetic acid / (2S)-2-amino-N-(2-aminoethyl)-4-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanamide (1:1)



10

15 mg (0.018 mmol) of Intermediate C64 were dissolved in 4 ml of 2,2,2-trifluoroethanol. 15 ml (0.110 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 2 h. 32 mg (0.110 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were then added, and the reaction was concentrated under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 9.5 mg (77% of theory) of the title compound.

20

LC-MS (Method 1):  $R_t = 0.68$  min; MS (ESIpos):  $m/z = 556$   $(M+H)^+$ .

### Example 199

25 4-[(2-[[2-[(2S)-2-Amino-4-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino)-2-oxoethyl]amino]-3-[[{(2R)-2-amino-2-carboxyethyl]sulphonyl]-4-oxobutanoic acid / trifluoroacetic acid (1:1)



in 2 ml of 2,2,2-trifluoroethanol. 12 ml (0.088 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 30 min. 26 mg (0.088 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid were then added, and the solvent was evaporated  
5 under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 8.3 mg (99% of theory) of the title compound as a regioisomer mixture in a ratio of 87:13.

10

LC-MS (Method 5):  $R_t = 2.3$  min and 2.43 min; MS (ESIpos):  $m/z = 832$  (M+H)<sup>+</sup>.

15

<sup>1</sup>H NMR main isomer: (500 MHz, DMSO-*d*<sub>6</sub>):  $\delta = 8.7$  (m, 1H), 8.5 (m, 2H), 8.1 (m, 1H), 7.6 (m, 1H), 7.5 (s, 1H) 7.4-7.15 (m, 6H), 6.9-7.0 (m, 1H), 6.85 (s, 1H), 5.61 (s, 1H), 4.9 and 5.2 (2d, 2H), 4.26 and 4.06 (2d, 2H), 3.5-3.8 (m, 5H), 3.0-3.4 (m, 5H), 2.75-3.0 (m, 3H), 2.58 and 2.57 (dd, 1H), 0.77 and 1,5 (2m, 2H), 0.81 (s, 9H).

20

Alternatively, the regioisomeric title compounds were prepared as follows:

25

To this end, first L-cysteine was converted with 1-([2-(trimethylsilyl)ethoxy]carbonyl)oxy)pyrrolidine-2,5-dione in DMF in the presence of *N,N*-diisopropylethylamine into *N*-{[2-(trimethylsilyl)ethoxy]carbonyl}-L-cysteine.

30

35

55 mg (0.068 mmol) of Intermediate F104 and 36 mg (0.136 mmol) of *N*-{[2-(trimethylsilyl) ethoxy]carbonyl}-L-cysteine were dissolved in 15 ml of DMF, and the mixture was stirred at RT for 20 h. The mixture was then concentrated and the residue was purified by preparative HPLC. The appropriate fractions were combined and the solvents were evaporated under reduced  
35 pressure, and the residue was then dissolved in 15 ml of THF/water 1:1. 131  $\mu$ l of a 2M aqueous lithium hydroxide solution were added and the reaction was stirred at RT for 1 h. The reaction was then neutralized with a 1M hydrochloric acid, the

solvent was evaporated under reduced pressure and the residue was purified by preparative HPLC. This gave 37 mg (50% of theory) of the regioisomeric protected intermediates as a colourless foam.

5

LC-MS (Method 5):  $R_t = 3.33$  min and 3.36 min; MS (ESIpos):  $m/z = 976$  (M+H)<sup>+</sup>.

In the last step, 25 mg of this intermediate were dissolved in 3 ml of 2,2,2-trifluoroethanol. 12.5 ml (0.092 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 4 h. 27 mg (0.092 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid were then added, and the solvent was evaporated under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 18.5 mg (85% of theory) of the title compound as a regioisomer mixture in a ratio of 21:79.

LC-MS (Method 5):  $R_t = 2.37$  min and 3.44 min; MS (ESIpos):  $m/z = 832$  (M+H)<sup>+</sup>.

The targeted preparation of the individual regioisomers of the title compounds was carried out as follows:

25

#### Example 199-2

4-[(2-{{[2-((2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-oxoethyl)amino]-2-[[2R)-2-amino-2-carboxyethyl]sulphonyl]-4-oxobutanoic acid / trifluoroacetic acid (1:1)

First, methyl L-cysteinate was converted with 1-([2-(trimethylsilyl)ethoxy]carbonyl)oxy)pyrrolidine-2,5-dione in DMF in the presence of *N,N*-diisopropylethylamine into methyl *N*-{[2-(trimethylsilyl)ethoxy]carbonyl}-L-cysteinate.



53 mg (0.251 mmol) of commercially available 3-bromo-4-methoxy-4-oxobutanoic acid and 70 mg (0.251 mmol) of methyl N-{{[2-(trimethylsilyl)ethoxy]carbonyl}-L-cysteinate were dissolved in 5 ml of DMF, and aqueous sodium bicarbonate solution was added while monitoring the pH. After stirring 15 min of stirring at RT, the mixture was adjusted to pH=4.3 with acetic acid and the reaction was concentrated. The residue was purified by preparative HPLC. Combination of the appropriate fractions and evaporation of the solvents under reduced pressure gave 72 mg (70% of theory) of 4-methoxy-3-{{[(2R)-3-methoxy-3-oxo-2-{{[2-(trimethylsilyl)ethoxy]carbonyl}amino)propyl]sulphanyl}-4-oxobutanoic acid.

LC-MS (Method 1):  $R_t = 0.93$  min; MS (ESIpos):  $m/z = 410$  (M+H)<sup>+</sup>.

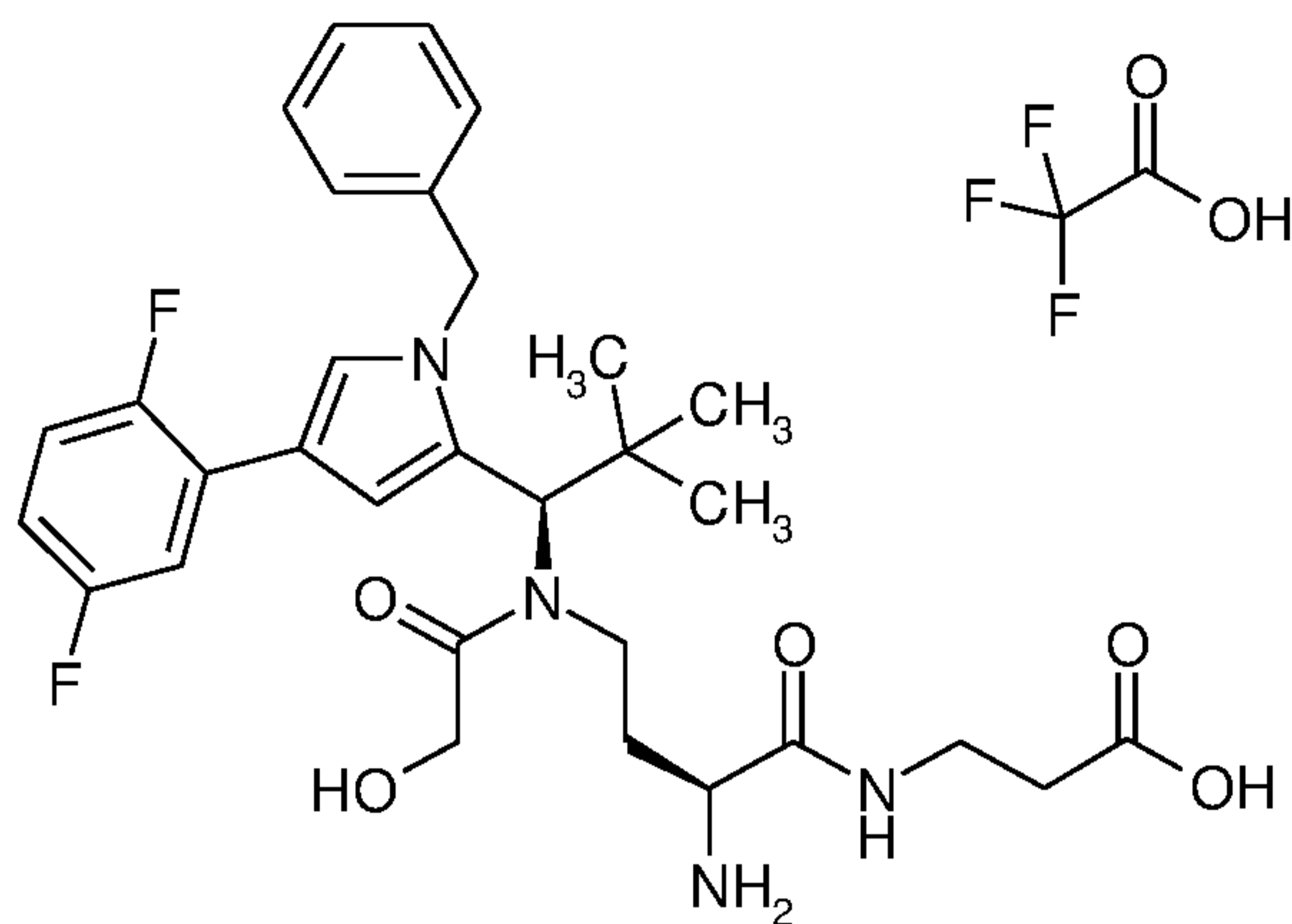
This intermediate was coupled in the presence of HATU with Intermediate C66 and then deprotected completely as described above first with lithium hydroxide in methanol and then with zinc chloride. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 2 mg of the title compound.

LC-MS (Method 1):  $R_t = 0.78$  min; MS (ESIpos):  $m/z = 832$  (M+H)<sup>+</sup>.

Isomer 1 can be prepared in an analogous manner.

### Example 200

N-{{(2S)-2-Amino-4-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-alanine / trifluoroacetic acid (1:1)



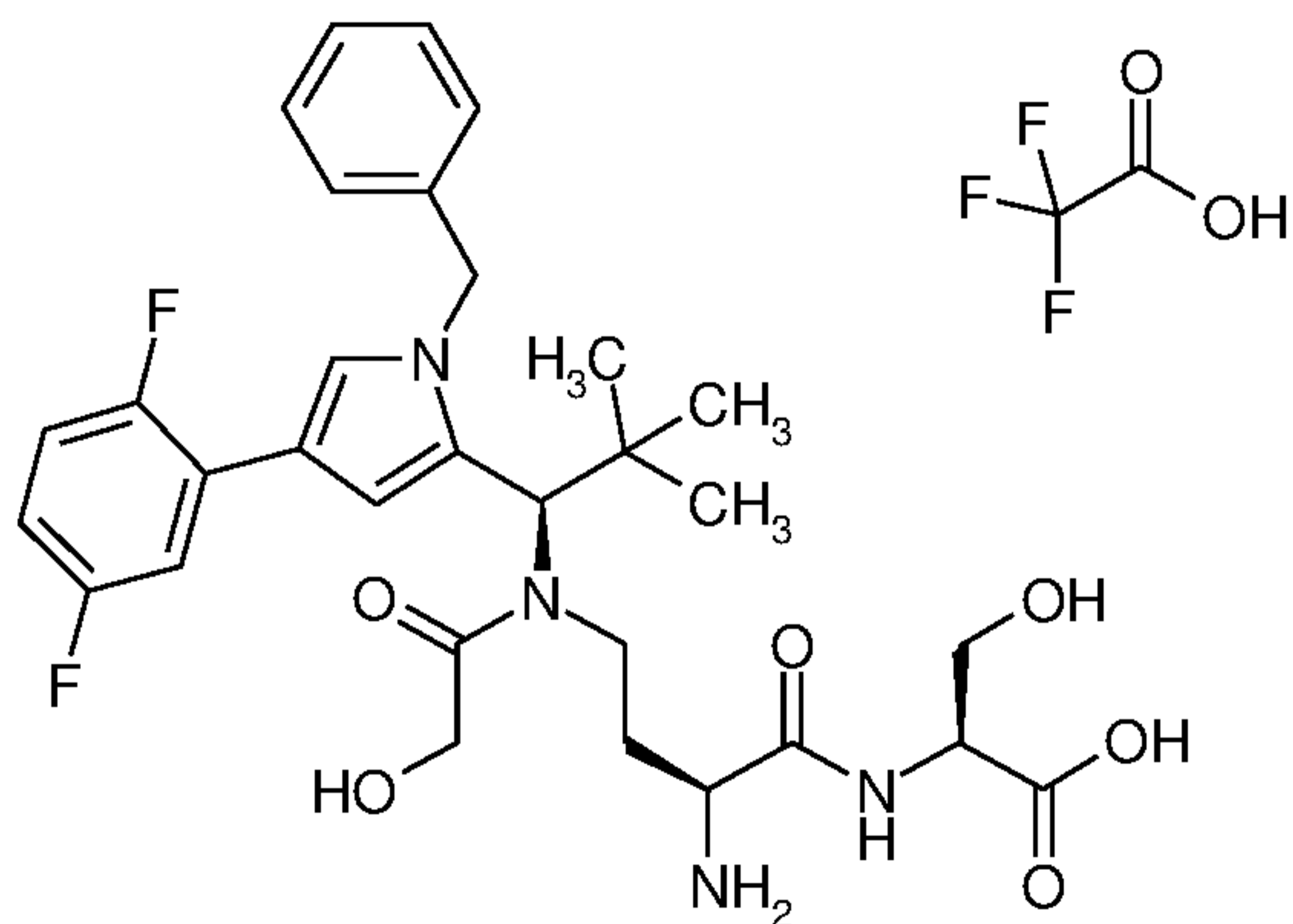
10 mg (0.014 mmol) of Intermediate C61 were dissolved in 3 ml  
 of 2,2,2-trifluoroethanol. 11 ml (0.082 mmol) of zinc chloride  
 5 were added, and the reaction was stirred at 50°C for 30 min. 24  
 mg (0.082 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid  
 were then added, and the solvent was evaporated under reduced  
 pressure. The residue was purified by preparative HPLC.  
 Concentration of the appropriate fractions and lyophilization  
 10 of the residue from acetonitrile/water gave 4.2 mg (40% of  
 theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.79$  min; MS (ESIpos):  $m/z = 585$  (M+H)<sup>+</sup>.

### 15 Example 201

N-{(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl}-  
 L-serine / trifluoroacetic acid (1:1)

20



20 mg (0.03 mmol) of Intermediate C58 and 8.5 mg (0.037 mmol)

of benzyl L-serinate hydrochloride (1:1) were taken up in 5 ml of DMF, and 17 mg (0.046 mmol) of HATU and 21  $\mu$ l of *N,N*-diisopropylethylamine were added. After stirring at RT for 10 minutes, the mixture was concentrated and the residue was purified by preparative HPLC. This gave 12.5 mg (49% of theory) of the intermediate. LC-MS (Method 1):  $R_t$  = 1.42 min; MS (ESIpos):  $m/z$  = 835 (M+H)<sup>+</sup>.

12.5 mg (0.015 mmol) of this intermediate were dissolved in 10 ml of ethanol, palladium on carbon (10%) was added and the mixture was hydrogenated at RT with hydrogen at standard pressure for 30 min. The catalyst was filtered off and the solvents were evaporated under reduced pressure giving, after lyophilization of the residue from acetonitrile/water, 7.5 mg (67% of theory) of the intermediate.

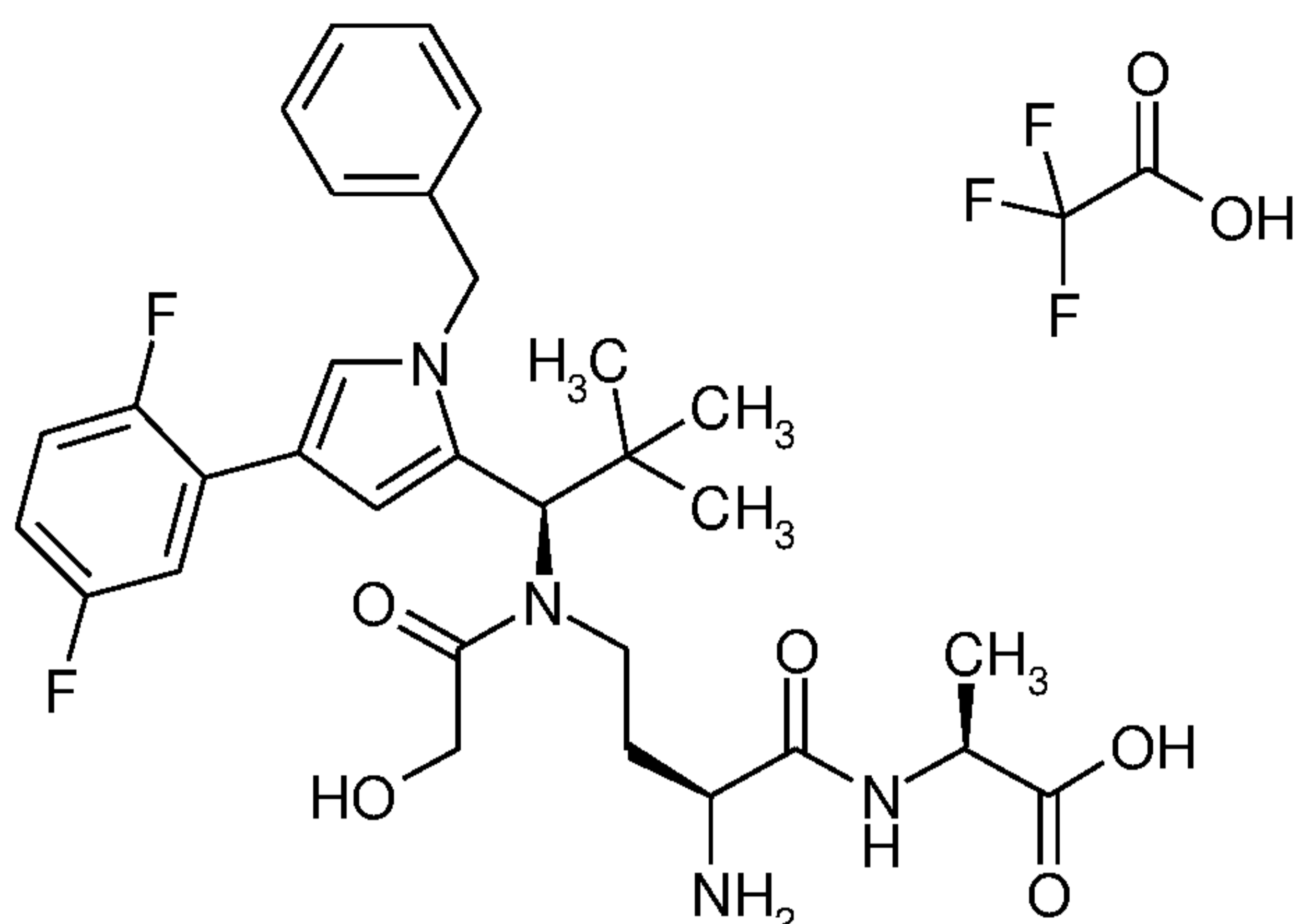
LC-MS (Method 1):  $R_t$  = 1.28 min; MS (ESIpos):  $m/z$  = 745 (M+H)<sup>+</sup>.

7.5 mg (0.01 mmol) of this intermediate were dissolved in 3 ml of 2,2,2-trifluoroethanol. 8 ml (0.06 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 4.5 h. 17.7 mg (0.06 mmol) of ethylenediamine-*N,N,N',N'*-tetraacetic acid were then added, and the solvent was evaporated under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 4.2 mg (58% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.75 min; MS (ESIpos):  $m/z$  = 601 (M+H)<sup>+</sup>.

### Example 202

*N*-{(2*S*)-2-Amino-4-[(1*R*)-1-[1-benzyl-4-(2,5-difluorophenyl)-1*H*-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-*L*-alanine / trifluoroacetic acid (1:1)



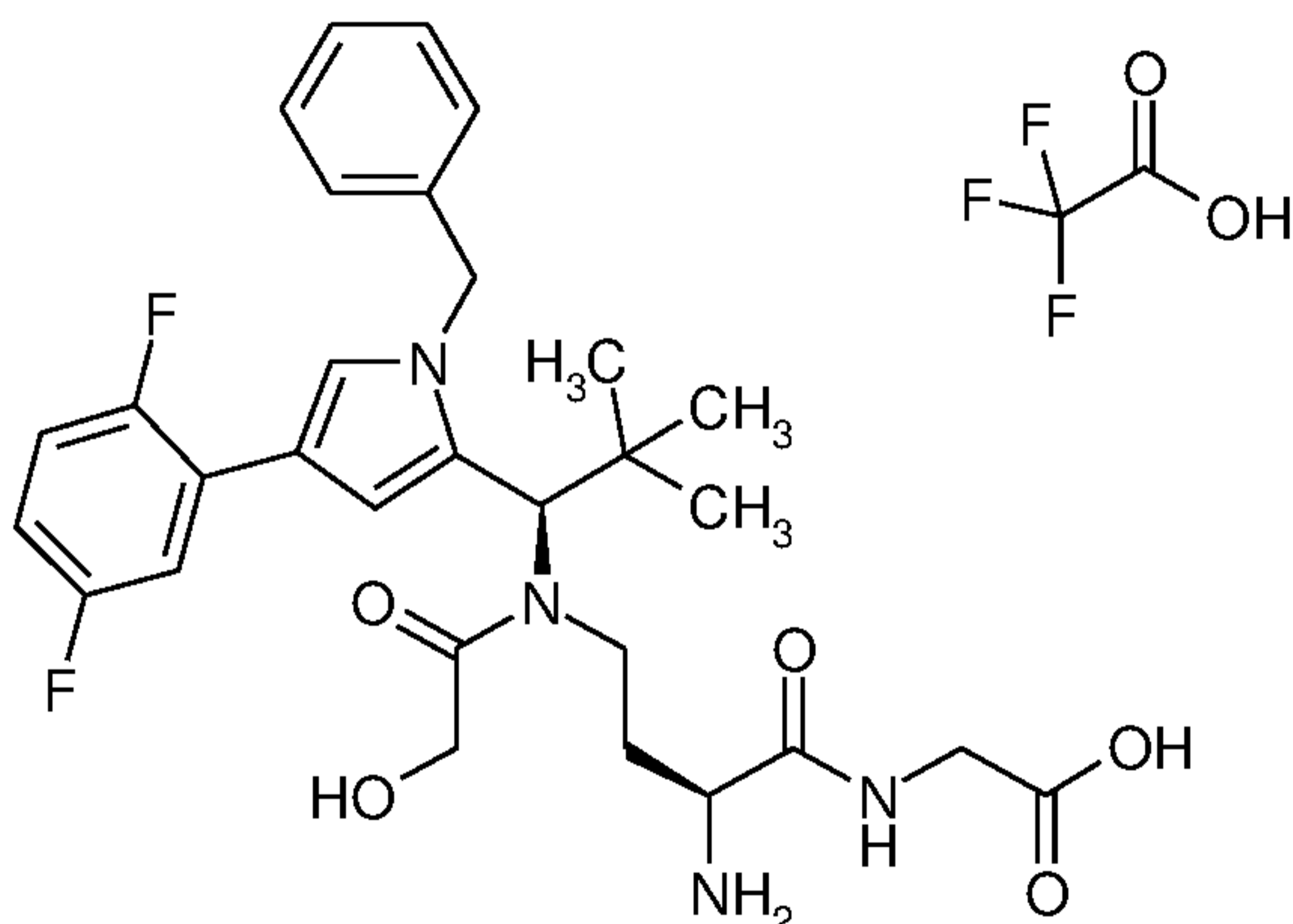
The title compound was prepared analogously to Example 201 from Intermediate C58 and benzyl L-alaninate hydrochloride (1:1).

5

LC-MS (Method 1):  $R_t = 0.82$  min; MS (ESIpos):  $m/z = 585$  (M+H)<sup>+</sup>.

### Example 203

10 N-{(2S)-2-Amino-4-[[{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}glycine /  
trifluoroacetic acid (1:1)

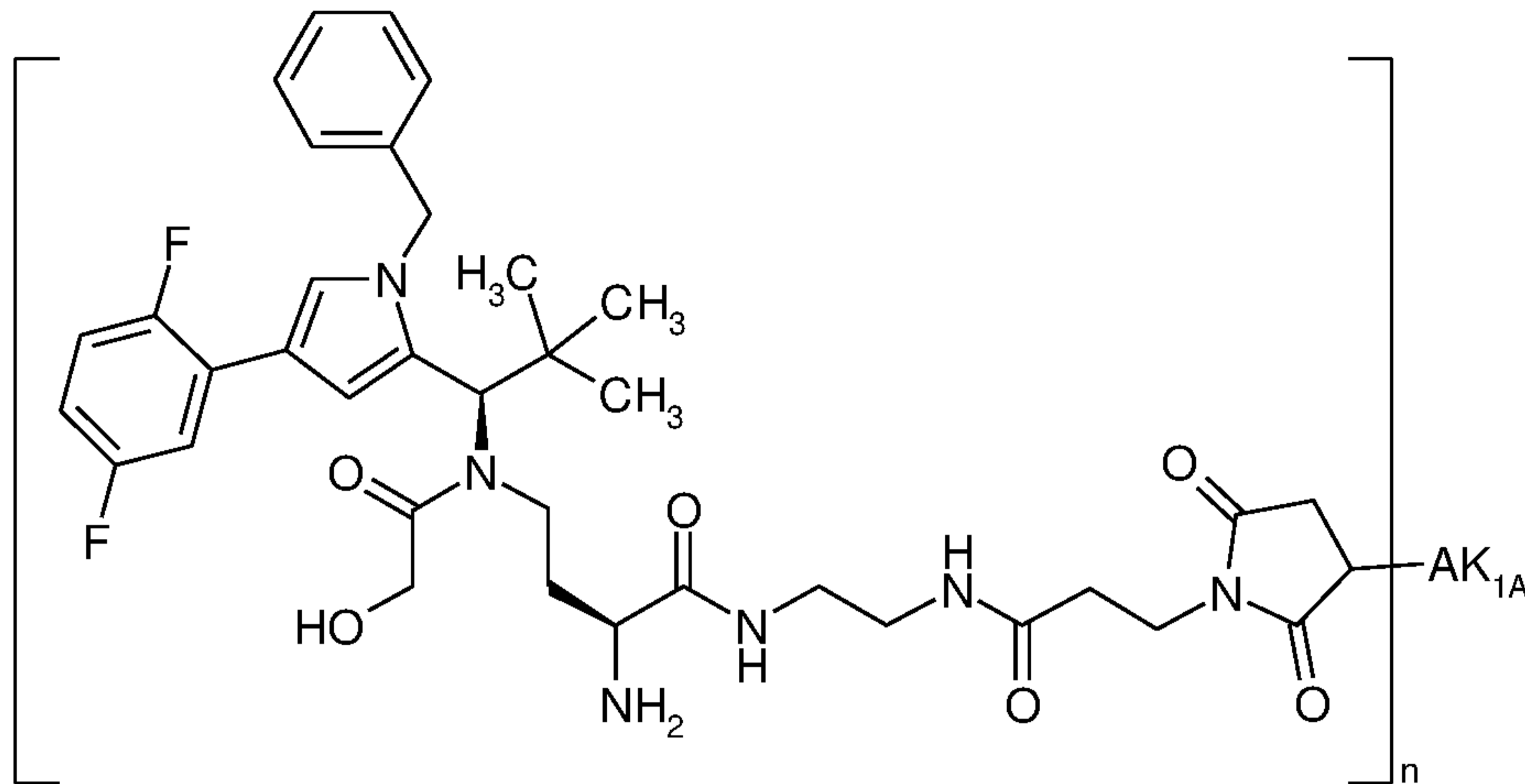


15

The title compound was prepared analogously to Example 201 from Intermediate C58 and benzyl glycinate hydrochloride (1:1).

20 LC-MS (Method 1):  $R_t = 0.82$  min; MS (ESIpos):  $m/z = 571$  (M+H)<sup>+</sup>.

### Example 204A



Here, 5 mg of cetuximab in PBS (c=23.1 mg/ml) were used for coupling with Intermediate F204, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.88 mg/ml

Drug/mAb ratio: 2.6

#### Example 204B

15

Here, 50 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F204, and the reaction was, after  
 Sephadex purification, concentrated by ultracentrifugation, rediluted with PBS and concentrated again. Some of the ADC may  
 20 also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 12.66 mg/ml

25 Drug/mAb ratio: 3.5

#### Example 204E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.5 mg/ml) were

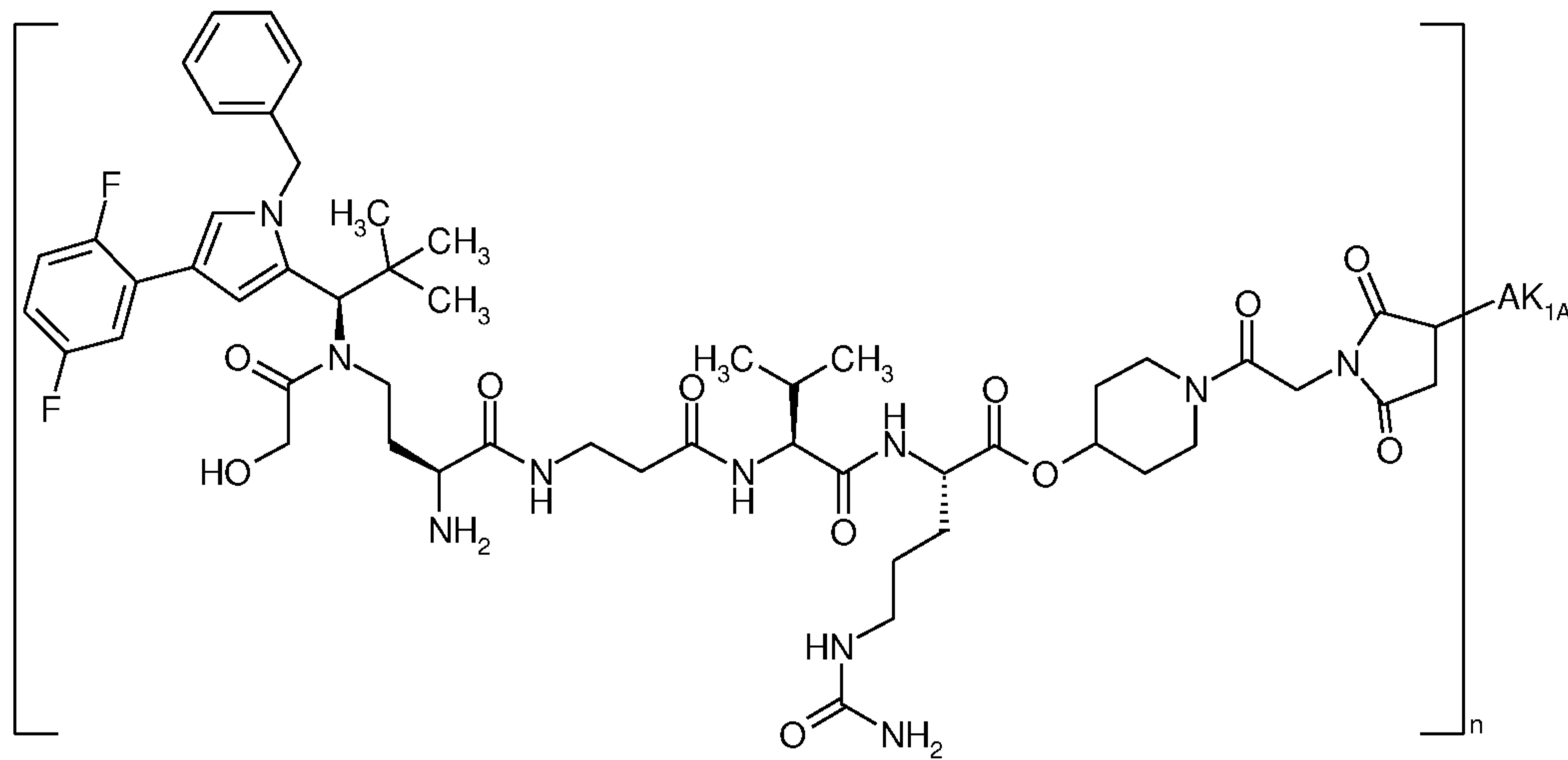
used for coupling with Intermediate F204, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.65 mg/ml

Drug/mAb ratio: 3.5

10

### Example 205A



15 Here, 5 mg of cetuximab in PBS (c=23.1 mg/ml) were used for coupling with Intermediate F205, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.99 mg/ml

Drug/mAb ratio: 3.2

25

### Example 205B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F205, and the reaction was, after

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 0.96 mg/ml

Drug/mAb ratio: 2.6

### 10 Example 205E

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F205, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

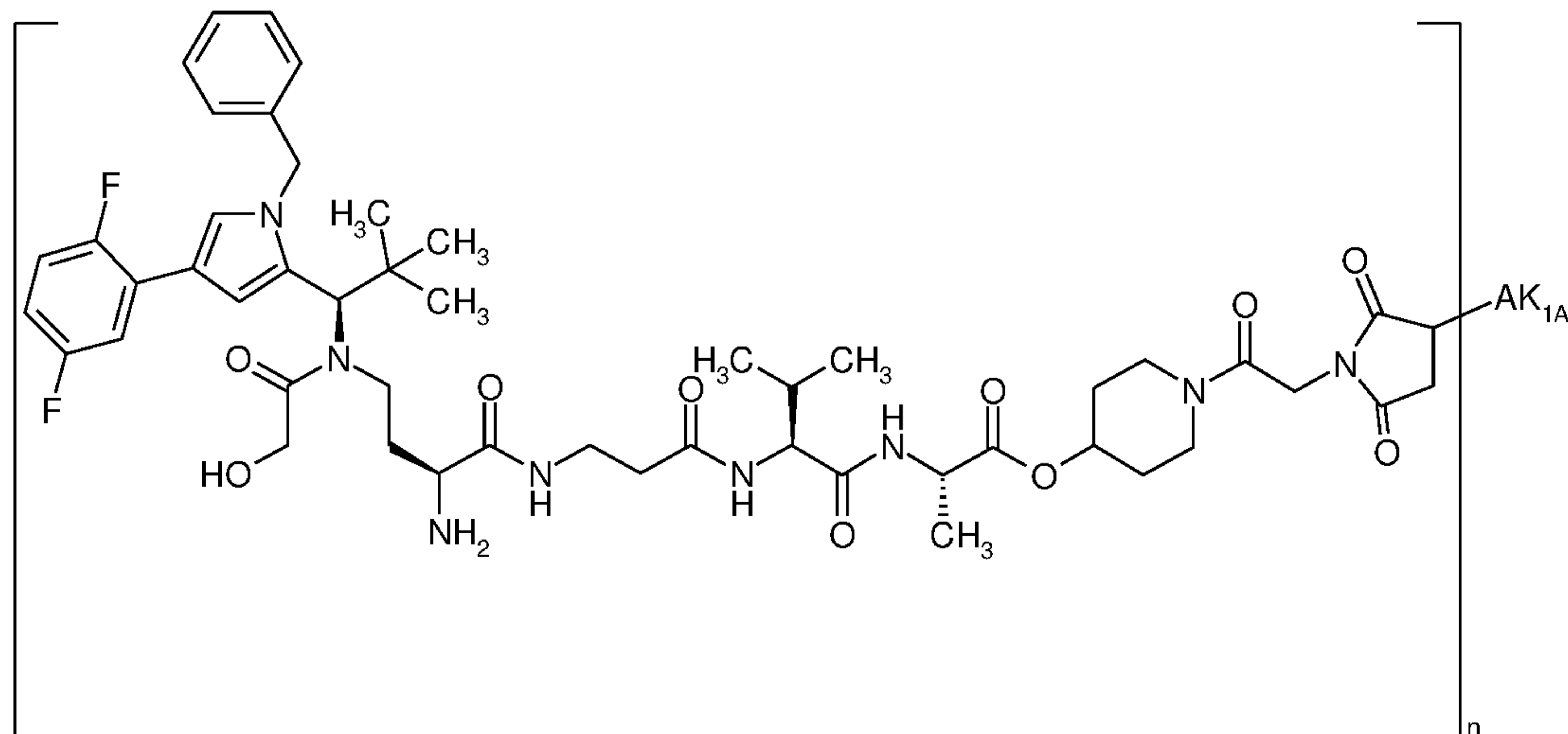
15

Protein concentration: 1.94 mg/ml

20

Drug/mAb ratio: 3.6

### Example 206A



25

Here, 5 mg of cetuximab in PBS (c=23.1 mg/ml) were used for coupling with Intermediate F206, and the reaction was, after

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 1.73 mg/ml

Drug/mAb ratio: 2.6

#### 10 Example 206B

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F206, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 1.21 mg/ml

20

Drug/mAb ratio: 2.0

#### Example 206E

Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F206, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

30

Protein concentration: 1.84 mg/ml

Drug/mAb ratio: 2.8

35

#### Example 207A





Here, 5 mg of trastuzumab in PBS (c=13.5 mg/ml) were used for coupling with Intermediate F207, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 2.39 mg/ml

10 Drug/mAb ratio: 2.8

#### Example 207I

Here, 5.0 mg of nimotuzumab in PBS (c=13.1 mg/ml) were used for coupling with Intermediate F207, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

20

Protein concentration: 1.92 mg/ml

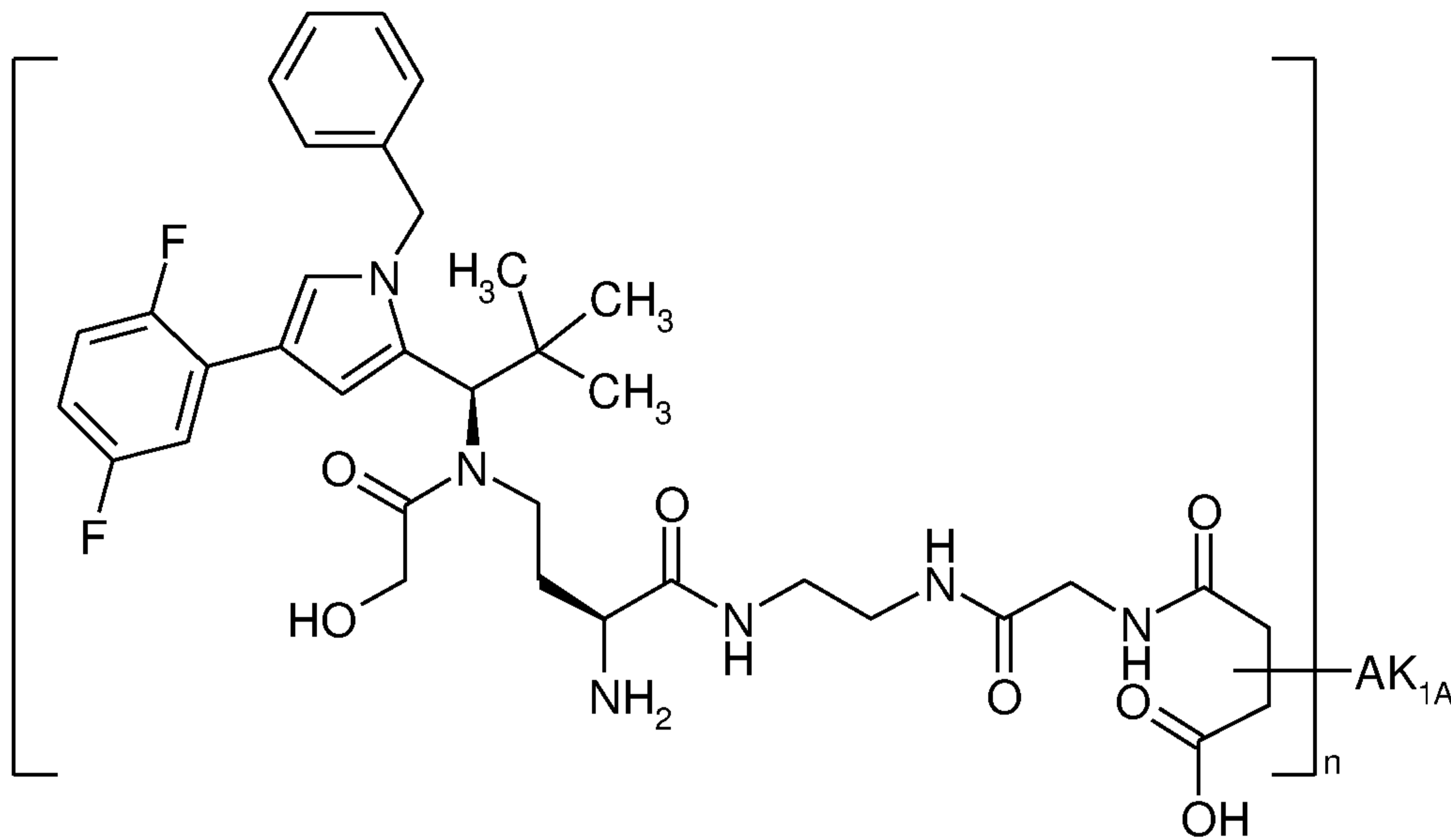
Drug/mAb ratio: 3.5

#### Example 207H

Here, 5.0 mg of panitumumab in PBS (c = 13.6 mg/ml) were used for coupling with Intermediate F207. The reduction time with TCEP was 4 h and the stirring time for the ADC coupling was 20 h. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

35 Protein concentration: 2.0 mg/ml

Drug/mAb ratio: 1.9

Example 208A

5 Under argon, a solution of 0.344 mg of TCEP in 100  $\mu$ l of PBS buffer was added to 60 mg of cetuximab in 5494  $\mu$ l of PBS (c=10.92 mg/ml). The reaction was stirred at RT for 30 min, and 2.582 mg (0.003 mmol) of Intermediate F104 dissolved in 600  $\mu$ l of DMSO were then added. After a further 120 min of stirring at RT, the  
 10 reaction was diluted with 1306  $\mu$ l of PBS buffer which had been adjusted to pH 8 beforehand.

This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was diluted  
 15 with PBS buffer pH 8 to a total volume of 14 ml. This solution was stirred under argon at RT overnight and then concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and reconcentrated again. The ADC batch obtained was characterized  
 20 as follows:

Protein concentration: 13.36 mg/ml

Drug/mAb ratio: 1.8

25

For this ADC preparation, a proportion of 94% was determined for the ring-opened succinamide form.

**Example 208B**

Under argon, 60 mg of anti-TWEAKR AK-1 in 3225 µl of PBS (c=18.6  
5 mg/ml) were diluted with 775 µl of PBS buffer, and a solution  
of 0.344 mg of TCEP in 100 µl of PBS buffer was then added. The  
reaction was stirred at RT for 30 min, and 2.582 mg (0.003 mmol)  
of Intermediate F104 dissolved in 600 µl of DMSO were then added.  
After a further 120 min of stirring at RT, the reaction was  
10 diluted with 300 µl of PBS buffer which had been adjusted to pH  
8 beforehand.

This solution was then applied to PD 10 columns (Sephadex® G-  
25, GE Healthcare) which had been equilibrated with PBS buffer  
15 pH 8 and was eluted with PBS buffer pH 8. The eluate was diluted  
with PBS buffer pH 8 to a total volume of 14 ml. This solution  
was stirred under argon at RT overnight and then concentrated  
by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and  
reconcentrated again. The ADC batch obtained was characterized  
20 as follows:

Protein concentration: 14.95 mg/ml

Drug/mAb ratio: 3.2

25

**Example 208I**

Under argon, a solution of 0.344 mg of TCEP in 100 µl of PBS  
buffer was added to 60 mg of nimotuzumab in 4587 µl of PBS  
30 (c=13.1 mg/ml). The reaction was stirred at RT for 30 min, and  
2.582 mg (0.003 mmol) of Intermediate F104 dissolved in 600 µl  
of DMSO were then added. After a further 120 min of stirring at  
RT, the reaction was diluted with 2213 µl of PBS buffer which  
had been adjusted to pH 8 beforehand.

35

This solution was then applied to PD 10 columns (Sephadex® G-  
25, GE Healthcare) which had been equilibrated with PBS buffer  
pH 8 and was eluted with PBS buffer pH 8. The eluate was diluted

with PBS buffer pH 8 to a total volume of 14 ml. This solution was stirred under argon at RT overnight and then concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and reconcentrated again. The ADC batch obtained was characterized  
5 as follows:

Protein concentration: 14.79 mg/ml

Drug/mAb ratio: 3.1

10

For this ADC preparation, a proportion of 91% was determined for the ring-opened succinamide form.

#### Example 208K

15

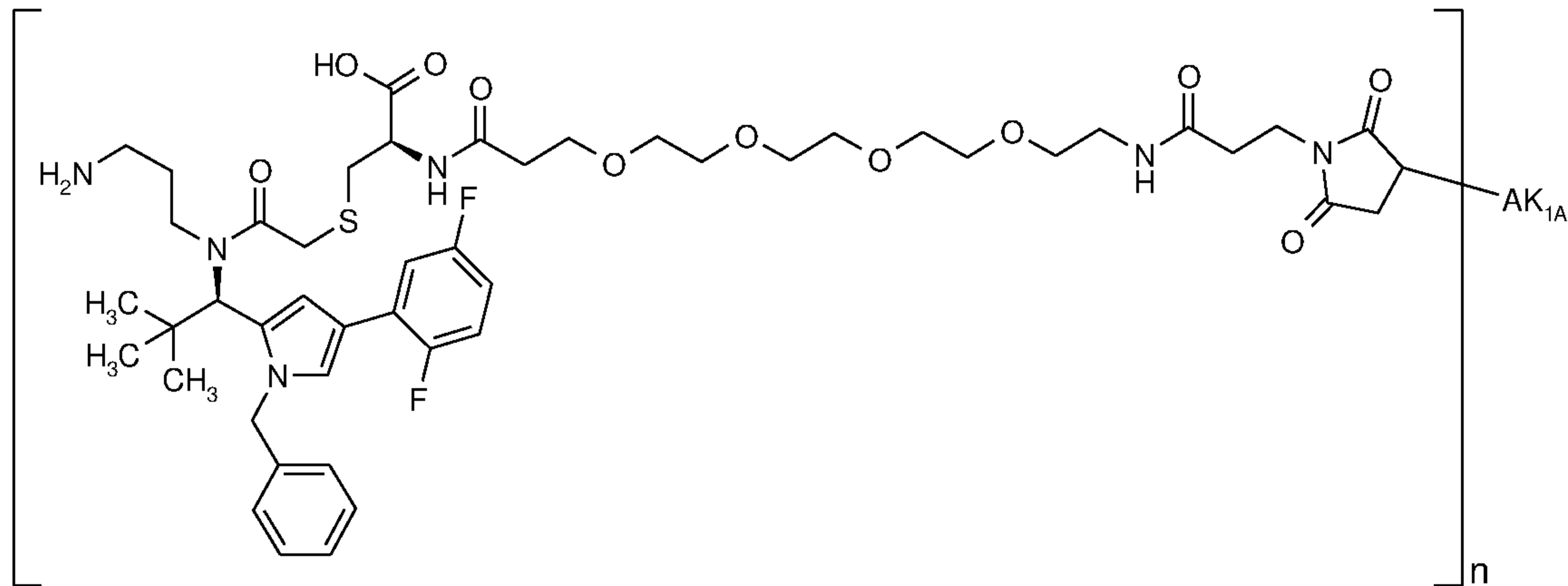
Under argon, a solution of 0.23 mg of TCEP in 67 µl of PBS buffer was added to 40 mg of anti-TWEAKR AK-2 in 2759 µl of PBS (c=14.5 mg/ml). The reaction was stirred at RT for 30 min, and 1.72 mg (0.002 mmol) of Intermediate F104 dissolved in 400 µl of DMSO  
20 were then added. After a further 120 min of stirring at RT, the reaction was diluted with 1774 µl of PBS buffer which had been adjusted to pH 8 beforehand.

This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-  
25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was diluted with PBS buffer pH 8 to a total volume of 14 ml. This solution was stirred under argon at RT overnight and then concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and  
30 reconcentrated again. The ADC batch obtained was characterized as follows:

Protein concentration: 11.66 mg/ml

35 Drug/mAb ratio: 3.1

#### Example 209A



Here, 5.0 mg of cetuximab in PBS ( $c=21.32$  mg/ml) were used for coupling with Intermediate F209, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.75 mg/ml

Drug/mAb ratio: 2.4

#### Example 209B

15

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS ( $c=18.60$  mg/ml) were used for coupling with Intermediate F209, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be  
 20 present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.30 mg/ml

25 Drug/mAb ratio: 2.1

#### Example 209E

Here, 5.0 mg of trastuzumab antibody in PBS ( $c=13.50$  mg/ml) were used for coupling with Intermediate F209, and the reaction was,  
 30

after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 2.03 mg/ml

Drug/mAb ratio: 2.3

#### 10 Example 209H

Here, 5.0 mg of panitumumab antibody in PBS (c=70.5 mg/ml) were used for coupling with Intermediate F209, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 1.91 mg/ml

20

Drug/mAb ratio: 1.5

#### Example 209I

Here, 5.0 mg of nimotuzumab antibody in PBS (c=13.1 mg/ml) were used for coupling with Intermediate F209, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

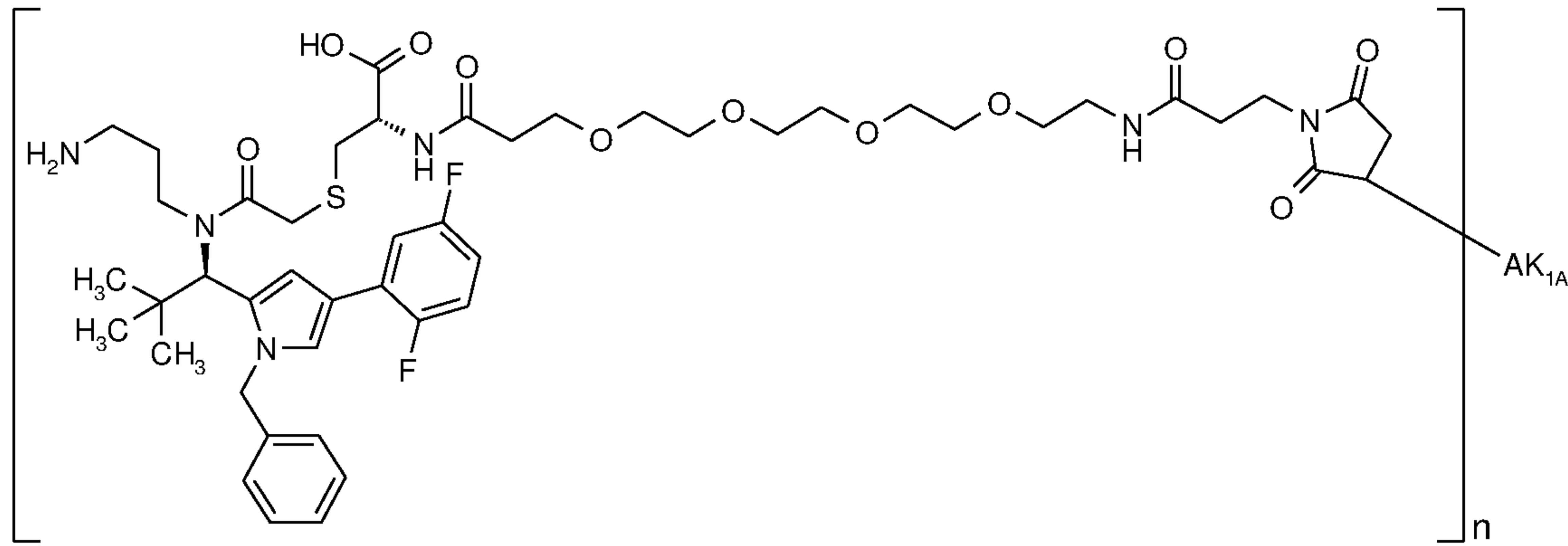
30

Protein concentration: 1.81 mg/ml

Drug/mAb ratio: 2.5

35

#### Example 210A



Here, 5.0 mg of cetuximab in PBS (c=21.32 mg/ml) were used for coupling with Intermediate F210, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.92 mg/ml

Drug/mAb ratio: 2.9

#### Example 210B

15

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.60 mg/ml) were used for coupling with Intermediate F210, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be  
 20 present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.41 mg/ml

25 Drug/mAb ratio: 2.5

#### Example 210E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were  
 30 used for coupling with Intermediate F210, and the reaction was, after Sephadex purification, concentrated by



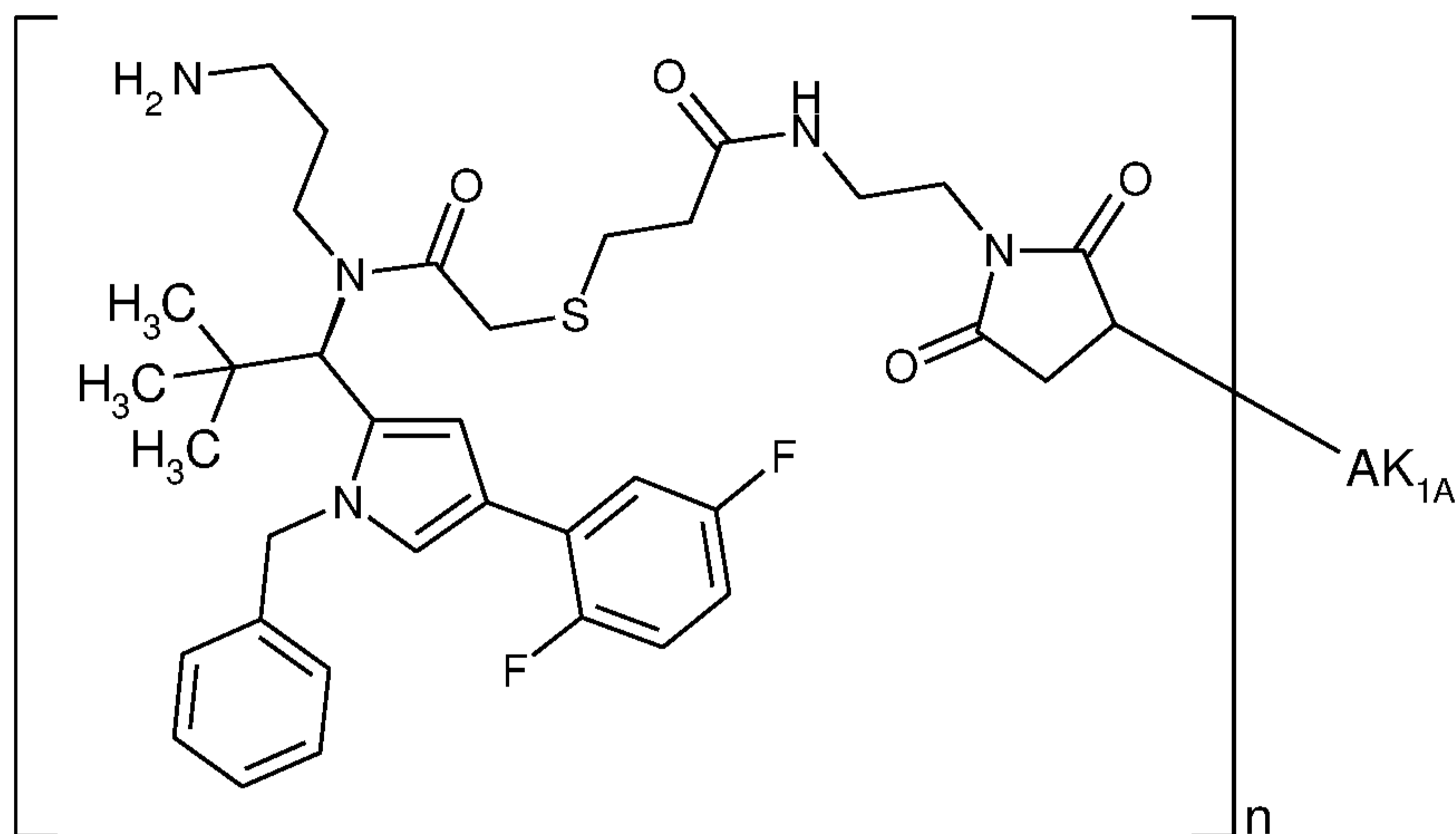
ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5 Protein concentration: 1.79 mg/ml

Drug/mAb ratio: 2.8

**Example 211A**

10



15 Here, 5.0 mg of cetuximab in PBS (c=12.33 mg/ml) were used for coupling with Intermediate F211, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

20 Protein concentration: 1.82 mg/ml

Drug/mAb ratio: 2.0

**Example 211B**

25

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=34.42 mg/ml) were used for coupling with Intermediate F211, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be

present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.52 mg/ml

5

Drug/mAb ratio: 2.4

### Example 211E

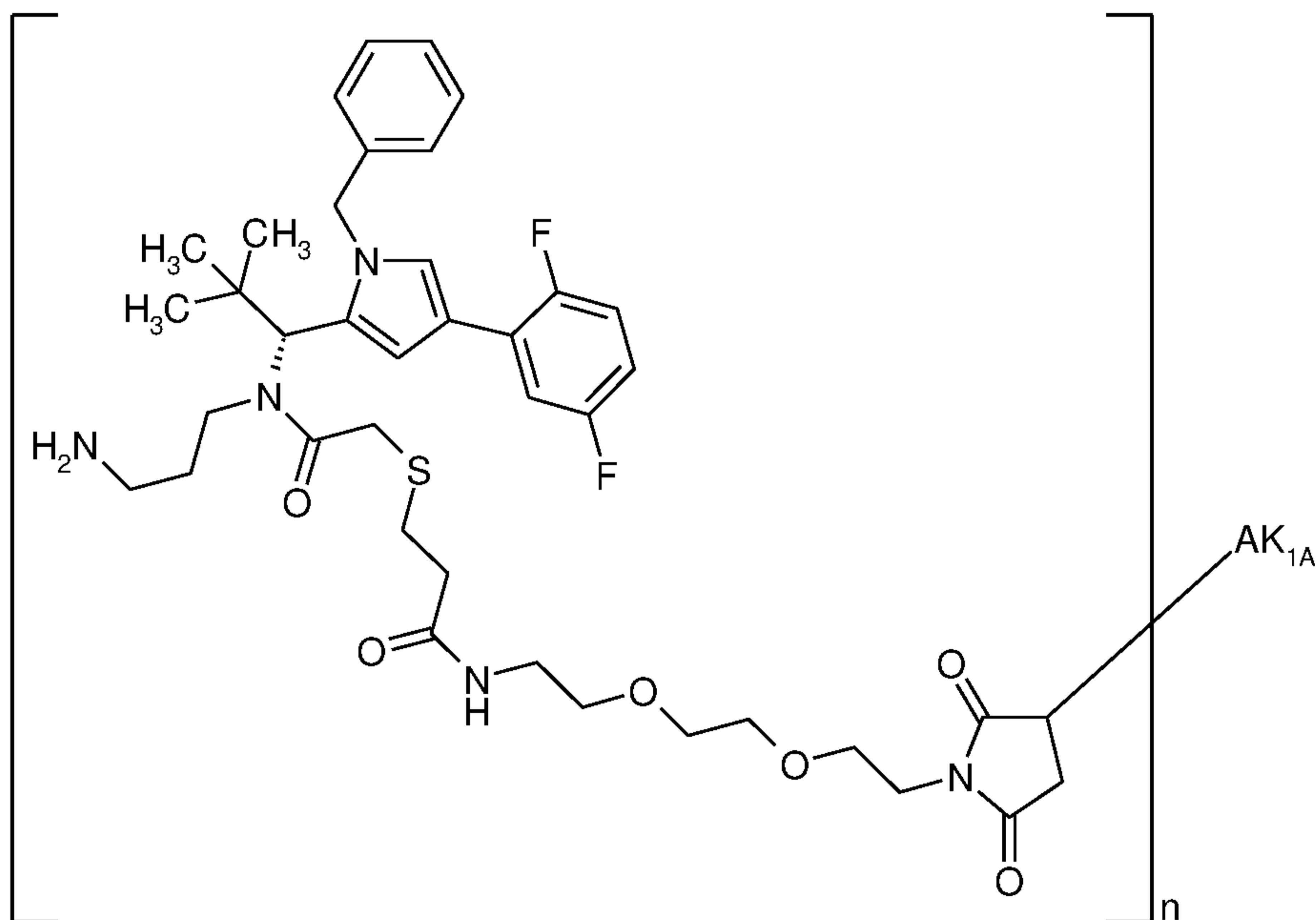
10 Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F211, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides  
15 attached to the antibody.

Protein concentration: 1.84 mg/ml

Drug/mAb ratio: 2.4

20

### Example 212A



25 Here, 5.0 mg of cetuximab in PBS (c=11.3 mg/ml) were used for coupling with Intermediate F212, and the reaction was, after

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 1.94 mg/ml

Drug/mAb ratio: 3.3

#### 10 Example 212B

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F212, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 0.85 mg/ml

20

Drug/mAb ratio: 2.0

#### Example 212E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F212, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

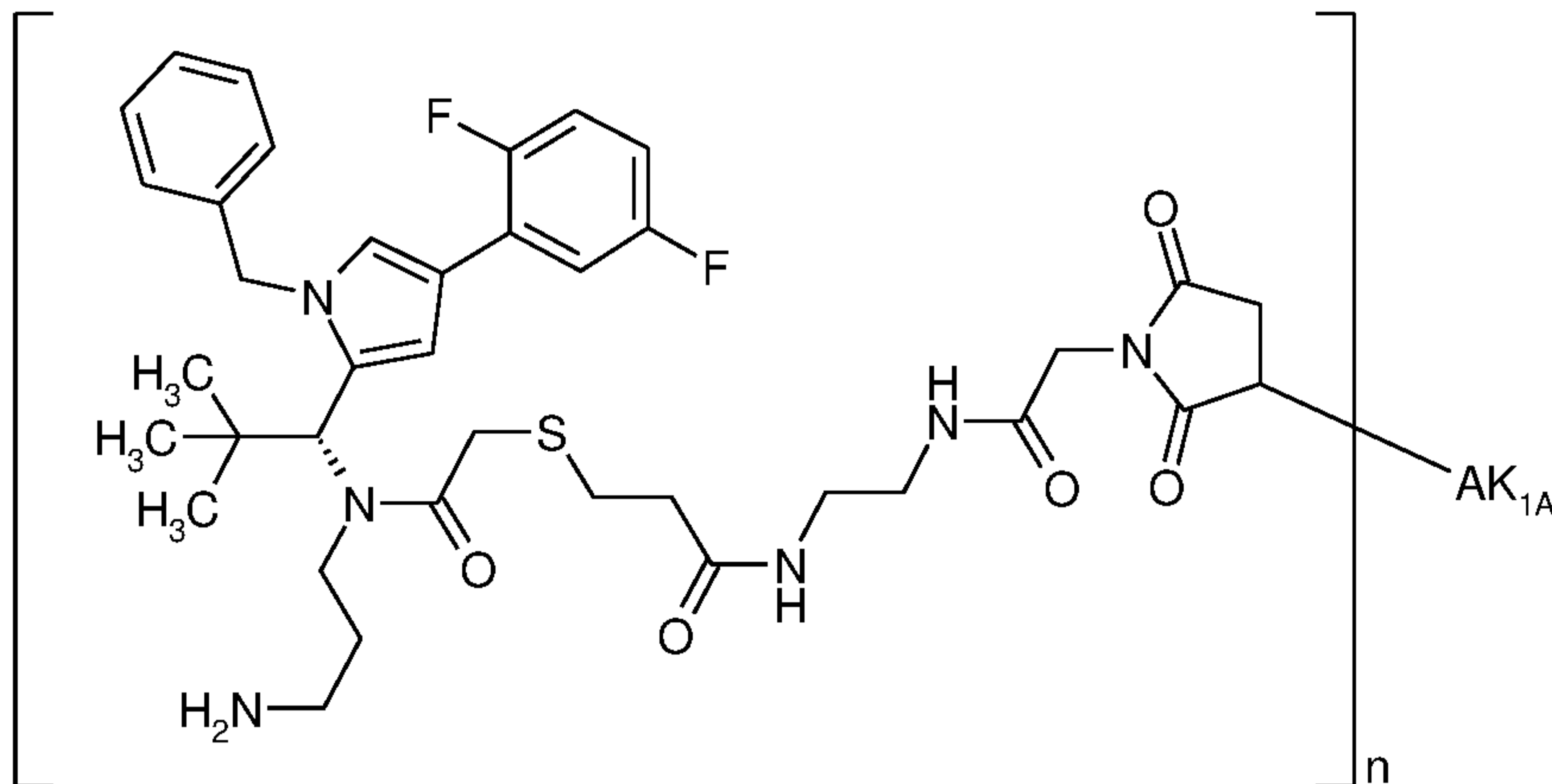
30

Protein concentration: 1.55 mg/ml

Drug/mAb ratio: 2.0

35

#### Example 213A



Here, 5.0 mg of cetuximab in PBS (c=21.32 mg/ml) were used for coupling with Intermediate F213, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 1.83 mg/ml

Drug/mAb ratio: 2.4

### Example 213B

15

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F213, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be  
 20 present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.4 mg/ml

25 Drug/mAb ratio: 2.3

### Example 213E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were  
 30 used for coupling with Intermediate F213, and the reaction was,

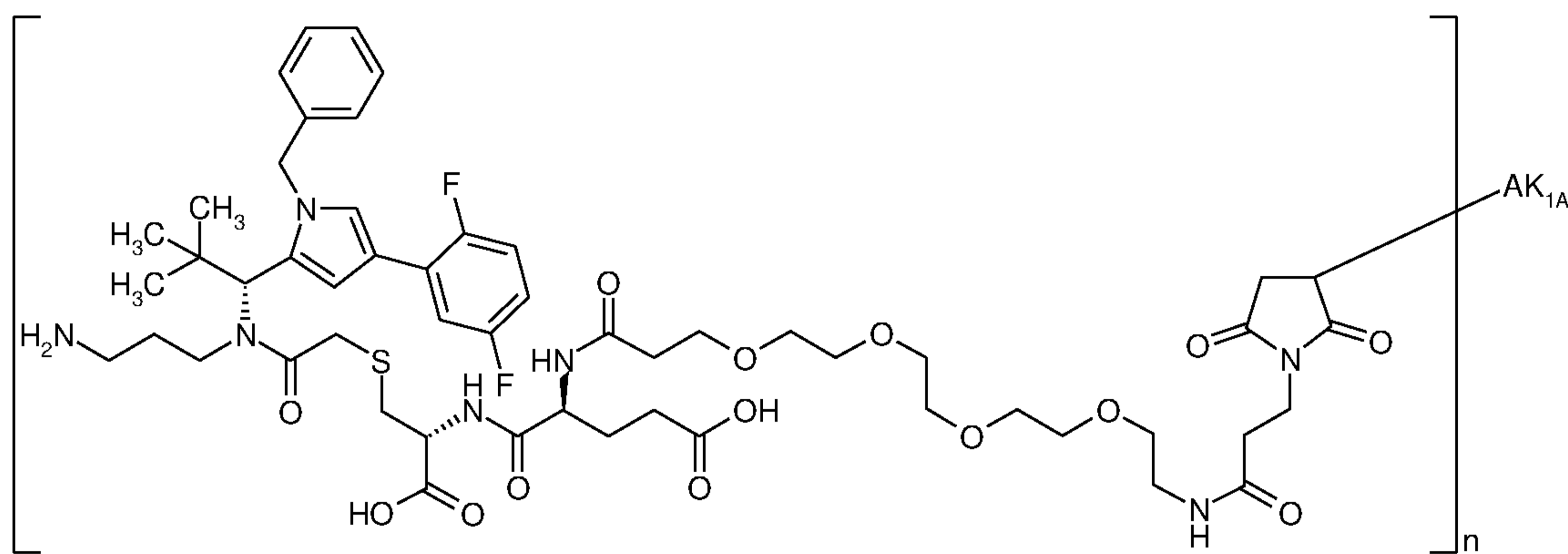
after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

Protein concentration: 1.94 mg/ml

Drug/mAb ratio: 2.5

### 10 Example 214A



Here, 5.0 mg of cetuximab in PBS (c=15.21 mg/ml) were used for coupling with Intermediate F214, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

20

Protein concentration: 2.00 mg/ml

Drug/mAb ratio: 2.6

### 25 Example 214B

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F214, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides

30

attached to the antibody.

Protein concentration: 2.01 mg/ml

5 Drug/mAb ratio: 2.6

### Example 214E

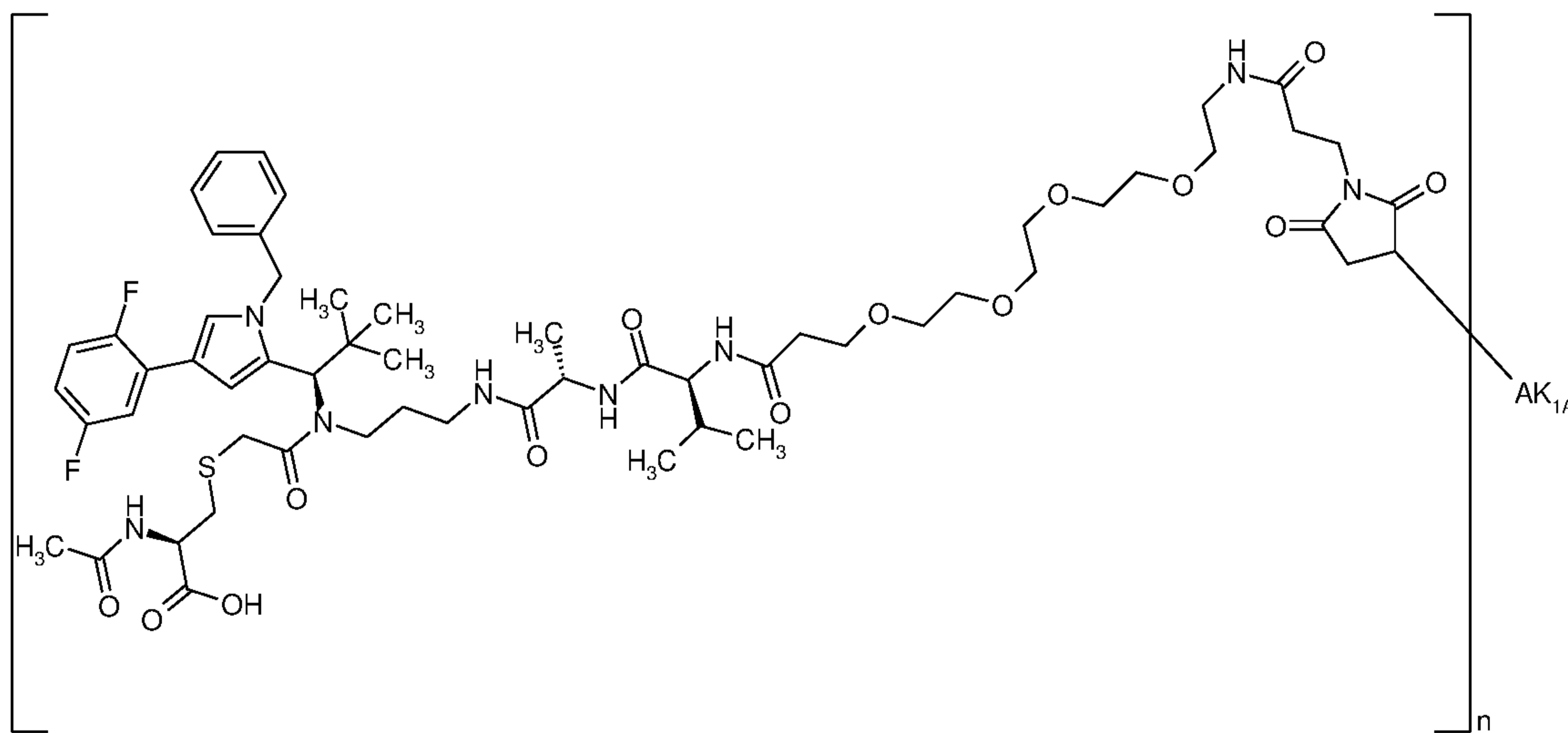
10 Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F214, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 1.86 mg/ml

Drug/mAb ratio: 2.7

### 20 Example 215A



25 Here, 5.0 mg of cetuximab in PBS (c=15.21 mg/ml) were used for coupling with Intermediate F215, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the

antibody.

Protein concentration: 1.99 mg/ml

5 Drug/mAb ratio: 2.7

#### Example 215B

10 Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F215, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 1.64 mg/ml

Drug/mAb ratio: 2.8

#### 20 Example 215E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F215, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

25

Protein concentration: 1.84 mg/ml

30

Drug/mAb ratio: 2.8

#### Example 215H

35 Here, 5.0 mg of panitumumab antibody in PBS (c=70.5 mg/ml) were used for coupling with Intermediate F215, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be

present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.86 mg/ml

5

Drug/mAb ratio: 1.4

### Example 215I

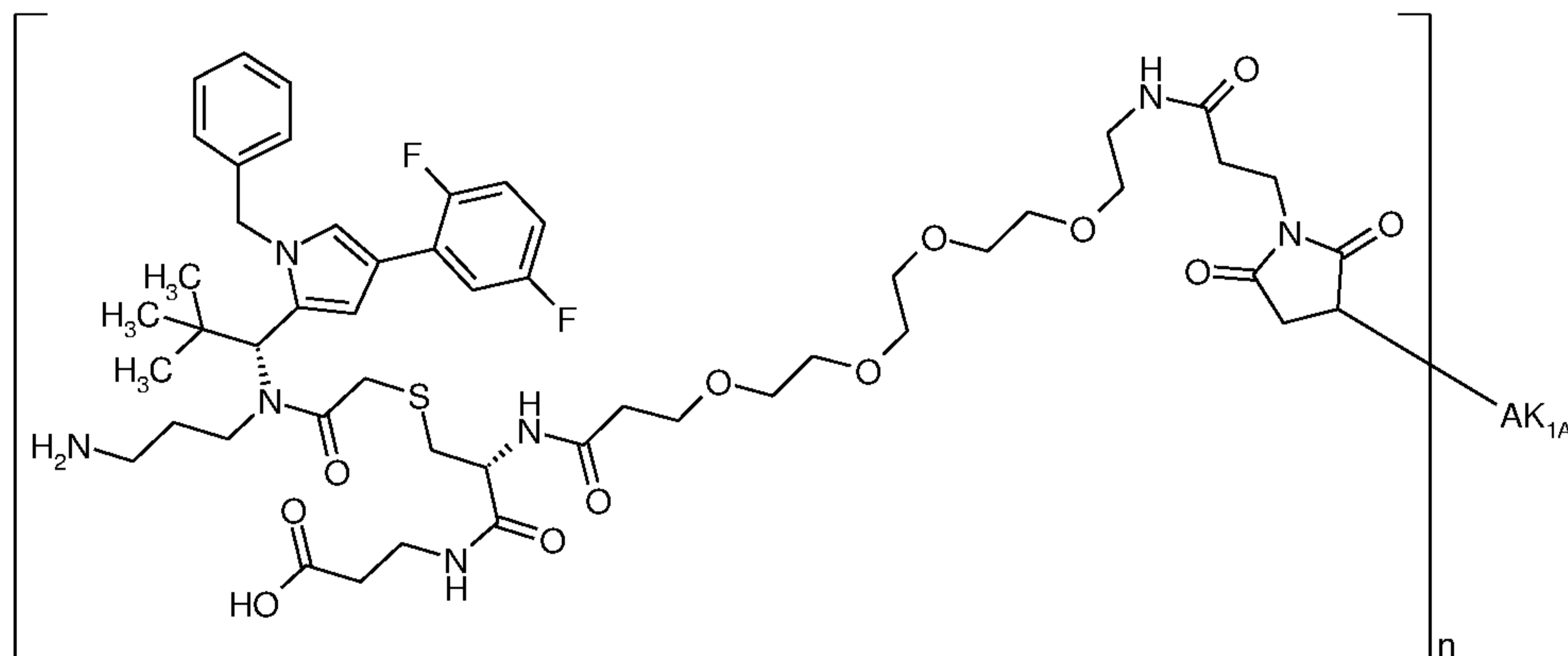
10 Here, 5.0 mg of nimotuzumab in PBS (c=13.1 mg/ml) were used for  
coupling with Intermediate F215, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS. Some of the ADC may also be present in the  
15 form of the hydrolysed open-chain succinamides attached to the  
antibody.

Protein concentration: 1.83 mg/ml

Drug/mAb ratio: 2.6

20

### Example 216A



25 Here, 5.0 mg of cetuximab in PBS (c=15.21 mg/ml) were used for  
coupling with Intermediate F216, and the reaction was, after  
Sephadex purification, concentrated by ultracentrifugation and  
rediluted with PBS. Some of the ADC may also be present in the  
form of the hydrolysed open-chain succinamides attached to the  
30 antibody.



Protein concentration: 1.97 mg/ml

Drug/mAb ratio: 2.8

5 **Example 216B**

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F216, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.94 mg/ml

15

Drug/mAb ratio: 2.5

**Example 216E**

20 Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F216, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

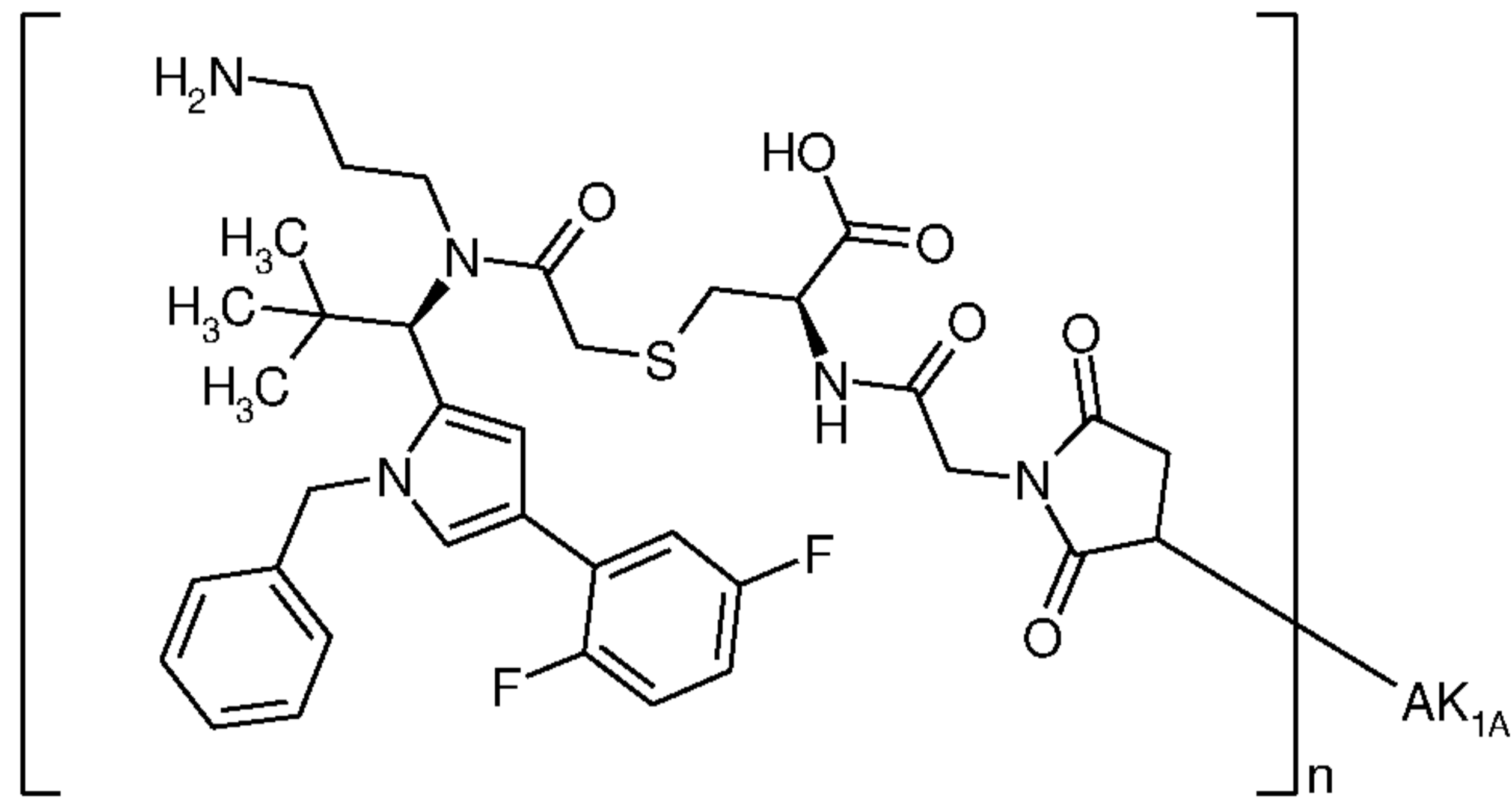
25

Protein concentration: 1.90 mg/ml

Drug/mAb ratio: 1.7

30

**Example 217A**



Here, 5.0 mg of cetuximab in PBS (c=15.21 mg/ml) were used for coupling with Intermediate F217, and the reaction was, after  
 5 Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

10 Protein concentration: 2.05 mg/ml

Drug/mAb ratio: 2.7

#### Example 217B

15

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F217, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be  
 20 present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.44 mg/ml

25 Drug/mAb ratio: 2.4

#### Example 217E

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F217, and the reaction was, after  
 30 Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be

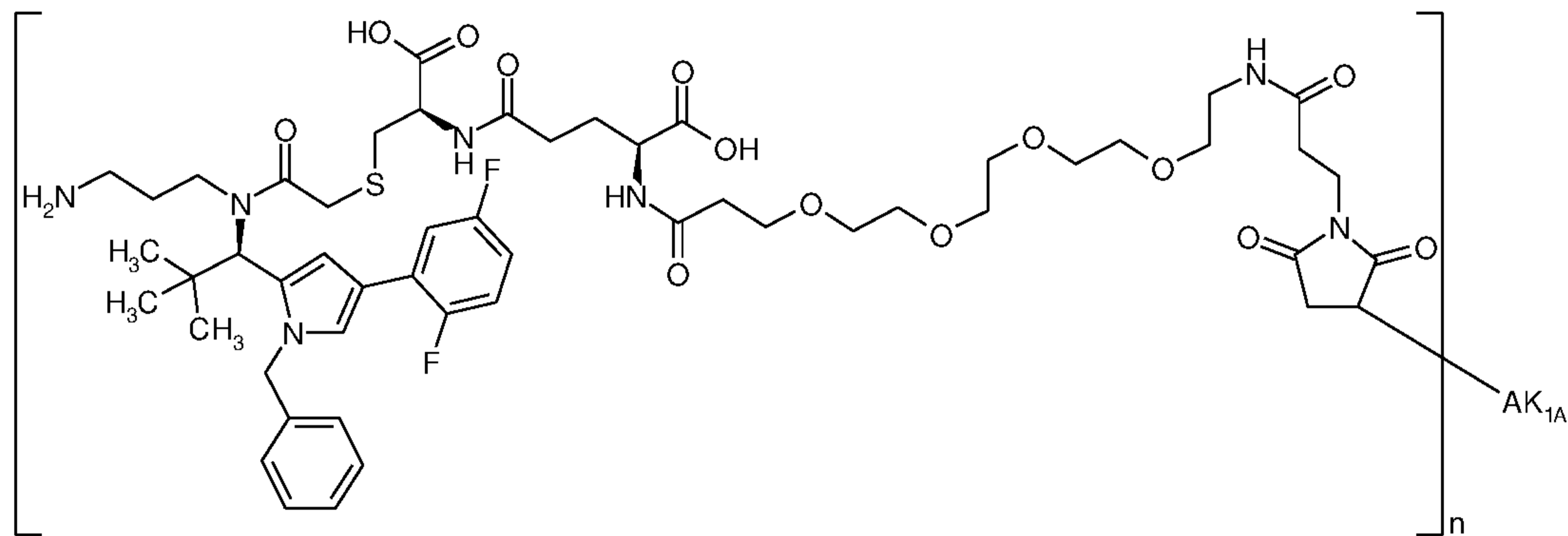
present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.85 mg/ml

5

Drug/mAb ratio: 2.8

### Example 218A



10

Here, 5.0 mg of cetuximab in PBS (c=15.21 mg/ml) were used for coupling with Intermediate F218, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

15

Protein concentration: 2.05 mg/ml

20

Drug/mAb ratio: 3.0

### Example 218B

Here, 5.0 mg of anti-TWEAKR AK-1 antibody in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F218, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

30

Protein concentration: 1.95 mg/ml

Drug/mAb ratio: 2.9

5 **Example 218E**

Here, 5.0 mg of trastuzumab antibody in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F218, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.84 mg/ml

15

Drug/mAb ratio: 2.7

**Example 218H**

20 Here, 5.0 mg of panitumumab antibody in PBS (c = 20 mg/ml) were used for coupling with Intermediate F218. The time for the reduction with TCEP was increased to 4 h and stirring time for the ADC coupling was increased to 20 h. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted. Some of the ADC may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

Protein concentration: 1.33 mg/ml

30

Drug/mAb ratio: 0.8

**Example 218I**

35 Here, 5.0 mg of nimotuzumab antibody in PBS (c=13.8 mg/ml) were used for coupling with Intermediate F218, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted. Some of the ADC may also be

present in the form of the hydrolysed open-chain succinamides attached to the antibody.

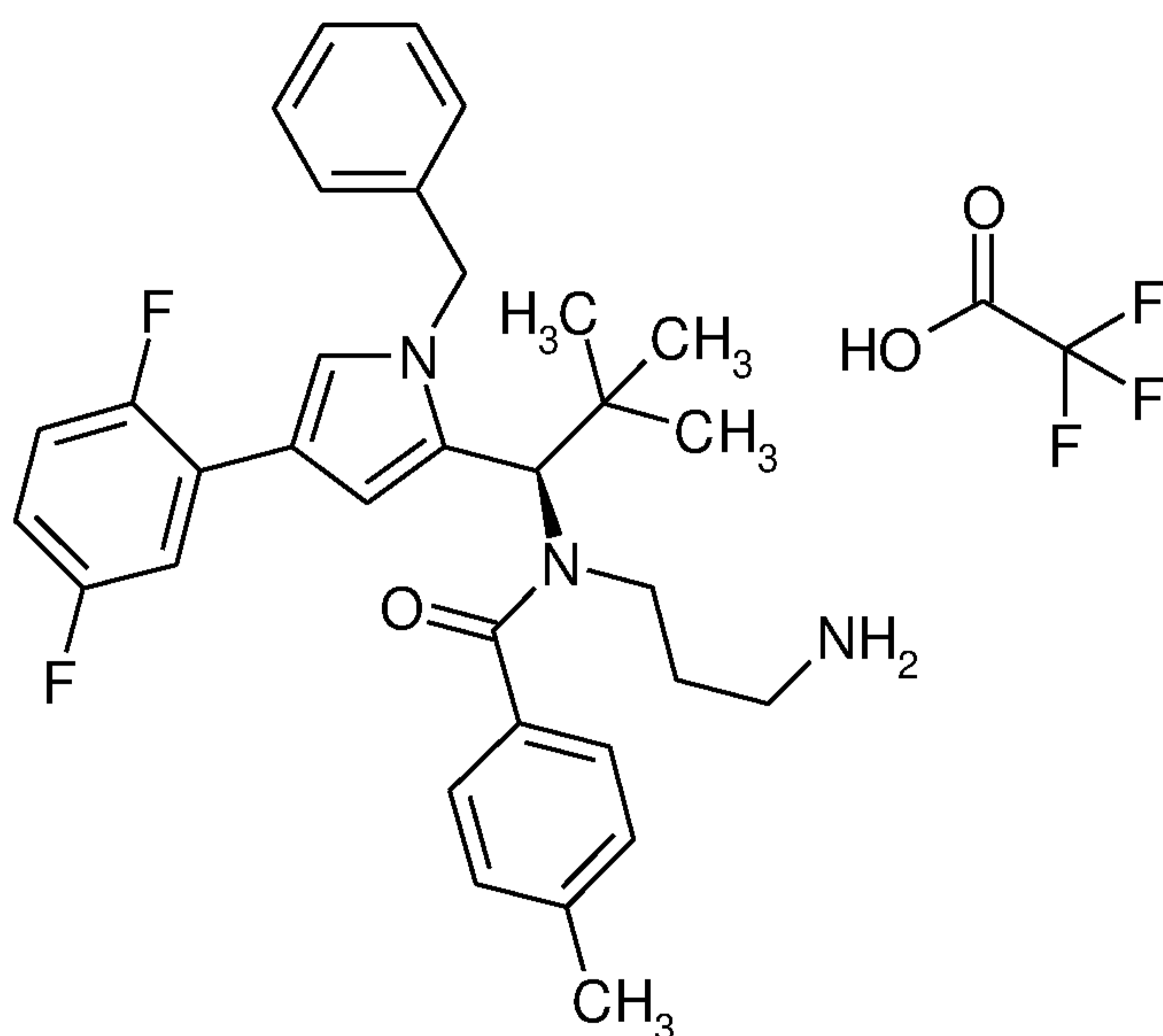
Protein concentration: 1.48 mg/ml

5

Drug/mAb ratio: 3.0

### Example 219

10 Trifluoroacetic acid / N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-methylbenzamide (1:1)



15

70.0 mg (0.09 mmol) of 9H-fluoren-9-ylmethyl-[3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino)propyl]carbamate (Intermediate C67) were initially charged in 3.0 ml of dichloromethane, and 31.3 mg (0.31 mmol) of triethylamine and 31.8 mg (0.21 mmol) of 4-methylbenzoyl chloride were added. The reaction mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and the residue was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 33.4 mg (48% of theory) of the compound 9H-fluoren-9-

20

25

ylmethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](4-methylbenzoyl)amino]propyl}carbamate.

5 LC-MS (Method 2):  $R_t = 11.91$  min; MS (ESIpos):  $m/z = 774$  (M+Na)<sup>+</sup>.

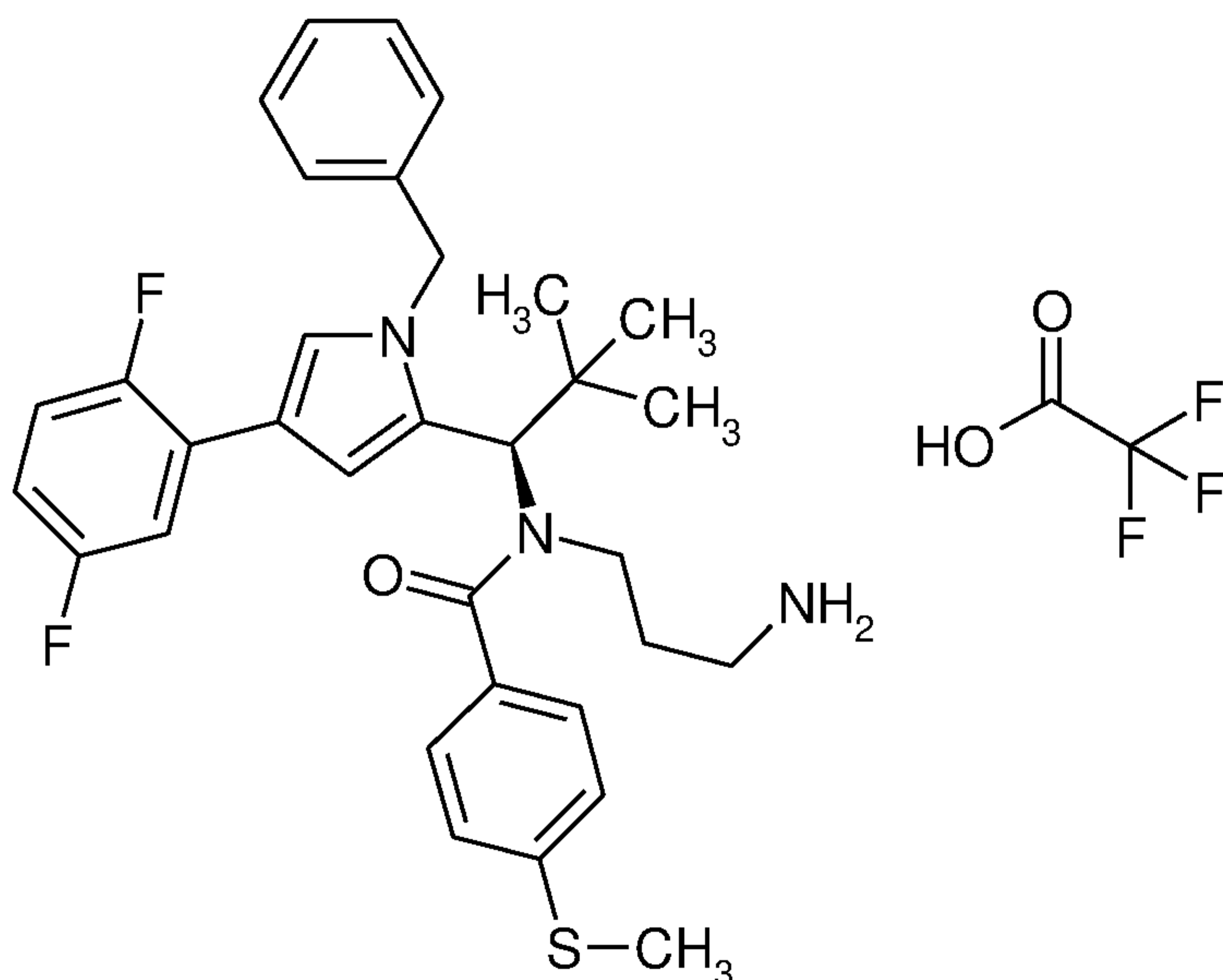
33.0 mg (0.04 mmol) of 9H-fluoren-9-ylmethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](4-methylbenzoyl)amino]propyl}carbamate in 1.0 ml of DMF were stirred with 20.0 mg (0.23 mmol) of morpholine overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 9.6 mg (34% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 1.01$  min; MS (ESIpos):  $m/z = 530$  (M+H)<sup>+</sup>.

### Example 220

20

Trifluoroacetic acid / N-(3-aminopropyl)-N-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]-4-(methylsulphanyl)benzamide (1:1)



25

50.0 mg (0.07 mmol) of 9H-fluoren-9-ylmethyl-[3-((1R)-1-[1-

benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino)propyl]carbamate (Intermediate C67) were initially charged in 2.0 ml of dichloromethane, and 22.3 mg (0.22 mmol) of triethylamine and 27.5 mg (0.15 mmol) of 4-(methylsulphanyl)benzoyl chloride were added. The reaction mixture was stirred at 40°C for 4 h, another 10.2 mg (0.10 mmol) of triethylamine and 27.5 mg (0.15 mmol) of 4-(methylsulphanyl)benzoyl chloride were added and the mixture was stirred at RT overnight. Another 14.9 mg (0.15 mmol) of triethylamine and 27.5 mg (0.15 mmol) of 4-(methylsulphanyl)benzoyl chloride were then added, and the mixture was stirred at 40°C for 2 h. The mixture was diluted with ethyl acetate and the organic phase was washed three times with water and once with saturated NaCl solution. The organic phase was dried over magnesium sulphate and concentrated under reduced pressure. The residue was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 39.8 mg (76% of theory) of the compound 9H-fluoren-9-ylmethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[4-(methylsulphanyl)benzoyl]amino)propyl]carbamate.

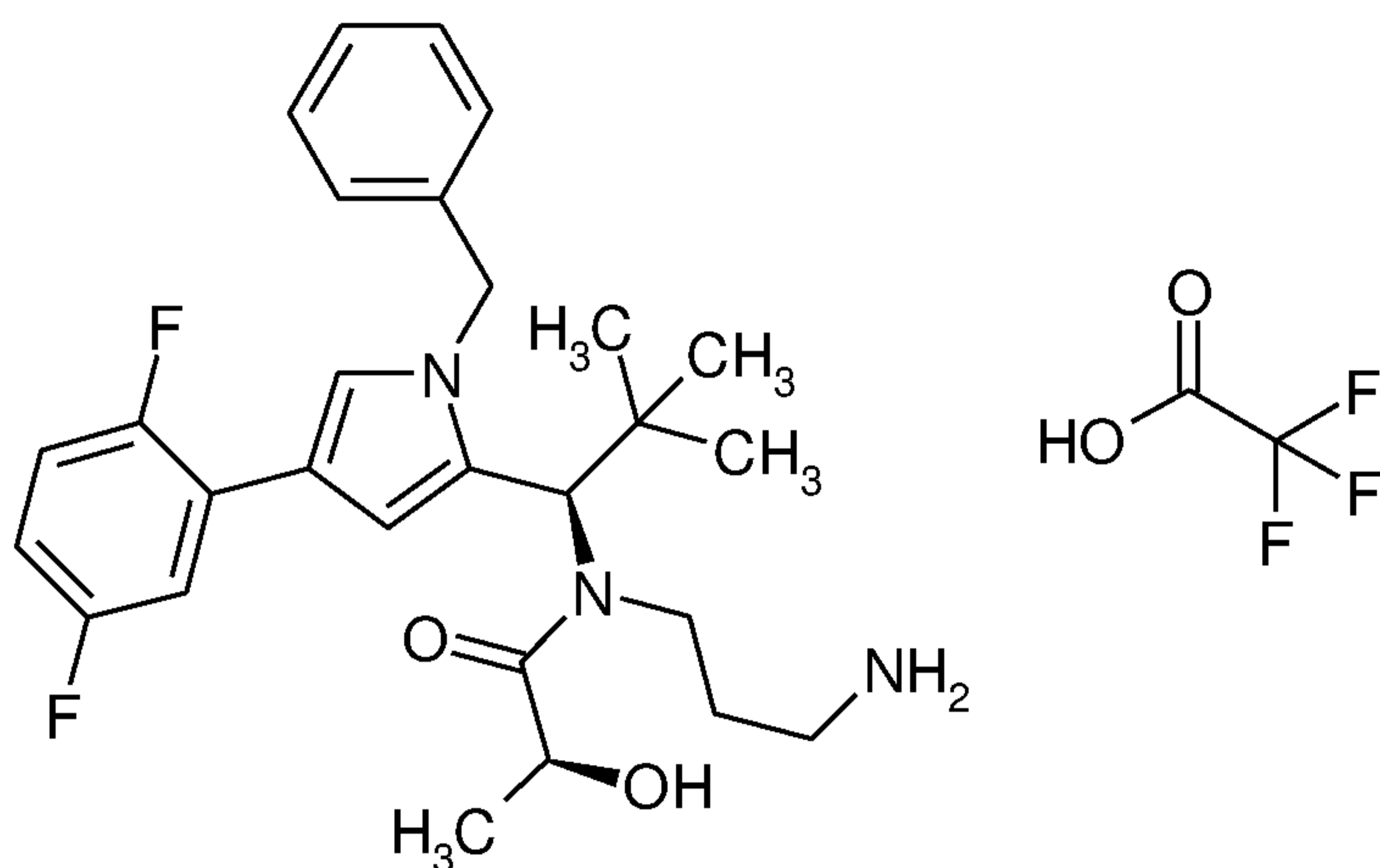
LC-MS (Method 1):  $R_t$  = 1.59 min; MS (ESIpos):  $m/z$  = 785 (M+H)<sup>+</sup>.

18.0 mg (0.02 mmol) of 9H-fluoren-9-ylmethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[4-(methylsulphanyl)benzoyl]amino)propyl]carbamate in 1.0 ml of DMF were stirred with 10.0 mg (0.12 mmol) of morpholine overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 6.2 mg (40% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.99 min; MS (ESIpos):  $m/z$  = 562 (M+H)<sup>+</sup>.

Example 221

Trifluoroacetic acid / (2S)-N-(3-aminopropyl)-N-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2-hydroxypropanamide (1:1)



40.0 mg (0.06 mmol) of 9H-fluoren-9-ylmethyl-[3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino)propyl]carbamate (Intermediate C67) were initially charged in 2.0 ml of dichloromethane, and 9.6 mg (0.10 mmol) of triethylamine and 14.3 mg (0.10 mmol) of (2S)-1-chloro-1-oxopropan-2-yl acetate were added. The reaction mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and the residue was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 39.7 mg (84% of theory) of the compound (2S)-1-[[1-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)](3-[[9H-fluoren-9-ylmethoxy]carbonyl]amino)propyl]amino]-1-oxopropan-2-yl acetate.

25

LC-MS (Method 1):  $R_t$  = 1.51 min; MS (ESIpos):  $m/z$  = 748 (M+H)<sup>+</sup>.

37.0 mg (0.05 mmol) of (2S)-1-[[1-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)](3-[[9H-



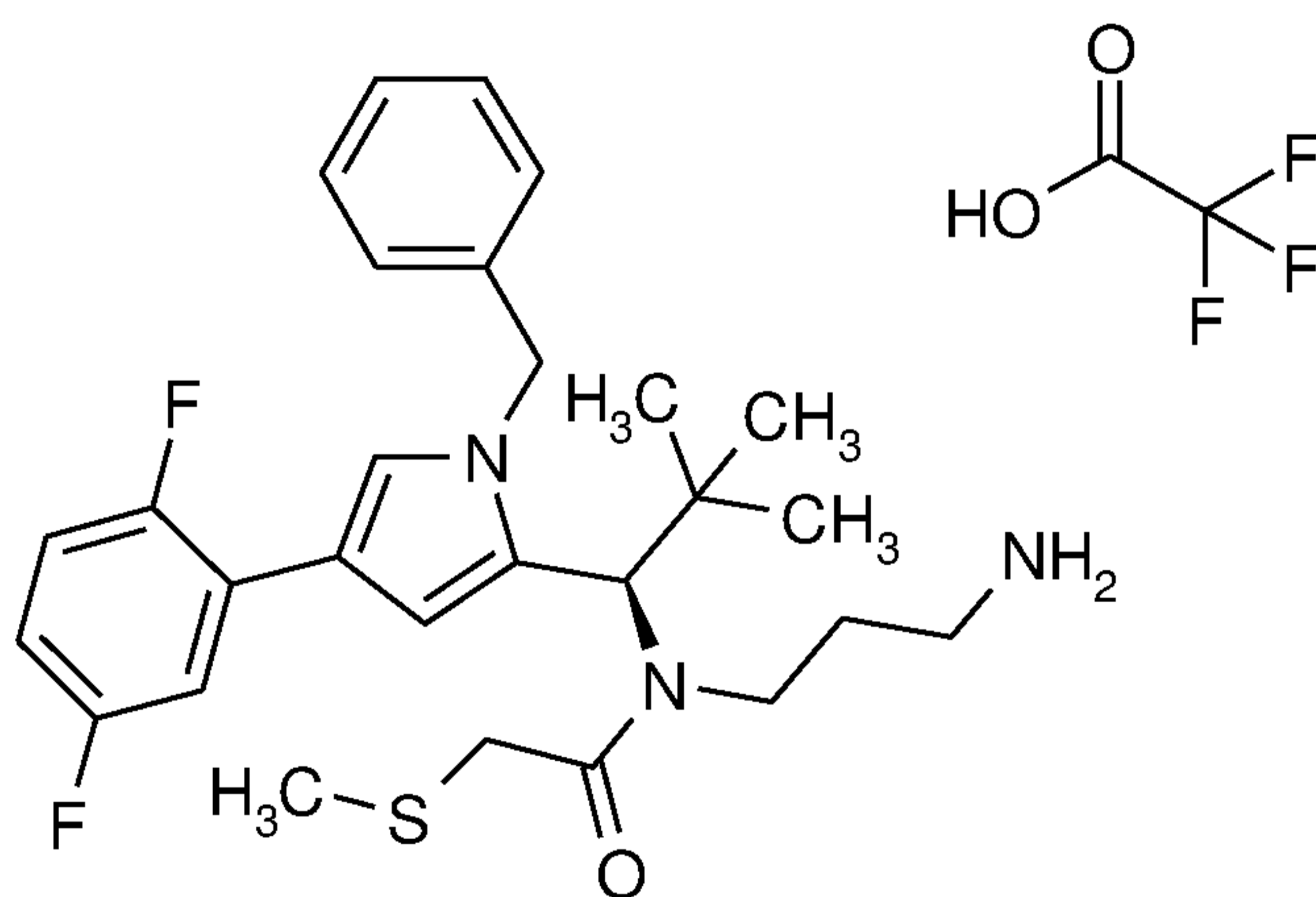
fluoren-9-ylmethoxy) carbonyl] amino] propyl) amino]-1-oxopropan-  
 2-yl acetate in 1.0 ml of DMF were stirred with 0.1 ml of  
 morpholine and 3 drops of water at 50°C for 10 h. Another 0.1  
 ml of morpholine and 0.1 ml of water were added, and the mixture  
 5 was stirred at 50°C for 10 h. After addition of 20.5 mg (0.15  
 mmol) of potassium carbonate and 72 h of stirring at RT, 0.1 ml  
 of 1N NaOH solution was added and the mixture was stirred at RT  
 overnight. The reaction mixture was purified directly by  
 preparative RP-HPLC (column: Reprosil 125x30; 10µ, flow rate:  
 10 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated  
 under reduced pressure and the residue was dried under high  
 vacuum. This gave 20.5 mg (69% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.89$  min; MS (ESIpos):  $m/z = 484$  (M+H)<sup>+</sup>.

15

### Example 222

Trifluoroacetic acid / N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-  
 (2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-  
 20 (methylsulphonyl)acetamide (1:1)



70.0 mg (0.11 mmol) of 2-(trimethylsilyl)ethyl-3-[(1R)-1-[1-  
 25 benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl](chloroacetyl)amino]propyl} carbamate  
 (Intermediate C 70) were initially charged in 3.0 ml of DMF.  
 15.5 mg (0.22 mmol) of sodium methanethiolate

30 were added, and the reaction mixture was stirred at 50°C for 2

h. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 60.0 mg  
5 (84% of theory) of the compound 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)}[(methylsulphonyl)acetyl]amino)propyl]carbamate .

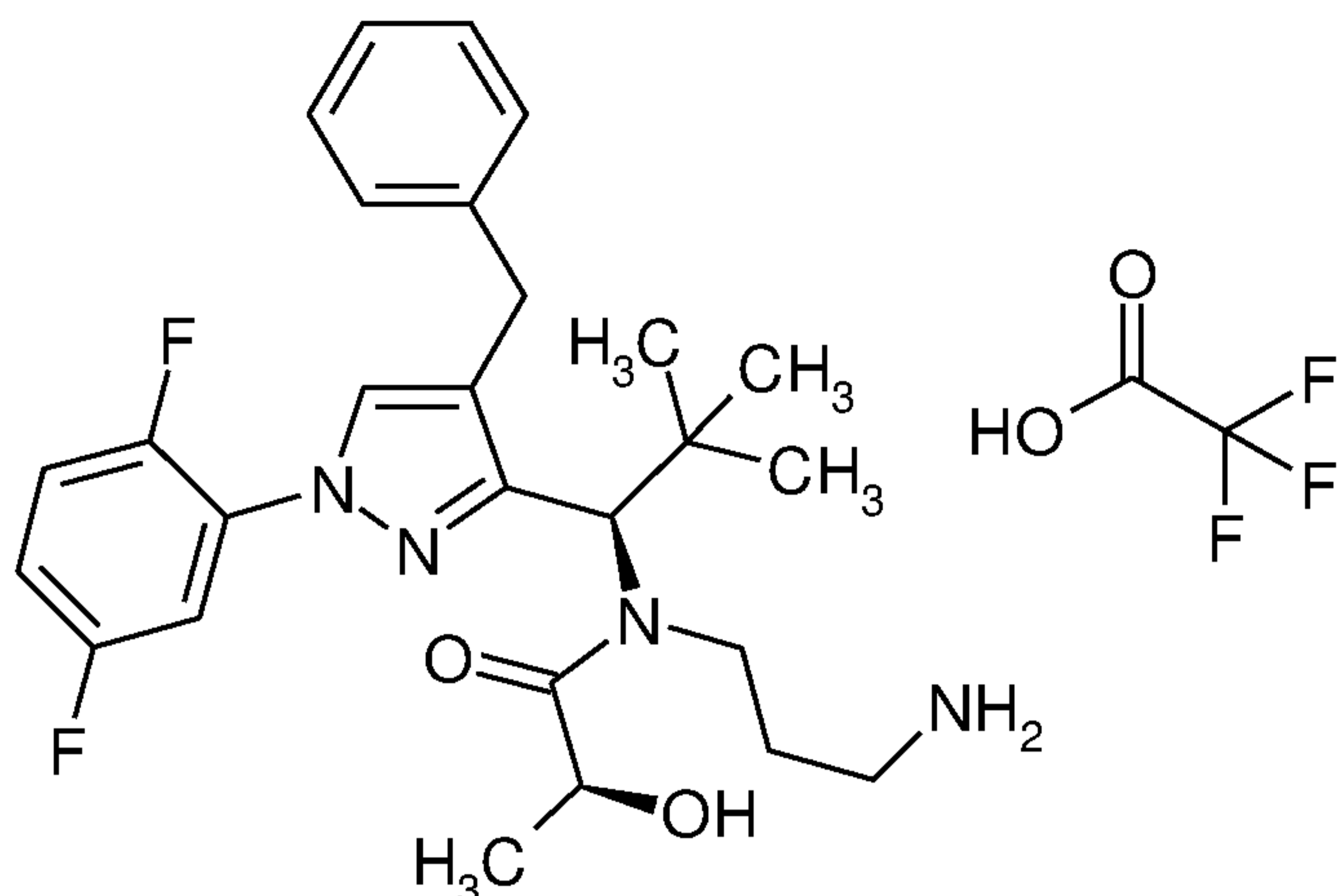
10 LC-MS (Method 1):  $R_t = 1.50$  min; MS (ESIpos):  $m/z = 644$  (M+H)<sup>+</sup>.

40.0 mg (0.06 mol) of 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)}[(methylsulphonyl)acetyl]amino)propyl]carbamate  
15 were dissolved in 2.0 ml of trifluoroethanol, and 21.2 mg (0.16 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C overnight. 45.4 mg (0.01 mmol) of ethylenediamin-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was  
20 then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 34.6 mg (91% of theory) of the title compound.

25 LC-MS (Method 1):  $R_t = 0.92$  min; MS (ESIpos):  $m/z = 500$  (M+H)<sup>+</sup>.

### Example 223

30 Trifluoroacetic acid / (2S)-N-(3-aminopropyl)-N-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)-2-hydroxypropanamide (1:1)



40.0 mg (0.08 mmol) of tert-butyl [3-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)amino]propyl]carbamate were dissolved in 2.0 ml of dichloromethane, and 19.7 mg (0.20 mmol) of triethylamine and 29.4 mg (0.20 mmol) of (2S)-1-chloro-1-oxopropan-2-yl acetate were added. The reaction mixture was stirred overnight. The solvent was evaporated under reduced pressure and the residue was purified by prep. RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 21.2 mg (43% of theory) of the compound (2S)-1-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl){3-[(tert-butoxycarbonyl)amino]propyl}amino)-1-oxopropan-2-yl acetate.

LC-MS (Method 1):  $R_t$  = 1.46 min; MS (ESIpos):  $m/z$  = 627 (M+H)<sup>+</sup>.

21.2 mg (0.03 mmol) of (2S)-1-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl){3-[(tert-butoxycarbonyl)amino]propyl}amino)-1-oxopropan-2-yl acetate were dissolved in 1.0 ml of dichloromethane, 77.1 mg (0.68 mmol) of trifluoroacetic acid were added and the mixture was stirred at RT for 2 h. Two more times, a further 77.1 mg (0.68 mmol) of trifluoroacetic acid were added, and in each case the mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and the residue was repeatedly co-distilled with dichloromethane and then dried under high vacuum. The residue, comprising the substance trifluoroacetic acid / (2S)-

1-[(3-aminopropyl){(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}amino]-1-oxopropan-2-yl acetate (1:1), was reacted further without further purification.

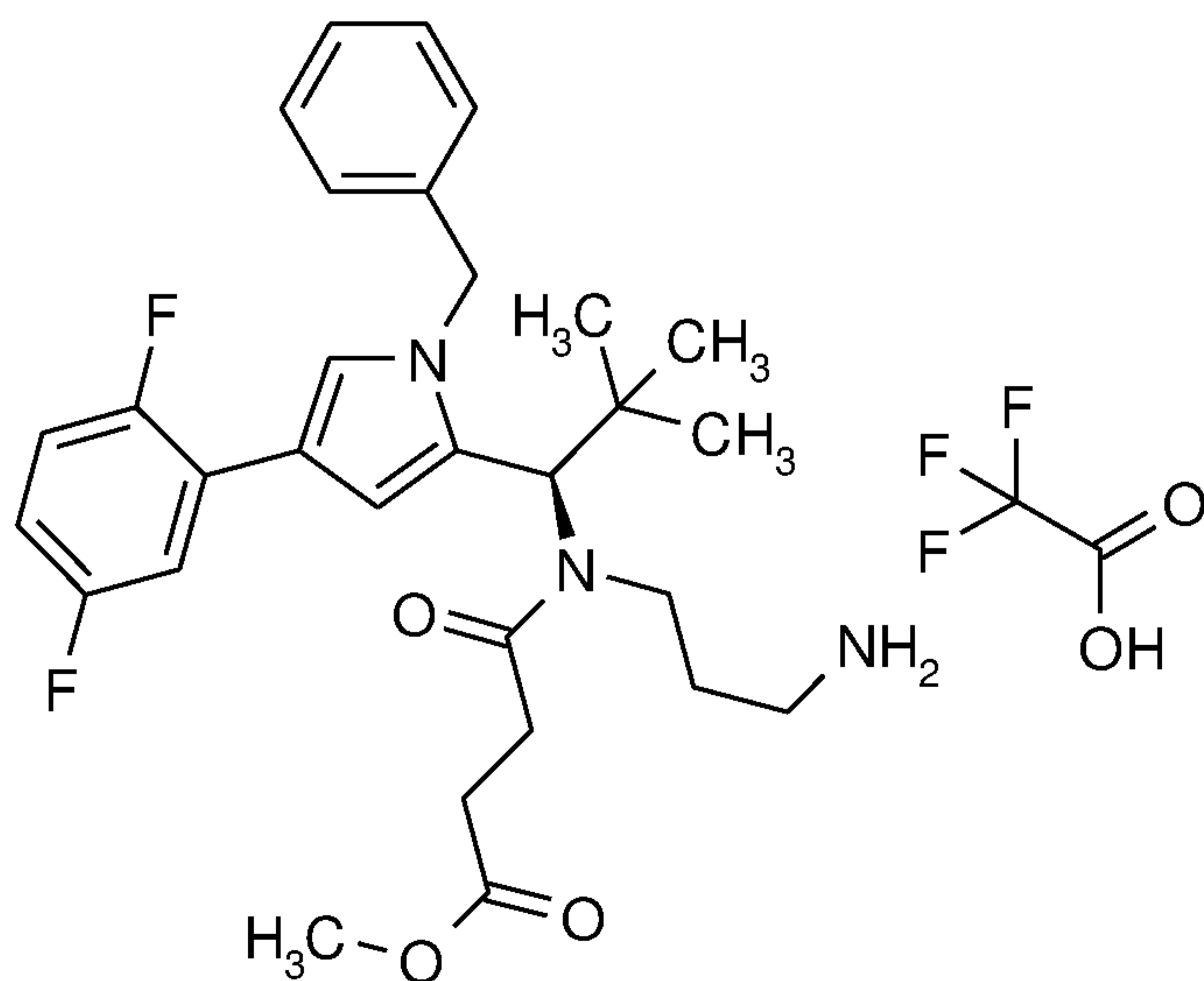
5 LC-MS (Method 1):  $R_t = 0.92$  min; MS (ESIpos):  $m/z = 527$  (M+H)<sup>+</sup>.

26.5 mg (0.04 mmol) of trifluoroacetic acid / (2S)-1-[(3-aminopropyl){(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}amino]-1-oxopropan-2-yl acetat (1:1) were dissolved in THF/methanol/water (1.0 ml/1.0 ml/0.05 ml), and 17.2 mg of potassium carbonate were added. The reaction mixture was stirred at RT overnight. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprisil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water; 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 17.3 mg (70% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.91$  min; MS (ESIpos):  $m/z = 485$  (M+H)<sup>+</sup>.  
20

#### Example 224

Trifluoroacetic acid / methyl 4-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-4-oxobutanoate (1:1)  
25



60.0 mg (0.11 mmol) of 2-(trimethylsilyl)ethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino]propyl]carbamate (see synthesis of Intermediate C11) were dissolved in 1.0 ml of dichloromethane, and 19.6 mg (0.25 mmol) of pyridine and 35.8 mg (0.24 mmol) of methyl 4-chloro-4-oxobutanoate were added. The reaction mixture was stirred at 40°C overnight. Another 19.6 mg (0.25 mmol) of pyridine and 35.8 mg (0.24 mmol) of methyl 4-chloro-4-oxobutanoate were added and the mixture was stirred at 40°C overnight. The solvent was evaporated under reduced pressure and the residue was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 16.1 mg (22% of theory) of the compound methyl 11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silapentadecan-15-oate.

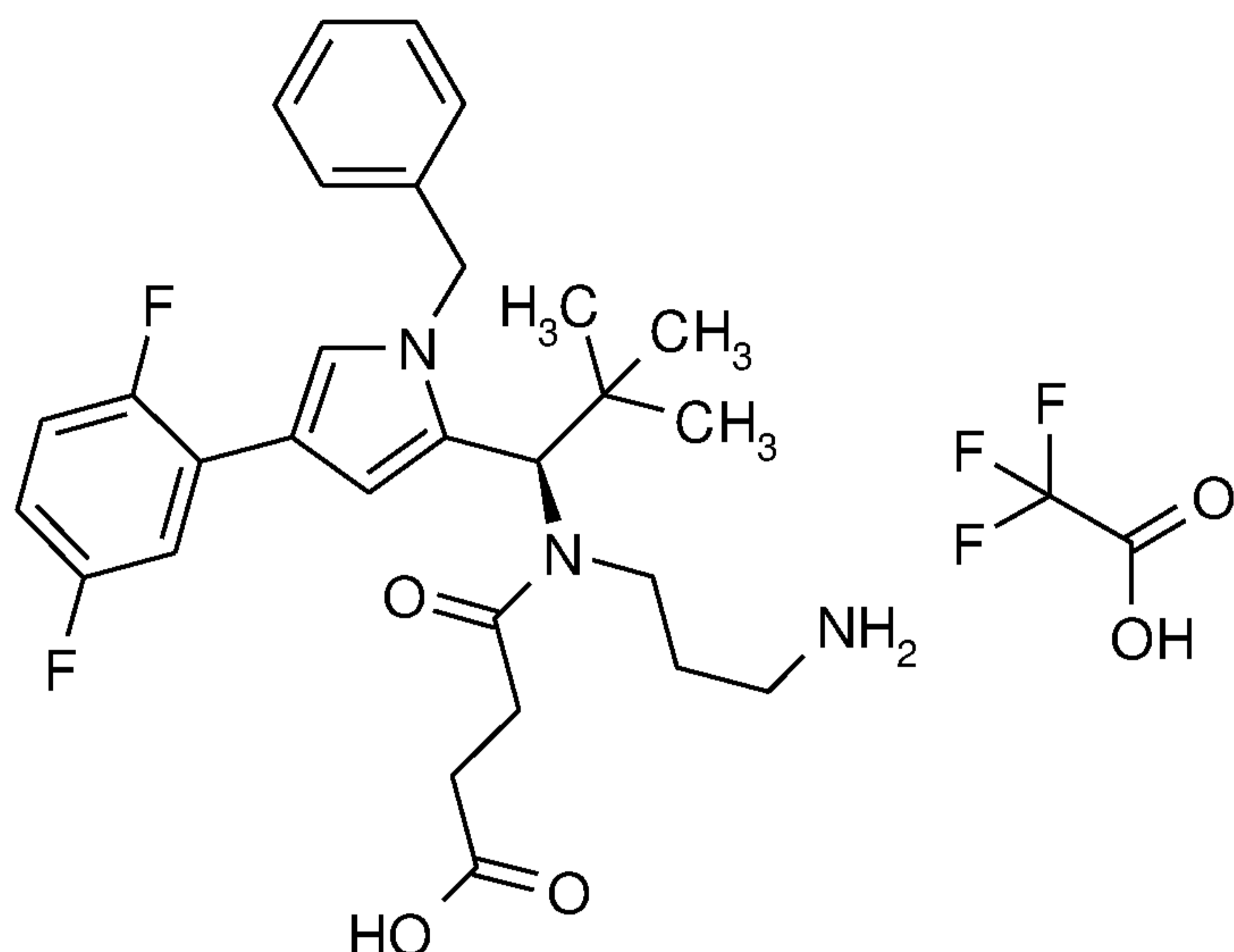
LC-MS (Method 1):  $R_t$  = 1.52 min; MS (ESIpos):  $m/z$  = 670 (M+H)<sup>+</sup>.

16.1 mg (0.02 mmol) of methyl 11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silapentadecan-15-oate were dissolved in 1.0 ml of trifluoroethanol, and 16.4 mg (0.12 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C for 5 h. 35.1 mg (0.12 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 11.1 mg (72% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.93 min; MS (ESIpos):  $m/z$  = 526 (M+H)<sup>+</sup>.

### Example 225

4-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-4-oxobutanoic acid / trifluoroacetic acid (1:1)



5

9.7 mg (0.02 mmol) of trifluoroacetic acid / methyl 4-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-4-oxobutanoate (1:1) (Example 224) were initially charged in THF/methanol/water (1.0 ml/0.2 ml/0.04 ml), and 1.3 mg (0.03 mmol) of lithium hydroxide monohydrate were added. The reaction mixture was stirred at RT overnight. A further 1.3 mg (0.03 mmol) of lithium hydroxide monohydrate were added, and the mixture was stirred at RT overnight. 3.6 mg (0.06 mmol) of HOAc were added and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water; 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 5.4 mg (57% of theory) of the title compound.

20

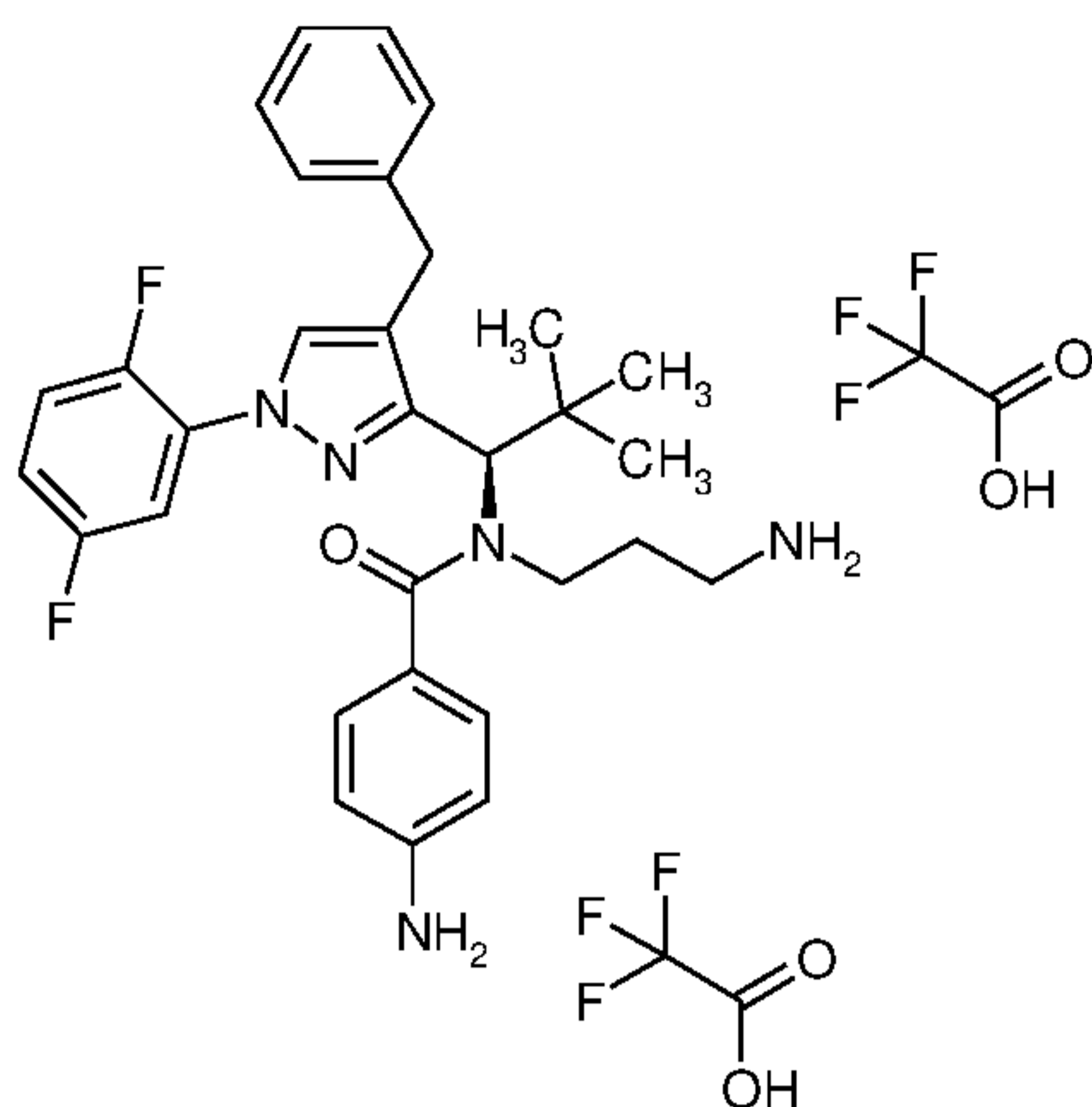
LC-MS (Method 1):  $R_t$  = 0.90 min; MS (ESIpos):  $m/z$  = 512 (M+H)<sup>+</sup>.

### Example 226

25

(2R)-22-[(3R/S)-3-[[2R)-2-Amino-2-carboxyethyl]sulphonyl]-2,5-dioxopyrrolidin-1-yl]-2-[[2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-





50.0 mg (0.10 mol) of tert-butyl [3-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl)amino]propyl carbamate (Intermediate C68) were initially charged in dichloromethane, and 54.9 mg (0.22 mmol) of tert-butyl [4-(chlorocarbonyl)phenyl]carbamate (Intermediate L71) and 22.7 mg (0.22 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight, and another 54.9 mg (0.22 mmol) of tert-butyl [4-(chlorocarbonyl)phenyl]carbamate (Intermediate L71) and 22.7 mg (0.22 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and the residue was purified by prep. RP-HPLC (column: Reprosil 250x40; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 26.2 mg (37% of theory) of the compound.

tert-Butyl [3-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl){4-[(tert-butoxycarbonyl)amino]benzoyl}amino]propyl carbamate.

LC-MS (Method 1):  $R_t$  = 5.34 min; MS (ESIpos):  $m/z$  = 732 (M+H)<sup>+</sup>.

26.2 mg (0.04 mmol) of tert-butyl [3-((1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl){4-[(tert-butoxycarbonyl)amino]benzoyl}amino]propyl carbamate were dissolved in 2.0 ml of dichloromethane, and 204.1 mg (1.79 mmol) of TFA were added. The reaction mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and



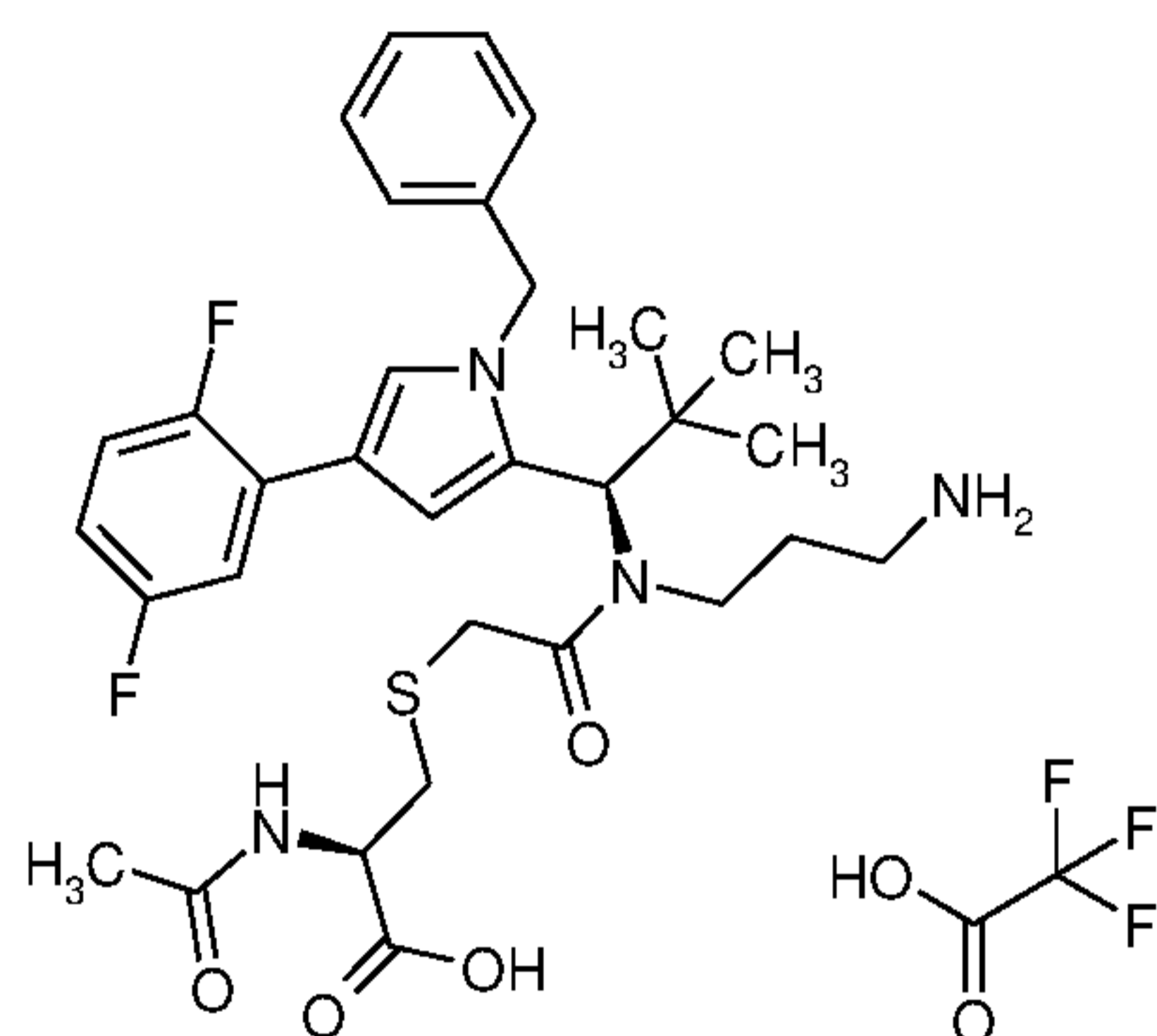
the residue was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 3.4 mg (13% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.92 min; MS (ESIpos):  $m/z$  = 532 (M+H)<sup>+</sup>.

### Example 228

10

N-Acetyl-S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-L-cysteine / trifluoroacetic acid (1:1)



15

50.0 mg (0.08 mol) of 2-(trimethylsilyl)ethyl-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(chloroacetyl)amino]propyl}carbamate (Intermediate C70) were suspended in 0.30 ml of water with 66.43 mg of sodium bicarbonate. A solution of 144.47 (0.95 mmol) of 1,8-diazabicyclo(5.4.0)undec-7-ene and 51.62 mg (0.32 mmol) of N-acetyl-L-cysteine in 3.0 ml of isopropanol was added to the suspension. The reaction mixture was stirred at 50°C for 2.5 h. Water (0.1% TFA) was added and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 55.2 mg (92% of theory) of the compound

N-acetyl-S-(11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-

30

cysteine.

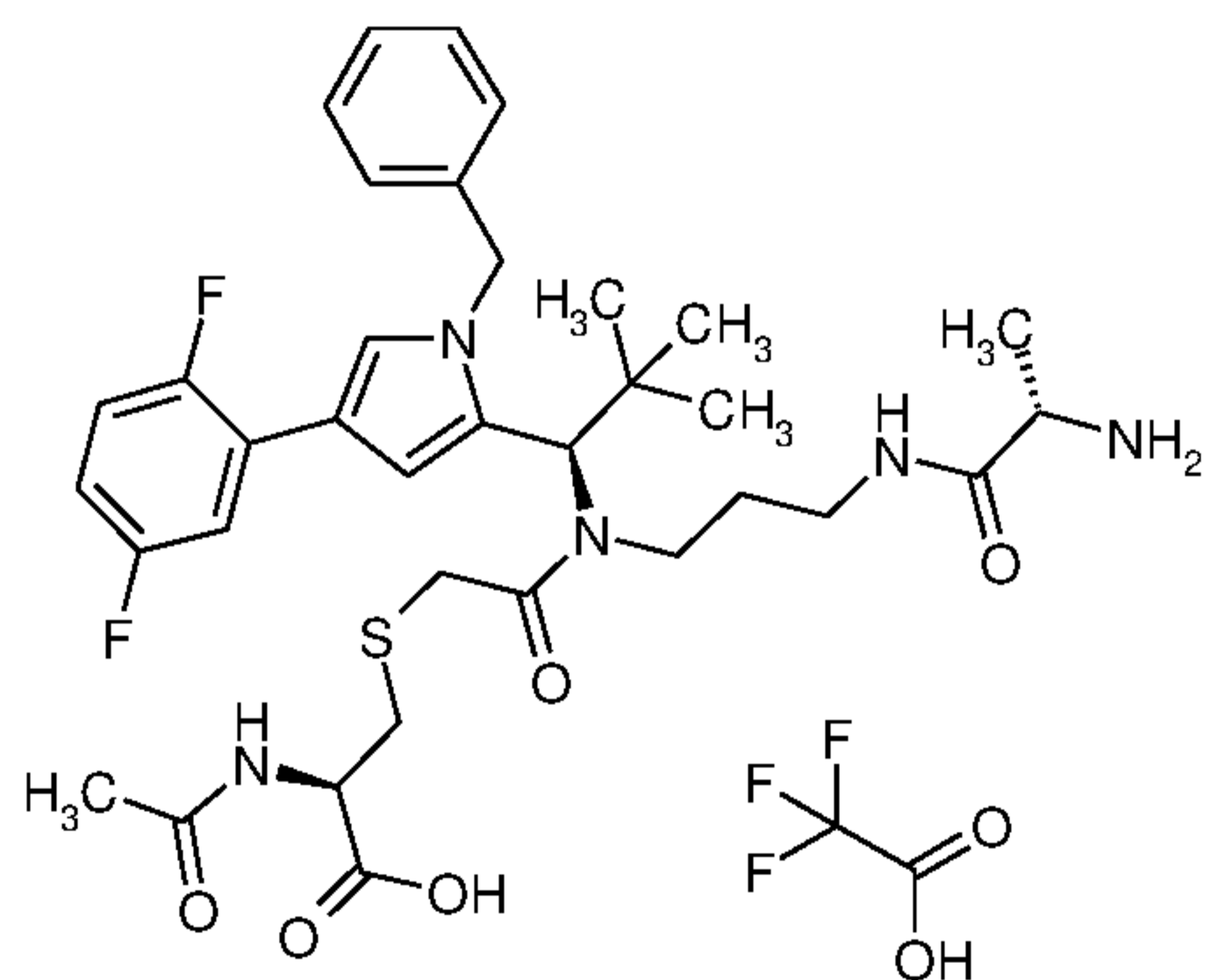
LC-MS (Method 1):  $R_t = 1.41$  min; MS (ESIpos):  $m/z = 759$  (M+H)<sup>+</sup>.

5 53.1 mg (69.96  $\mu$ mol) of N-acetyl-S-(11-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)-L-cysteine were dissolved in 5.0 ml of trifluoroethanol, and 57.2 mg (419.76  $\mu$ mol) of zinc dichloride were added. The reaction mixture was stirred at 50°C for 4 h. 122.67 mg (0.42 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). This gave 32.5 mg (64% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.90$  min; MS (ESIpos):  $m/z = 615$  (M+H)<sup>+</sup>.

## 20 Example 229

N-Acetyl-S-[2-((3-(L-alanyl-amino)propyl)((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino)-2-oxoethyl]-L-cysteine / trifluoroacetic acid (1:1)



30.3 mg (41.58  $\mu$ mol) of N-acetyl-S-{2-[(3-aminopropyl)((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino]-2-oxoethyl}-L-cysteine / trifluoroacetic acid (1:1) (Example 228) were dissolved in 1.5 ml of DMF, and

8.4 mg (83.15  $\mu$ mol) of 4-methylmorpholine and 14.65 mg (45.73  $\mu$ mol) of 2,5-dioxopyrrolidin-1-yl-N-[(benzyloxy)carbonyl]-L-alaninate were added. The reaction mixture was stirred at RT overnight, and another 8.4 mg (83.15  $\mu$ mol) of 4-methylmorpholine were then added. Once more, the reaction mixture was stirred at RT overnight. 10.0 mg (0.17 mmol) of acetic acid were added and the reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 31.2 mg (92% of theory) of the compound N-acetyl-S-[2-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)[3-((N-[(benzyloxy)carbonyl]-L-alanyl)amino)propyl]amino)-2-oxoethyl]-L-cysteine.

LC-MS (Method 1):  $R_t$  = 1.22 min; MS (ESIpos):  $m/z$  = 820 (M+H)<sup>+</sup>.

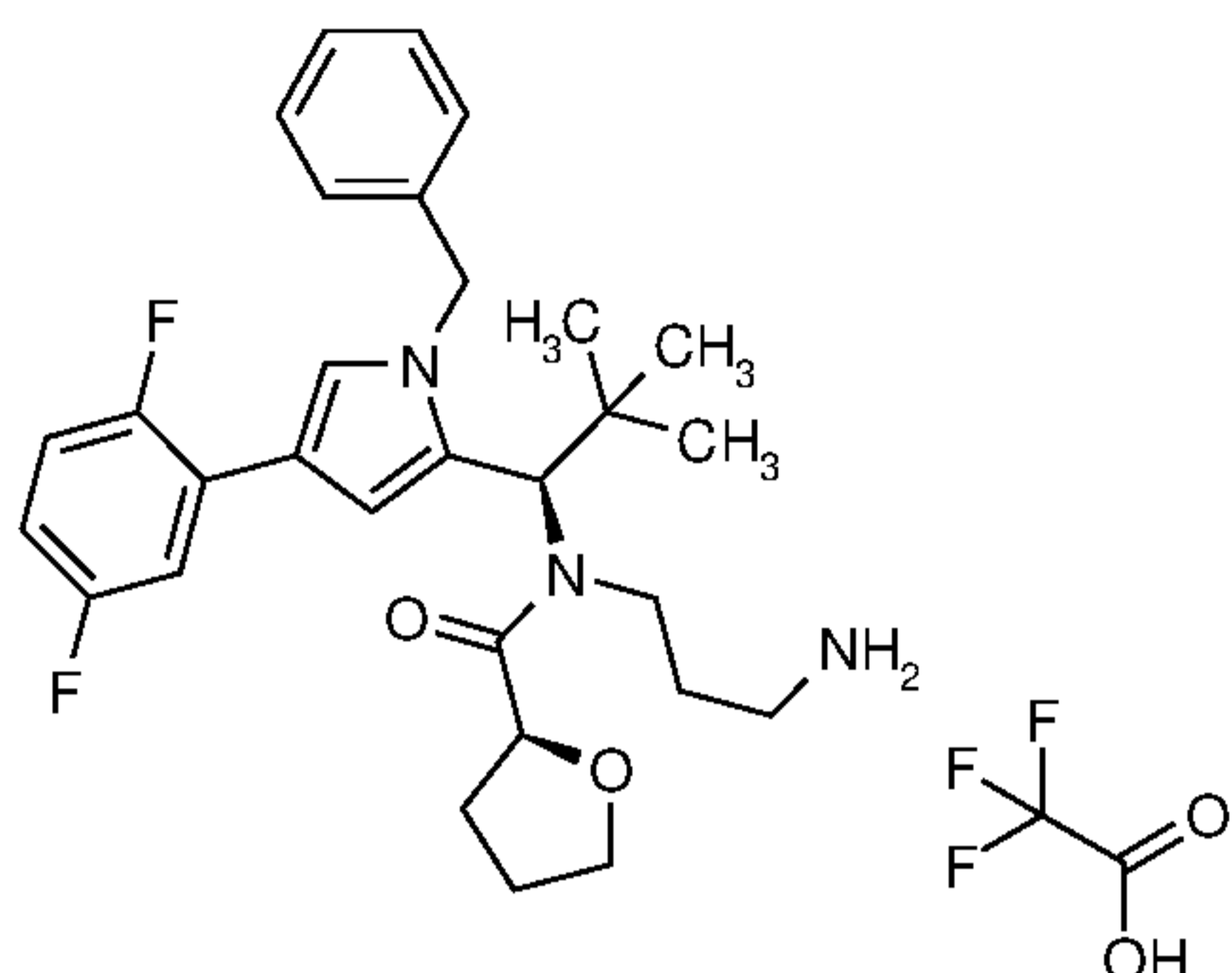
28.6 mg (0.04 mmol) of N-acetyl-S-[2-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)[3-((N-[(benzyloxy)carbonyl]-L-alanyl)amino)propyl]amino)-2-oxoethyl]-L-cysteine were dissolved in 5.0 ml of ethanol, and 2.9 mg of palladium on activated carbon (10%) were added. The reaction mixture was hydrogenated at standard pressure and RT overnight. The mixture was filtered through Celite and the filter cake was washed with a mixture of ethanol. The solvents were evaporated under reduced pressure and the residue was purified by preparative RP-HPLC (column: Reprosil 250x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was lyophilized. This gave 17.4 mg (62% of theory) of the title compound.

LC-MS (Method 1):  $R_t$  = 0.85 min; MS (ESIpos):  $m/z$  = 686 (M+H)<sup>+</sup>.

### 35 Example 230

Trifluoroacetic acid / (2S)-N-(3-aminopropyl)-N-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-

dimethylpropyl}tetrahydrofuran-2-carboxamide (1:1)



- 5 100.0 mg (0.16 mmol) of 9H-fluoren-9-ylmethyl-[3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino)propyl]carbamate (Intermediate C67) were dissolved in 5.0 ml of dichloromethane with 71.9 mg (0.71 mmol) of triethylamine and added dropwise to a solution of freshly prepared (2S)-tetrahydrofuran-2-carbonyl chloride (preparation: 10 54.7 mg (0.39 mmol) of (2S)-tetrahydrofuran-2-carboxylic acid were initially charged in 0.7 ml of toluene, 0.04 ml of thionyl chloride were added and the mixture was stirred at 90°C for 1 h. After cooling, the crude reaction solution was reacted 15 further). The reaction mixture was stirred at RT overnight. The solvents were evaporated under reduced pressure and the residue was purified by preparative RP-HPLC (column: Reprisil 125x30; 10µ, flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried 20 under high vacuum. This gave 44.3 mg (38% of theory) of the compound 9H-fluoren-9-ylmethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[(2S)-tetrahydrofuran-2-ylcarbonyl]amino)propyl]carbamate.
- 25 LC-MS (Method 1):  $R_t = 1.52$  min; MS (ESIpos):  $m/z = 776$  ( $M+HCOOH-H$ )<sup>-</sup>.

- 26.0 mg (0.04 mmol) of 9H-fluoren-9-ylmethyl [3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}[(2S)-tetrahydrofuran-2-ylcarbonyl]amino)propyl]carbamate were dissolved in 2.6 ml of DMF, and 0.26 ml of morpholine were added. The reaction mixture
- 30

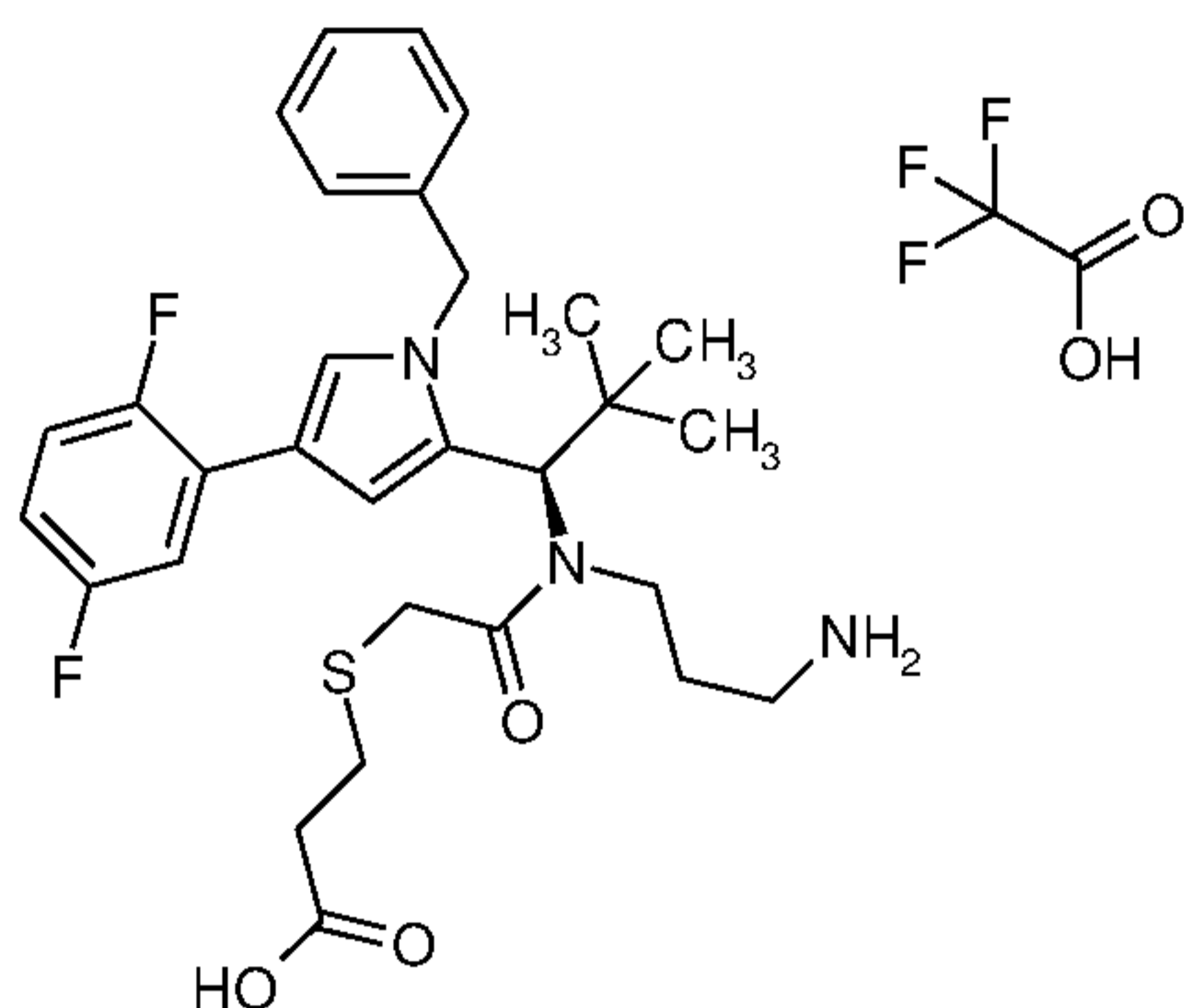
was stirred at RT for 3 h. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). This gave 11.7 mg (53% of theory) of the title compound.

5

LC-MS (Method 1):  $R_t$  = 0.93 min; MS (ESIpos):  $m/z$  = 510 (M+H)<sup>+</sup>.

### Example 231

10 3-({2-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulphonyl)propanoic acid / trifluoroacetic acid (1:1)



15

53.9 mg (0.08 mmol) of 11-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2,2-dimethyl-6,12-dioxo-5-oxa-14-thia-7,11-diaza-2-silaheptadecan-17-oic acid (Intermediate C69) were dissolved in 4.0 ml of trifluoroethanol, and 31.4 mg (0.23 mmol) of zinc dichloride were added. The reaction mixture was stirred at 50°C overnight. 67.3 mg (0.23 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). This gave 34.2 mg (66% of theory) of the title compound.

20

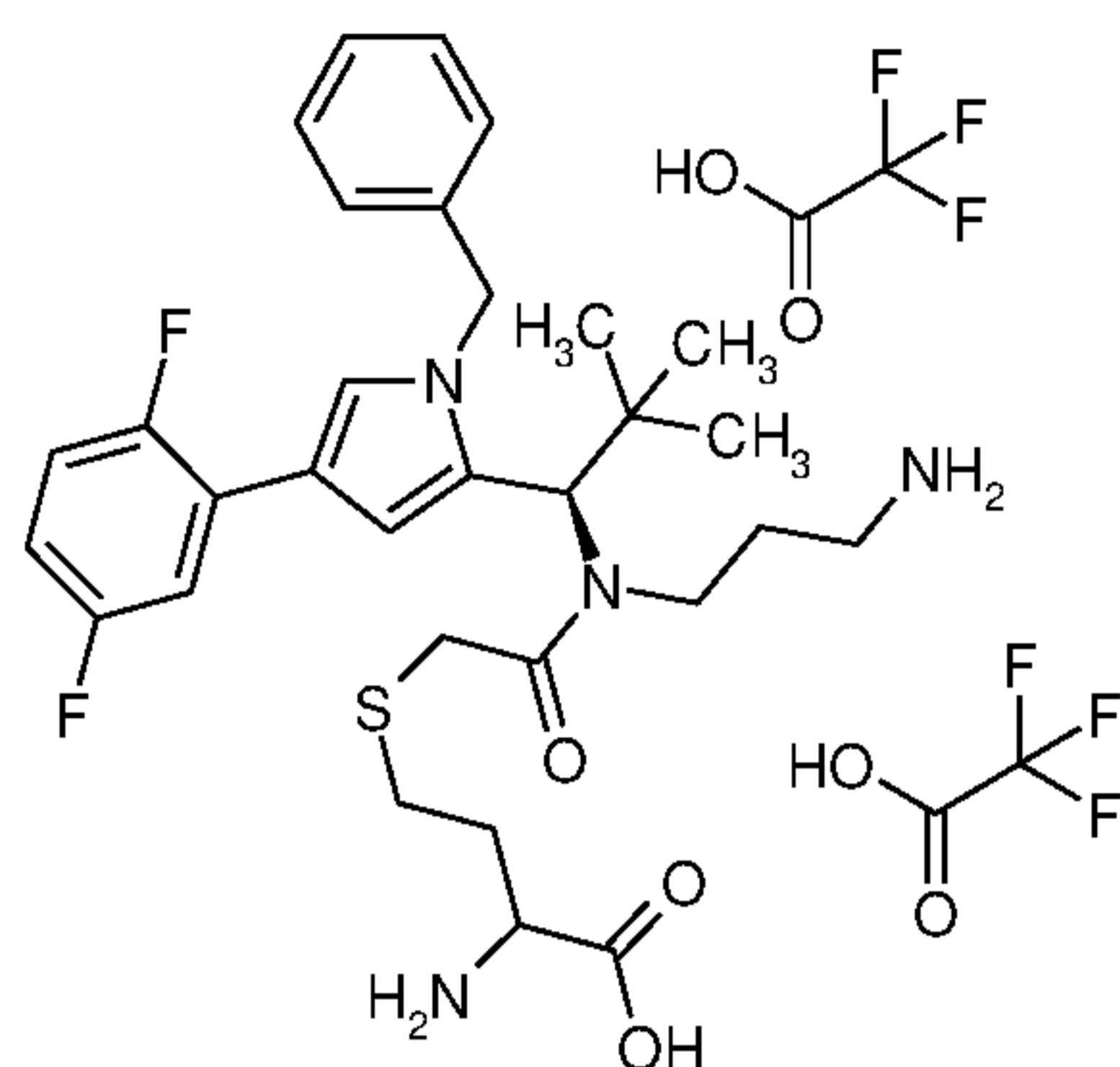
25

30

LC-MS (Method 1):  $R_t$  = 0.91 min; MS (ESIpos):  $m/z$  = 558 (M+H)<sup>+</sup>.

### Example 232

S-{2-[(3-Aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}homocysteine / trifluoroacetic acid (1:2)



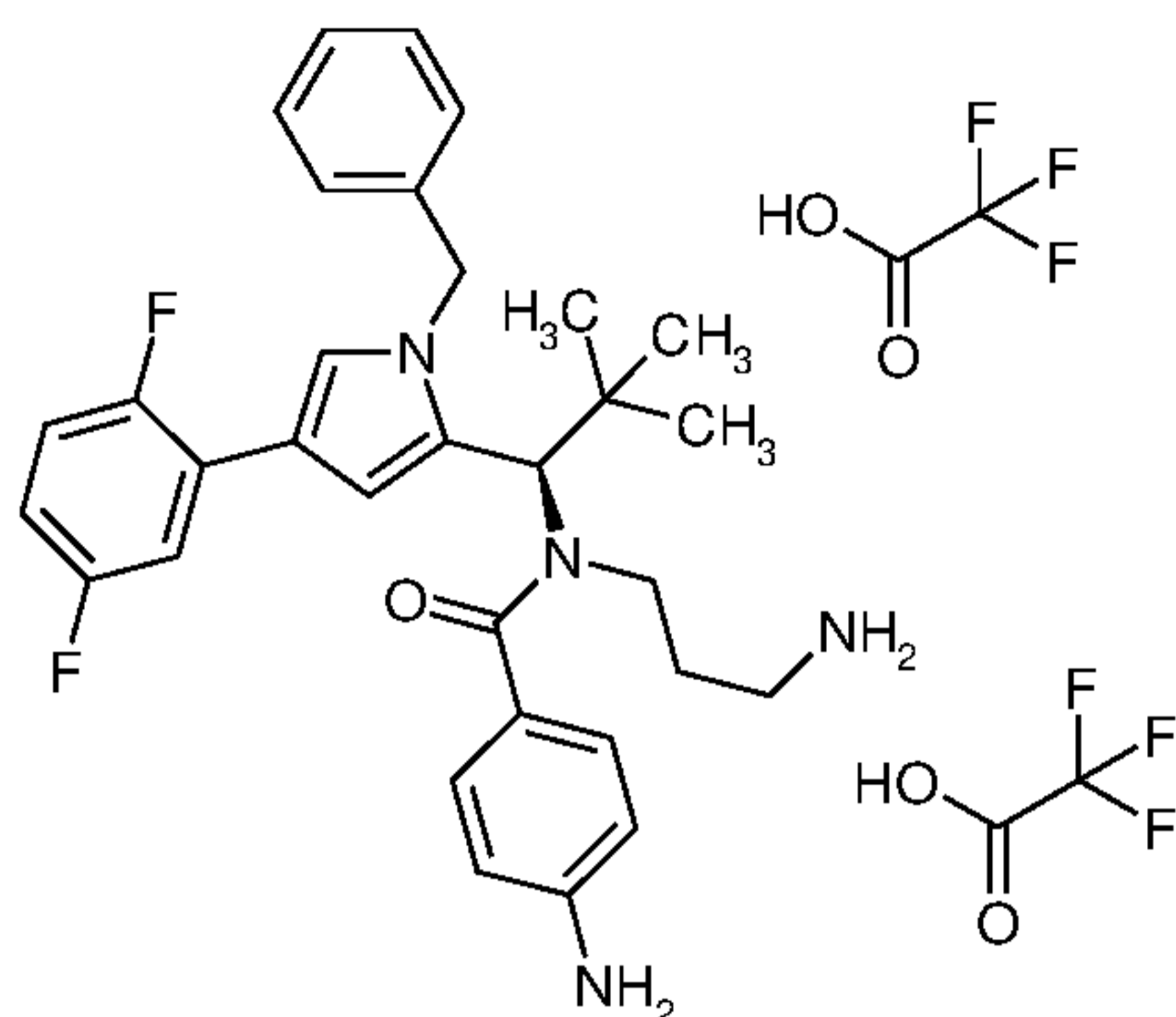
5

40.0 mg (47.3  $\mu\text{mol}$ ) of S-(11-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]-2,2-dimethyl-6,12-dioxo-5-oxa-7,11-diaza-2-silatridecan-13-yl)homocysteine (Intermediate C11) were dissolved in 3.0 ml of trifluoroethanol, and 38.7 mg (0.28  $\mu\text{mol}$ ) of zinc dichloride were added. The reaction mixture was stirred at 50°C for 6 h. 83 mg (0.28 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were added, the reaction mixture was stirred for 10 min and water (0.1% TFA) was then added. Purification was carried out directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). This gave 32.4 mg (84% of theory) of the title compound.

20 LC-MS (Method 1):  $R_t$  = 0.81 min; MS (ESIpos):  $m/z$  = 587 (M+H)<sup>+</sup>.

### Example 233

25 Trifluoroacetic acid / 4-amino-N-(3-aminopropyl)-N-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]benzamide (2:1)



73.0 mg (0.12 mmol) of 9H-fluoren-9-ylmethyl-[3-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl)amino]propyl}carbamate (Intermediate C67) and 27.8 mg (0.15 mmol) of 4-nitrobenzoyl chloride were dissolved in 2.0 ml of dichloromethane, and 15.2 mg (0.15 mmol) of triethylamine were added. The reaction mixture was stirred at RT overnight. The same amount of 4-nitrobenzoyl chloride and triethylamine was added again, and the reaction mixture was stirred at RT overnight. The solvent was evaporated under reduced pressure and the residue was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). This gave 39.3 mg (44% of theory) of the compound 9H-fluoren-9-ylmethyl {3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]}(4-nitrobenzoyl)amino]propyl}carbamate.

LC-MS (Method 1):  $R_t$  = 1.54 min; MS (ESIpos):  $m/z$  = 783 (M+H)<sup>+</sup>.

39.3 mg (0.05 mmol) of fluoren-9-ylmethyl-{3-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]}(4-nitrobenzoyl)amino]propyl}carbamate were dissolved in 2.0 ml of ethanol, 3.9 mg of palladium hydroxide on activated carbon (20%) were added and the mixture was hydrogenated at standard pressure overnight. The reaction mixture was filtered through a paper filter and the filter cake was washed with ethanol. The solvent was evaporated under reduced pressure. The residue was purified by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min, MeCN/water, 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum.





**Example 234B**

Under argon, a solution of 0.029 mg of TCEP in 50 µl of PBS  
5 buffer was added to 5 mg of anti-TWEAKR AK-1 in 269 µl of PBS  
(c=18.6 mg/ml). The reaction was stirred at RT for 30 min, and  
0.2 mg (0.00027 mmol) of Intermediate F85 dissolved in 50 µl of  
DMSO were then added. After a further 90 min of stirring at RT,  
10 the reaction was diluted with 2130 µl of PBS buffer which had  
been adjusted to pH 8 beforehand.

This solution was then applied to PD 10 columns (Sephadex® G-  
25, GE Healthcare) which had been equilibrated with PBS buffer  
pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred  
15 under argon at RT overnight and then concentrated by  
ultracentrifugation and rediluted with PBS buffer (pH 7.2).  
Under these conditions, some of the ADSs may also be present in  
the ring-closed form. The ADC batch obtained was characterized  
as follows:

20 Protein concentration: 1.57 mg/ml

Drug/mAb ratio: 3.7

**Example 234I**

Under argon, a solution of 0.029 mg of TCEP in 50 µl of PBS  
buffer was added to 5 mg of nimotuzumab in 500 µl of PBS (c=10  
mg/ml). The reaction was stirred at RT for 30 min, and 0.2 mg  
30 (0.00027 mmol) of Intermediate F85 dissolved in 50 µl of DMSO  
were then added. After a further 90 min of stirring at RT, the  
reaction was diluted with 1900 µl of PBS buffer which had been  
adjusted to pH 8 beforehand.

35 This solution was then applied to PD 10 columns (Sephadex® G-  
25, GE Healthcare) which had been equilibrated with PBS buffer  
pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred  
under argon at RT overnight and then concentrated by

ultracentrifugation and rediluted with PBS buffer (pH 7.2). Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

5

Protein concentration: 1.93 mg/ml

Drug/mAb ratio: 3.6

#### 10 Example 234H

Under argon, a solution of 0.029 mg of TCEP in 50 µl of PBS buffer was added to 5 mg of panitumumab in 500 µl of PBS (c=10 mg/ml). The reaction was stirred at RT for 4 h, and 0.2 mg  
15 (0.00027 mmol) of Intermediate F85 dissolved in 50 µl of DMSO were then added. After a further 120 min of stirring at RT, the reaction was diluted with 1900 µl of PBS buffer which had been adjusted to pH 8 beforehand.

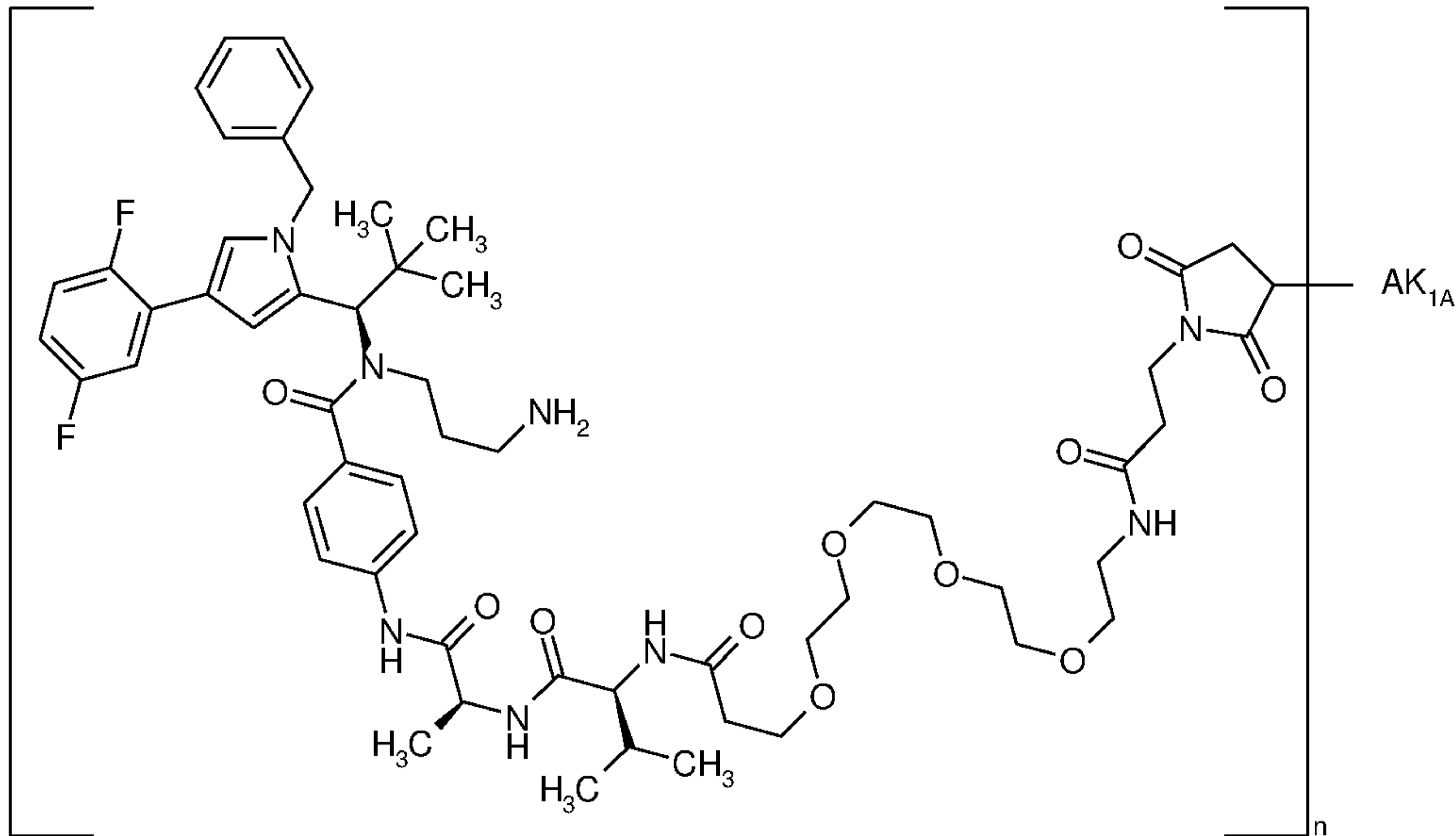
20 This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2).  
25 Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

30

Protein concentration: 2.12 mg/ml

Drug/mAb ratio: 1.4

#### Example 235A



Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS buffer was added to 5 mg of cetuximab in 329  $\mu$ l of PBS (c=15.2 mg/ml), and the mixture was then diluted with 2021  $\mu$ l of PBS buffer which had been adjusted to pH 8 beforehand. The reaction was stirred at RT for 1 h, and 0.28 mg of Intermediate F235 dissolved in 50  $\mu$ l of DMSO were then added. After a further 90 min of stirring at RT, the reaction was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

Protein concentration: 1.5 mg/ml

Drug/mAb ratio: 2.4

#### Example 235B

Analogously to Example 235A, 5 mg of anti-TWEAKR AK-1 in 269  $\mu$ l of PBS (c=18.6 mg/ml) were diluted with PBS buffer pH 8 to a concentration of 2 mg/ml and coupled with Intermediate F235.

Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

5

Protein concentration: 0.56 mg/ml

Drug/mAb ratio: 1.0

#### 10 **Example 235E**

Analogously to Example 235A, 5 mg of trastuzumab in 370 µl of PBS (c=14.5 mg/ml) were diluted with PBS buffer pH 8 to a concentration of 2 mg/ml and coupled with Intermediate F235.

15

Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

20

Protein concentration: 1.67 mg/ml

Drug/mAb ratio: 2.9

#### **Example 235I**

25

Analogously to Example 235A, 5 mg of nimotuzumab in 382 µl of PBS (c=13.08 mg/ml) were diluted with PBS buffer pH 8 to a concentration of 2 mg/ml and coupled with Intermediate F235.

30

Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

Protein concentration: 1.81 mg/ml

35

Drug/mAb ratio: 2.8

#### **Example 235H**

Analogously to Example 235A, 5 mg of panitumumab in 74  $\mu$ l of PBS (c=67.4 mg/ml) were diluted with PBS buffer pH 8 to a concentration of 2 mg/ml and coupled with Intermediate F235.

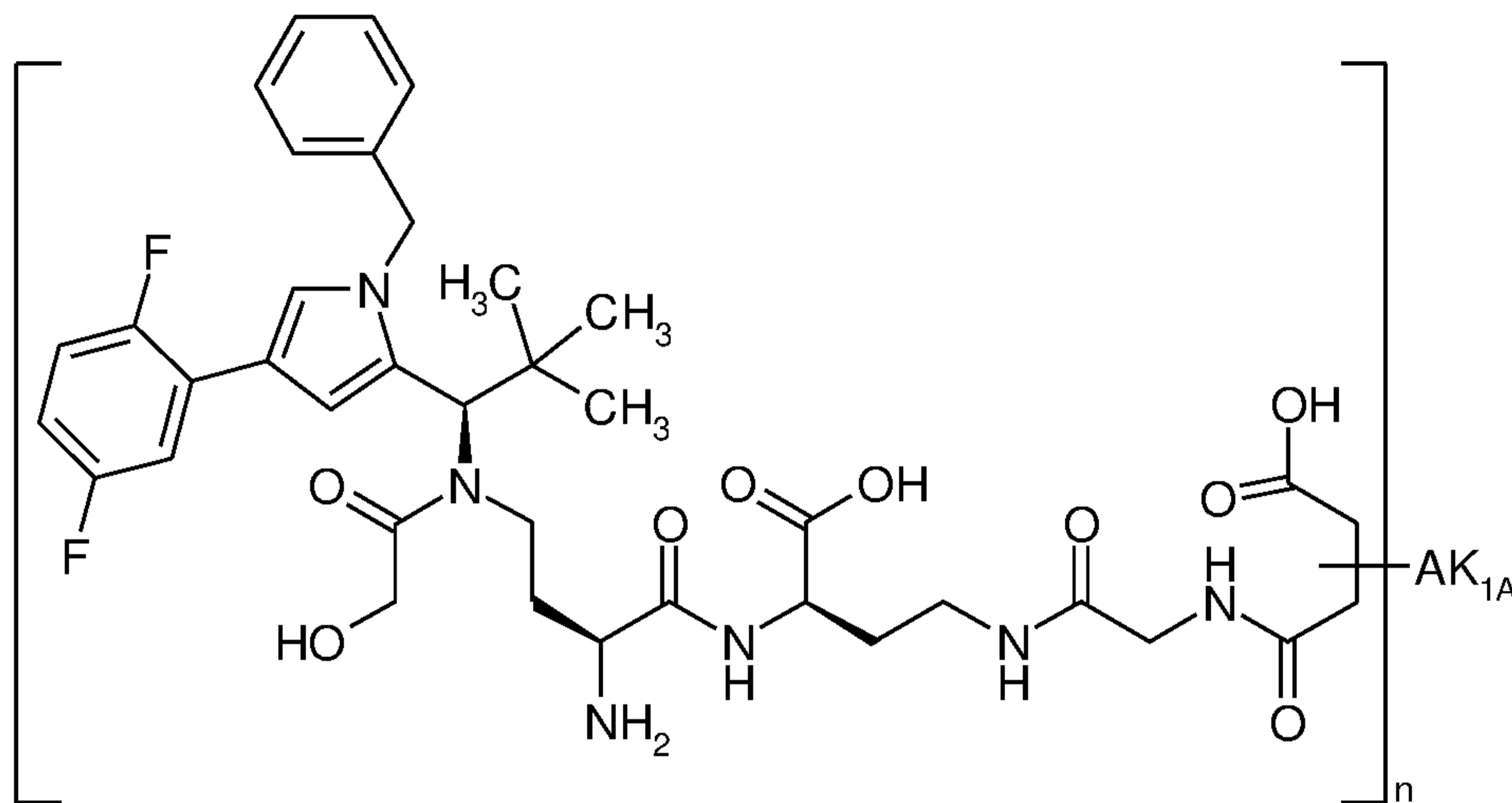
5 Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

10 Protein concentration: 1.14 mg/ml

Drug/mAb ratio: 0.9

**Example 236A**

15



Analogously to Example 234A, 5 mg of cetuximab in 500  $\mu$ l of PBS (c= 10 mg/ml) were coupled with Intermediate F236. Under these  
 20 conditions, some of the ADCs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.85 mg/ml

25

Drug/mAb ratio: 3.6

**Example 236B**

Analogously to Example 234A, 5 mg of anti-TWEAKR AK-1 in 500  $\mu$ l of PBS (c= 10 mg/ml) were coupled with Intermediate F236. Under these conditions, some of the ADCs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.67 mg/ml

Drug/mAb ratio: 3.5

10

### Example 236E

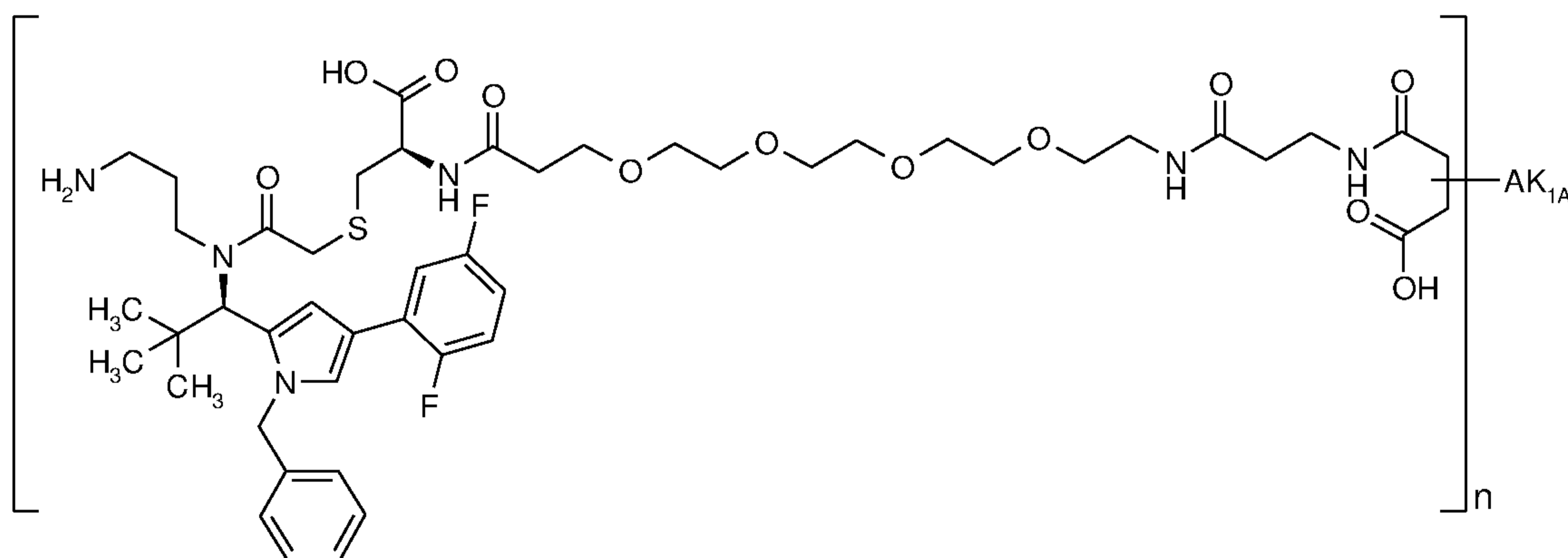
Analogously to Example 234A, 5 mg of trastuzumab in 500  $\mu$ l of PBS (c= 10 mg/ml) were coupled with Intermediate F236. Under these conditions, some of the ADCs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.67 mg/ml

20

Drug/mAb ratio: 3.9

### Example 237A



25

Under argon, a solution of 0.344 mg of TCEP in 100  $\mu$ l of PBS buffer was added to 60 mg of cetuximab in 4000  $\mu$ l of PBS (c=15 mg/ml). The reaction was stirred at RT for 1 h, and 3.04 mg (0.003 mmol) of Intermediate F104 dissolved in 600  $\mu$ l of DMSO were then added. After a further 90 min of stirring at RT, the

30

reaction was diluted to 15 ml with PBS buffer which had been adjusted to pH 8 beforehand.

This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. This solution was stirred under argon at RT overnight and then concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and reconcentrated again. Under these conditions, some of the ADCs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 13.78 mg/ml

Drug/mAb ratio: 4.8

#### **Example 237B**

Under argon, a solution of 0.287 mg of TCEP in 500  $\mu$ l of PBS buffer was added to 50 mg of anti-TWEAKR AK-1 in 2688  $\mu$ l of PBS (c=18.6 mg/ml), and the mixture was then diluted with 8812  $\mu$ l of PBS buffer which had been adjusted to pH 8 beforehand. The reaction was stirred at RT for 1 h, and 2,894 mg (0.003 mmol) of Intermediate F104 dissolved in 500  $\mu$ l of DMSO were then added. After a further 90 min of stirring at RT, the reaction was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. This solution was stirred under argon at RT overnight and then concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and reconcentrated again. Under these conditions, some of the ADCs may also be present in the ring-closed form. For this batch, the proportion of the ring-opened form was determined directly after the synthesis to be 23%. The ADC batch obtained was characterized as follows:

35

Protein concentration: 11.1 mg/ml

Drug/mAb ratio: 3.3

Example 237I

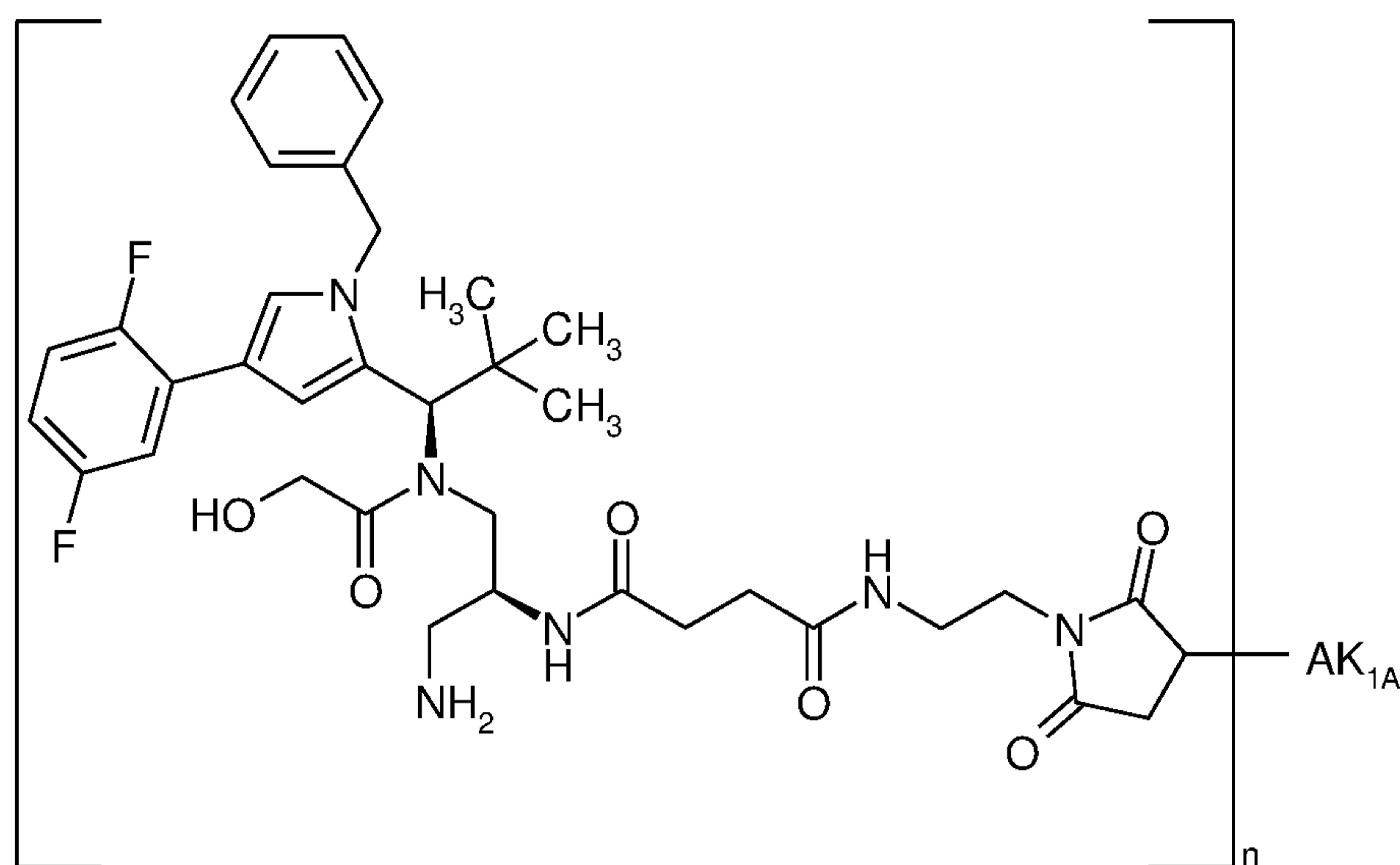
Analogously to Example 237A, 60 mg of nimotuzumab in 4587  $\mu$ l of  
 5 PBS (c= 13.08 mg/ml) were coupled with Intermediate F209. Under  
 these conditions, some of the ADSs may also be present in the  
 ring-closed form. The ADC batch obtained was characterized as  
 follows:

10 Protein concentration: 15.88 mg/ml

Drug/mAb ratio: 4.0

Example 238A

15



Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS  
 buffer was added to 5 mg of cetuximab in 500  $\mu$ l of PBS (c=10  
 20 mg/ml). The reaction was stirred at RT for 30 min, and 0.22 mg  
 (0.00027 mmol) of Intermediate F238 dissolved in 50  $\mu$ l of DMSO  
 were then added. After a further 90 min of stirring at RT, the  
 reaction was diluted with 1900  $\mu$ l of PBS buffer which had been  
 adjusted to pH 8 beforehand.

25

This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-  
 25, GE Healthcare) which had been equilibrated with PBS buffer



pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

Protein concentration: 1.89 mg/ml

10 Drug/mAb ratio: 2.9

#### **Example 238B**

15 Analogously to Example 238A, 5 mg of anti-TWEAKR AK-1 in 500 µl of PBS (c= 10 mg/ml) were coupled with Intermediate F238. Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

20 Protein concentration: 1.66 mg/ml

Drug/mAb ratio: 3.1

25 Under these coupling conditions, 26% of the ADC were present in the form of the hydrolysed open-chain succinamides attached to the antibody.

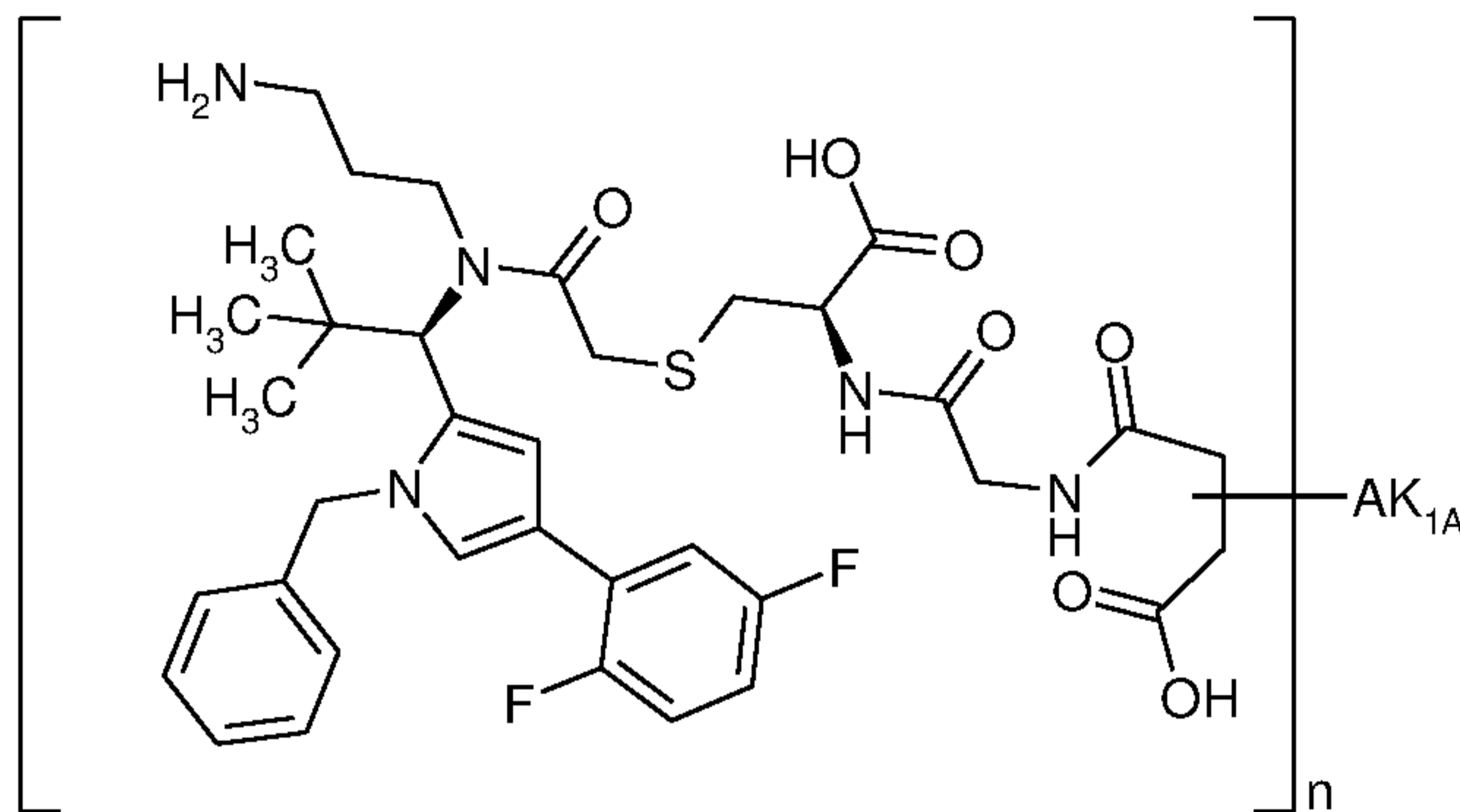
#### **Example 238E**

30 Analogously to Example 238A, 5 mg of trastuzumab in 500 µl of PBS (c= 10 mg/ml) were coupled with Intermediate F238. Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody. The ADC batch obtained was characterized as follows:

35

Protein concentration: 1.84 mg/ml

Drug/mAb ratio: 3.5

**Example 239A**

5

Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS buffer was added to 5 mg of cetuximab in 458  $\mu$ l of PBS (c=10.92 mg/ml). The reaction was diluted with 1892  $\mu$ l of PBS buffer which had been adjusted to pH 8 beforehand and stirred at RT for 10 1 h. 0.19 mg (0.00027 mmol) of Intermediate F217 dissolved in 100  $\mu$ l of DMSO were then added. After a further 90 min of stirring at RT, the reaction was applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate 15 was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

20

Protein concentration: 1.86 mg/ml

Drug/mAb ratio: 2.8

25 **Example 239B**

Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS buffer was added to 5 mg of anti-TWEAKR AK-1 in 269  $\mu$ l of PBS (c=18.6 mg/ml). The reaction was diluted with 2081  $\mu$ l of PBS 30 buffer which had been adjusted to pH 8 beforehand and stirred at RT for 1 h. 0.19 mg (0.00027 mmol) of Intermediate F217

dissolved in 100 µl of DMSO were then added. After a further 90 min of stirring at RT, the reaction was applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate  
5 was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

10

Protein concentration: 1.28 mg/ml

Drug/mAb ratio: 2.6

15 **Example 239I**

Under argon, a solution of 0.029 mg of TCEP in 50 µl of PBS buffer was added to 5 mg of nimotuzumab in 382 µl of PBS (c=13.1 mg/ml). The reaction was diluted with 1968 µl of PBS buffer  
20 which had been adjusted to pH 8 beforehand and stirred at RT for 1 h. 0.19 mg (0.00027 mmol) of Intermediate F217 dissolved in 100 µl of DMSO were then added. After a further 90 min of stirring at RT, the reaction was applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with  
25 PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized  
30 as follows:

Protein concentration: 1.32 mg/ml

Drug/mAb ratio: 2.9

35

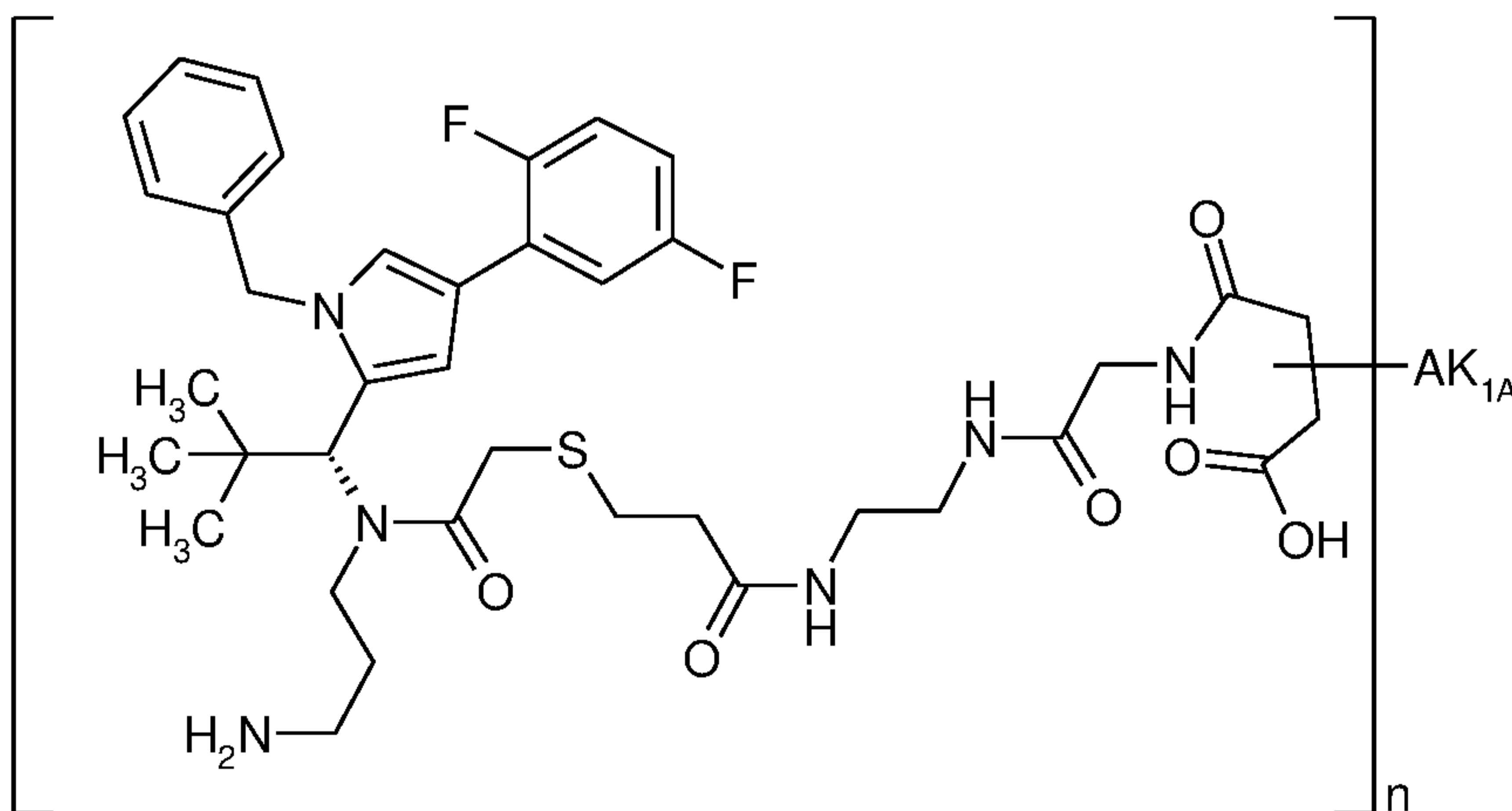
For this ADC preparation, a proportion of 89% was determined for the ring-opened succinamide form.

**Example 239H**

Under argon, a solution of 0.048 mg of TCEP in 83  $\mu$ l of PBS buffer was added to 5 mg of panitumumab in 74  $\mu$ l of PBS (c=67.5 mg/ml). The reaction was diluted with 2243  $\mu$ l of PBS buffer which had been adjusted to pH 8 beforehand and stirred at RT for 4 h. 0.19 mg (0.00027 mmol) of Intermediate F217 dissolved in 100  $\mu$ l of DMSO were then added. The reaction was stirred at RT overnight. The solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.33 mg/ml

Drug/mAb ratio: 2.1

**Example 240A**

25

Under argon, a solution of 0.29 mg of TCEP in 500  $\mu$ l of PBS buffer was added to 50 mg of cetuximab in 4579  $\mu$ l of PBS (c=10.92 mg/ml). The reaction was diluted with 7421  $\mu$ l of PBS buffer which had been adjusted to pH 8 beforehand and stirred at RT for

1 h. 1.4 mg (0.0027 mmol) of Intermediate F213 dissolved in 500  
µl of DMSO were then added. After a further 90 min of stirring  
at RT, the reaction was applied to PD 10 columns (Sephadex<sup>®</sup> G-  
25, GE Healthcare) which had been equilibrated with PBS buffer  
5 pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred  
under argon at RT overnight and then concentrated by  
ultracentrifugation and rediluted with PBS buffer (pH 7.2).  
Under these conditions, some of the ADSs may also be present in  
the ring-closed form. The ADC batch obtained was characterized  
10 as follows:

Protein concentration: 15.63 mg/ml

Drug/mAb ratio: 2.7

15

#### **Example 240B**

Analogously to Example 239A, 5 mg of anti-TWEAKR AK-1 in 500 µl  
of PBS (c= 10 mg/ml) were coupled with Intermediate F213. Under  
20 these conditions, some of the ADSs may also be present in the  
ring-closed form. The ADC batch obtained was characterized as  
follows:

Protein concentration: 0.9 mg/ml

25

Drug/mAb ratio: 2.0

#### **Example 240E**

30 Analogously to Example 239A, 5 mg of trastuzumab in 500 µl of  
PBS (c= 10 mg/ml) were coupled with Intermediate F213. Under  
these conditions, some of the ADSs may also be present in the  
ring-closed form. The ADC batch obtained was characterized as  
follows:

35

Protein concentration: 1.4 mg/ml

Drug/mAb ratio: 2.1



ultracentrifugation and rediluted with PBS.

Protein concentration: 1.39 mg/ml

5 Drug/mAb ratio: 3.9

**Example 241E**

10 Here, 5 mg of trastuzumab in PBS (c=10 mg/ml) were used for coupling with Intermediate F241. After TCEP reduction, coupling with the antibody was carried out with stirring overnight, followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

15

Protein concentration: 1.83 mg/ml

Drug/mAb ratio: 4.7

20 **Example 241I**

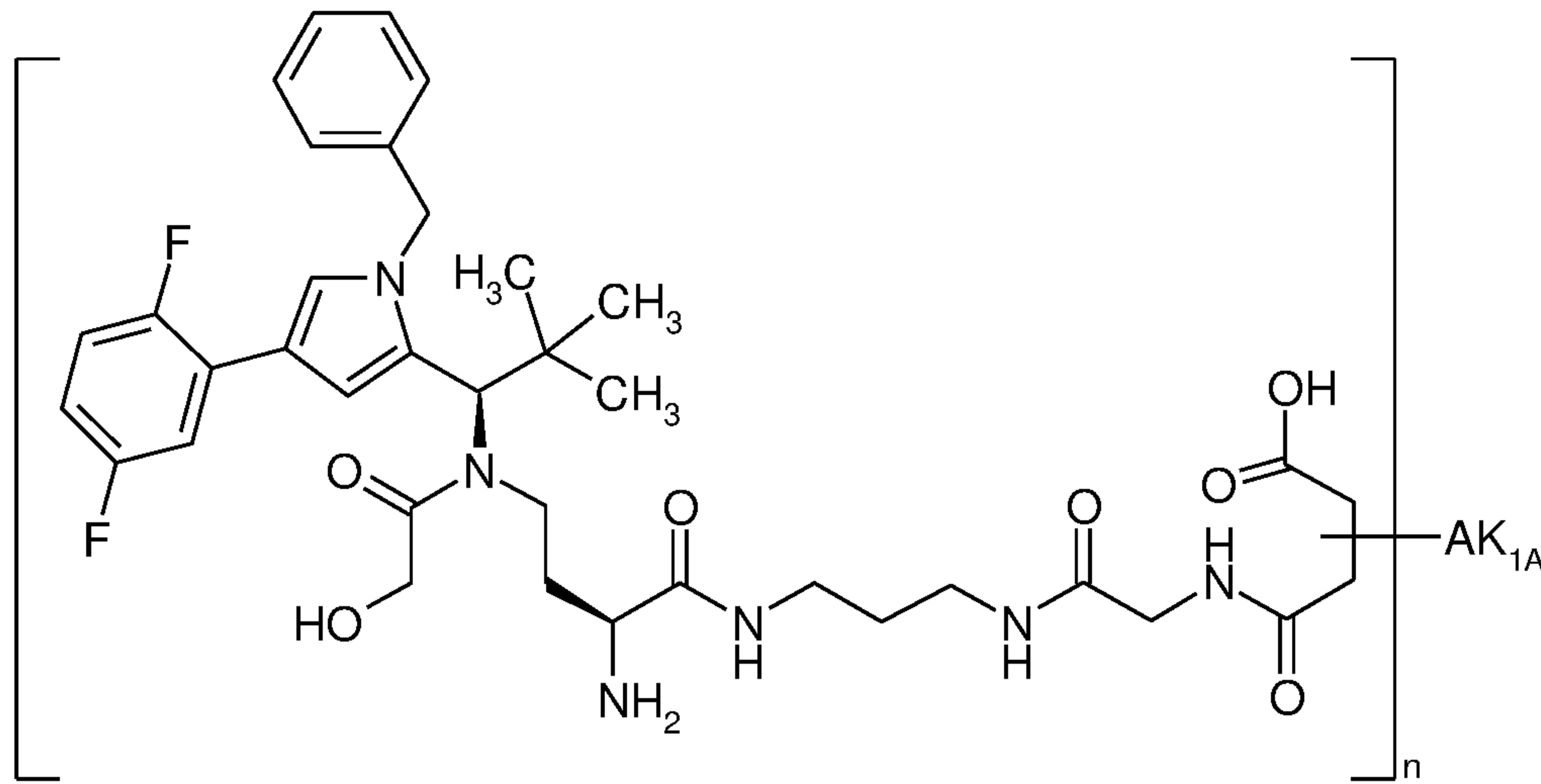
Here, 5 mg of nimotuzumab in PBS (c=10 mg/ml) were used for coupling with Intermediate F241. After TCEP reduction, coupling with the antibody was carried out with stirring overnight,  
25 followed by further work-up by Sephadex purification. After Sephadex purification, the reaction was concentrated by ultracentrifugation and rediluted with PBS.

30

Protein concentration: 1.6 mg/ml

Drug/mAb ratio: 4.0

**Example 242A**



Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS  
 buffer was added to 5 mg of cetuximab in 500  $\mu$ l of PBS (c=10  
 5 mg/ml). The reaction was stirred at RT for 30 min, and 0.22 mg  
 (0.00027 mmol) of Intermediate F242 dissolved in 50  $\mu$ l of DMSO  
 were then added. After a further 90 min of stirring at RT, the  
 reaction was diluted with 1900  $\mu$ l of PBS buffer which had been  
 adjusted to pH 8 beforehand.

10

This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-  
 25, GE Healthcare) which had been equilibrated with PBS buffer  
 pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred  
 under argon at RT overnight and then concentrated by  
 15 ultracentrifugation and rediluted with PBS buffer (pH 7.2).  
 Under these conditions, some of the ADSs may also be present in  
 the ring-closed form. The ADC batch obtained was characterized  
 as follows:

20 Protein concentration: 1.92 mg/ml

Drug/mAb ratio: 2.7

### Example 242B

25

As described in Example 242A, 5 mg of anti-TWEAKR AK-1 in 500  
 $\mu$ l of PBS (c= 10 mg/ml) were coupled with 0.22 mg of Intermediate  
 F242. Under these conditions, some of the ADSs may also be



present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.54 mg/ml

5

Drug/mAb ratio: 3.1

#### Example 242E

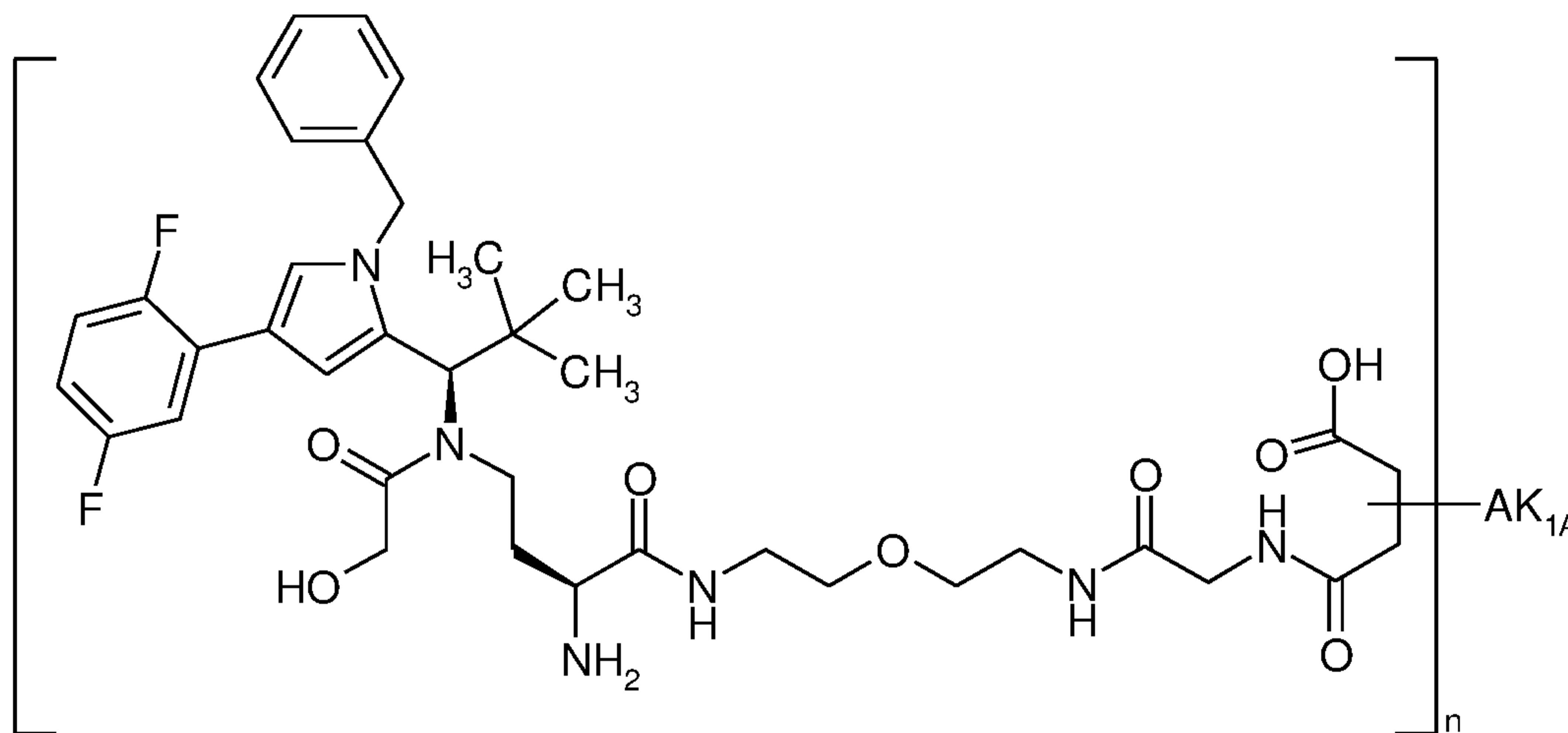
10 As described in Example 242A, 5 mg of trastuzumab in 500  $\mu$ l of PBS (c= 10 mg/ml) were coupled with 0.22 mg of Intermediate F242. Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

15

Protein concentration: 1.76 mg/ml

Drug/mAb ratio: 3.6

#### 20 Example 243A



25 Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS buffer was added to 5 mg of cetuximab in 500  $\mu$ l of PBS (c=10 mg/ml). The reaction was stirred at RT for 30 min, and 0.23 mg (0.00027 mmol) of Intermediate F243 dissolved in 50  $\mu$ l of DMSO were then added. After a further 90 min of stirring at RT, the reaction was diluted with 1900  $\mu$ l of PBS buffer which had been

adjusted to pH 8 beforehand.

This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.92 mg/ml

Drug/mAb ratio: 2.9

#### **Example 243B**

As described in Example 243A, 5 mg of anti-TWEAKR AK-1 in 500  $\mu$ l of PBS (c= 10 mg/ml) were coupled with 0.23 mg of Intermediate F243. Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.63 mg/ml

Drug/mAb ratio: 3.1

#### **Example 243E**

As described in Example 243A, 5 mg of trastuzumab in 500  $\mu$ l of PBS (c= 10 mg/ml) were coupled with 0.23 mg of Intermediate F243. Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

Protein concentration: 1.8 mg/ml

Drug/mAb ratio: 3.5



Drug/mAb ratio: 3.5

**Example 244B**

5

Here, 5.0 mg of anti-TWEAK AK-1 in PBS (c=18.60 mg/ml) were used for coupling with Intermediate F244, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.

10

Protein concentration: 1.65 mg/ml

Drug/mAb ratio: 3.5

15 **Example 244E**

Here, 5.0 mg of trastuzumab in PBS (c=13.50 mg/ml) were used for coupling with Intermediate F244, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and

20

rediluted with PBS.  
Protein concentration: 1.91 mg/ml

Drug/mAb ratio: 3.5

25

**Example 244I**

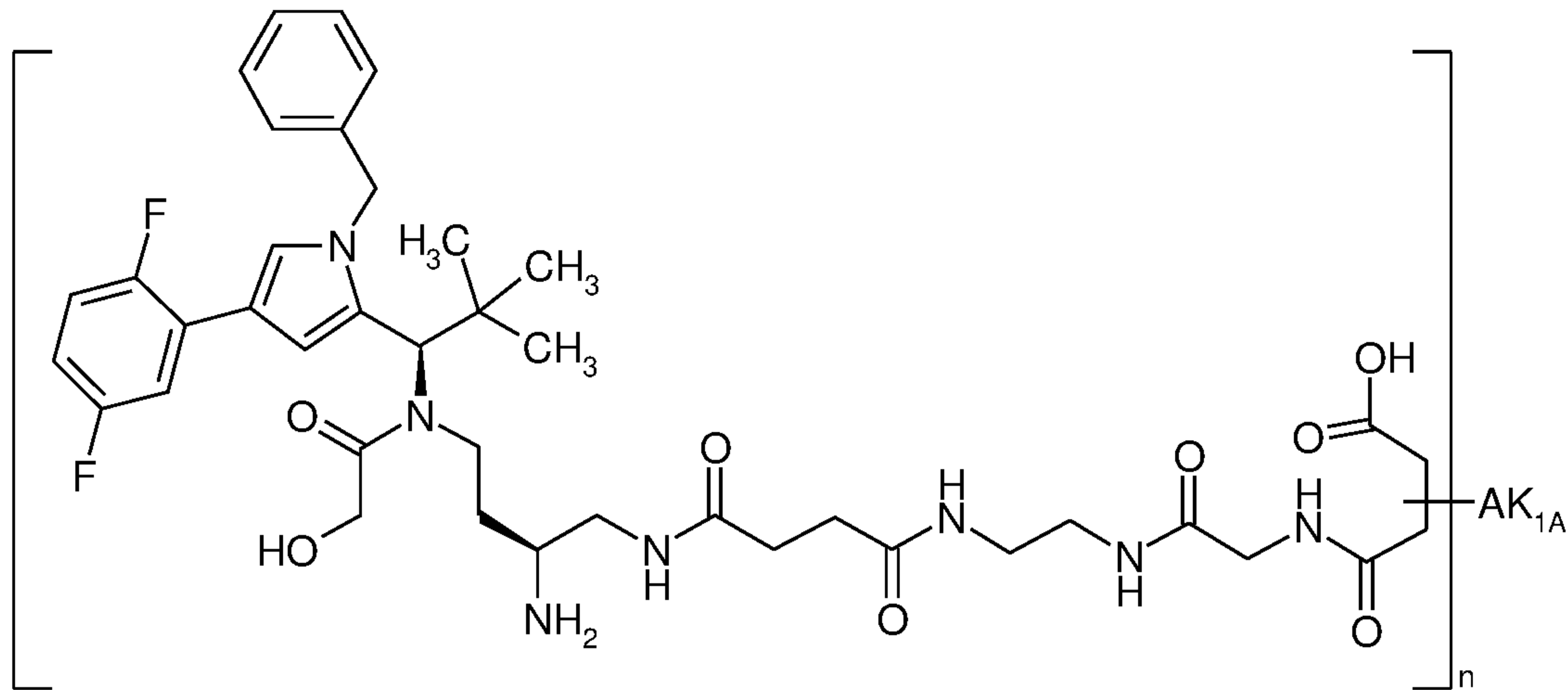
Here, 5.0 mg of nimotuzumab in PBS (c=13.08 mg/ml) were used for coupling with Intermediate F244, and the reaction was, after

30

Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS.  
Protein concentration: 1.91 mg/ml

35 Drug/mAb ratio: 2.9

**Example 245A**



Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS buffer was added to 5 mg of cetuximab in 500  $\mu$ l of PBS (c=10 mg/ml). The reaction was stirred at RT for 30 min, and 0.24 mg (0.00027 mmol) of Intermediate F245 dissolved in 50  $\mu$ l of DMSO were then added. After a further 90 min of stirring at RT, the reaction was diluted with 1900  $\mu$ l of PBS buffer which had been adjusted to pH 8 beforehand.

10

This solution was then applied to PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) which had been equilibrated with PBS buffer pH 8 and was eluted with PBS buffer pH 8. The eluate was stirred under argon at RT overnight and then concentrated by ultracentrifugation and rediluted with PBS buffer (pH 7.2). Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was characterized as follows:

20 Protein concentration: 1.69 mg/ml

Drug/mAb ratio: 2.4

### Example 245B

25

As described in Example 245A, 5 mg of anti-TWEAKR AK-1 in 500  $\mu$ l of PBS (c= 10 mg/ml) were coupled with 0.24 mg of Intermediate F245. Under these conditions, some of the ADSs may also be present in the ring-closed form. The ADC batch obtained was

characterized as follows:

Protein concentration: 1.51 mg/ml

5 Drug/mAb ratio: 2.5

**Example 245E**

As described in Example 245A, 5 mg of trastuzumab in 500 µl of  
 10 PBS (c= 10 mg/ml) were coupled with 0.24 mg of Intermediate  
 F245. Under these conditions, some of the ADSs may also be  
 present in the ring-closed form. The ADC batch obtained was  
 characterized as follows:

15 Protein concentration: 1.8 mg/ml

Drug/mAb ratio: 3.5

**Example 245I**

20 As described in Example 245A, 5 mg of nimotuzumab in 500 µl of  
 PBS (c= 10 mg/ml) were coupled with 0.24 mg of Intermediate  
 F245. Under these conditions, some of the ADSs may also be  
 present in the ring-closed form. The ADC batch obtained was  
 25 characterized as follows:

Protein concentration: 1.65 mg/ml

Drug/mAb ratio: 2.3

30

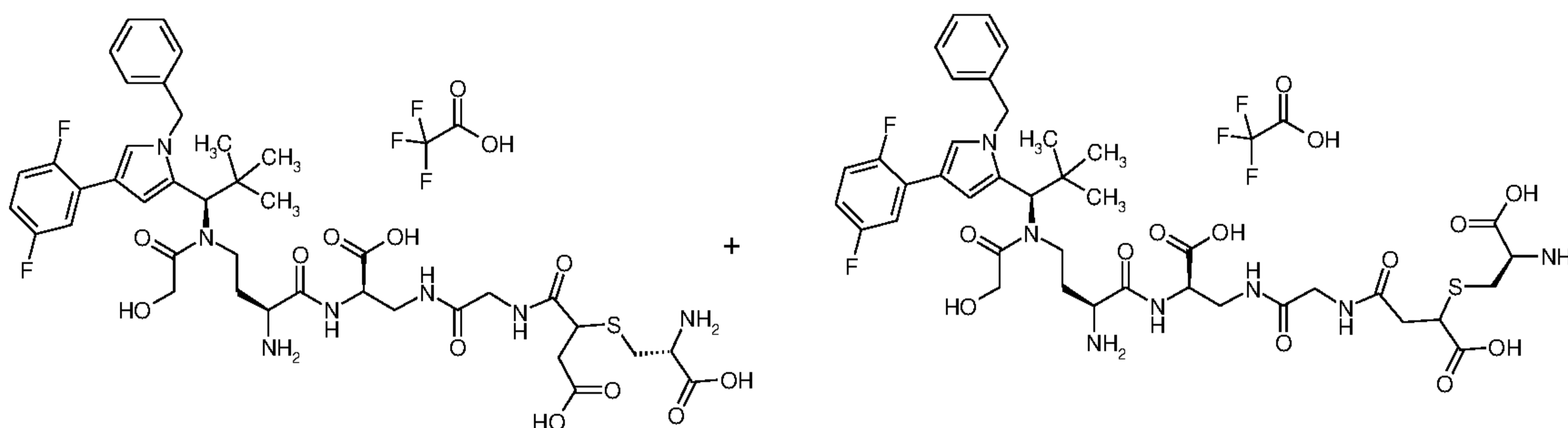
**Example 246**

4-[(2-[[ (2R)-2-((2S)-2-Amino-4-[[ (1R)-1-[1-benzyl-4-(2,5-  
 difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 35 dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-2-  
 carboxyethyl]amino]-2-oxoethyl)amino]-3-[[ (2R)-2-amino-2-  
 carboxyethyl]sulphonyl]-4-oxobutanoic acid / trifluoroacetic  
 acid

and

4-[(2-[[[(2R)-2-({(2S)-2-amino-4-[[[(1R)-1-[1-benzyl-4-(2,5-  
 5 difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-2-  
 carboxyethyl]amino]-2-oxoethyl)amino]-2-[[[(2R)-2-amino-2-  
 carboxyethyl]sulphonyl]-4-oxobutanoic acid / trifluoroacetic  
 acid (1:1)

10



First, L-cysteine was converted with 1-([2-  
 (trimethylsilyl)ethoxy]carbonyl)oxy)pyrrolidine-2,5-dione in  
 15 DMF in the presence of *N,N*-diisopropylethylamine into *N*-{[2-  
 (trimethylsilyl)ethoxy]carbonyl}-L-cysteine.

11 mg (0.013 mmol) of Intermediate F193 and 8 mg (0.016 mmol)  
 of *N*-{[2-(trimethylsilyl)ethoxy]carbonyl}-L-cysteine were  
 20 dissolved in 3 ml of DMF, and the mixture was stirred at RT for  
 20 h. The mixture was then concentrated and the residue was  
 purified by preparative HPLC.

The appropriate fractions were combined and the solvents were  
 25 evaporated under reduced pressure, and the residue was then  
 dissolved in 2 ml of THF/water 1:1. 19  $\mu$ l of a 2M aqueous lithium  
 hydroxide solution were added and the reaction was stirred at  
 RT for 1 h. Another 19  $\mu$ l of the 2M aqueous lithium hydroxide  
 solution were then added and the reaction was stirred at RT  
 30 overnight. The mixture was then neutralized with a 1M  
 hydrochloric acid, the solvent was evaporated under reduced  
 pressure and the residue was purified by preparative HPLC. This  
 gave 4.1 mg (38% of theory) of the regioisomeric protected

intermediates as a colourless foam.

LC-MS (Method 1):  $R_t = 1.03$  min (broad); MS (ESIpos):  $m/z = 1020$  (M+H)<sup>+</sup>.

5

In the last step, 4.1 mg (0.004 mmol) of this intermediate were dissolved in 3 ml of 2,2,2-trifluoroethanol. 3 ml (0.022 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 1 h. 6 mg (0.022 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid and 2 ml of a 0.1% strength aqueous trifluoroacetic acid were then added, and the solvent was evaporated under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 5 mg (quant.) of the title compound as a regioisomer mixture in a ratio of 20:80.

10

15

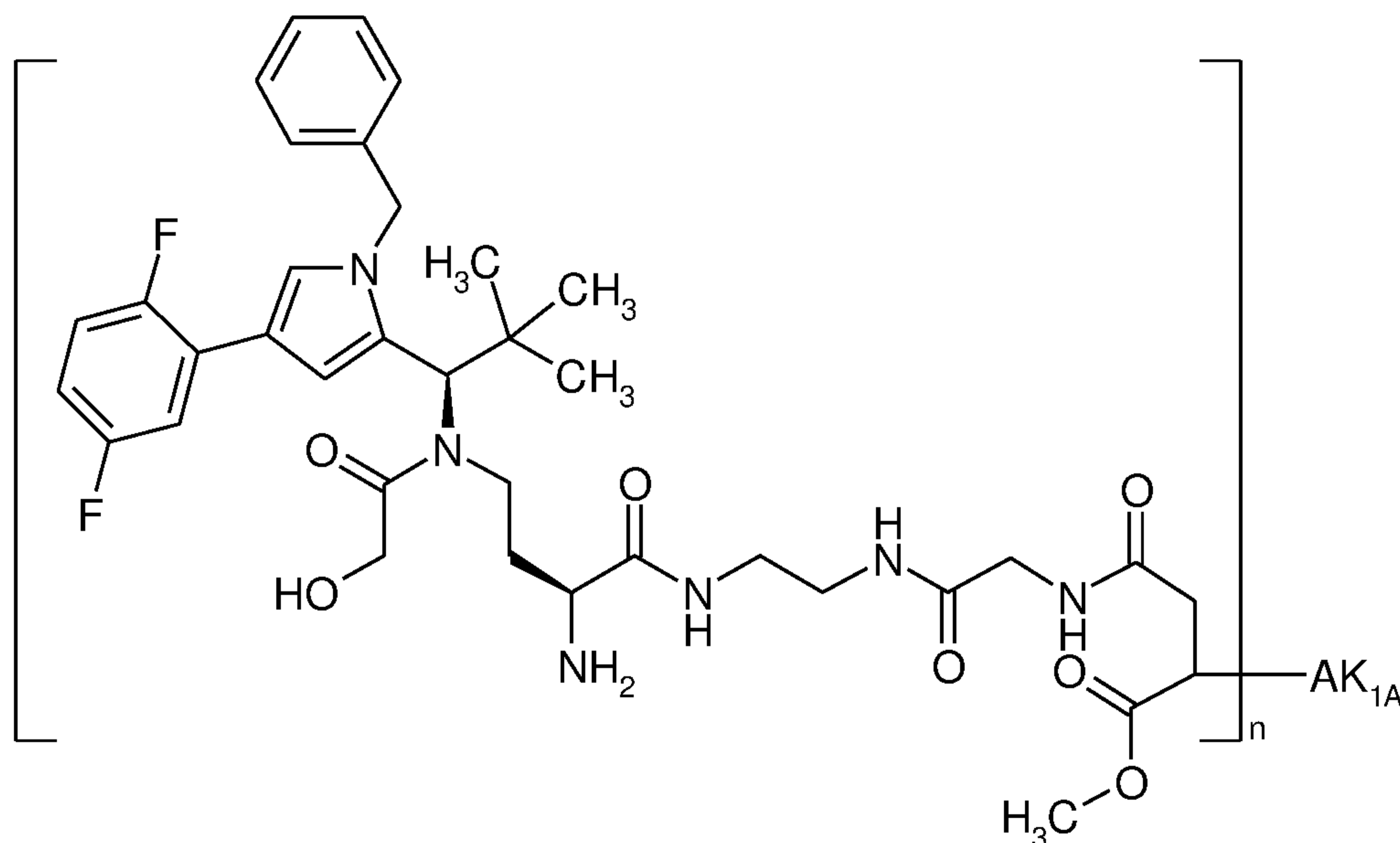
LC-MS (Method 1):  $R_t = 0.78$  min (broad); MS (ESIpos):  $m/z = 876$  (M+H)<sup>+</sup>.

20

LC-MS (Method 5):  $R_t = 2.36$  min and 2.39 min; MS (ESIpos):  $m/z = 876$  (M+H)<sup>+</sup>.

### Example 247A

25





Under argon, a solution of 0.029 mg of TCEP in 50  $\mu$ l of PBS buffer was added to 5 mg of cetuximab in 500  $\mu$ l of PBS (c=10 mg/ml). The reaction was stirred at RT for 30 min, and 0.264 mg (0.27  $\mu$ mol) of Intermediate F247 dissolved in 50  $\mu$ l of DMSO were then added. After 20 h of stirring at RT, the reaction was diluted with 1.9 ml of PBS buffer and eluted through PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) using PBS buffer. The eluate was then concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and reconcentrated again. The ADC batch obtained was characterized as follows:

Protein concentration: 1.66 mg/ml

Drug/mAb ratio: 2.2

#### **Example 247B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=10 mg/ml) were used for coupling with Intermediate F247, and coupling and work-up were carried out as described in Example 247A.

Protein concentration: 1.49 mg/ml

Drug/mAb ratio: 2.6

#### **Example 247E**

Here, 5 mg of trastuzumab in PBS (c=10 mg/ml) were used for coupling with Intermediate F247, and coupling and work-up were carried out as described in Example 247A.

Protein concentration: 1.67 mg/ml

Drug/mAb ratio: 2.3

#### **Example 247I**

Here, 5 mg of nimotuzumab in PBS (c=10 mg/ml) were used for

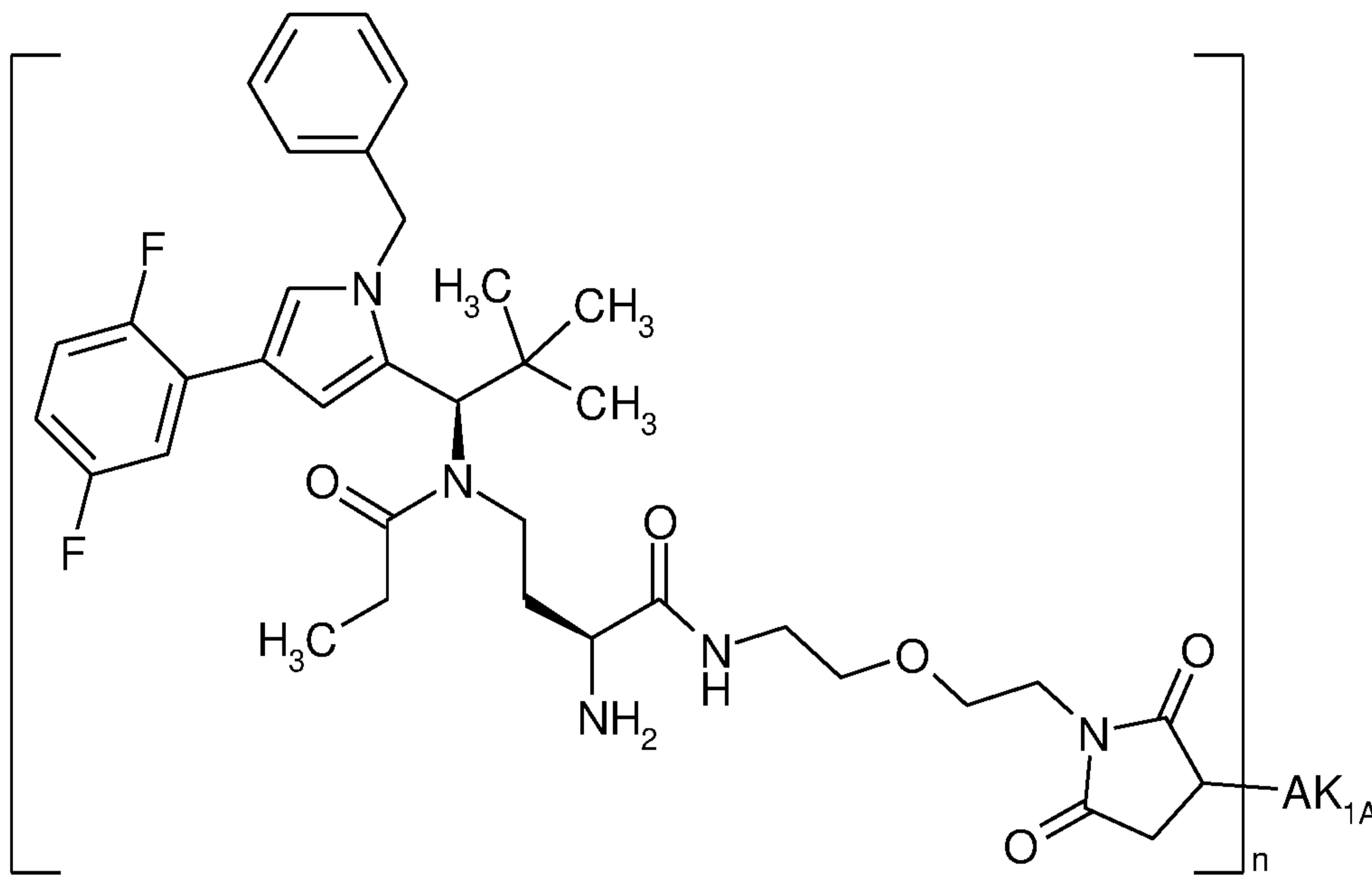
coupling with Intermediate F247, and coupling and work-up were carried out as described in Example 247A.

Protein concentration: 1.62 mg/ml

5

Drug/mAb ratio: 2.4

### Example 248A



10

Here, analogously to Example 5A, 5 mg of cetuximab in PBS (c=10.92 mg/ml) were used for coupling with Intermediate F248, and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

20

Protein concentration: 2.06 mg/ml

Drug/mAb ratio: 3.8

### Example 248B

25 Here, analogously to Example 5B, 5 mg of anti-TWEAKR AK-1 in PBS (c=18.6 mg/ml) were used for coupling with Intermediate F248,

and the reaction was, after Sephadex purification, concentrated by ultracentrifugation and rediluted with PBS. Some of the ADC prepared in this manner may also be present in the form of the hydrolysed open-chain succinamides attached to the antibody.

5

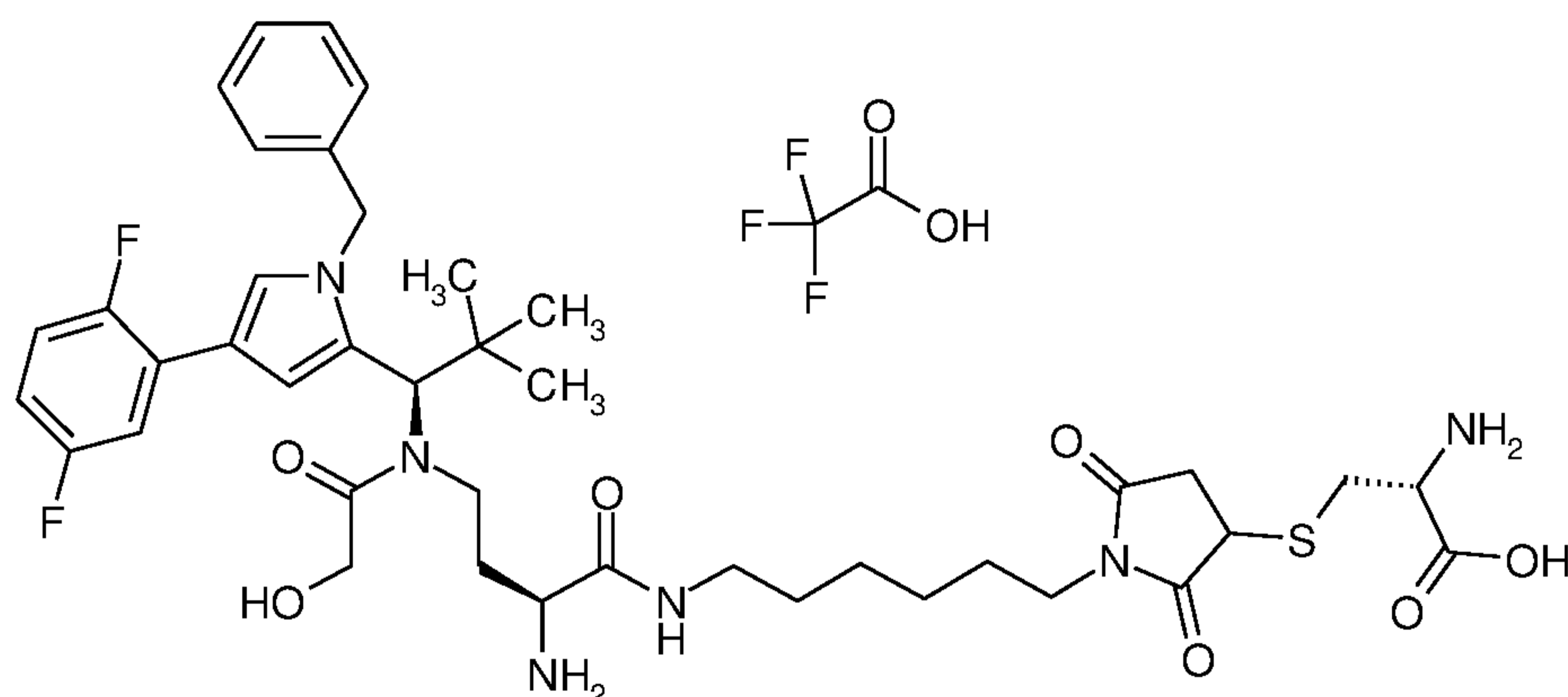
Protein concentration: 1.84 mg/ml

Drug/mAb ratio: 4.1

10 **Example 249**

S-{1-[6-({(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)hexyl]-2,5-dioxopyrrolidin-3-yl}-L-cysteine/trifluoroacetic acid (1:1)

15



11 mg (14  $\mu\text{mol}$ ) of Intermediate F179 were taken up in 2.2 ml of DMF, and 3.3 mg (27  $\mu\text{mol}$ ) of L-cysteine were added. The reaction mixture was stirred at RT for 3 h and then concentrated under reduced pressure. The residue was purified by preparative HPLC. The appropriate fractions were concentrated, giving, after lyophilization of the residue from acetonitrile/water, 7.3 mg (58% of theory) of the title compound as a colourless foam.

25

LC-MS (Method 4):  $R_t = 1.04$  min; MS (EIpos):  $m/z = 813$   $[M+H]^+$ .

**Example 250**

30

4-{{2-({(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-

difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-3-[[ (2R)-2-amino-2-carboxyethyl]sulphanyl]-4-oxobutanoic acid / trifluoroacetic acid (1:1)

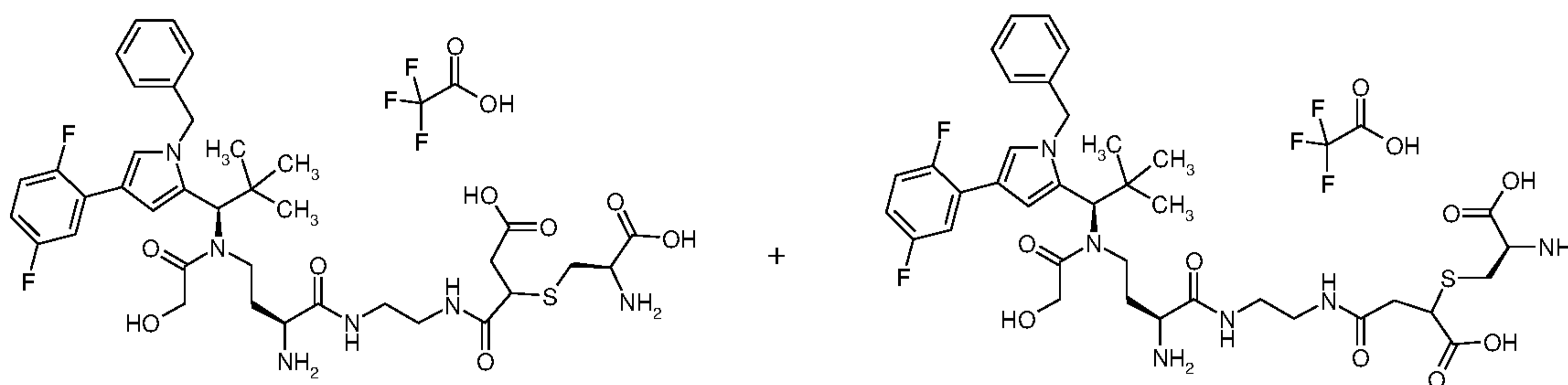
5

and

4-{{2-({(2S)-2-amino-4-[[ (1R)-1-[1-benzyl-4-(2,5-

difluorophenyl)-1H-pyrrol-2-yl]-2,2-

10 dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]amino}-2-[[ (2R)-2-amino-2-carboxyethyl]sulphanyl]-4-oxobutanoic acid / trifluoroacetic acid (1:1)



15

10 mg (0.013 mmol) of Intermediate F85 and 5.3 mg (0.02 mmol) of N-{{2-(trimethylsilyl) ethoxy}carbonyl}-L-cysteine were dissolved in 3 ml of DMF, and the mixture was stirred at RT for 3 days. The mixture was then concentrated and the residue was purified by preparative HPLC. The appropriate fractions were combined and the solvents were evaporated under reduced pressure, and the residue was then dissolved in 2 ml of THF/water 1:1. 9  $\mu$ l of a 2M aqueous lithium hydroxide solution were added and the reaction was stirred at RT for 1 h. The reaction was then adjusted to a pH of ~3 with a 1M hydrochloric acid, the solvent was evaporated under reduced pressure and the residue was purified by preparative HPLC. This gave 3 mg (24% of theory over 2 steps) of the regioisomeric protected intermediates as a colourless foam.

30

LC-MS (Method 5):  $R_t$  = 3.39 min and 3.43 min; MS (ESIpos):  $m/z$  = 919 (M+H)<sup>+</sup>.

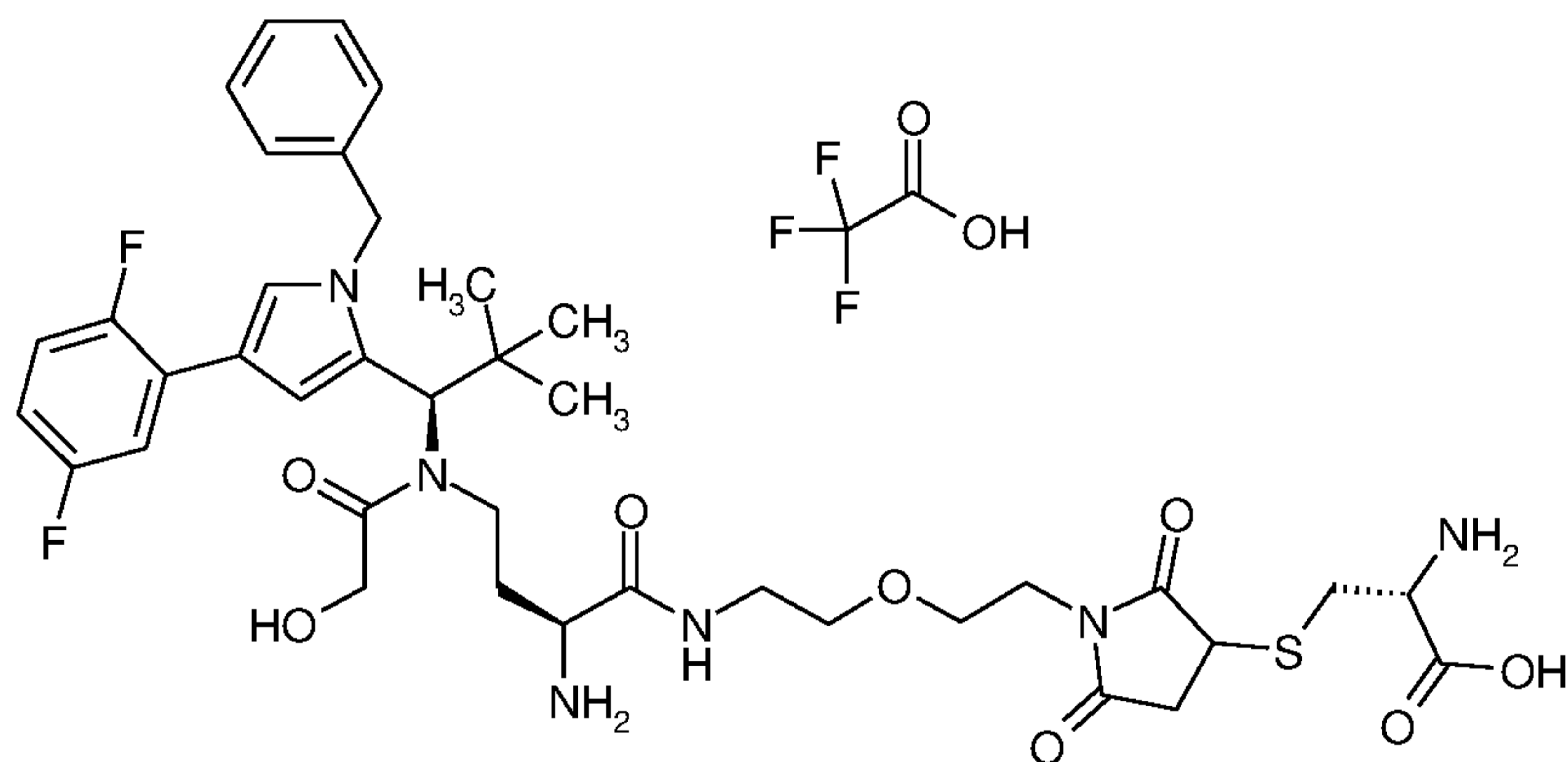
In the last step, 3 mg (0.0033 mmol) of this intermediate were

dissolved in 3 ml of 2,2,2-trifluoroethanol. 2.2 ml (0.016 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 3.5 h. 4.8 mg (0.016 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were then added, and the solvent was evaporated under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 1 mg (33% of theory) of the title compound as a regioisomer mixture in a ratio of 43:34. The isomer mixture comprised 23% of a further isomer (RT = 2.51).

LC-MS (Method 5):  $R_t$  = 2.57 min and 2.62 min; MS (ESIpos):  $m/z$  = 775 (M+H)<sup>+</sup>.

### 15 Example 251

S-(1-{2-[2-({(2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethoxy]ethyl}-2,5-dioxopyrrolidin-3-yl)-L-cysteine / trifluoroacetic acid (1:1)



25 3 mg (4  $\mu$ mol) of Intermediate F248 were taken up in 2 ml of DMF, and 0.9 mg (8  $\mu$ mol) of L-cysteine were added. The reaction mixture was stirred at RT for 18 h and then concentrated under reduced pressure. The residue was purified by preparative HPLC. The appropriate fractions were concentrated, giving, after lyophilization of the residue from acetonitrile/water, 1.1 mg (32% of theory) of the title compound as a white solid.

LC-MS (Method 1):  $R_t = 0.78$  min; MS (EIpos):  $m/z = 801$   $[M+H]^+$ .

### Example 252

5

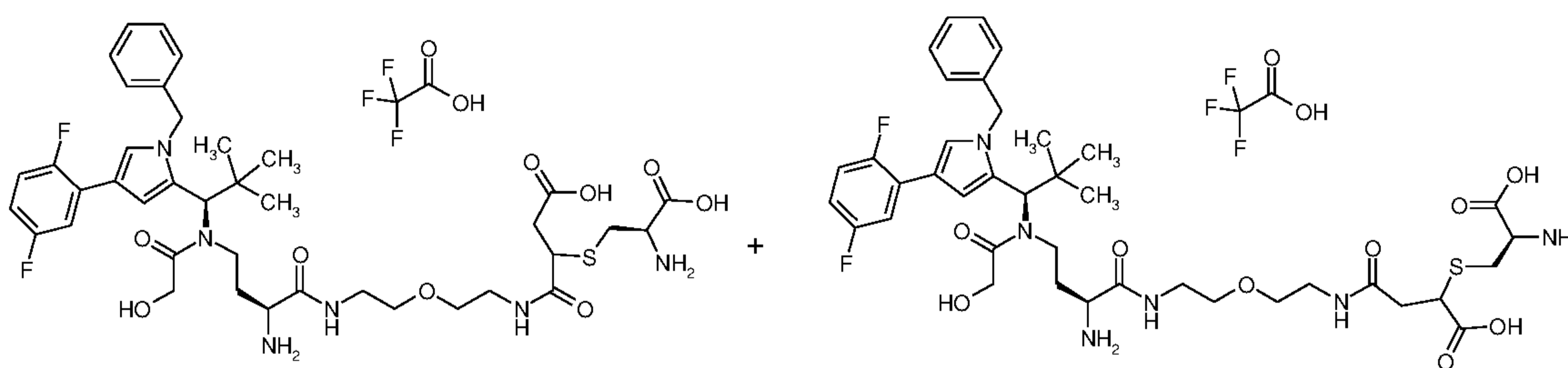
(3R,7S)-7-Amino-17-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-3-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-4-glycoloyl-2,2-dimethyl-8,16-dioxo-12-oxa-4,9,15-triazanonadecan-19-oic acid / trifluoroacetic acid (1:1)

10

and

(3R,7S)-7-amino-18-{[(2R)-2-amino-2-carboxyethyl]sulphonyl}-3-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-4-glycoloyl-2,2-dimethyl-8,16-dioxo-12-oxa-4,9,15-triazanonadecan-19-oic acid / trifluoroacetic acid (1:1)

15



20 8 mg (0.010 mmol) of the protected intermediate of Intermediate F248 and 5.1 mg (0.02 mmol) of N-{[2-(trimethylsilyl)ethoxy]carbonyl}-L-cysteine were dissolved in 3 ml of DMF, and the mixture was stirred at RT for 18 h and then treated in an ultrasonic bath for 2 h. The mixture was then concentrated and

25 the residue was purified by preparative HPLC. The appropriate fractions were combined and the solvents were evaporated under reduced pressure, and the residue was then dissolved in 2 ml of THF/water 1:1. 15  $\mu$ l of a 2M aqueous lithium hydroxide solution were added and the reaction was stirred at RT for 15 min. The

30 reaction was then adjusted to a pH of  $\sim$ 3 with a 1M hydrochloric acid, diluted with 20 ml of sodium chloride solution and extracted twice with 20 ml of ethyl acetate. The organic phase was dried over magnesium sulphate and concentrated, and the residue was lyophilized from acetonitrile/water. This gave 8.4

mg (78% of theory over 2 steps) of the regioisomeric protected intermediates as a colourless foam.

LC-MS (Method 1):  $R_t$  = 1.44 min and 3.43 min; MS (ESIpos):  $m/z$   
5 = 1107 (M+H)<sup>+</sup>.

In the last step, 8 mg (0.007 mmol) of this intermediate were dissolved in 5 ml of 2,2,2-trifluoroethanol. 9.8 ml (0.072 mmol) of zinc chloride were added, and the reaction was stirred at  
10 50°C for 1.5 h. Ethylenediamine-N,N,N',N'-tetraacetic acid were then added, and the solvent was evaporated under reduced pressure. The residue was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 4 mg (59% of theory)  
15 of the title compound as a regioisomer mixture in a ratio of 31:67.

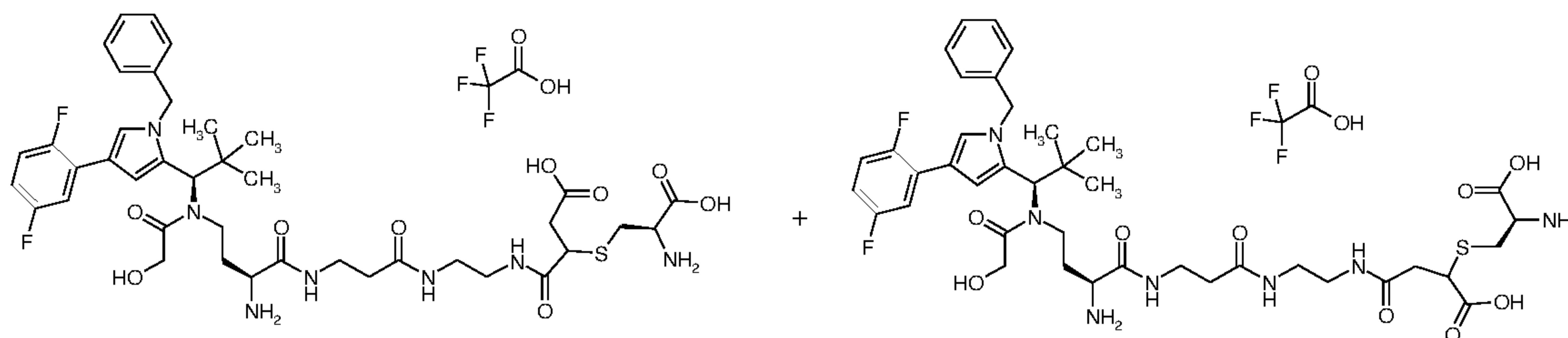
LC-MS (Method 1):  $R_t$  = 0.79 min and 0.81 min; MS (ESIpos):  $m/z$   
20 = 819 (M+H)<sup>+</sup>.

### Example 253

4-({2-[(N-((2S)-2-Amino-4-[(1R)-1-[1-benzyl-4-(2,5-  
difluorophenyl)-1H-pyrrol-2-yl])-2,2-  
25 dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-  
alanyl)amino]ethyl}amino)-3-[[ (2R)-2-amino-2-  
carboxyethyl]sulphonyl]-4-oxobutanoic acid / trifluoroacetic  
acid (1:1)

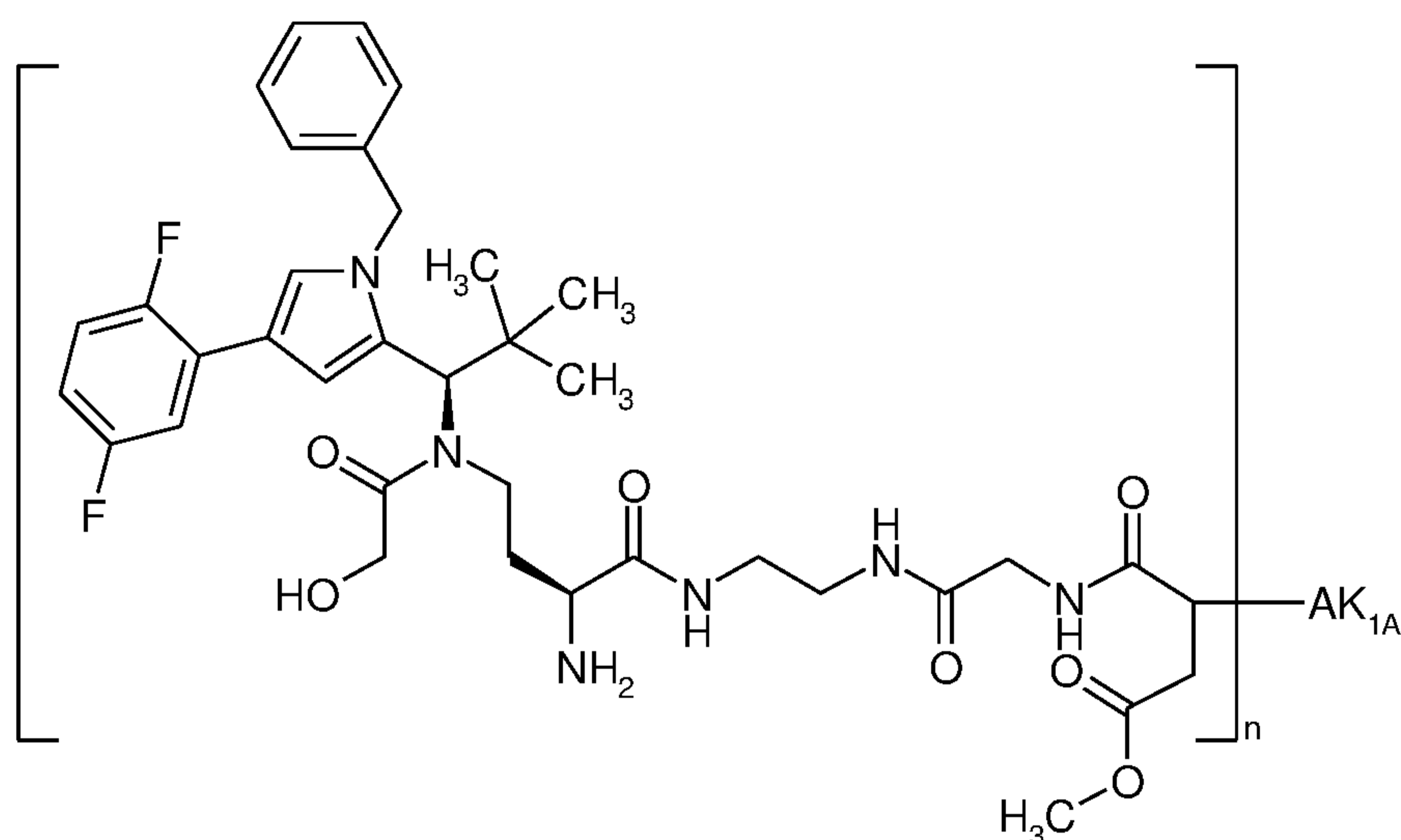
30 and

4-({2-[(N-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-  
difluorophenyl)-1H-pyrrol-2-yl])-2,2-  
dimethylpropyl}(glycoloyl)amino]butanoyl}-beta-  
35 alanyl)amino]ethyl}amino)-2-[[ (2R)-2-amino-2-  
carboxyethyl]sulphonyl]-4-oxobutanoic acid / trifluoroacetic  
acid (1:1)



The isomeric title compounds were prepared analogously to Example 250 from Intermediate F84 and N- $\{[2-(\text{trimethylsilyl})\text{ethoxy}]\text{carbonyl}\}$ -L-cysteine.

### Example 254A



10

Under argon, a solution of 0.029 mg of TCEP in 50  $\mu\text{l}$  of PBS buffer was added to 5 mg of cetuximab in 500  $\mu\text{l}$  of PBS ( $c=10$  mg/ml). The reaction was stirred at RT for 30 min, and 0.264 mg (0.27  $\mu\text{mol}$ ) of Intermediate F254 dissolved in 50  $\mu\text{l}$  of DMSO were then added. After 20 h of stirring at RT, the reaction was diluted with 1.9 ml of PBS buffer and eluted through PD 10 columns (Sephadex<sup>®</sup> G-25, GE Healthcare) using PBS buffer. The eluate was then concentrated by ultracentrifugation, rediluted with PBS buffer (pH 7.2) and reconcentrated again. The ADC batch obtained was characterized as follows:

20

Protein concentration: 1.74 mg/ml

Drug/mAb ratio: 2.2



**Example 254B**

Here, 5 mg of anti-TWEAKR AK-1 in PBS (c=10 mg/ml) were used for  
 5 coupling with Intermediate F254, and coupling and work-up were  
 carried out as described in Example 254A.

Protein concentration: 1.8 mg/ml

10 Drug/mAb ratio: 2.5

**Example 254E**

Here, 5 mg of trastuzumab in PBS (c=10 mg/ml) were used for  
 15 coupling with Intermediate F254, and coupling and work-up were  
 carried out as described in Example 254A.

Protein concentration: 1.74 mg/ml

20 Drug/mAb ratio: 2.4

**Example 254I**

Here, 5 mg of nimotuzumab in PBS (c=10 mg/ml) were used for  
 25 coupling with Intermediate F254, and coupling and work-up were  
 carried out as described in Example 254A.

Protein concentration: 1.73 mg/ml

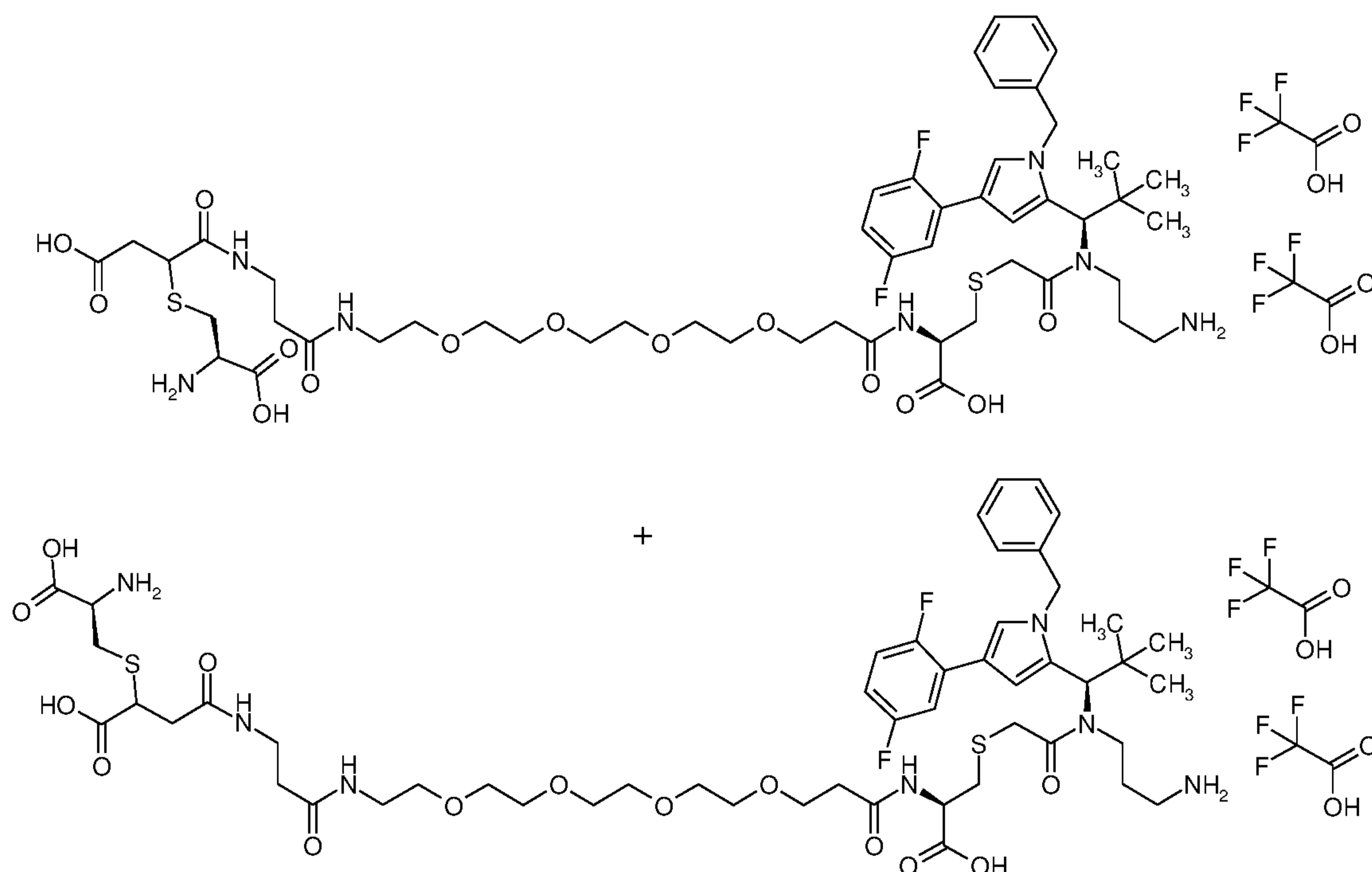
30 Drug/mAb ratio: 2.0

**Example 255**

(2R,28R)-28-Amino-2-[(2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-  
 35 (2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl}amino]-2-oxoethyl)sulphonyl)methyl]-25-  
 (carboxymethyl)-4,20,24-trioxo-7,10,13,16-tetraoxa-26-thia-  
 3,19,23-triazanonacosan-1,29-dioic acid / trifluoroacetic acid

(1:2) and

(1R,28R,34R)-1-amino-33-(3-aminopropyl)-34-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-35,35-dimethyl-6,10,26,32-tetraoxo-14,17,20,23-tetraoxa-3,30-dithia-7,11,27,33-tetraazahexatriacontan-1,4,28-tricarboxylic acid / trifluoroacetic acid (1:2)



10

20 mg (0.018 mmol) of R-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}-N-[19-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-L-cysteine / trifluoroacetic acid (1:1) (Intermediate F209) and 9.78 mg (0.036 mmol) of N-{[2-(trimethylsilyl)ethoxy]carbonyl}-L-cysteine were dissolved in 2 ml of DMF, and the mixture was stirred at RT for 18 h. The reaction mixture was concentrated under reduced pressure. The residue (47.7 mg) was dissolved in 3 ml of THF/water 1:1. 0.08 ml of a 2M aqueous lithium hydroxide solution were added and the reaction was stirred at RT for 1 hour. The reaction was then adjusted to a pH of ~7 using 9.26 mg (0.15 mmol) of acetic acid. The reaction mixture was purified directly by preparative RP-HPLC (column: Reprosil 125x30; 10 $\mu$ , flow rate: 50 ml/min,

15

20

25

MeCN/water; 0.1% TFA). The solvents were evaporated under reduced pressure and the residue was dried under high vacuum. This gave 15.3 mg (29% over 2 steps) of the regioisomeric protected intermediates.

5

LC-MS (Method 6):  $R_t$  = 12.26 min and 12.30 min; MS (ESIpos):  $m/z$  = 1254 (M+H)<sup>+</sup>.

In the last step, 15.3 mg (0.01 mmol) of this intermediate were dissolved in 2 ml of 2,2,2-trifluoroethanol. 6.1 ml (0.05 mmol) of zinc chloride were added, and the reaction was stirred at 50°C for 2 h. 13.1 mg (0.05 mmol) of ethylenediamine-N,N,N',N'-tetraacetic acid were then added, and the product was purified by preparative HPLC. Concentration of the appropriate fractions and lyophilization of the residue from acetonitrile/water gave 11.9 mg (79.5%) of the title compound as a regioisomer mixture.

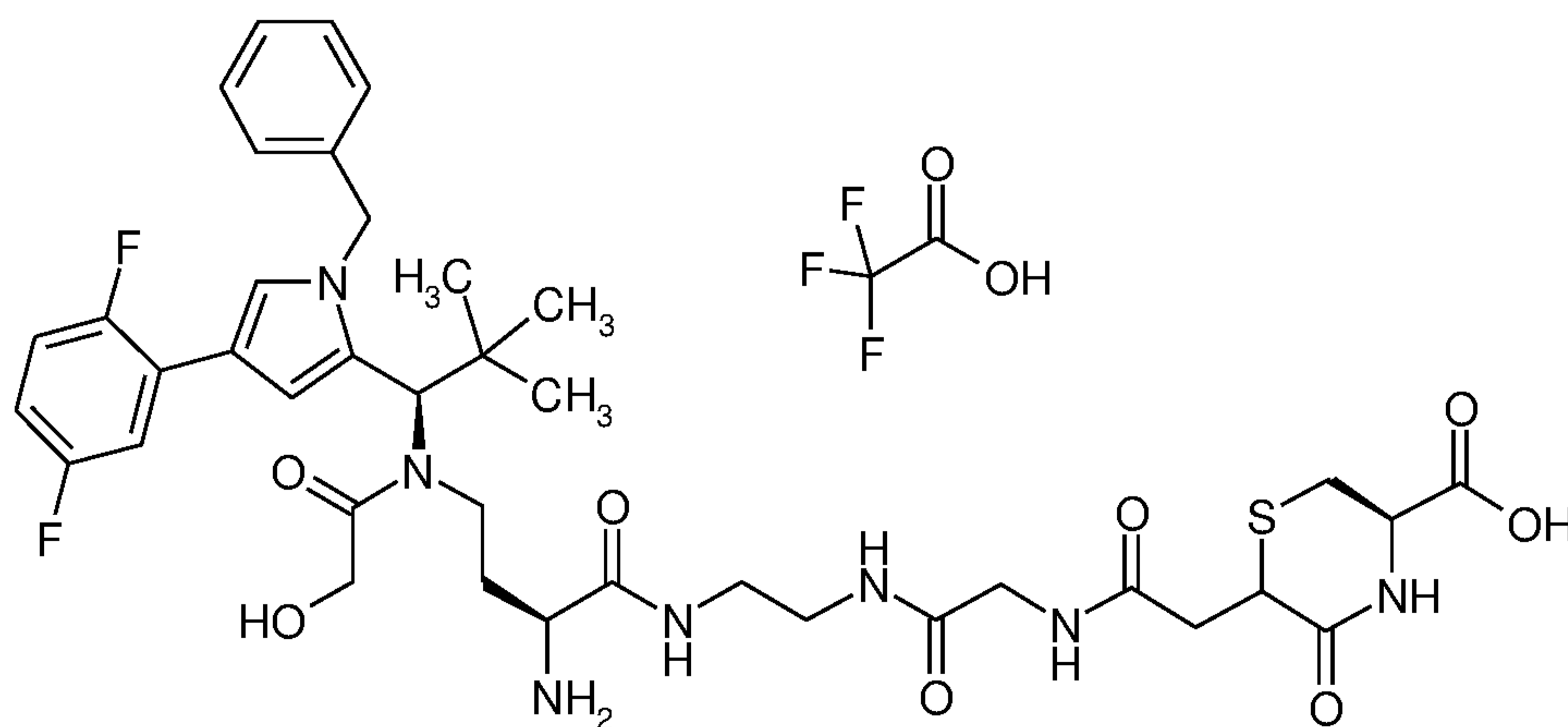
15

LC-MS (Method 1):  $R_t$  = 0.85 min; MS (ESIpos):  $m/z$  = 1110 (M+H)<sup>+</sup>.

## 20 Example 256

(3R)-6-{(11S,15R)-11-Amino-15-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-14-glycoloyl-16,16-dimethyl-2,5,10-trioxo-3,6,9,14-tetraazaheptadec-1-yl}-5-oxothiomorpholine-3-carboxylic acid / trifluoroacetic acid (1:1)

25



30 4 mg (0.004 mmol) of the compound from Example 135 were dissolved

in 4 ml of THF/water, and 48 µl of a 2-molar aqueous lithium hydroxide solution were added. The reaction was stirred at RT for 1 h and then concentrated and purified by preparative HPLC. Combination, concentration and lyophilization of the appropriate  
5 fractions from acetonitrile/water gave 2.4 mg (60% of theory) of the title compound.

LC-MS (Method 1):  $R_t = 0.86$  min; MS (EIpos):  $m/z = 814$   $[M+H]^+$ .

10 **C: Assessment of biological efficacy**

The biological activity of the compounds according to the invention can be shown in the assays described below:

15 m. C-1a Determination of the cytotoxic effects of the ADCs directed against TWEAKR

The analysis of the cytotoxic effects of the anti-TWEAKR-ADCs was carried out with various cell lines:

20

NCI-H292: human mucoepidermoid lung carcinoma cells, ATCC-CRL-1848, standard medium: RPMI 1640 (Biochrom; #FG1215, stab. glutamine) + 10% FCS (Biochrom; #S0415), TWEAKR-positive, EGFR-positive.

25

BxPC3: human pancreas carcinoma cells, ATCC-CRL-1687, standard medium: RPMI 1640 (Biochrom; #FG1215, stab. glutamine) + 10% FCS (Biochrom; #S0415), TWEAKR-positive.

30

KPL4: human breast carcinoma cells, standard medium: RPMI 1640 + GlutaMAX I + 10% FBS, cell bank, Bayer Pharma AG (identity checked and confirmed on 19.7.2012 at DSMZ), Berlin, ERBB2-positive.

35

The cells are cultivated by a standard method, as indicated in the American Tissue Type Collection (ATCC) for the respective cell lines.

## MTT assay

The test is carried out by detaching the cells with a solution of Accutase in PBS (Biochrom AG #L2143), pelleting, resuspending  
5 in culture medium, counting and sowing into a 96-well culture plate with white bottom (Costar #3610) (1000-2000 cells in 100  $\mu$ l/well depending on the cell used) and incubating in an incubator at 37°C and 5% carbon dioxide. After 48 h, the antibody drug conjugates are added in 10  $\mu$ l of culture medium in  
10 concentrations of from  $10^{-5}$ M to  $10^{-13}$ M to the cells (triplicates) and incubated in an incubator at 37°C and 5% carbon dioxide. After 72 h, the proliferation is detected using the MTT assay (ATCC, Manassas, Virginia, USA, catalogue No. 30-1010K). At the end of the selected incubation time, the MTT reagent is incubated  
15 with the cells for 4 h, followed by lysis of the cells overnight by addition of the detergent. The dye formed was detected at 570 nm. The proliferation of cells which were not treated with test substance but were otherwise identically treated was defined as the 100% figure.

20

## CTG assay

The cells were cultivated according to the standard method using the growth media listed under C-1. The test was carried out by  
25 detaching the cells with a solution of trypsin (0.05%) and EDTA (0.02%) in PBS (Biochrom AG #L2143), pelleting, resuspending in culture medium, counting and sowing into a 96-well culture plate with white bottom (Costar #3610) (at 75  $\mu$ l/well, the following cell numbers are per well: NCI-H292: 2500 cells/well, BxPC3 2500  
30 cells/well) and incubating in an incubator at 37°C and 5% carbon dioxide. After 24 h, the antibody drug conjugates were added in 25  $\mu$ l of culture medium (concentrated four-fold) to the cells to give final antibody drug conjugate concentrations of  $3 \times 10^{-7}$  M to  $3 \times 10^{-11}$  M on the cells (triplicates). The cells were then  
35 incubated in an incubator at 37°C and 5% carbon dioxide. On a parallel plate, the cell activity at the start of the active compound treatment (day 0) was determined using the Cell Titer Glow (CTG) luminescent cell viability assay (Promega #G7573 and

#G7571). To this end, per cell batch 100  $\mu$ l of the substrate were added, the plates were then covered with aluminium foil, shaken on the plate shaker at 180 rpm for 2 minutes, allowed to stand on the laboratory bench for 8 minutes and then measured using a luminometer (Victor X2, Perkin Elmer). The substrate detects the ATP content of the living cells generating a luminescence signal whose intensity is directly proportional to the viability of the cells. After 72 h of incubation with the antibody drug conjugates, the viability of these cells was then also determined using the Cell Titer Glow luminescent cell viability assay as described above. From the data measured, the IC<sub>50</sub> of the growth inhibition was calculated in comparison to day 0 using the DRC (Dose Response Curve) analysis spreadsheets and a 4-parameter fit. The DRC analysis spreadsheet is a biobook spreadsheet developed by Bayer Pharma AG and Bayer Business Services on the IDBS E-WorkBook Suite platform (IDBS: ID Business Solutions Ltd., Guildford, UK).

The Tables 1a, 1b and 1c list the IC<sub>50</sub> values of representative working examples using the anti-TWEAKR antibody from this assay:

Table 1a

<b>Example</b>	<b>BxPC3 IC<sub>50</sub> [M] CTG</b>	<b>NCI-H292 IC<sub>50</sub> [M] CTG</b>
58b	5.32E-09	5.76E-09
82b	4.26E-09	6.00E-07
83b	4.68E-09	1.97E-08
84b	1.00E-09	8.58E-10
85b	2.03E-09	8.84E-10
86b	2.96E-09	8.68E-09
87b	2.16E-09	1.33E-09
88b	2.0E-08	8.34E-09
89b	1.87E-09	6.00E-07
90b	2.12E-08	8.49E-09
91b	9.98E-09	3.07E-09

Table 1b

Example	BxPC3 IC <sub>50</sub> [M] CTG	NCI-H292 IC <sub>50</sub> [M] CTG	Example	BxPC3 IC <sub>50</sub> [M] CTG	NCI-H292 IC <sub>50</sub> [M] CTG
104b	5.88E-10	1.44E-10	126b	6.00E-07	6.00E-07
106b	2.43E-09	6.00E-07	127b	3.31E-08	1.94E-09
107b	4.19E-09	4.52E-09	129b	6.00E-07	6.00E-07
109b	9.26E-08	1.61E-07	142b	1.41E-09	6.92E-10
112b	6.00E-07	6.00E-07	143b	1.04E-07	4.33E-09
113b	2.48E-08	7.36E-08	144b	1.17E-08	8.29E-08
115b	2.43E-07	6.00E-07	146b	4.85E-08	1.05E-08
116b	1.01E-08	6.00E-07			
117b	6.00E-07	6.00E-07			
118b	6.00E-07	6.00E-07	153b	1.44E-09	1.17E-08
119b	8.90E-09	9.14E-10	155b	9.44E-10	5.09E-10
121b	6.00E-07	6.00E-07	156b	1.39E-09	6.00E-07
122b	6.00E-07	1.03E-08			
124b	6.00E-07	6.00E-07			
125b	2.99E-08	3.58E-09			

Table 1c

<b>Example</b>	<b>BxPC3 IC<sub>50</sub> [M] CTG</b>	<b>NCI-H292 IC<sub>50</sub> [M] CTG</b>	<b>Example</b>	<b>BxPC3 IC<sub>50</sub> [M] CTG</b>	<b>NCI-H292 IC<sub>50</sub> [M] CTG</b>
163b	3.80E-09	1.21E-08	211b	5.94E-09	3.75E-09
164b	8.54E-09	4.10E-09	212b	3.65E-08	3.31E-08
165b	9.14E-10	8.95E-10	213b	6.02E-09	2.16E-09
166b	3.72E-09	9.27E-08	214b	3.86E-09	3.82E-09
168b	9.12E-10	5.78E-09	216b	7.15E-10	5.28E-10
169b	2.04E-09	3.19E-09	217b	5.60E-09	1.33E-08
171b	6.00E-07	4.25E-09	234b	8.17E-09	1.82E-09
173b	1.39E-09	1.07E-09	236b	1.93E-07	2.38E-07
174b	5.27E-09	6.00E-07	237b	1.05E-09	3.74E-10
175b	9.58E-09	6.00E-07	238b	6.00E-07	6.03E-08
176b	3.84E-09	6.00E-07	239b	2.52E-08	7.04E-09
177b	2.00E-08	5.11E-08	240b	1.13E-08	8.69E-09
178b	1.78E-08	3.78E-08	241b	1.10E-08	1.15E-09
179b	2.29E-08	6.00E-07	242b	8.54E-09	1.21E-09
180b	4.39E-09	2.34E-09	243b	9.99E-09	1.18E-09
192b	6.29E-09	6.00E-07	244b	4.63E-08	3.28E-08
193b	1.85E-09	8.98E-09	245b	1.05E-08	1.30E-09
194b	3.96E-	2.92E-09	247b	6.19E-10	3.63E-10



	09				
195b	4.07E-09	2.95E-09	248b	1.53E-08	1.28E-09
196b	5.73E-09	1.58E-09	254b	3.46E-09	3.37E-10
204b	6.42E-09	1.25E-09			
205b	1.45E-09	5.17E-08			
206b	3.89E-09	3.01E-08			
207b	1.36E-09	3.33E-10			
208b	2.55E-09	6.98E-10			
209b	7.07E-10	1.42E-09			
210b	8.53E-10	1.89E-08			

The activity data reported relate to the working examples described in the present experimental section, with the drug/mAB ratios indicated. The values may possibly deviate for different drug/mAB ratios. The IC<sub>50</sub> values are means of several independent experiments or individual values. The action of the TWEAKR antibody drug conjugates was selective for the respective isotype control comprising the respective linker and toxophor.

Table 2 below lists the IC<sub>50</sub> values of representative working examples with the cetuximab antibody from the MTT assay:

Table 2

<b>Example</b>	<b>NCI-H292 IC<sub>50</sub> [M] MTT</b>
35a	5.15E-09
36a	1.65E-10

37a	1.31E-10
82a	4.37E-09
83a	1.07E-09
84a	9.43E-10
85a	3.40E-10
86a	1.05E-08
87a	4.34E-09
88a	1.32E-09
89a	3.78E-09
90a	2.35E-09
91a	1.45E-10
96	2.09E-09
98	1.65E-09
99	8.51E-10
102	3.29E-08
104a	1.85E-10
104h	4.60E-10
104i	1.47E-09
106a	7.89E-11
107a	4.02E-11
109a	2.63E-10
112a	1.80E-10
113a	1.97E-10
115a	2.39E-10
116a	8.99E-11
117a	1.09E-10
118a	8.33E-09
119a	8.39E-10
121a	8.78E-11
122a	2.01E-10
124a	5.75E-11
125a	9.11E-11
126a	8.38E-09
127a	5.97E-10
129a	1.70E-09
131	5.00E-07
132	5.00E-07

133	1.55E-07
134	5.00E-07
135	5.00E-07
136	5.00E-07
137	5.00E-07
138	5.00E-07
139	1.34E-07
141	1.61E-08
142a	4.95E-11
142h	6.50E-10
142i	3.62E-09
143a	1.21E-10
146a	6.44E-11
153a	2.40E-12
155a	7.34E-11
156a	3.74E-11
157	5.00E-07
158	2.22E-08
160	5.00E-07
161	3.55E-08
162	1.21E-07
163a	7.24E-10
163h	6.03E-10
164a	1.29E-10
165a	3.26E-11
166a	1.08E-10
168a	1.23E-10
168h	1.57E-10
169a	1.86E-10
171a	5.05E-10
173a	3.16E-11
174a	5.41E-10
175a	4.72E-10
176a	2.19E-10
177a	1.87E-11
178a	1.99E-10
179a	1.51E-09

180a	2.32E-10
181	7.72E-09
182	1.20E-09
183	3.29E-08
184	2.78E-08
185	1.20E-08
186	9.29E-09
187	1.98E-09
188	9.28E-10
189	8.76E-09
190	5.42E-09
191	9.36E-10
192a	5.18E-10
193a	3.10E-10
193h	2.52E-10
194a	6.17E-11
195a	8.27E-10
196a	3.67E-09
197	7.35E-09
198	1.95E-08
199	3.23E-07
199-2	
200	1.40E-07
201	>5.00E-07
202	4.22E-07
203	>5.00E-07
204a	3.60E-09
205a	6.64E-10
206a	1.37E-09
207a	1.05E-10
207h	2.39E-09
207i	6.87E-10
208a	1.16E-08
208i	8.22E-09
209a	2.02E-11
209h	1.35E-09
209i	1.37E-09

210a	1.30E-11
211a	4.71E-10
212a	4.43E-10
213a	1.95E-11
214a	6.83E-10
215a	5.20E-10
215h	5.18E-10
215i	1.58E-08
216a	1.12E-10
217a	1.27E-09
218a	3.44E-10
218h	9.42E-10
218i	1.37E-08
219	3.48E-09
220	1.97E-10
221	1.00E-10
222	3.30E-09
223	7.70E-12
224	3.37E-12
225	1.51E-09
226	2.22E-07
227	1.82E-09
228	5.36E-09
229	5.00E-07
230	1.21E-11
231	9.18E-10
232	1.55E-08
233	9.63E-10
234a	2.10E-09
234h	7.28E-09
234i	6.97E-09
235a	4.6E-08
235h	3.04E-10
235i	1.45E-08
236a	5.98E-09
237a	1.51E-09
237i	5.94E-09

238a	3.49E-10
239a	7.78E-11
239h	1.65E-09
239i	6.48E-09
240a	3.66E-11
241a	5.91E-11
241i	5.80E-10
242a	8.26E-11
243a	3.37E-10
243i	3.22E-09
244a	3.70E-10
244i	4.50E-09
245a	5.30E-11
245i	2.50E-08
246	9.38E-09
247a	<1.00E-10
247i	1.53E-10
248a	
249	2.66E-08
252	
254a	
254i	

The activity data reported relate to the working examples described in the present experimental section, with the drug/mAB ratios indicated. The values may possibly deviate for different drug/mAB ratios. The IC50 values are means of several independent experiments or individual values. The action of the cetuximab antibody drug conjugates was selective for the respective isotype control comprising the respective linker and toxophor.

5

Table 3 below lists the IC50 values of representative working examples with the trastuzumab antibody from the MTT assay:

Table 3

15

<b>Example</b>	<b>KPL4</b>
----------------	-------------

	<b>IC<sub>50</sub> [M]</b> <b>MTT</b>
82e	9.53E-08
83e	5.81E-09
104e	2.09E-10
106e	5.00E-05
118e	5.0E-07
126e	5.0E-07
127e	1.22E-09
129e	5.0E-07
142e	1.53E-07
143e	5.0E-07
146e	5.0E-07
156e	5.00E-07
164e	5.85E-10
165e	7.44E-11
166e	5.00E-07
168e	3.50E-10
169e	6.27E-08
173e	8.07E-10
175e	5.00E-07
176e	2.55E-08
179e	1.21E-07
180e	4.13E-10
192e	5.00E-07
193e	1.23E-09
194e	1.55E-08
195e	7.27E-10
196e	5.49E-10
204e	1.66E-10
205e	3.24E-10
206e	1.13E-08
207e	1.55E-10
210e	5.00E-07
211e	1.06E-07
212e	5.03E-09
213e	5.00E-07

214e	8.48E-09
215e	1.89E-10
216e	9.64E-11
217e	1.47E-10
218e	1.67E-09
235e	8.0E-08
236e	8.40E-08
238e	2.30E-08
240e	1.90E-11
241e	1.90E-10
242e	6.288E-11
243e	2.92E-10
244e	8.30E-09
245e	4.48E-10
247e	1.43E-10
254e	1.09E-10

The activity data reported relate to the working examples described in the present experimental section, with the drug/mAB ratios indicated. The values may possibly deviate for different drug/mAB ratios. The IC50 values are means of several independent experiments or individual values. The action of the trastuzumab antibody drug conjugates was selective for the respective isotype control comprising the respective linker and toxophor.

5

#### C-1b Determination of the inhibition of the kinesin spindle protein KSP/ Eg5 by selected examples

10

The motor domain of the human kinesin spindle protein KSP /Eg5 (from tebu-bio/ Cytoskeleton Inc, No. 027EG01-XL) is incubated at a concentration of 10 nM with 50 µg/ml taxol- (from Sigma No. T7191-5MG) stabilized microtubuli (bovine or porcine, from tebu-bio/ Cytoskeleton Inc) for 5 min at RT in 15 mM PIPES, pH 6.8 (5 mM MgCl<sub>2</sub> and 10mM DTT, from Sigma). The freshly prepared mixture is aliquoted into a 384-well MTP. The inhibitors to be examined at concentrations of 1.0 x 10<sup>-6</sup> M to 1.0 x 10<sup>-13</sup> M and ATP (final concentration 500 µM, from Sigma) are then added.

15

20



Incubation is at RT for 2 h. ATPase activity is detected by detecting the inorganic phosphate formed using malachite green (from Biomol). After additon of the reagent, the assay was incubated at RT for 50 minutes prior to detection of the absorption at a wavelength of 620 nm. Monastrol and Ispinesib (from Adooq A10486) are used as positive control. The individual data of the dose-activity curve are eight-fold determinations. The IC50 values are means of three independent experiments. The 100% control was the sample which had not been treated with inhibitors.

Table 4 below summarizes the IC50 values of representative working examples from the assay described: In an exemplary manner, they confirm the high potency at the target of the toxophors and ADC methods described.

Table 4

<b>Examples</b>	<b>KSP assay IC<sub>50</sub> [M]</b>
96	7.80E-09
98	1.80E-09
99	n.d.
102	4.70E-09
131	1.24E-09
132	1.65E-09
133	4.13E-09
134	4.35E-09
135	2.04E-09
136	3.28E-10
137	7.78E-10
138	1.01E-9
139	1.82E-09
141	9.02E-10
157	9.61E-09
158	6.20E-10
160	6.81E-10
161	5.17E-09

162	2.58E-09
181	7.90E-09
182	1.04E-08
183	6.21E-09
184	1.43E-08
185	1.80E-08
186	2.53E-08
187	1.41E-08
188	1.01E-08
189	1.12E-08
190	9.61E-09
191	1.26E-08
197	4.17E-09
198	1.07E-08
199	6.00E-09
199-2	9.09E-10
200	1.08E-10
201	1.79E-10
202	2.56E-10
203	2.30E-10
219	1.13E-09
220	1.45E-09
221	1.12E-10
222	1.58E-09
223	2.89E-10
224	1.96E-10
225	4.25E-10
226	1.49E-09
227	1.78E-09
228	3.07E-09
229	4.44E-09
230	3.41E-09
231	1.67E-09
232	2.20E-09
233	3.88E-09
246	2.69E-09
249	9.15E-10

250	3.26E-09
251	2.71E-10
252	4.57E-10
256	1.64E-08

The activity data reported relate to the working examples described in the present experimental section.

## 5 C-2 Internalisation assay

Internalisation is a key process which enables specific and efficient provision of the cytotoxic payload in antigen-expressing cancer cells via antibody drug conjugates (ADC). This process is monitored via fluorescent labelling of specific TWEAKR antibodies and an isotype control antibody (M014). First, the fluorescent dye is conjugated to lysines of the antibody. Conjugation is carried out using a two-fold molar excess of CypHer 5E mono NHS ester (Batch 357392, GE Healthcare) at pH 8.3. After the coupling, the reaction mixture is dialysed at 4°C (sSlide-A-Lyser Dialysis Cassettes MWCD 10kD, from Pierce) overnight to remove excess dye and to adjust the pH, and the protein solution is then concentrated (VIVASPIN 500, from Sartorius stedim biotec). Determination of the dye load of the antibody is by spectrophotometric analysis (NanoDrop) and subsequent calculation ( $D: P = A_{\text{dye}} \epsilon_{\text{protein}} : (A_{280} - 0.16A_{\text{dye}}) \epsilon_{\text{dye}}$ ). The dye load of the TWEAKR antibody examined here and the isotype control were of a comparable order. In cell binding assays, it is confirmed that the conjugation did not lead to a change in the affinity of the antibody.

The labelled antibodies are used for the internalisation assay. Prior to the start of the treatment, the cells ( $2 \times 10^4$ /well) are sown in 100 µl medium in a 96-well MTP (fat, black, clear bottom No 4308776, from Applied Biosystems). After 18 h of incubation at 37°C/5%CO<sub>2</sub>, the medium is replaced and labelled anti-TWEAKR antibodies are added in different concentrations (10, 5, 2.5, 1, 0.1 µg/ml). The same treatment protocol is applied to the labelled isotype control (negative control). The

chosen incubation times are 0, 0.25 h, 0.5 h, 1 h, 1.5 h, 2 h, 3 h, 6 h and 24 h. The fluorescence measurement is carried out using the InCellanalyser 1000 (from GE Healthcare). This was followed by kinetic evaluation via measurement of the parameters  
5 granule counts/cell and totale granule intensity/cell.

Following binding to the TWEAKR, TWEAKR antibodies were examined for the internalisation ability. For this purpose, cells with different TWEAKR expression levels were chosen. A target-  
10 mediated specific internalisation was observed with the TWEAKR antibodies, whereas the isotype control showed no internalisation.

### C-3 In vitro tests for determining cell permeability

15

The cell permeability of a substance can be investigated by means of in vitro testing in a flux assay using Caco-2 cells [M.D. Troutman and D.R. Thakker, *Pharm. Res.* 20 (8), 1210-1224 (2003)]. For this purpose, the cells were cultured for 15-16  
20 days on 24-well filter plates. For the determination of permeation, the respective working example was applied in a HEPES buffer to the cells either apically (A) or basally (B) and incubated for 2 hours. After 0 hours and after 2 hours, samples were taken from the cis and trans compartments. The samples were  
25 separated by HPLC (Agilent 1200, Böblingen, Germany) using reverse phase columns. The HPLC system was coupled via a Turbo Ion Spray Interface to a Triple Quadropol mass spectrometer API 4000 (Applied Biosystems Applera, Darmstadt, Germany). The permeability was evaluated on the basis of a Papp value, which  
30 was calculated using the formula published by Schwab et al. [D. Schwab et al., *J. Med. Chem.* 46, 1716-1725 (2003)]. A substance was classified as actively transported when the ratio of  $P_{app}$  (B-A) to  $P_{app}$  (A-B) (efflux ratio) was  $>2$  or  $<0.5$ .

35 Of critical importance for toxophores which are released intracellularly is the permeability from B to A [ $P_{app}$  (B-A)] and the ratio of  $P_{app}$  (B-A) to  $P_{app}$  (A-B) (efflux ratio): the lower this permeability, the slower the active and passive transport

processes of the substance through the monolayer of Caco-2 cells. If additionally the efflux ratio does not indicate any active transport, the substance may, following intracellular release, remain longer in the cell. Hence, there is also more  
 5 time available for interaction with the biochemical target (in this case: kinesin spindle protein, KSP / Eg5).

Table 5 below sets out permeability data for representative working examples from this assay:

10

Table 5

Working Example	$P_{app}$ (B-A) [nm/s]	Efflux ratio
92	4	3
96	285	16
98	213	16
99	303	8.7
131	9	10
132	1.6	1
133	4.3	6
134	2.9	0.5
135	7.8	4
136	9.1	8.5
137	17	19
138	1.7	1.6
139	21	7.7
141	30	35
157	7.7	7.2
158	5.6	2.2
160	1.6	1
161	1.6	2
162	2	2.9
197	93	81
198	519	34
199	4.8	6.4
226	3	0.2
232	49	14

246	1.4	1.3
251	21	19
252	20	26

C-4 In vitro tests for determining the substrate properties for P-glycoprotein (P-gp)

5 Many tumour cells express transporter proteins for drugs, and this frequently accompanies the development of resistance towards cytostatics. Substances which are not substrates of such transporter proteins, such as P-glycoprotein (P-gp) or BCRP, for example, could therefore exhibit an improved activity profile.

10

The substrate properties of a substance for P-gp (ABCB1) were determined by means of a flux assay using LLC-PK1 cells which overexpress P-gp (L-MDR1 cells) [A.H. Schinkel *et al.*, *J. Clin. Invest.* 96, 1698-1705 (1995)]. For this purpose, the LLC-PK1  
15 cells or L MDR1 cells were cultured on 96-well filter plates for 3-4 days. For determination of the permeation, the respective test substance, alone or in the presence of an inhibitor (such as ivermectin or verapamil, for example), was applied in a HEPES buffer to the cells either apically (A) or basally (B) and  
20 incubated for 2 hours. After 0 hours and after 2 hours, samples were taken from the cis and trans compartments. The samples were separated by HPLC using reverse phase columns. The HPLC system was coupled via a Turbo Ion Spray Interface to a Triple Quadropol mass spectrometer API 3000 (Applied Biosystems Applera,  
25 Darmstadt, Germany). The permeability was evaluated on the basis of a Papp value, which was calculated using the formula published by Schwab *et al.* [D. Schwab *et al.*, *J. Med. Chem.* 46, 1716-1725 (2003)]. A substance was classified as P-gp substrate when the efflux ratio of  $P_{app} (B-A)$  to  $P_{app} (A-B)$  was  $>2$ .

30

As further criteria for the evaluation of the P-gp substrate properties, the efflux ratios in L-MDR1 and LLC-PK1 cells or the efflux ratio in the presence or absence of an inhibitor may be compared. If these values differ by a factor of more than 2, the  
35 substance in question is a P-gp substrate.

C-5 PharmacokineticsC5a: Identification of the ADC metabolites after internalisation  
5 in vitro

Description of the method:

Internalisation studies with immunoconjugates are carried out  
10 to analyse metabolites formed intracellularly. To this end,  
human lung tumour cells NCI H292 ( $3 \times 10^5$ /well) are sown in 6-well  
plates and incubated overnight ( $37^\circ\text{C}$ , 5%  $\text{CO}_2$ ). The cells are  
treated with 10  $\mu\text{g}/\text{ml}$  of the ADC to be examined. Internalisation  
is carried out at  $37^\circ\text{C}$  and 5%  $\text{CO}_2$ . At various time points (0, 4,  
15 24, 48, 72 h), cell samples are taken for further analysis.  
First, the supernatants (about 5 ml) are harvested and, after  
centrifugation (2 min, RT, 1000 rpm Heraeus Variofuge 3.0R),  
stored at  $-80^\circ\text{C}$ . The cells are washed with PBS and detached with  
Accutase, and the cell number is determined. After another  
20 washing, a defined number of cells ( $2 \times 10^5$ ) is treated with 100  
 $\mu\text{l}$  of lysis buffer (Mammalian Cell Lysis Kit (Sigma MCL1) and  
incubated with continuous shaking (Thermomixer, 15 min,  $4^\circ\text{C}$ , 650  
rpm) in Protein LoBind tubes (Eppendorf Cat. No. 0030 108.116).  
After the incubation, the lysate is centrifuged (10 min,  $4^\circ\text{C}$ ,  
25 12000 g, Eppendorf 5415R) and the supernatant is harvested. The  
supernatant obtained is stored at  $-80^\circ\text{C}$ . All samples are then  
analysed as follows.

Measurement of the compounds in the culture supernatant or cell  
30 lysate is carried out after precipitation of the proteins with  
methanol or acetonitrile by high-pressure liquid chromatography  
(HPLC) coupled to a triple-quadrupole mass spectrometer (MS).

For work-up of 50  $\mu\text{l}$  of culture supernatant/cell lysate, 150  $\mu\text{l}$   
35 of precipitation reagent (generally acetonitrile) are added and  
the mixture is shaken for 10 seconds. The precipitation reagent  
contains an internal standard (ISTD) in a suitable concentration  
(generally in the range of 20-100 ng/ml). After 3 minutes of

centrifugation at 16000 g, the supernatant is transferred into an autosampler vial, made up with 500 µl of a buffer suitable for the mobile phase and shaken again.

- 5 The two matrix samples are then measured using the HPLC-coupled triple-quadrupol mass spectrometer API6500 from AB SCIEX Deutschland GmbH.

10 For calibration, concentrations of 0.5–2000 µg/l are added to plasma samples. The detection limit (LOQ) is about 2 µg/l. The linear range extends from 2 to 1000 µg/l.

15 For calibration of the tumour samples, concentrations of 0.5–200 µg/l are added to the supernatant of untreated tumours. The detection limit is 4 µg/l. The linear range extends from 4 to 200 µg/l.

Quality controls for testing validity contain 5 and 50 µg/l.

- 20 NCI-H292 cells were incubated with in each case 10 µg/ml of the ADCs from Examples 104b, 119b, 155b, 165b and 173b. After 72 h, the cells were washed with PBS, lysed and deep-frozen (-80°C). Using the method described above, the cell lysates and cell culture supernatants were worked up, and after extraction the  
25 following metabolites were identified and quantified:

Incubated ADC Example	Isolated metabolite	Metabolite concentration in the cell lysate [µg/l]	Metabolite concentration in the supernatant [µg/l]
104b	199	11.6	< 1
104b	198	< 2	< 2
104b	135	< 0.2	< 0.2
104b	197	< 1	< 1
104b	256	<0.2	<0.2
119b	158	14	< 0.2
155b	162	10	< 0.2



165b	162	8.8	< 0.2
173b	161	18	< 0.2

NCI-H292 cells were incubated with in each case 10 µg/ml of the ADCs from Examples 179a, 226a, 85b and 208b. After 72 h, the cells were washed with PBS, lysed and deep-frozen (-80°C). Using the method described above, the cell lysates and cell culture supernatants were worked up, and after extraction the following metabolites were identified and quantified:

Incubated ADC Example	Isolated metabolite	Metabolite concentration in the cell lysate [µg/l]	Metabolite concentration in the supernatant [µg/l]
179a	249	< 1	5.6
226a	226	<1	9.1
226a	255	<1	2.5
85b	139	1.5	3.2
85b	250	7.4	1.8
208b	199	13.5	2.3
208b	198	<1	<1

#### 10 C5b: Identification of the ADC metabolites *in vivo*

After i.v. administration of 3-30 mg/kg of different ADCs, the plasma and tumour concentrations of the ADCs and any metabolites occurring can be measured, and the pharmacokinetic parameters such as clearance (CL), area under the curve (AUC) and half-times ( $t_{1/2}$ ) can be calculated.

#### Analysis for quantification of any metabolites occurring

20 Measurement of the compounds in plasma and tumour is carried out after precipitation of the proteins with methanol or acetonitrile by high-pressure liquid chromatography (HPLC) coupled to a triple-quadrupole mass spectrometer (MS).

For work-up of 50 µl of plasma, 250 µl of precipitation reagent (generally acetonitrile) are added and the mixture is shaken for 10 seconds. The precipitation reagent contains an internal standard (ISTD) in a suitable concentration (generally in the  
5 range of 20-100 ng/ml). After 3 minutes of centrifugation at 16000 g, the supernatant is transferred into an autosampler vial, made up with 500 µl of a buffer suitable for the mobile phase and shaken again.

10 During the work-up of a tumour, the latter is treated with 3 times the amount of extraction buffer. The extraction buffer contains 50 ml of Tissue Protein Extraction Reagent (Pierce, Rockford, IL), two pellets of Complete-Protease-Inhibitor-Cocktail (Roche Diagnostics GmbH, Mannheim, Germany) and  
15 phenylmethylsulphonyl fluoride (Sigma, St. Louis, MO) in a final concentration of 1 mM. The sample is homogenized twice for 20 minutes in a Tissuelyser II (Qiagen), at maximum stroke number. 50 µl of the homogenate are transferred into an autosampler vial and made up with 150 µl of methanol including ISTD. After 3  
20 minutes of centrifugation at 16000 g, 10 µl of the supernatant are made up with 180 µl of a buffer suitable for the mobile phase and shaken again. The tumour sample is then ready for measuring.

25 The two matrix samples are then measured using the HPLC-coupled triple-quadrupol mass spectrometer API6500 from AB SCIEX Deutschland GmbH.

For calibration, concentrations of 0.5 - 2000 µg/l are added to  
30 plasma samples. The detection limit (LOQ) is about 2 µg/l. The linear range extends from 2 to 1000 µg/l.

For calibration of the tumour samples, concentrations of 0.5 - 2000 µg/l are added to the supernatant of untreated tumours. The  
35 detection limit is 5 µg/l. The linear range extends from 5 to 200 µg/l.

Quality controls for testing validity contain 5 and 50 µg/l, in

plasma additionally 500 µg/l.

Following administration of 10 mg/kg of the ADCs from Examples 119b and 104b in the control groups from the xenograft models with NCI-H292, the mice were sacrificed after 24 h, blood was removed and the tumours were isolated. Using the method described above, the plasma and tumour samples were worked up, and after extraction the following metabolites were identified and quantified:

10

ADC administered (Example Number)	Isolated metabolite (Example Number)	Metabolite concentration in the tumour [µg/l]	Metabolite concentration in the plasma [µg/l]
119b	158	155	< 5
104b	199	147	< 5
104b	198	27.4	< 5
104b	135	5.23	< 5
104b	197	7.75	< 5

#### Analysis for quantification of the antibodies used

The antibody part of the ADCs was determined using a ligand binding assay (ELISA) as total IgG concentration in plasma samples and tumour lysates. Here, the sandwich ELISA format was used. This ELISA had been qualified and validated for the determination in plasma and tumour samples. The ELISA plates were coated with anti-human goat IgG Fc antibodies. After incubation with the sample, the plates were washed and incubated with a detector conjugate of simian anti-human IgG(H+L) antibody and horseradish peroxidase (HRP). After a further washing step, the HRP substrate was added to OPD and the colour development was monitored via absorption at 490 nm. Standard samples having a known IgG concentration were fitted using a 4-parameter equation. Within the lower (LLOQ) and upper (ULOQ) quantification limits, the unknown concentrations were determined by interpolation.

C-6 Activity test *in vivo*

The activity of the conjugates according to the invention was tested, for example, using xenograft models. The person skilled in the art is familiar with methods in the prior art which allow the activity of the compounds according to the invention to be tested (see, for example, WO 2005/081711; Polson et al., Cancer Res. 2009 Mar 15;69(6):2358-64). To this end, a tumour cell line expressing the target molecule of the binder was implanted into rodents (for example mice). A conjugate according to the invention, an isotype control conjugate, a control antibody or isotonic saline was then administered to the implant animals. The administration took place once or more than once. Following an incubation time of several days, the size of the tumour was determined by comparing conjugate-treated animals and the control group. The conjugate-treated animals displayed a smaller tumour size.

**C-6a. Growth inhibition / regression of experimental tumours in the mouse**

Human tumour cells expressing the antigen for the antibody drug conjugate are inoculated subcutaneously into the flank of immunosuppressed mice, for example NMRi nude or SCID mice. 1-10 million cells are detached from the cell culture, centrifuged and resuspended in medium or medium / matrigel. The cell suspension is injected under the skin of the mouse.

Within a few days, a tumour grows. Treatment is commenced after the tumour is established, at a tumour size of approximately 40 mm<sup>2</sup>. To examine the effect on larger tumours, treatment may be initiated only at a tumour size of 50-100 mm<sup>2</sup>.

Treatment with ADCs is carried out via the intravenous route into the tail vein of the mouse. The ADC is administered in a volume of 5 ml/kg.

The treatment protocol depends on the pharmacokinetics of the antibody. As standard, treatment takes place three times in succession every fourth day. For a quick assessment, a protocol with a single treatment may be employed. However, the treatment  
5 may also be continued, or a second cycle of three treatment days may follow at a later time.

As standard, 8 animals are used per treatment group. In addition to the groups to which the active substances are administered,  
10 one group is treated as control group only with the buffer, according to the same protocol.

During the experiment, the tumour area is measured regularly in two dimensions (length / width) using a caliper. The tumour  
15 area is determined as length x width. The ratio of the mean tumour area of the treatment group to that of the control group is stated as T/C area.

When after the end of the treatment all groups of the experiment  
20 are terminated at the same time, the tumours can be removed and weighed. The ratio of the mean tumour weights of the treatment group to that of the control group is stated as T/C weight.

**C-6c. Efficacy of the anti-EGFR antibody drug conjugates in  
25 the NCI-H292 tumour model on single treatment**

1 million NCI-H292 cells are inoculated subcutaneously into the flank of female NMRI-nude mice (Janvier). At a tumour size of ~37 mm<sup>2</sup> on day 9, the animals are treated once with a dose of 3  
30 mg/kg of ADC or 2.5 mg/kg toxophor A (N-(3-aminopropyl)-N-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-imidazol-2-yl]-2,2-dimethylpropyl)-2-hydroxyacetamide) or toxophor B (N-[(3R)-3-amino-4-fluorobutyl]-N-((1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-imidazol-2-yl]-2,2-dimethylpropyl)-2-  
35 hydroxyacetamide). After the treatment, the tumour growth is monitored up to day 25.

Treatment with the antibody drug conjugates according to

Examples 35A and 02A results in a marked inhibition of the growth of the tumours compared to the control group. During the treatment with 3 mg/kg, there is a regression of most tumours. Table 7 shows the T/C values determined for tumour weights and tumour area on day 25. Here, the antibody drug conjugate 35A shows better efficacy than cetuximab and than toxophor A or toxophor B.

Table 7:

10

Example	Dosage	T/C weight	T/C area
Toxophor A	2.5 mg/kg	0.82	0.87
Toxophor B	2.5 mg/kg	0.40	0.57
cetuximab	3 mg/kg	0.47	0.54

**C-6d. Efficacy of anti-EGFR and anti-TWEAKR antibody drug conjugates in the NCI-H292 tumour model on single treatment**

15 1 million NCI-H292 cells are inoculated subcutaneously into the flank of female NMRI-nude mice (Janvier). At a tumour size of 37 mm<sup>2</sup> on day 7, treatment is carried out once intravenously at a dose of 3 or 10 mg/kg of antibody drug conjugate. After the treatment, the tumour growth is monitored up to day 24.

20

Single treatment with the anti-TWEAKR antibody drug conjugate 02B results in a marked and long-lasting inhibition of growth of the tumours compared to the control group and the unconjugated anti-TWEAKR antibody. Table 8 shows the T/C values determined for tumour weights and tumour area on day 24.

25

Table 8:

Example	Dosage	T/C weight	T/C area
91A	3mg/kg	n.d.	0.78
anti-TWEAKR antibody	10mg/kg	0.99	1.01
cetuximab	3mg/kg	n.d.	0.67

nimotuzumab	3mg/kg	n.d.	0.95
237i	3 mg/kg	n.d.	0.47
208i	3 mg/kg	n.d.	0.69

**C-6f. Efficacy of the anti-TWEAKR antibody drug conjugates in the NCI-H292 tumour model on single treatment (2 independent experiments)**

5

In both experiments, 1 million NCI-H292 cells are inoculated subcutaneously into the flank of female NMRI-nude mice (Janvier). At a tumour size of 45 mm<sup>2</sup> on day 10 (experiment 1) or day 8 (experiment 2), treatment is carried out once intravenously at a dose of 3 mg/kg of antibody drug conjugate. After the treatment, the tumour growth is monitored up to day 18 (Experiment 1) or day 24 (Experiment 2).

Single treatment with the anti-TWEAKR antibody drug conjugates 02B, 07B and 08B in Experiment 1 results in a marked inhibition of growth of the tumours compared to the control group and the unconjugated anti-TWEAKR antibody. Table 10 (Experiment 1) shows the T/C values determined for tumour weights and tumour area on day 18. Single treatment with the anti-TWEAKR antibody drug conjugates 155B, 173B, 165B and 085B in Experiment 2 likewise results in a marked inhibition of growth of the tumours compared to the control group and the respective isotype control (not shown). Table 11 (Experiment 2) shows the T/C values determined for the tumour area on day 24.

25

Table 10:

Example	Dosage	T/C weight	T/C area
91B	10 mg/kg	0.52	0.58
anti-TWEAKR antibody	10 mg/kg	0.68	0.84

Table 11:

30

Example	Dosage	T/C weight	T/C area
155B	10 mg/kg	/	0.32
173B	10 mg/kg	/	0.32
165B	10 mg/kg	/	0.30
085B	10 mg/kg	/	0.24

**C-6g. Efficacy of the anti-TWEAKR antibody drug conjugates in the A375 tumour model on repeat treatment**

5 5 million A375 (human melanoma) cells are inoculated subcutaneously into the flank of female NMRI-nude mice (Janvier). At a tumour size of 41 mm<sup>2</sup> on day 10, treatment is initiated intravenously at a dosage of 10 mg/kg (day 10, 14, 18). After the treatment, the tumour growth is monitored up to  
10 day 22.

Treatment with the antibody drug conjugates according to Examples 38B and 104B results in a marked inhibition of the growth of the tumours compared to the control group. Table 10  
15 shows the T/C values determined for tumour weights and tumour area on day 22. Treatment with the respective control conjugate (isotype antibody against an irrelevant antigen) results in a markedly weaker tumour growth-inhibiting action. Treatment with the unconjugated antibodies likewise results in an in some cases  
20 weaker inhibition of the growth of the tumours.

Table 10:

Example	Dosage	T/C weight	T/C area
anti-TWEAKR antibody	10 mg/kg	0.32	0.52
isotype control for 104B	10 mg/kg	0.94	0.96
anti-TWEAKR antibody	10 mg/kg	0.32	0.52
104B	10 mg/kg	0.14	0.29



C-6h. **Efficacy of the anti-TWEAKR antibody drug conjugates in the LoVo tumour model on repeat treatment**

5 million LoVo (human colon carcinoma) cells are inoculated subcutaneously into the flank of female NMRI-nude mice (Janvier). At a tumour size of 43 mm<sup>2</sup> on day 7, treatment is initiated intravenously at a dosage of 10 mg/kg (day 7, 11, 15). After the treatment, the tumour growth is monitored up to day 45.

10

Treatment with the antibody drug conjugates according to Examples 07B, 87B and 104B results in a marked inhibition of the growth of the tumours compared to the control group. Table 11 shows the T/C values determined for the tumour area on day 45.

15

Treatment with the control conjugate for 07B (isotype antibody against an irrelevant antigen) results in a markedly weaker tumour growth-inhibiting action.

Table 11:

20

Example	Dosage	T/C area
104B	10 mg/kg	0.43
87B	10 mg/kg	0.52

D. **Working examples of pharmaceutical compositions**

The compounds according to the invention can be converted to pharmaceutical formulations as follows:

25

**i.v. solution:**

The compound according to the invention is dissolved in a concentration below the saturation solubility in a physiologically acceptable solvent (e.g. isotonic saline solution, D-PBS, or a formulation with glycine and sodium chloride in citrate buffer with addition of polysorbate 80). The solution is subjected to sterile filtration and dispensed into sterile and pyrogen-free injection vessels.

35

**i.v. solution:**

The compounds according to the invention can be converted to the administration forms mentioned. This can be accomplished in a manner known per se by "mixing with" or "dissolving in" inert, non-toxic, pharmaceutically suitable excipients (e.g. buffer substances, stabilizers, solubilizers, preservatives). The following, for example, may be present: amino acids (glycine, histidine, methionine, arginine, lysine, leucine, isoleucine, threonine, glutamic acid, phenylalanine and others), sugars and related compounds (glucose, saccharose, mannitol, trehalose, sucrose, mannose, lactose, sorbitol), glycerol, sodium salts, potassium, ammonium salts and calcium salts (e.g. sodium chloride, potassium chloride or disodiumhydrogenphosphate and many others), acetate/acetic acid buffer systems, phosphate buffer systems, citric acid and citrate buffer systems, trometamol (TRIS and TRIS salts), Polysorbates (e.g. Polysorbate 80 and Polysorbate 20), Poloxamers (e.g. Poloxamer 188 and Poloxamer 171), Macrogols (PEG derivatives, e.g. 3350), Triton X-100, EDTA salts, glutathione, albumins (e.g. human), urea, benzyl alcohol, phenol, chlorocresol, metacresol, benzalkonium chloride and many others.

**Lyophilizate for subsequent conversion into an i.v., s.c. or i.m. solution:**

Alternatively the compounds of the invention may be converted into a stable lyophilizate (possibly with the aid of abovementioned excipients) and, before being administered, reconstituted with a suitable solvent (e.g. injection-grade water, isotonic saline solution) and administered.

**Working examples of anti-TWEAKR antibodies**

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All examples were carried out using standard methods known to the person skilled in the art, unless described here in detail. Routine methods of molecular biology of the examples that follow

can be carried out as described in standard laboratory textbooks such as Sambrook et al., *Molecular Cloning: a Laboratory Manual*, 2. Edition; Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., 1989.

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#### AK EXAMPLE 1: Antibody preparation using an antibody library

A complete human phage display library (Hoet RM et al, *Nat Biotechnol* 2005;23(3):344-8) was employed to isolate TWEAKR-specific human monoclonal antibodies of the present invention by protein panning (Hoogenboom H.R., *Nat Biotechnol* 2005;23(3):1105-16), where dimeric Fc-fused extracellular domains of human and murine TWEAKR were immobilized as target.

15 **Table AK-1:** List of recombinant antigens used for antibody selection

Nomenclature	Description	SEQ ID NO
TPP-599	HUMAN-TNFRSF12Aaa28-80-hIgG1-Fc	138
TPP-601	MURIN-TNFRSF12Aaa28-80-hIgG1-Fc	137

The antigens were biotinylated using an about 2-fold molar excess of biotin-LC-NHS (Pierce; Cat. No. 21347) according to the instructions of the manufacturer and desalted using Zeba desalting columns (Pierce; Cat. No. 89889). Washed magnetic beads (DynaBeads) were incubated overnight with 200 nM biotinylated antigen at 4°C and blocked for 1 h at 4°C with blocking buffer (PBS with 3% BSA, 0.05% Tween-20). The blocked Fab phage library was added to the blocked TWEAKR beads (DynaBeads Streptavidin-M280 - Invitrogen 112-06D) and incubated at room temperature for 30 min. After stringent washing (3 x with blocking buffer and 9 x with PBS (150 mM NaCl; 8 mM Na<sub>2</sub>HPO<sub>4</sub>; 1.5 mM KH<sub>2</sub>PO<sub>4</sub>; adjusted to pH = 7.4-7.6) with 0.05% Tween-20), Fab phages binding specifically to biotinylated TWEAKR beads (DynaBeads Streptavidin- M280 - Invitrogen 112-06D) were resuspended in PBS and, for amplification, used directly for

30

infecting *Escherichia coli* strain TG1. In the second selection round, two murine TWEAKR (200 nM) were used to select for cross-reactive binders, and in the third selection round the concentration of human TWEAKR was reduced (100 nM) to increase the selection pressure for high-affinity binders.

11 different Fab phages were identified and the corresponding antibodies were cloned into a mammalian IgG expression vector which provided the missing CH2-CH3 domains not present in the soluble Fab. The resulting IgGs were expressed transiently in mammalian cells as described in Tom et al., Chapter 12 in *Methods Express: Expression Systems* edited by Micheal R. Dyson and Yves Durocher, Scion Publishing Ltd, 2007. Briefly, a CMV promoter-based expression plasmid was transfected into HEK293-6E cells and incubated in Fernbach bottles or Wave bags. Expression took place at 37°C for 5 to 6 days in F17 medium (Invitrogen). 1% Ultra-Low IgG FCS (Invitrogen) and 0.5 mM valproic acid (Sigma) were added as supplements 24 h after the transfection. The antibodies were purified by protein-A chromatography and characterized further by their binding affinity to soluble monomeric TWEAKR using ELISA and BIAcore analysis, as described in AK-Example 2.

**Table AK-2:** List of recombinant antigen used for the affinity measurement

Nomenclature	Description	Origin	Cat. No. (Fitzgerald Inc)	SEQ ID NO
TPP-2305	hTNFRSF12 amino acids a28-80	human	30R-AT080	168

To determine the cell binding characteristics of the anti-TWEAKR antibodies, binding to a number of cell lines (HT29, HS68, HS578) was examined by flow cytometry. The cells were suspended in dilutions of the antibodies (5 µg/ml) in FACS buffer and

incubated on ice for 1 h. A second antibody (PE goat-anti-human IgG, Dianova #109-115-098) was then added. After 1 h of incubation on ice, the cells were analysed by flow cytometry using an FACS array (BD Biosciences).

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NF-kappaB reporter gene assays were carried out to assess the agonistic activity of all 11 antibodies identified (human IgG1). HEK293 cells were transiently transfected with an NF-kappaB reporter construct (BioCat, Cat. No. LR-0051-PA) using 293fectin according to the instructions of the manufacturer. Transfected cells were sown in F17 media (serum-free; Invitrogen) at 37C, 5% CO2 into white polylysine-coated 384-well plates (BD). The next day, the cells were stimulated with various concentrations of purified antibodies for 6 h, and a luciferase assay was then carried out using standard methods.

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Internalisation was monitored via fluorescence labelling of anti-TWEAKR antibodies (CypHer 5E mono NHS ester; GE Healthcare). Prior to the treatment, HT29 cells were sown (2 x 10<sup>4</sup>/well) in 100 µl of medium in 96-well MTP plates (thick, black, transparent botton, No. 4308776, Applied Biosystems). After 18 h of incubation at 37°C/5%CO<sub>2</sub>, the medium was replaced and labelled anti-TWEAKR antibodies were added in different concentrations (10, 5, 2.5, 1, 0.1 µg/ml). The chosen incubation time was 0, 0.25, 0.5, 1, 1.5, 2, 3, 6 and 24h. Fluorescence measurement was carried out in an InCell-analyser 1000 (GE Healthcare).

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The antibody having the highest in vitro activity (TPP-883) was selected for further activity and affinity maturation.

30

### TPP-883

SEQ ID NO.71

35

AQDIQMTQSPATLSASVGDRTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGV

PSRFSGSGSGTDFLTISLQPEDFATYYCQQSYSSPGITFGPGTKVEIKRTVAAPSVFI

FPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSS

TLTLSKADYKHKLYACEVTHQGLSSPVTKSFNRGEC

5

SEQ ID NO.72

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMMWVRQAPGKGLEWVSYISPSGGKTHY

10 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDGYFDYFDYWGQTLTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKRVEPKSCDKTHTCPPCPAPELLGG

15

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREE

20 MTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRW

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

25 Amino acid sequences of the light (SEQ ID NO.71) and heavy (SEQ ID NO.72) chains of TPP-883; CDRs both of the heavy and the light chain are underlined.

30 Maturation was carried out in a first mutations collection round, followed by recombination of those amino acid modifications which increased affinity and activity most. For collecting mutations NNK (N = AGCT, K = G or T), randomization was carried out at the following individual amino acid positions by site-directed mutagenesis using synthetic oligonucleotides including NNK codon diversification (continuous amino acid nomenclature): S35, S36, Y37 and N39 in CDR-L1; A51, S53, S54, Q56 and S57 in CDR-L2; S92, Y93, S94, S95, G97 and I98 in CDR-L3; P31, Y32, P33, M34 and M35 in CDR-H1; Y50, S52, P53, S54, G56, K57 and H59 in CDR-H2; G99, G100, D101, G102, Y103, F104,

D105 and Y106 in CDR-H3. The DNA of all individual NNK saturation mutagenesis libraries was cloned into a mammalian IgG expression vector for activity maturation or into a phagemid vector for affinity maturation. Affinity maturation was carried out by phage panning. Washed magnetic beads (DynaBeads) were incubated overnight with 10 nM, 1 nM, 100 pM or 10 pM biotinylated antigen at 4°C and blocked for 1 h at 4°C with blocking buffer (PBS with 3% BSA, 0.05% Tween-20). The blocked Fab phage library was added in 10000-fold, 1000-fold or 100-fold excess, compared to the theoretical library complexity, to the blocked TWEAKR-DynaBeads and incubated at room temperature for 30 min. That means that 12 strategies were followed in total (4 antigen concentrations x 3 Fab phage titres). After stringent washing (3 x with blocking buffer and 9 x with PBS with 0.05% Tween-20), Fab phages binding specifically to biotinylated TWEAKR DynaBeads (DynaBeads Streptavidin- M280 - Invitrogen 112-06D) were resuspended in PBS and, for amplification, used directly for infecting *Escherichia coli* strain TG1. In selection round two, the concentration of human TWEAKR-Fc was reduced (1 nM, 100 pM, 10 pM and 1 pM), and the same Fab phage titre was used for all 12 strategies ( $4.4 \times 10^{11}$ ). For the expression of soluble Fab, the phagemid vector was digested with MluI to remove the gene-III membrane anchor sequence required for the Fab display on the phage, and the vector was re-ligated. 96 variants of each of the 12 selection pools were expressed as soluble Fabs and examined in an ELISA format. To this end, 2.5 nM biotinylated TWEAKR-Fc were antigen-coated, and binding of soluble Fabs was demonstrated using anti-c-Myc antibodies (Abcam ab62928). 7 single substitution variants (consecutive amino acid nomenclature) with improved binding to TWEAKR-Fc (Seq ID No 138) were demonstrated: S36G of CDR-L1, A51Q and S57K of CDR-L2, S94T and G97F of CDR-L3, M35I of CDR-H1 and G102T of CDR-H3. For the activity maturation, HEK293 cells were transfected with an NF-kappaB reporter (BioCat, Cat. No. LR-0051-PA). Transfected cells were sown in F17 media (serum-free; Invitrogen) in white, polylysine-coated 384-well plates (BD), and individual variants of the NNK-diversified position antibodies (human IgG1) libraries were expressed transiently in mammalian cells. The next day, NF-kappaB reporter

cells were stimulated with the individual NNK antibody variants expressed for 6 h, and a luciferase assay was then carried out using standard methods. 1 single substitution variant having improved agonistic activity was detected: G102T of CDR-H3. This variant was also obtained by affinity maturation, and there, too, it showed the highest enhancement of affinity. After mutation collection by affinity and activity screening, all 7 favourable individual substitutions (library complexity: 128 variants) were recombined into a recombination library. To this end, oligonucleotides were synthesized to introduce selected mutations or the corresponding wild type amino acid at each selected position. The library was established using successive rounds of overlap extension PCR. The final PCR product was ligated into a bacterial soluble Fab expression vector, and 528 variants were selected at random (~ 4-fold excess of the sample taken) for an equilibrium ELISA screen with soluble Fabs, as described above. In the end, 7 variants were selected based on increased affinity compared to the best single substitution variant G102T. The corresponding DNA of these was cloned into a mammalian IgG expression vector and examined for functional activity in the above-mentioned NF-kappaB reporter cell assay. Finally, the sequences obtained were compared with human germ line sequences, and deviations without any significant effect on the affinity and the efficacy were adapted. Antibodies having the sequences below were obtained by antibody library screening and by affinity and/or activity maturation:

**TPP-2090**

30 SEQ ID NO.1:

DIQMTQSPSSLSASVGDRVTITCRASQSISGYLNWYQQKPGKAPKLLIYQASSLQSGVPS

RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTSPFITFGQGTKVEIKRTVAAPSVFIFP

35

PSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTL

TLISKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC



SEQ ID NO.2:

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMIWVRQAPGKGLEWVSYISPSGGSTHY

5

ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLVTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

10 GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

15

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

20 Amino acid sequences of the light (SEQ ID NO.1) and heavy (SEQ ID NO.2) chains of TPP-2090; CDRs both of the heavy and the light chain are underlined.

**TPP-2149**

25

SEQ ID NO.11

DIQMTQSPATLSASVGRVTITCRASQSISGYLNWYQQKPGKAPKLLIYQASSLQSGVPS

30 RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTSPFITFGPGTKVEIKRTVAAPSVFIFP

PSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDYSLSTL

TLISKADYEKHKLYACEVTHQGLSSPVTKSFNRGEC

35

SEQ ID NO.12

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMIWVRQAPGKGLEWVSYISPSGGKTHY

ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLTVTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSQVHTFPAVLQSS

5

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

10 STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

15

Amino acid sequences of the light (SEQ ID NO.11) and heavy (SEQ ID NO.12) chains of TPP-2149; CDRs both of the heavy and the light chain are underlined.

20 **TPP-2093**

SEQ ID NO.21

DIQMTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYQASSLQSGVPS

25

RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTSPFITFGQGTKVEIKRTVAAPSVFIFP

PSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDESTYSLSTL

30 TLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO.22

EVQLLESQGGGLVQPGGSLRLSCAASGFTFSPYPMWVRQAPGKGLEWVSYISPSGGSTHY

35

ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLTVTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSQVHTFPAVLQSS

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

5

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

10 QQGNVFSCSVMHEALHNHYTQKSLSLSPG

Amino acid sequences of the light (SEQ ID NO.21) and heavy (SEQ ID NO.22) chains of TPP-2093; CDRs both of the heavy and the light chain are underlined.

15

**TPP-2148**

SEQ ID NO.31

20 DIQMTQSPATLSASVGRVTITCRASQSISSYLNWYQQKPGKAPKLLIYQASSLQSGVPS

RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTSPFITFGPGTKVEIKRTVAAPSVFIFP

PSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDESTYSLSTL

25

TLISKADYEKHKLYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO.32

30 EVQLLES GGGLVQPGGSLRLS CAASGFTFSPYPMWVRQAPGKGLEWVS YISPSGGKTHY

ADSVKGRFTISRDN SKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLVTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

35

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

5

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

Amino acid sequences of the light (SEQ ID NO.31) and heavy (SEQ  
ID NO.32) chains of TPP-2148; CDRs both of the heavy and the  
10 light chain are underlined.

**TPP-2084**

SEQ ID NO.41

15

DIQMTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGVPS

RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPGITFGQGTKVEIKRTVAAPSVFIFP

20

PSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDYSLSTL

TLISKADYEEKHKVYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO.42

25

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMWVRQAPGKGLEWVSYISPSGGSTHY

ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQTLTVSS

30

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

GLYSLSSVTVTPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

35

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

Amino acid sequences of the light (SEQ ID NO.41) and heavy (SEQ  
5 ID NO.42) chains of TPP-2084; CDRs both of the heavy and the  
light chain are underlined.

**TPP-2077**

10 SEQ ID NO.51

DIQMTQSPATLSASVGRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGVPS

15 RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSSPGITFGPGTKVEIKRTVAAPSVFIFP

PSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDYSLSTL

TLSKADYEKHKLYACEVTHQGLSSPVTKSFNRGEC

20 SEQ ID NO.52

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMWVRQAPGKGLEWVSYISPSGGKTHY

25 ADSVKGRFTISRDNKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLVTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

30 PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

35

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

Amino acid sequences of the light (SEQ ID NO.51) and heavy (SEQ

ID NO.52) chains of TPP-2077; CDRs both of the heavy and the light chain are underlined.

**TPP-1538**

5

SEQ ID NO.61

AQDIQMTQSPATLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGV

10 PSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSSPGITFGPGTKVEIKRTVAAPSVFI

FPPSDEQLKSGTASVCLLNFPYFVWYQFKVDNALQSGNSQESVTEQDSKDSTYSLS

TLTSLKADYEKHKLYACEVTHQGLSSPVTKSFNRGEC

15

SEQ ID NO.62

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMWVRQAPGKGLEWVSYISPSGGKTHY

20 ADSVKGRFTISRDNKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLVTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

25

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

30 LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRW

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

35 Amino acid sequences of the light (SEQ ID NO.61) and heavy (SEQ ID NO.62) chains of TPP-1538; CDRs both of the heavy and the light chain are underlined.

**TPP-1854**

SEQ ID NO.81

AQDIQMTQSPATLSASVGDRVTITCRASQSISGYLNWYQQKPGKAPKLLIYNASSLQSGV  
 5 PSRFSGSGSGTDFLTITSSLQPEDFATYYCQQSYTSPFITFGPGTKVEIKRTVAAPSVFI  
 FPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLS  
 10 TLTLKADYEKHKLYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO.82

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMIWRQAPGKGLEWVSYISPSGGKTHY  
 15 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQTLTVSS  
 ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSQVHTFPAVLQSS  
 20 GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG  
 PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN  
 STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE  
 25 LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRW  
 QQGNVFSCSVMHEALHNHYTQKSLSLSPG

30 Amino acid sequences of the light (SEQ ID NO.81) and heavy (SEQ  
 ID NO.82) chains of TPP-1854; CDRs both of the heavy and the  
 light chain are underlined.

**TPP-1853**

35

SEQ ID NO.91

AQDIQMTQSPATLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYNASSLQSGV

PSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTSPGITFGPGTKVEIKRTVAAPSVFI

FPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSS

5

TLTLSKADYEKHKLYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO.92

10 EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMMWVRQAPGKGLEWVSYISPSGGKTHY

ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLTVTVSS

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSQGVHTEFPAVLQSS

15

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

20 STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRW

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

25

Amino acid sequences of the light (SEQ ID NO.91) and heavy (SEQ ID NO.92) chains of TPP-1853; CDRs both of the heavy and the light chain are underlined.

30 **TPP-1857**

SEQ ID NO.101

AQDIQMTQSPATLSASVGRVTITCRASQSISGYLNWYQQKPGKAPKLLIYNASSLQSGV

35

PSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTSPGITFGPGTKVEIKRTVAAPSVFI

FPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSS



TLTLSKADYEEKHKLYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO.102

5

EVQLLESGGGLVQPGGSLRLSCAASGFTFSSPYPMWVRQAPGKGLEWVSYISPSGGKTHY

ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLTVTVSS

10 ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

15

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

20 QQGNVFSCSVMHEALHNHYTQKSLSLSPG

Amino acid sequences of the light (SEQ ID NO.101) and heavy (SEQ ID NO.102) chains of TPP-1857; CDRs both of the heavy and the light chain are underlined.

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**TPP-1858**

SEQ ID NO.111

30 AQDIQMTQSPATLSASVGRVTITCRASQSISSYLNWYQQKPGKAPKLLIYNASSLQSGV

PSRFGSGSGTDFTLTISSLQPEDFATYYCQQSYTSPFITFGPGTKVEIKRTVAAPSVFI

FPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLS

35

TLTLSKADYEEKHKLYACEVTHQGLSSPVTKSFNRGEC

SEQ ID NO.112

EVQLLESGGGLVQPGGSLRLSCAASGFTFSPYPMMWVRQAPGKGLEWVSYISPSGGKTHY

ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGGDTYFDYFDYWGQGLVTVSS

5

ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

GLYSLSSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG

10

PSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN

STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE

LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRW

15

QQGNVFSCSVMHEALHNHYTQKSLSLSPG

Amino acid sequences of the light (SEQ ID NO.111) and heavy (SEQ ID NO.112) chains of TPP-1858; CDRs both of the heavy and the light chain are underlined.

20

## EXAMPLE 2: Biochemical characteristics of the antibodies

25 *Determination of binding affinities by Biacore analysis:*

Binding affinities of anti-TWEAKR antibodies were examined using surface plasmon resonance analysis on a Biacore T100 instrument (GE Healthcare Biacore, Inc.). The antibodies were immobilized on a CM5 sensor chip using an indirect capture reagent, anti-human IgG(Fc). Reagents of the "Human Antibody Capture Kit" (BR-1008-39, GE Healthcare Biacore, Inc.) were used as described by the manufacturer. Anti-TWEAKR antibodies were injected at a concentration of 10 µg/ml at 10 µl/min for 10 sec.

35

**Table AK-3:** List of recombinant antigen (TWEAKR) used for affinity measurement

Nomenclature	Description	Origin	Cat. (Fitzgerald Inc)	No.	SEQ ID NO
TPP-2305	hTNFRSF12 amino acids a28-80	human	30R-AT080		168

**Table AK-4:** List of antibodies used for the affinity measurement

Nomenclature	Description	SEQ ID NO	
		Light chain	Heavy chain
P3G5 (TPP-2195)	murine IgG2a	121	122
P4A8 (TPP-1324)	human IgG1	125	126
P2D3 (TPP-2196)	murine IgG2a	131	132
136.1 (TPP-2194)	murine IgG2a	123	124
PDL-192 (TPP-1104)	human IgG1	127	128
18.3.3 (TPP-2193)	murine IgG2a	129	130
TPP-883	human IgG1	71	72
TPP-1538	human IgG1	61	62
TPP-2077	human IgG1	51	52
TPP-2084	human IgG1	41	42
TPP-2148	human IgG1	31	32
TPP-2093	human IgG1	21	22
TPP-2149	human IgG1	11	12
TPP-2090	human IgG1	1	2

5 **Table AK-5:** List of commercially available antibodies used for the affinity measurement

Nomenclature	Description	Cat. No. (Abcam)
ITEM-1	murine IgG1	ab21359
ITEM-4	murine IgG1	ab21127

Various concentrations (200 nM, 100 nM, 50 nM, 25 nM, 12.5 nM, 6.25 nM, 3.12 nM, 1.56 nM) of purified recombinant human TWEAKR protein (TPP-2305, SEQ ID NO:168) in HEPES-EP buffer (GE Healthcare Biacore, Inc.) were injected over immobilised anti-TWEAKR antibodies at a flow rate of 60  $\mu$ l/min for 3 minutes, the dissociation time being 5 minutes. Sensorgrams were generated after in-line reference cell correction, followed by subtraction of the buffer sample. The dissociation constant ( $K_D$ ) was calculated based on the ratio of association ( $k_{on}$ ) and dissociation ( $k_{off}$ ) constants, obtained by fitting sensorgrams using a 1:1 first order binding model.

**Table AK-6:** Monovalent  $K_D$  values of anti-TWEAKR antibodies measured using Biacore with TWEAKR protein (TPP-2305 (SEQ ID NO: 168)).

	$k_a$ (1/MS)	$k_d$ (1/s)	$K_D$ (nM)
TPP-883	4.40E+06	9.10E-01	205.9
TPP-1538	4.20E+06	1.10E-01	27.6
TPP-2077	3.00E+06	8.60E-02	28.9
TPP-2084	4.20E+06	1.10E-01	27.6
TPP-2148	5.10E+06	1.30E-01	24.5
TPP-2093	4.10E+06	9.00E-02	22.1
TPP-2149	8.40E+06	1.00E-01	12.1
TPP 2090	9.10E+06	1.10E-01	12.4
PDL-192 (TPP-1104)	1.00E+07	3.80E-02	3.7
136.1 (TPP-2194)	3.84E+07	3.24E-02	0.8
18.3.3 (TPP-2193)	1.64E+07	2.85E-02	1.7
P4A8 (TPP-1324)	1.20E+06	2.70E-03	2.3
P3G5 (TPP-2195)	2.31E+06	1.22E-03	0.5
P2D3 (TPP-2196)	1.32E+06	5.64E-04	0.4
ITEM-1	3.80E+06	1.10E-02	2.9
ITEM-4	2.80E+06	2.00E-03	0.7

It was determined that the antibodies of the invention bind TWEAKR with moderate affinity ( $K_D$  10 - 200 nM), whereas some

comparative antibodies (e.g. PDL-192 (TPP-1104), 136.1 (TPP-2194), 18.3.3 (TPP-2193), P4A8 (TPP-1324), P3G5 (TPP-2195), P2D3 (TPP-2196), ITEM-1, ITEM-4) show high-affinity binding (0.7 - 3.7 nM). The sequences of the variable domains of the antibodies PDL-192, 136.1, 18.3.3, P4A8, P3G5 and P2D3 were obtained from the patent literature WO2009/020933 and WO2009/140177, and the sequences coding for the constant region of human IgG1 and murine IgG2 were added, resulting in full-length IgGs PDL-192 (TPP-1104), 136.1 (TPP-2194), 18.3.3 (TPP-2193), P4A8 (TPP-1324), P3G5 (TPP-2195), P2D3 (TPP-2196). The range of the affinities measured in this study agrees well with published data: for PDL-192, 18.3.3 and 136.1, KD values of 5.5, 0.2 and 0.7 nM have been published (WO2009/020933); for P4A8 2.6 nM (WO2009/140177). For comparison: the native ligand TWEAK binds TWEAKR with a  $K_D$  value of 0.8 - 2.4 nM (Immunity. 2001 Nov;15(5):837-46; Biochem J. 2006 Jul 15;397(2):297-304; Arterioscler Thromb Vasc Biol. 2003 Apr 1;23(4):594-600).

As a result, it can be recorded that the antibodies of the invention (TPP-883, TPP-1538, TPP-2077, TPP-2084, TPP-2148, TPP-2093, TPP-2149 and TPP-2090) bind TWEAKR with moderate affinity ( $K_D$  10 - 200 nM).

*Characterization of the binding epitope of TPP-2090 using N- and C-terminally truncated variants of the TWEAKR ectodomain:*

The alignment of the cysteine-rich domain of TWEAKR (amino acids 34-68) of different species (Figure 1-Alignment) shows that it is well conserved in all 6 species analysed. PDL-192 binds depending on R56 (WO2009/020933: Figure 2B) and does therefore not bind to rat, pig and mouse TWEAKR. TPP-2090 binds depending on the conserved amino acid D47, and therefore binds to all species shown.

In a first approach to characterizing the binding epitope of the antibodies mentioned above, a N- and C-terminally truncated mutant of the TWEAKR ectodomain was generated and examined for its ability to bind the various anti-TWEAKR antibodies. Amino

acids 28 to 33 were deleted N-terminally and amino acids 69 to 80 were deleted C-terminally, such that the cysteine-rich domains with disulphide bridges between Cys36-Cys49, Cys52-Cys67 and Cys55-Cys64 remained intact (compare Figure 2). Both  
 5 constructs, the full ectodomain 28-80 including N- and C-terminus and the truncated ectodomain 34-68, were expressed and purified as Fc fusion proteins TPP-2202 and TPP-2203, respectively.

10 To analyse the binding, 1 µg/ml of the corresponding dimeric TWEAKR Fc construct was coated, and 0.3 µg/ml and 0.08 µg/ml of biotinylated IgG were used as soluble binding partner. Detection was carried out using streptavidin-HRP and Amplex Red substrate. IgGs were biotinylated using an about 2-fold molar excess of  
 15 biotin-LC-NHS (Pierce; Cat. No. 21347) according to the instructions of the manufacturer and desalted using Zeba desalting columns (Pierce; Cat. No. 89889). At all concentrations used of the soluble ligand, the antibodies of the present invention displayed saturated binding to both  
 20 constructs, whereas the antibodies P4A8(TPP-1324), P3G5(TPP-2195) and Item-4 showed saturated binding only to the full-length ectodomain, but worsened binding to the N- and C-terminally truncated constructs (Figure 3 & Figure 4). This shows that the binding epitope of the antibodies of the present  
 25 invention is located in the cysteine-rich domain between amino acids 34-68. To analyse whether the N-terminus or the C-terminus of the TWEAKR ectodomain is required for P4A8(TPP-1324) and P3G5(TPP-2195) binding, a monomeric ectodomain having the C-terminal deletion of amino acids 69 to 80 was generated. Binding  
 30 of P4A8(TPP-1324) and P3G5(TPP-2195) to the C-terminally truncated TWEAKR ectodomain is likewise worsened, whereas the antibodies of the present invention show saturated binding (Figure 5).

35 **Table AK-9:** List of recombinant antigens used in the ELISA analysis for epitope profiling

Nomenclature	Description	SEQ ID NO
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TPP-2202	TWEAKR-ECD-28-80-hIgGFc-His	139
TPP-2203	TWEAKR-ECD-34-68-hIgGFc-His	140
TPP-1984	hTNFRSF12 amino acids 28-68-CT-His	141

**Table AK-10:** List of antibodies used in the ELISA analysis for epitope profiling

Nomenclature	Description	SEQ ID NO	
		Light chain	Heavy chain
P3G5 (TPP-2195)	murine IgG2a	121	122
P4A8 (TPP-1324)	human IgG1	125	126
136.1 (TPP-2194)	murine IgG2a	123	124
PDL-192 (TPP-1104)	human IgG1	127	128
TPP 2090	human IgG1	1	2
TPP-2084	human IgG1	41	42

5

Thus, the binding epitope of TPP-2090, TPP-2084, PDL-192 (TPP-1104) and 136.1 (TPP-2194) in the cysteine-rich domain and the binding epitope of P4A8 (TPP-1324) and P3G5 (TPP-2195) are located at least partially outside of the cysteine-rich domain.

10

#### *Effect of TWEAKR-Fc muteins on the antibody affinity*

To examine the binding characteristics of the antibodies of the invention in more detail, certain muteins of TWEAKR suggested to be of relevance for the activity of known agonistic antibodies (WO2009/140177) were investigated. To this end, the full-length ectodomain (amino acids 28-80) having the individual amino acid substitutions below were expressed and purified as Fc fusion proteins: T33Q; S40R; W42A; M50A; R56P; H60K; L65Q.

20

**Table AK-11:** List of recombinant proteins used in the ELISA

analysis for mutein binding

Nomenclature	Description	SEQ ID NO
TPP-1990	hTNFRSF12 amino acids a28-80-L65Q-hIgG1-Fc	142
TPP-1989	hTNFRSF12 amino acids a28-80-H60K-hIgG1-Fc	143
TPP-2683	hTNFRSF12 amino acids a28-80-R56P-hIgG1-Fc	144
TPP-1988	hTNFRSF12 amino acids a28-80-M50A-hIgG1-Fc	145
TPP-1985	hTNFRSF12 amino acids a28-80-W42A-hIgG1-Fc	146
TPP-1987	hTNFRSF12 amino acids a28-80-S40R-hIgG1-Fc	147
TPP-1986	hTNFRSF12 amino acids a28-80-T33Q-hIgG1-Fc	148
TPP-599	hTNFRSF12 amino acids a28-80-hIgG1-Fc	138

To obtain dose-reaction data, the different TWEAKR-Fc muteins were coated at a low concentration (62 ng/ml) onto a 384-well Maxisorb ELISA plate, and a serial 2-fold dilution of biotinylated IgG beginning with a concentration of 100 nM was used as a soluble binding partner. Detection was carried out using streptavidin-HRP and Amplex Red. The IgGs examined were TPP-2090 and TPP-2084 of the present invention, PDL-192, 136.1 and 18.3.3 of WO2009/020933, P4A8 and P3G5 of WO2009/140177, and ITEM-1 and ITEM-4 of Nakayama et al [Biochem Biophys Res Com 306: 819-825].

**Table AK-12:** List of antibodies used in the ELISA analysis for mutein binding

Nomenclature	Description	SEQ ID NO	
		Light chain	Heavy chain
P3G5 (TPP-2195)	murine IgG2a	121	122



P4A8 (TPP-1324)	human IgG1	125	126
136.1 (TPP-2194)	murine IgG2a	123	124
PDL-192 (TPP-1104)	human IgG1	127	128
18.3.3 (TPP-2193)	murine IgG2a	129	130
TPP 2090	human IgG1	1	2
TPP-2084	human IgG1	41	42

**Table AK-13:** List of commercially available antibodies used in the ELISA for mutein binding

Nomenclature	Description	Cat. No. (Abcam)
ITEM-1	murine IgG1	ab21359
ITEM-4	murine IgG1	ab21127

5

IgGs were biotinylated using an about 2-fold molar excess of biotin-LC-NHS (Pierce; Cat. No. 21347) according to the instructions of the manufacturer and desalted using Zeba desalting columns (Pierce; Cat. No. 89889). The dose-reaction data were fitted and the IC<sub>50</sub>s were determined. To illustrate the results, a table was generated; "-" marks IC<sub>50</sub>s over 50 nM, "+" marks IC<sub>50</sub>s in the range from 1 to 150 pM.

10

**Table AK-14:** Effect of muteins on antibody binding

15

	T33Q	S40R	W42A	M50A	R56P	H60K	L65Q	WT
TPP-2084	+	+	-	+	+	+	+	+
TPP-2090	+	+	-	+	+	+	+	+
PDL-192 (TPP-1104)	+	+	-	+	-	+	+	+
136.1 (TPP-2194)	+	+	-	+	-	+	+	+
18.3.3 (TPP-2193)	+	+	-	+	-	+	+	+

P4A8 (TPP-1324)	+	+	-	+	+	+	+	+
P3G5 (TPP-2195)	+	+	-	+	+	+	+	+
ITEM1	+	+	-	+	-	+	+	+
ITEM4	+	+	-	+	+	-	+	+

As already published, ITEM-4 shows worsened binding to the H60K mutein [WO2009/140177: Figure 23F] and PDL-192 to the R56P mutein [WO2009/020933: Figure 22B]. In contrast to published data, ITEM-1 shows worsened binding to R56P, and all antibodies to W42A [WO2009/140177: Figure 23E, Figure 23F]. These differences can be explained by the methods chosen.

In contrast to ITEM-1, ITEM-4, PDL-192, 136.1 and 18.3.3, the antibodies of the present invention bind independently of all substitutions except for W42A.

*Alanine scan of the cysteine-rich domain:*

An alanine scan of the cysteine-rich domain (amino acids 34-68) was carried out in order to locate the binding site of the antibodies of the invention. Figure 6 shows that N- and C-terminally truncated variants of the full-length ectodomain of TWEAKR do not worsen binding of the antibodies of the invention. Accordingly, the binding epitope is located in the cysteine-rich domain. The substitutions below were introduced into the TWEAKR(34-68) Fc construct: S37A, R38A, S40A, S41A, W42A, S43A, D45A, D47A, K48A, D51A, S54A, R56A, R58A, P59A, H60A, S61A, D62A, F63A und L65A.

**Table AK-15:** List of TWEAKR mutein constructs for the alanine scan of the cysteine-rich domain

Nomenclature	Description	SEQ ID NO
TPP-2203	TweakR-ECD-34-68-hIgGFc-His	140
TPP-2625	TweakR-ECD-34-68-hIgGFc-His-L65A	149

TPP-2624	TweakR-ECD-34-68-hIgGFc-His-F63A	150
TPP-2623	TweakR-ECD-34-68-hIgGFc-His-D62A	151
TPP-2622	TweakR-ECD-34-68-hIgGFc-His-S61A	152
TPP-2621	TweakR-ECD-34-68-hIgGFc-His-H60A	153
TPP-2620	TweakR-ECD-34-68-hIgGFc-His-P59A	154
TPP-2619	TweakR-ECD-34-68-hIgGFc-His-R58A	155
TPP-2618	TweakR-ECD-34-68-hIgGFc-His-R56A	156
TPP-2617	TweakR-ECD-34-68-hIgGFc-His-S54A	157
TPP-2616	TweakR-ECD-34-68-hIgGFc-His-D51A	158
TPP-2615	TweakR-ECD-34-68-hIgGFc-His-K48A	159
TPP-2614	TweakR-ECD-34-68-hIgGFc-His-D47A	160
TPP-2613	TweakR-ECD-34-68-hIgGFc-His-D45A	161
TPP-2612	TweakR-ECD-34-68-hIgGFc-His-S43A	162
TPP-2611	TweakR-ECD-34-68-hIgGFc-His-W42A	163
TPP-2610	TweakR-ECD-34-68-hIgGFc-His-S41A	164
TPP-2609	TweakR-ECD-34-68-hIgGFc-His-S40A	165
TPP-2608	TweakR-ECD-34-68-hIgGFc-His-R38A	166
TPP-2607	TweakR-ECD-34-68-hIgGFc-His-S37A	167

These TWEAKR(34-68) Fc muteins were expressed in HEK293 cells.

To obtain dose-reaction data, IgGs were coated at a concentration of 1 µg/ml onto a 384-well Maxisorp ELISA plate, and a serial 2-fold dilution of the supernatant comprising the TWEAKR mutein was used as soluble binding partner. Detection was carried out using anti-HIS-HRP and Amplex Red. The IgGs examined were TPP-2090 of the present invention, PDL-192 of WO2009/020933 and P4A8 of WO2009/140177.

**Table AK-16:** List of antibodies used for the alanine scan of the cysteine-rich domain

Nomenclature	Description	SEQ ID NO	
		Light chain	Heavy chain
P4A8 (TPP-1324)	human IgG1	125	126
PDL-192 (TPP-1104)	human IgG1	127	128
TPP 2090	human IgG1	1	2

To assess the relevance of the TWEAKR mutein for binding to various IgGs, a correlation blot at a certain mutein concentration was prepared. By way of example, Figure 6 shows the correlation blots for the 8-fold diluted supernatants of the TWEAKR expression broth, with PDL-192 (TPP-1104) on the X axis and TPP-2090 on the Y axis. The blot shows that binding of TPP-2090 was worsened by substitution D47A, and binding of PDL-192 (TPP-1104) was worsened by substitution R56A. Binding to P4A8 (TPP-1324) was demonstrated for none of the constructs, which agrees with the results obtained above (Figure 6). Thus, the P4A8 epitope is localized at least partially outside of the cysteine-rich domain. The dependencies identified for certain TWEAKR amino acids for antibody interaction correlates with the agonistic activity determined for these antibodies. The native ligand TWEAK shows an effective activation of the TWEAKR and binds depending on leucine 46 in the cysteine-rich domain of TWEAKR (Pellegrini et al, FEBS 280:1818-1829). P4A8 displays a very low agonistic activity and interacts at least partially

with domains outside of the cysteine-rich domain of TWEAKR. PDL-192 displays a moderate agonistic activity and binds depending on R56 to the cysteine-rich domain, but opposite the TWEAK ligand site. TPP-2090 and TWEAK binding depends on D47 and L46, respectively, and they therefore bind to a similar binding site (Figure 7).

It can be concluded the the antibodies of the invention (e.g. TPP-2090) bind to TWEAKR in a manner depending on D47.

10

The dependencies identified for certain TWEAKR amino acids for antibody interaction correlates with the agonistic activity determined for these antibodies. The native ligand TWEAK shows an effective activation of the TWEAKR and binds depending on leucine 46 in the cysteine-rich domain of TWEAKR (Pellegrini et al, FEBS 280:1818-1829). P4A8 displays a very low agonistic activity and interacts at least partially with domains outside of the cysteine-rich domain of TWEAKR. PDL-192 displays a moderate agonistic activity and binds depending on R56 to the cysteine-rich domain, but opposite the TWEAK ligand site. Antibodies of the present invention (Example TPP-2090) bind in a manner depending on D47, and TWEAK binds in a manner depending on L46, and binds to a similar, but distinct, binding site (Figure 7). Accordingly, the antibodies of the present invention displaying strong agonistic activity bind to a novel epitope (D47-dependent) for antibodies associated with very high agonistic activity. It is interesting to note that Michaelson et al. (see page 369, left column in Michaelson JS et al, MAbs. 2011 Jul-Aug;3(4):362-75) gave an explanation for the fact that all agonistic antibodies examined by them have weaker agonistic activity than the natural ligand TWEAK. They conclude that reduced efficacy could be a function of the dimeric binding interaction of the antibodies with TWEAKR, with TWEAK probably entering into a trimeric interaction. It is therefore a surprising result that an antibody of the invention, in spite of its dimeric interaction with TWEAKR, has an even higher agonistic activity. This surprising activity is linked to the specific binding properties of the antibodies of the invention,

35

i.e. the specific binding to D47 of TWEAKR.

Further embodiments The scope of protection is solely defined by the claims.

5

1. Conjugate of a binder or derivative thereof with one or more active compound molecules, the active compound molecule being a kinesin spindle protein inhibitor attached to the binder via a linker L.

10

2. Conjugate according to Item 1 where the binder or derivative thereof is a binder peptide or protein or a derivative of a binder peptide or protein.

15

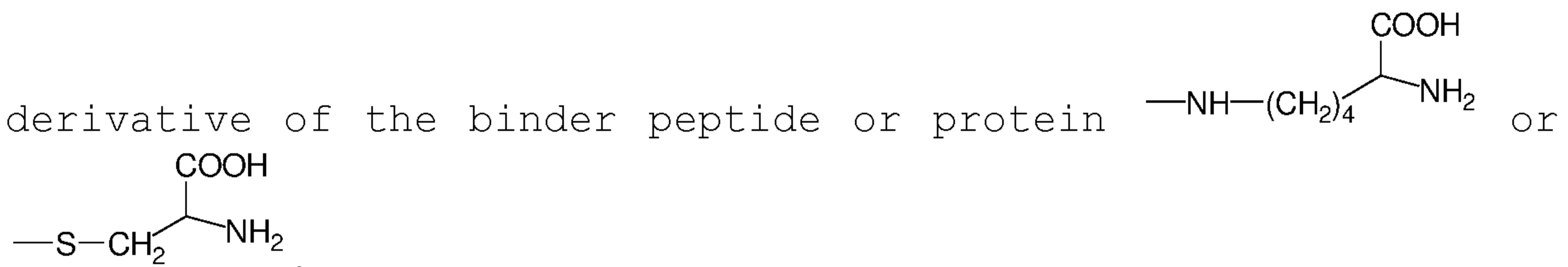
3. Conjugate according to Item 2 where each active compound molecule is attached to different amino acids of the binder peptide or protein or derivative thereof via the linker.

20

4. Conjugate according to one or more of the preceding items where the conjugate has on average 1.2 to 20 active compound molecules per binder.

25

5. Conjugate according to one or more of Items 2 to 4 where the binder peptide or protein represents an antibody or the derivative of the binder peptide or protein



30

6. Conjugate according to one or more of the preceding items where the binder binds to a cancer target molecule.

7. Conjugate according to Item 6 where the binder binds to an extracellular target molecule.

35

8. Conjugate according to Item 7 where the binder, after binding to the extracellular target molecule, is internalized and processed intracellularly (preferably lysosomally) by the

cell expressing the target molecule.

9. Conjugate according to one or more of Items 2 to 8 where the binder peptide or protein is a human, humanized or chimeric monoclonal antibody or an antigen-binding fragment thereof.

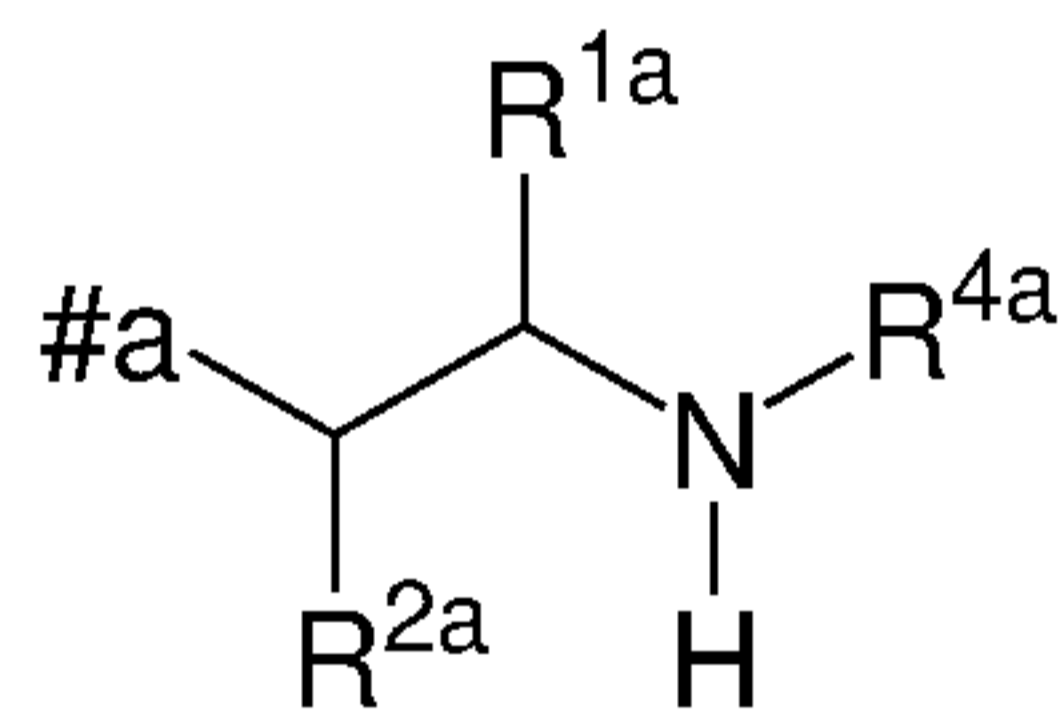
10. Conjugate according to Item 9 where the binder peptide or protein is an anti-HER2 antibody, an anti-EGFR antibody, an anti-TWEAKR antibody or an antigen-binding fragment thereof.

10

11. Conjugate according to Item 10 where the anti-TWEAKR antibody binds specifically to amino acid D in position 47 (D47) of TWEAKR (SEQ ID NO:169), preferably the anti-TWEAKR antibody TPP-2090.

15

12. Conjugate according to one or more of the preceding items where the kinesin spindle protein inhibitor has the substructure below:



20

where

\#a represents a bond to the remainder of the molecule;

25

$\text{R}^{1a}$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}$ , where Z represents -H,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $-\text{NHY}^3$ , halogen,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$ ,  $-(\text{CH}_2\text{CH}_2\text{O})_{0-3}-(\text{CH}_2)_{0-3}\text{Z}'$  or  $-\text{CH}(\text{CH}_2\text{W})\text{Z}'$ , and  $\text{Y}^3$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$  represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$ ,  $\text{COOH}$ ,  $-\text{NH}-\text{CO}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{NH}_2)\text{COOH}$  or  $-(\text{CO}-\text{NH}-\text{CHY}^4)_{1-3}\text{COOH}$ , where W represents H or OH, where  $\text{Y}^4$  independently of one another represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl which is optionally substituted by  $-\text{NHCONH}_2$ , or represents aryl or benzyl which are optionally

35

substituted by  $-\text{NH}_2$ ;

$\text{R}^{2a}$  and  $\text{R}^{4a}$  independently of one another represent H,  $-\text{CO}-\text{CHY}^4-$   
 $\text{NHY}^5$  or  $-(\text{CH}_2)_{0-3}\text{Z}$ , or  $\text{R}^{2a}$  and  $\text{R}^{4a}$  together (with formation of a  
 5 pyrrolidine ring) represent  $-\text{CH}_2-\text{CHR}^{10}-$  or  $-\text{CHR}^{10}-\text{CH}_2-$ , where  $\text{R}^{10}$   
 represents H,  $\text{NH}_2$ ,  $\text{COOH}$ ,  $\text{SO}_3\text{H}$ ,  $\text{SH}$  or  $\text{OH}$ ,

where Z represents  $-\text{H}$ ,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $-\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

10 where  $\text{Y}^1$  and  $\text{Y}^2$  independently of one another represent H,  $\text{NH}_2$  or  
 $-(\text{CH}_2)_{0-3}\text{Z}'$ , and  $\text{Y}^3$  represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where  $\text{Z}'$   
 represents H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  or  $\text{COOH}$ ;

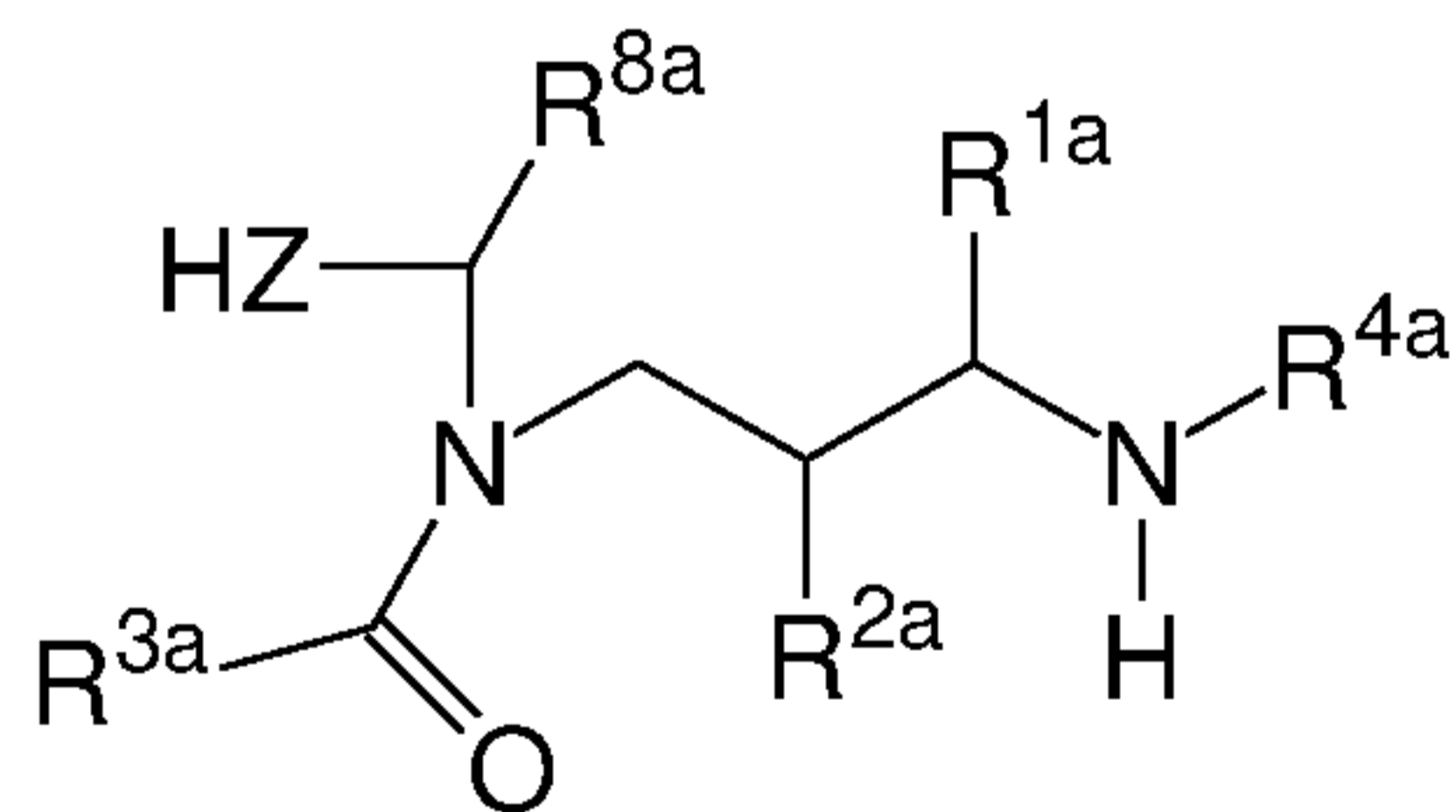
where  $\text{Y}^4$  independently of one another represents straight-chain  
 15 or branched  $\text{C}_{1-6}$  alkyl which is optionally substituted by  $-\text{NHCONH}_2$ , or represents aryl or benzyl which are optionally  
 substituted by  $-\text{NH}_2$ , and  $\text{Y}^5$  represents H or  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ , where  
 $\text{Y}^6$  represents straight-chain or branched  $\text{C}_{1-6}$ -alkyl;

20 where the kinesin spindle protein inhibitor is attached to the  
 linker by substitution of a hydrogen atom at  $\text{R}^{1a}$ ,  $\text{R}^{2a}$ ,  $\text{R}^{4a}$  or at  
 the pyrrolidine ring formed by  $\text{R}^{2a}$  and  $\text{R}^{4a}$ ,

and the salts, solvates and salts of the solvates thereof.

25

13. Conjugate according to one or more of the preceding items  
 where the kinesin spindle protein inhibitor is represented by  
 general formula (I):



30 (I)

where



R<sup>1a</sup> represents H or  $-(\text{CH}_2)_{0-3}\text{Z}$ , where Z represents -H,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $-\text{NHY}^3$ , halogen,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>,  $-(\text{CH}_2\text{CH}_2\text{O})_{0-3}$ ,  $-(\text{CH}_2)_{0-3}\text{Z}'$  or  $-\text{CH}(\text{CH}_2\text{W})\text{Z}'$ , and Y<sup>3</sup> represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub>, COOH,  $-\text{NH}-\text{CO}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{NH}_2)\text{COOH}$  or  $-(\text{CO}-\text{NH}-\text{CHY}^4)_{1-3}\text{COOH}$ , where W represents H or OH, where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by  $-\text{NHCONH}_2$ , or represents aryl or benzyl which are optionally substituted by  $-\text{NH}_2$ ;

R<sup>2a</sup> and R<sup>4a</sup> independently of one another represent H,  $-\text{CO}-\text{CHY}^4-\text{NHY}^5$  or  $-(\text{CH}_2)_{0-3}\text{Z}$ , or R<sup>2a</sup> and R<sup>4a</sup> together (with formation of a pyrrolidine ring) represent  $-\text{CH}_2-\text{CHR}^{10}-$  or  $-\text{CHR}^{10}-\text{CH}_2-$ , where R<sup>10</sup> represents H, SO<sub>3</sub>H, NH<sub>2</sub>, COOH, SH or OH,

where Z represents -H, halogen,  $-\text{OY}^3$ ,  $-\text{SY}^3$ ,  $-\text{NHY}^3$ ,  $-\text{CO}-\text{NY}^1\text{Y}^2$  or  $-\text{CO}-\text{OY}^3$ ,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , and Y<sup>3</sup> represents H or  $-(\text{CH}_2)_{0-3}\text{Z}'$ , where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by  $-\text{NHCONH}_2$ , or represents aryl or benzyl which are optionally substituted by  $-\text{NH}_2$ , and Y<sup>5</sup> represents H or  $-\text{CO}-\text{CHY}^6-\text{NH}_2$ , where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

R<sup>3a</sup> represents an optionally substituted alkyl, aryl, heteroaryl, heteroalkyl or heterocycloalkyl group,

preferably a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl- or C<sub>5-10</sub>-heterocycloalkyl group,

which may be substituted by 1-3 -OH groups, 1-3 halogen atoms,

1-3 halogenated alkyl groups (which may each have 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -  
 5  
 10 (CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH

(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl);

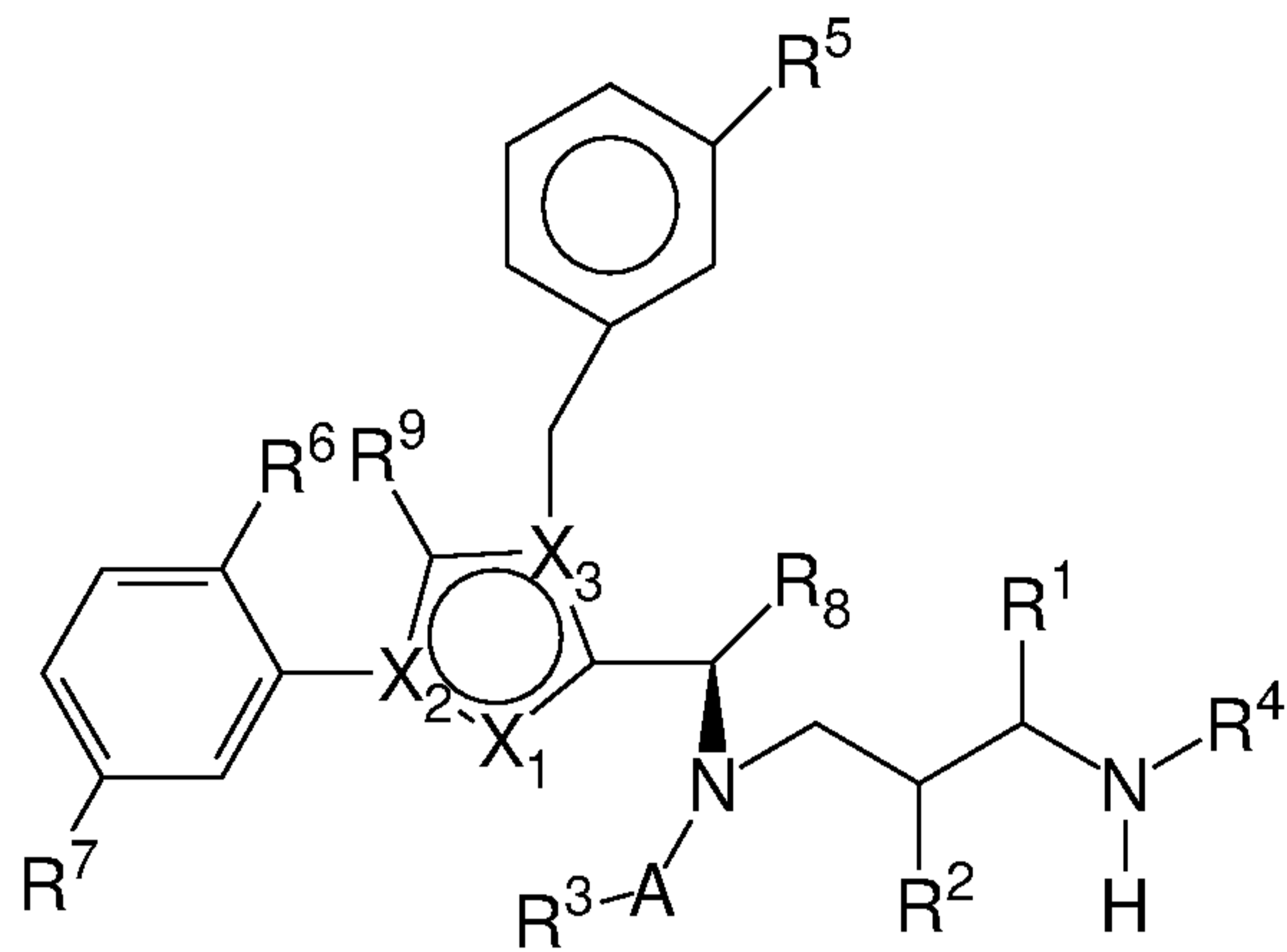
15 R<sup>8a</sup> represents C<sub>1-10</sub>-alkyl;

HZ represents a mono- or bicyclic heterocycle which may be substituted by one or more substituents selected from the group consisting of halogen, C<sub>1-10</sub>-alkyl groups, C<sub>6-10</sub>-aryl groups and  
 20 C<sub>6-10</sub>-aralkyl groups which may optionally be substituted by halogen;

where the kinesin spindle protein inhibitor is attached to the linker by substitution of a hydrogen atom at R<sup>1a</sup>, R<sup>2a</sup>, R<sup>3a</sup>, R<sup>4a</sup>,  
 25 R<sup>8a</sup> or at the pyrrolidine ring formed by R<sup>2a</sup> and R<sup>4a</sup>,

and the salts, solvates and salts of the solvates thereof.

14. Conjugate according to one or more of the preceding items  
 30 where the active compound molecule linker is represented by general formula (II):



(II)

where

5 X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C; or

X<sub>1</sub> represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represents N; or

X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C; or

10

X<sub>1</sub> represents CH, X<sub>2</sub> represents N and X<sub>3</sub> represents C

R<sup>1</sup> represents H, -L-#1 or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

15

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where W represents H or OH, where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>;

20

25 R<sup>2</sup> and R<sup>4</sup> independently of one another represent -L-#1, H, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents L-#1, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH or OH,

where Z represents -H, OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

A represents CO, SO, SO<sub>2</sub>, SO<sub>2</sub>NH or CNNH;

R<sup>3</sup> represents -L-#1 or an optionally substituted alkyl, aryl, heteroaryl, heteroalkyl or heterocycloalkyl group,

preferably -L-#1 or a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl, C<sub>6-10</sub>-aralkyl, C<sub>5-10</sub>-heteroalkyl, C<sub>1-10</sub>-alkyl-O-C<sub>6-10</sub>-aryl- or C<sub>5-10</sub>-heterocycloalkyl group,

which may be substituted by 1-3 -OH groups, 1-3 halogen atoms, 1-3 halogenated alkyl groups (which may each have 1-3 halogen atoms), 1-3 O-alkyl groups, 1-3 -SH groups, 1-3 -S-alkyl groups, 1-3 -O-CO-alkyl groups, 1-3 -O-CO-NH-alkyl groups, 1-3 -NH-CO-alkyl groups, 1-3 -NH-CO-NH-alkyl groups, 1-3 -S(O)<sub>n</sub>-alkyl groups, 1-3 -SO<sub>2</sub>-NH-alkyl groups, 1-3 -NH-alkyl groups, 1-3 -N(alkyl)<sub>2</sub> groups, 1-3 -NH<sub>2</sub> groups or 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z groups, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H, -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH

(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl);

R<sup>5</sup> represents -L-#1, H, F, NH<sub>2</sub>, NO<sub>2</sub>, halogen, SH or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, OY<sup>3</sup>, -SY<sup>3</sup>, halogen, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -

CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub> or  
 -(CH<sub>2</sub>)<sub>0-3</sub>Z', and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z', where Z'  
 5 represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where one of the substituents R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup> and R<sup>5</sup> represents -  
 L-#1,

10 L represents the linker and #1 represents the bond to the binder  
 or derivative thereof,

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano,  
 (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated)  
 15 C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy or  
 halogen,

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally  
 fluorinated) C<sub>4-10</sub>-cycloalkyl or optionally substituted oxetane;  
 20 and

R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

and the salts, solvates and salts of the solvates thereof.

25

15. Conjugate according to Item 14

where R<sup>1</sup> represents -L-#1, H, -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H,  
 -CO-NY<sup>1</sup>Y<sup>2</sup>, NHY<sup>3</sup>, OY<sup>3</sup>, -SY<sup>3</sup> or -CO-OY<sup>3</sup>,

30

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -  
 (CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup> represents H or -  
 (CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, NH<sub>2</sub>, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-  
 CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, where W represents H or OH,

35

where Y<sup>4</sup> independently of one another represents straight-chain  
 or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -  
 NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally  
 substituted by -NH<sub>2</sub>;

R<sup>2</sup> and R<sup>4</sup> independently of one another represent -L-#1, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> or H or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>-, where R<sup>10</sup> represents H, -L-#1, NH<sub>2</sub>,  
5 COOH, SH, OH or SO<sub>3</sub>H,

where Y<sup>4</sup> independently of one another represents straight-chain or branched C<sub>1-6</sub> alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally  
10 substituted by -NH<sub>2</sub>, and Y<sup>5</sup> represents H or -CO-CHY<sup>6</sup>-NH<sub>2</sub>, where Y<sup>6</sup> represents straight-chain or branched C<sub>1-6</sub>-alkyl;

A represents CO,

15 R<sup>3</sup> represents -(CH<sub>2</sub>)OH or -L-#1, and

R<sup>5</sup> represents -L-#1 or H,

where one of the substituents R<sup>1</sup>, R<sup>2</sup>, R<sup>3</sup>, R<sup>4</sup> and R<sup>5</sup> represents -  
20 L-#1.

16. Conjugate according to Item 14 or 15

where R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, C<sub>1-3</sub>-  
25 alkyl or halogen.

17. Conjugate according to one or more of Items 14 to 16

where R<sup>8</sup> represents C<sub>1-4</sub>-alkyl (preferably tert-butyl).  
30

18. Conjugate according to one or more of Items 14 to 17

where R<sup>9</sup> represents H.

35 19. Conjugate according to one or more of Items 14 to 18 where R<sup>6</sup> and R<sup>7</sup> represent F.

20. Conjugate according to one or more of the preceding items

where the linker -L- has one of the basic structures (i) to (iv) below:

- (i)  $-(CO)_m-SG1-L1-L2-$
- 5 (ii)  $-(CO)_m-L1-SG-L1-L2-$
- (iii)  $-(CO)_m-L1-L2-$
- 10 (iv)  $-(CO)_m-L1-SG-L2$

where m is 0 or 1, SG and SG1 are *in vivo* cleavable groups, L1 independently of one another represent organic groups not cleavable *in vivo*, and L2 represents a coupling group to the binder.

15

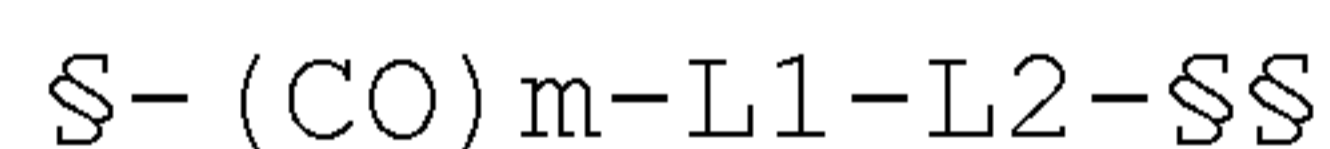
21. Conjugate according to Item 20 where the *in vivo* cleavable group SG is a 2-8 oligopeptide group, preferably a dipeptide group or a disulphide, a hydrazone, an acetal or an aminal and SG1 is a 2-8 oligopeptide group, preferably a dipeptide group.

20

22. Conjugate according to one or more of Items 2 to 21

where the linker is attached to a cysteine side chain or a cysteine residue and has the formula below:

25



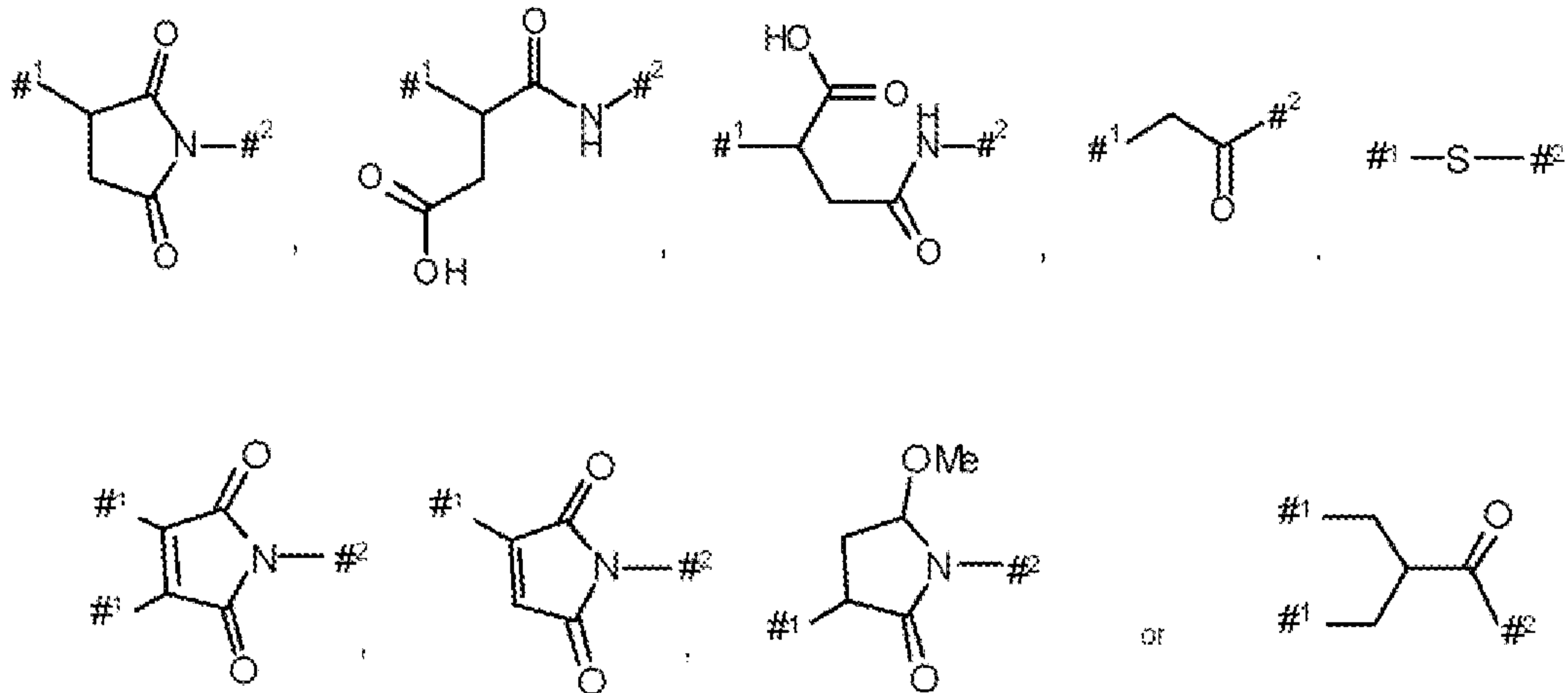
where

30 m is 0 or 1;

§ represents the bond to the active compound molecule and

35 §§ represents the bond to the binder peptide or protein, and

-L2- represents



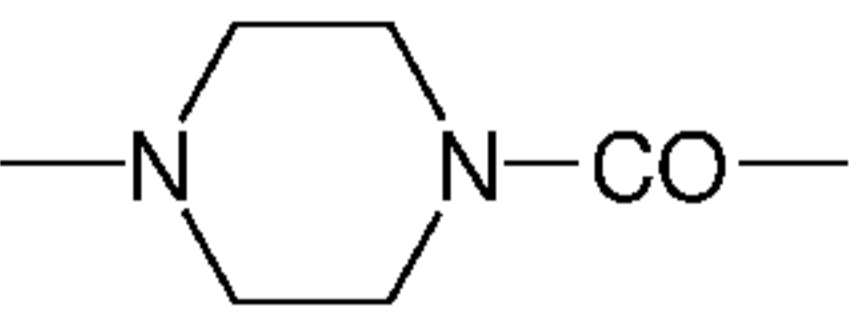
where

5 #<sup>1</sup> denotes the point of attachment to the sulphur atom of the binder,

#<sup>2</sup> denotes the point of attachment to group L<sup>1</sup>,

10 L<sup>1</sup> represents  $-(NR^{10})_n-(G1)_o-G2-$ ,

where R<sup>10</sup> represents H, NH<sub>2</sub> or C<sub>1</sub>-C<sub>3</sub>-alkyl;

G1 represents  $-NHCO-$  or  ;

15

n is 0 or 1;

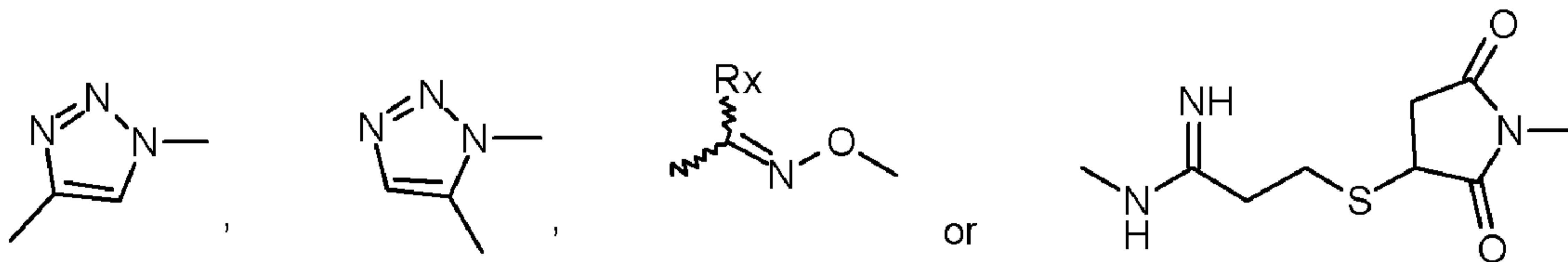
o is 0 or 1; and

20 G2 represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups  $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-NMe-$ ,  $-NHNH-$ ,  $-SO_2NHNH-$ ,  
 25  $-NHCO-$ ,  $-CONH-$ ,  $-CONHNH-$  and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S,  $-SO-$  or  $-SO_2-$  (preferably



$\text{—N—} \begin{array}{c} \diagup \\ \diagdown \end{array} \text{N—CO—}$ 
), where the side chains, if present, may be substituted by  $\text{—NHCONH}_2$ ,  $\text{—COOH}$ ,  $\text{—OH}$ ,  $\text{—NH}_2$ ,  $\text{NH—CNNH}_2$ , sulphonamide, sulphone, sulfoxide or sulphonic acid,

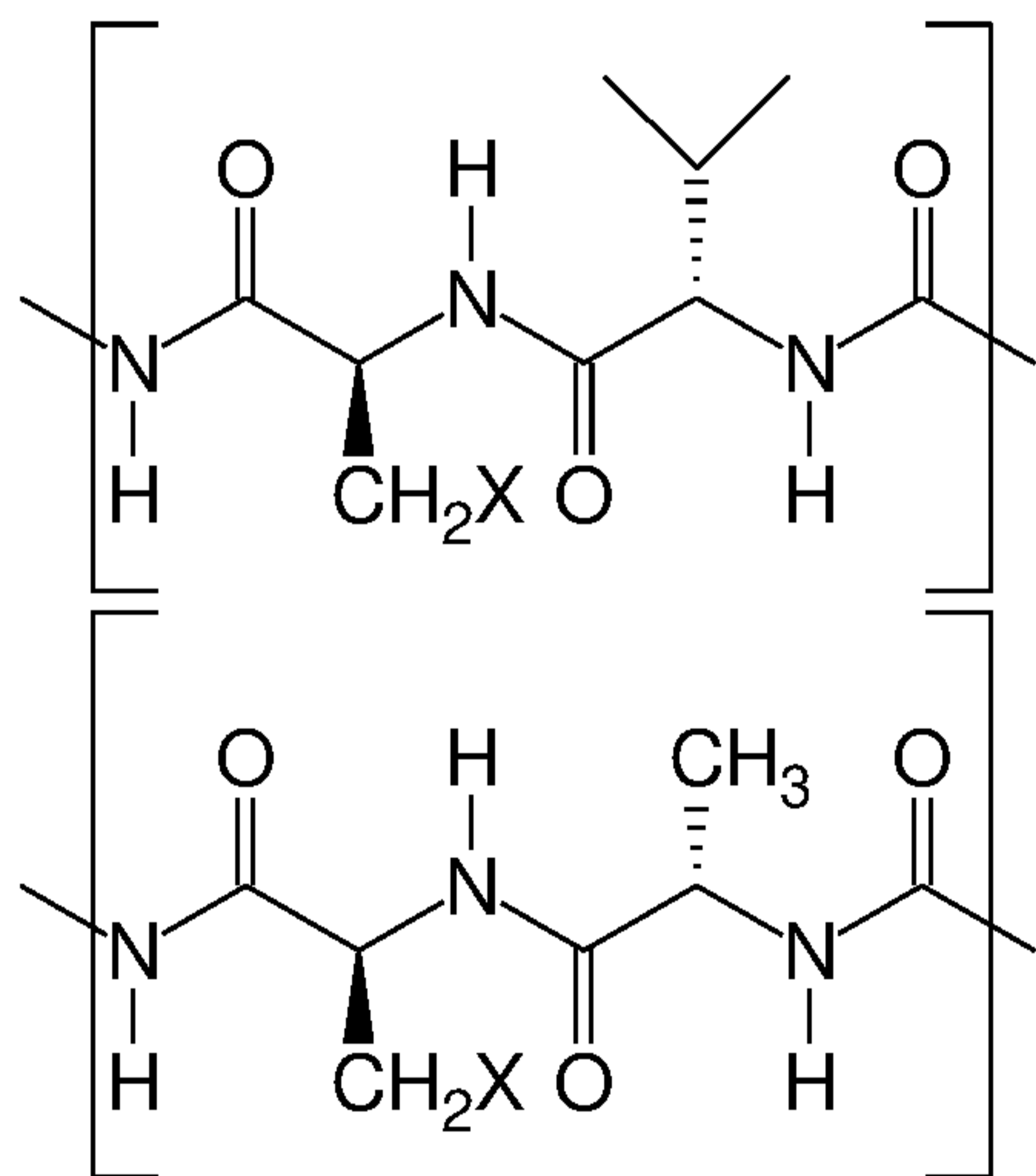
5 or represents one of the groups below:



where Rx represents H,  $\text{C}_1\text{—C}_3\text{—alkyl}$  or phenyl.

10

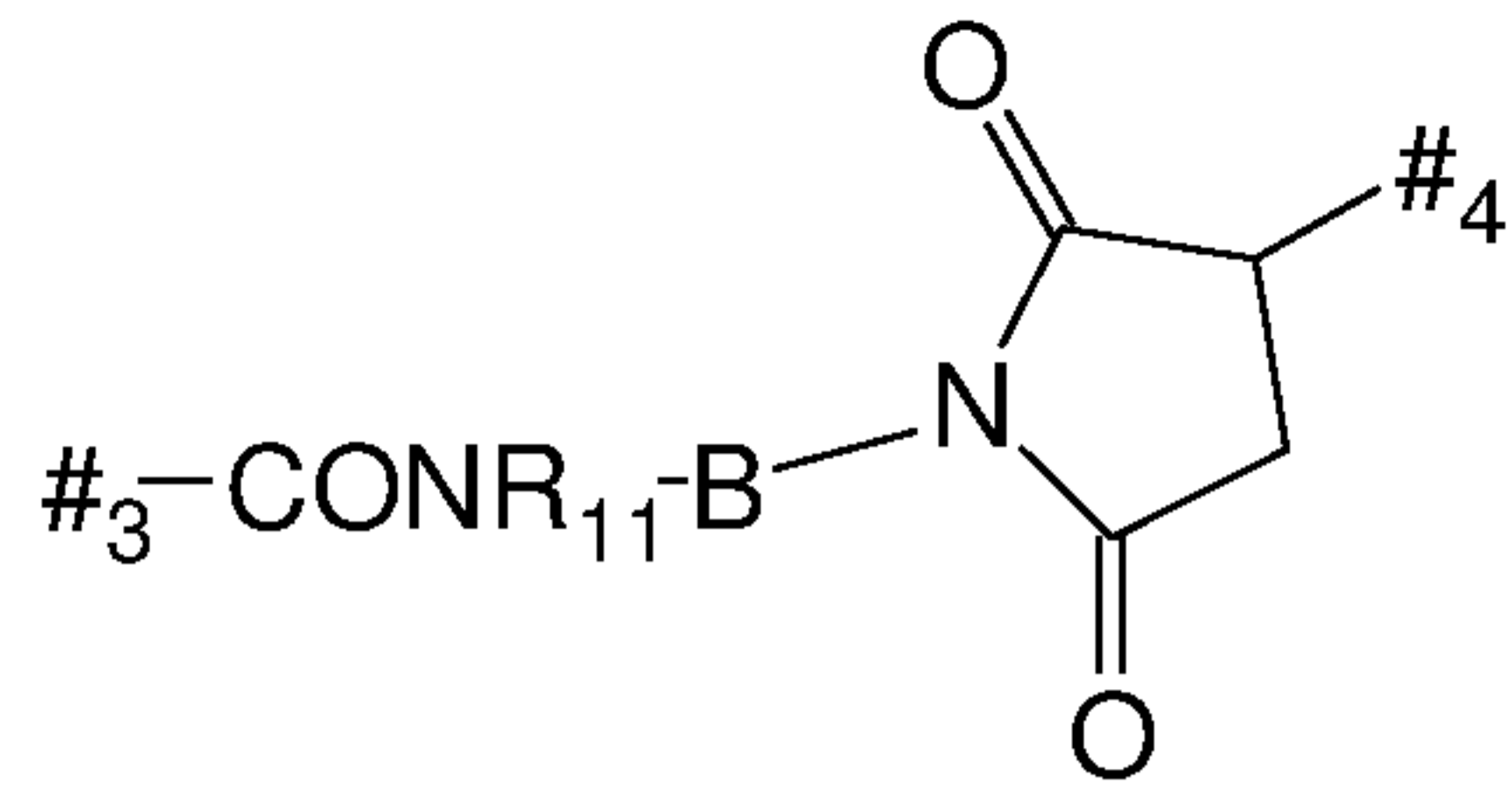
23. Conjugate according to Item 22 where the hydrocarbon chain is interrupted by one of the groups below:



15

where X represents H or a  $\text{C}_1\text{—C}_{10}\text{—alkyl}$  group which may optionally be substituted by  $\text{—NHCONH}_2$ ,  $\text{—COOH}$ ,  $\text{—OH}$ ,  $\text{NH}_2$ ,  $\text{—NH—CNNH}_2$ , sulphone, sulfoxide or sulphonic acid.

20 24. Conjugate according to Item 22 where the linker has the formula below:



where

5 #3 represents the bond to the active compound molecule,

#4 represents the bond to the binder peptide or protein,

R<sup>11</sup> represents H or NH<sub>2</sub>;

10

B represents  $-(\text{CH}_2)_x-(\text{X}^4)_y)_w-(\text{CH}_2)_z-$ ,

w = 0 to 20;

15 x = 0 to 5;

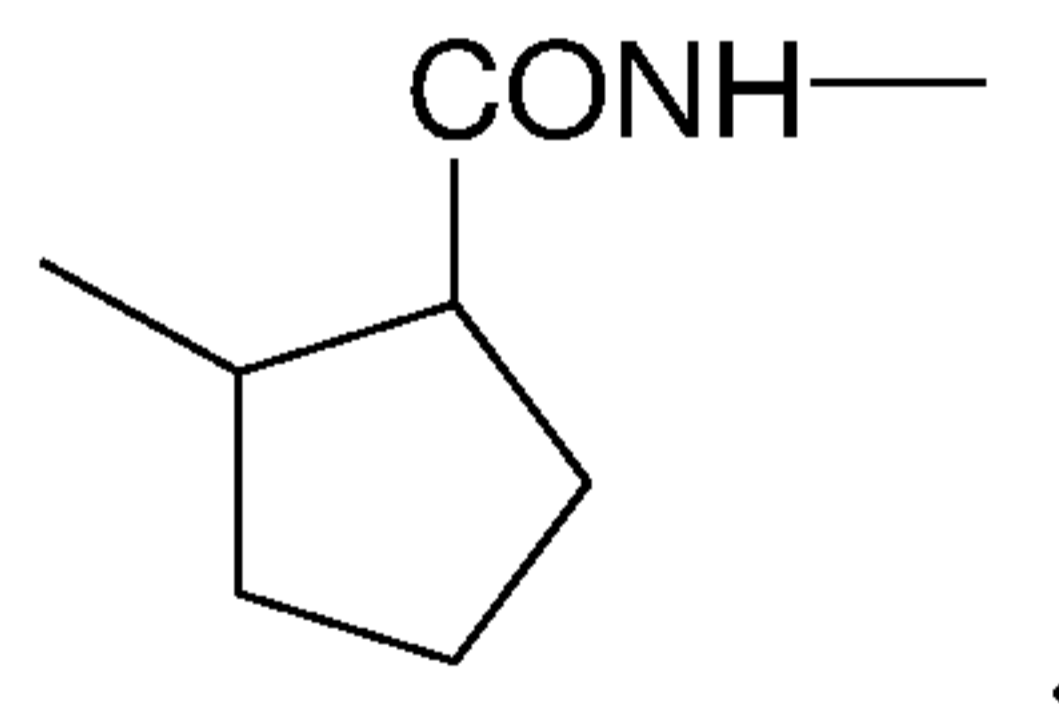
x = 0 to 5;

y = 0 or 1;

20

z = 0 to 5; and

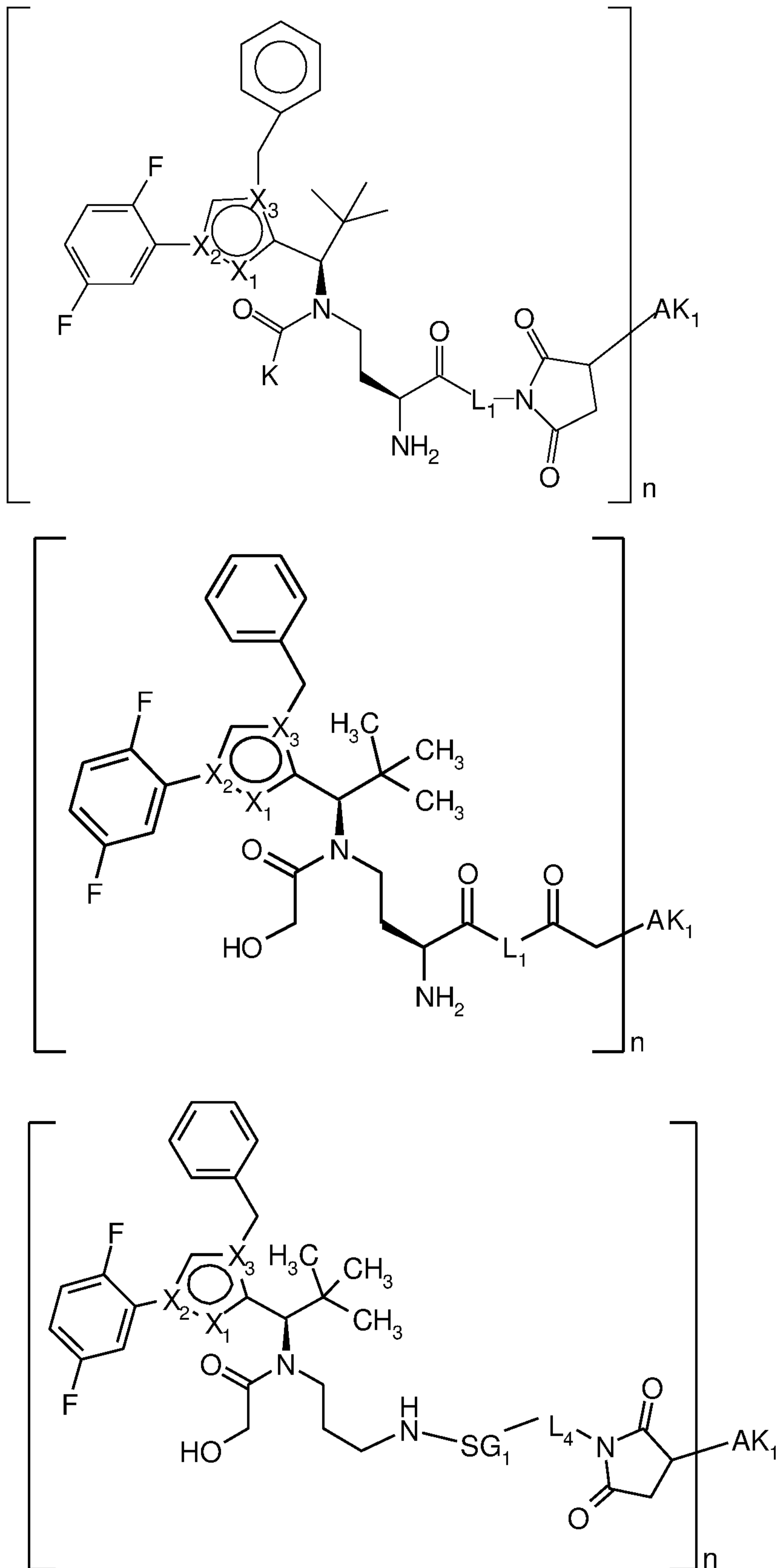
X<sup>4</sup> represents -O-, -CONH- or -NHCO-

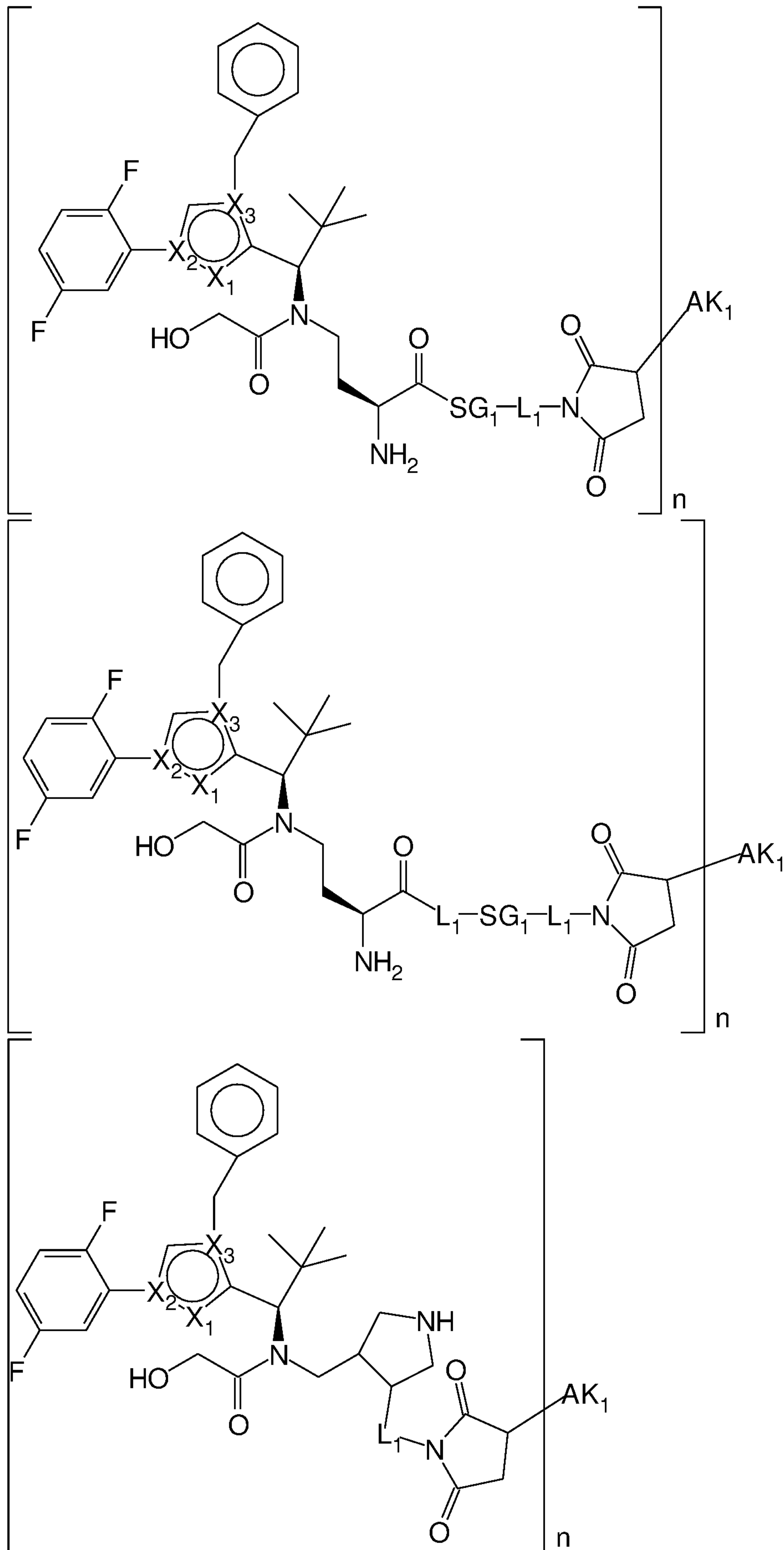


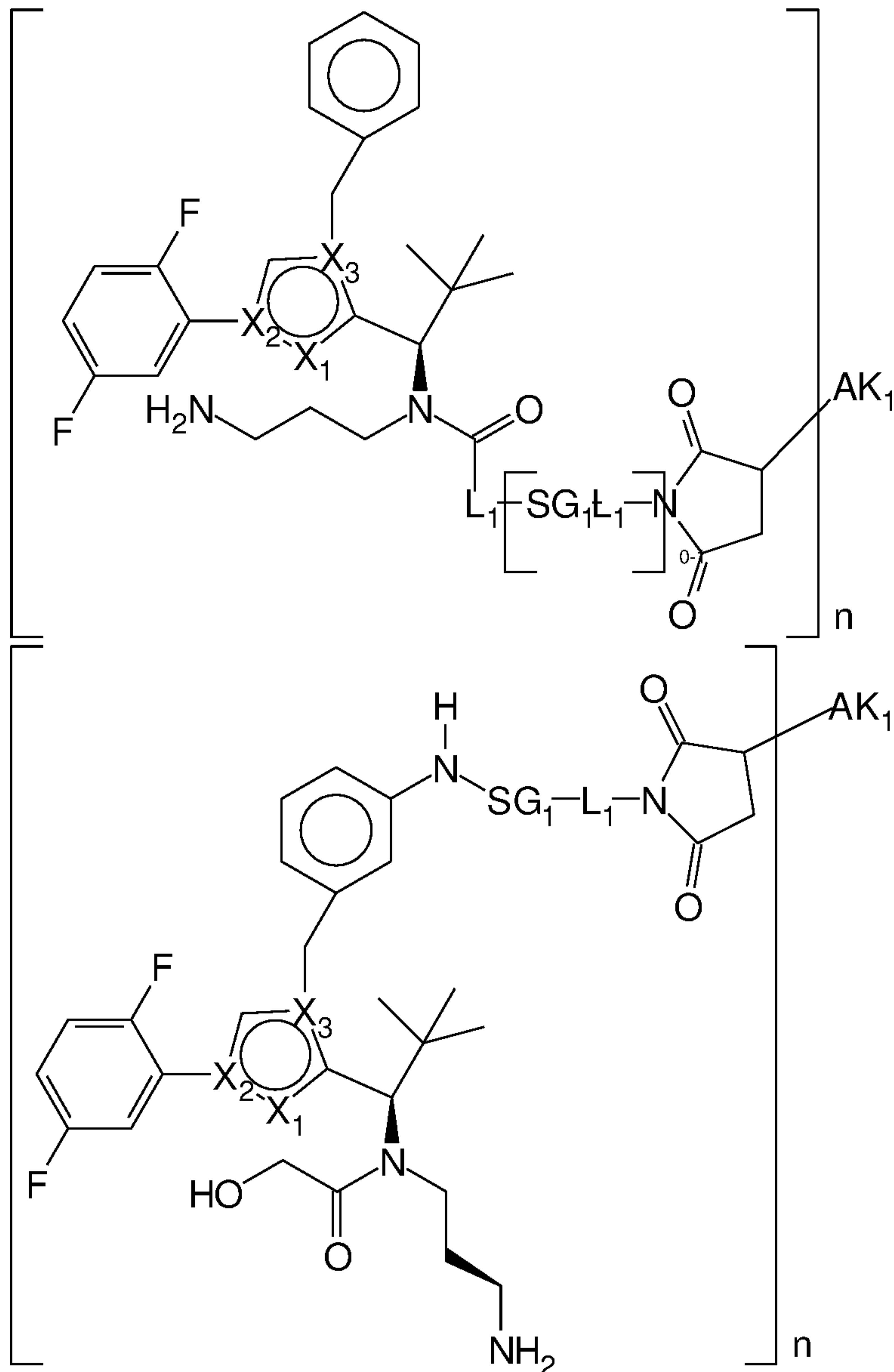
25 25. Conjugate according to Item 24 where R<sup>1</sup> or R<sup>4</sup> represents -L-#1.

26. Conjugate according to one or more of Items 21 to 25 where the conjugate has one of the formulae below:

30





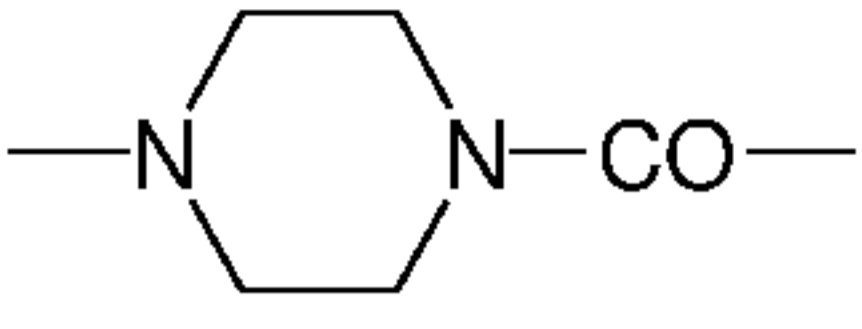


where

- 5  $X_1$ ,  $X_2$  and  $X_3$  have the same meaning as in Item 14,

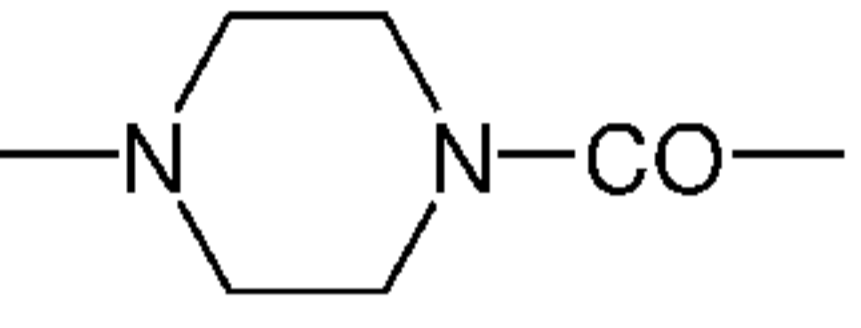
AK<sub>1</sub> represents a binder peptide or protein which is attached via a sulphur atom of the binder; n represents a number from 1 to 20; and L<sub>1</sub> represents an optionally branched hydrocarbon group having 1 to 70 carbon atoms, which represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-  
 10 , -CO-, -CONH-, -NHCO-, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- and  
 15

a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O

and S, or -SO- or -SO<sub>2</sub>- (preferably ) , where the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid,

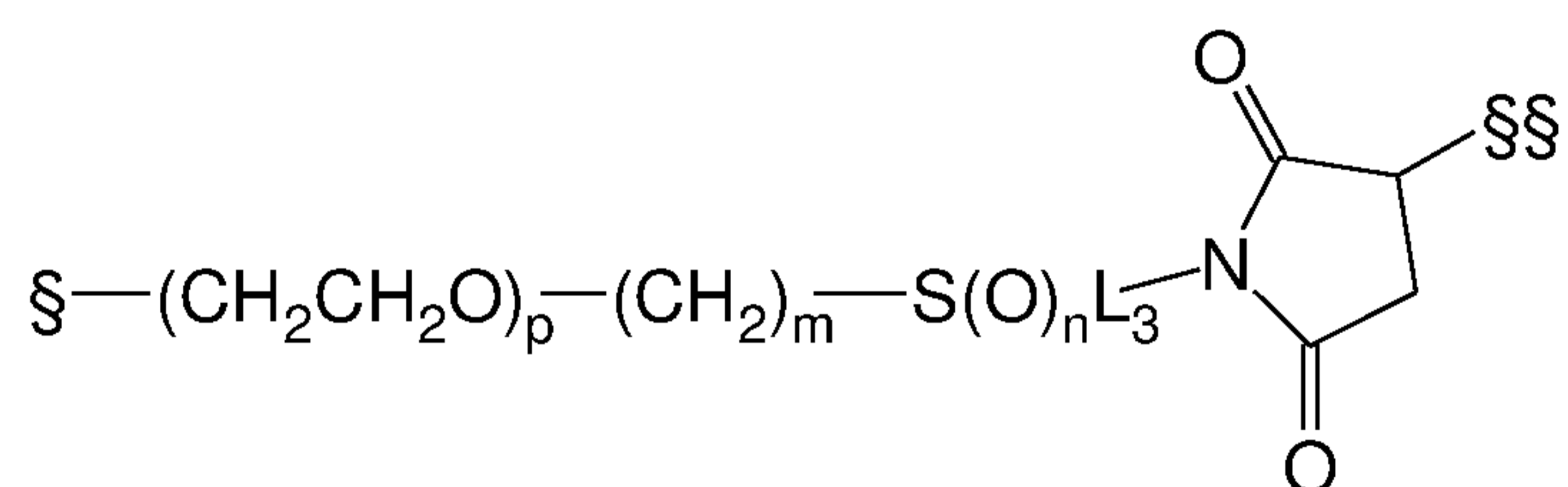
and SG1 is a 2-8 oligopeptide, preferably a dipeptide;

10 L4 is a single bond or a group -(CO)<sub>y</sub>-G4-, where y represents 0 or 1, and G4 represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more

15 of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, -SO- or -SO<sub>2</sub>- (preferably ) , where the side chains, if present,

20 may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid.

27. Conjugate according to one or more of Items 1 to 21 where the linker -L- is attached to a cysteine side chain or a cysteine residue and has the formula below:



where

30

§ represents the bond to the active compound molecule and

§§ represents the bond to the binder peptide or protein,

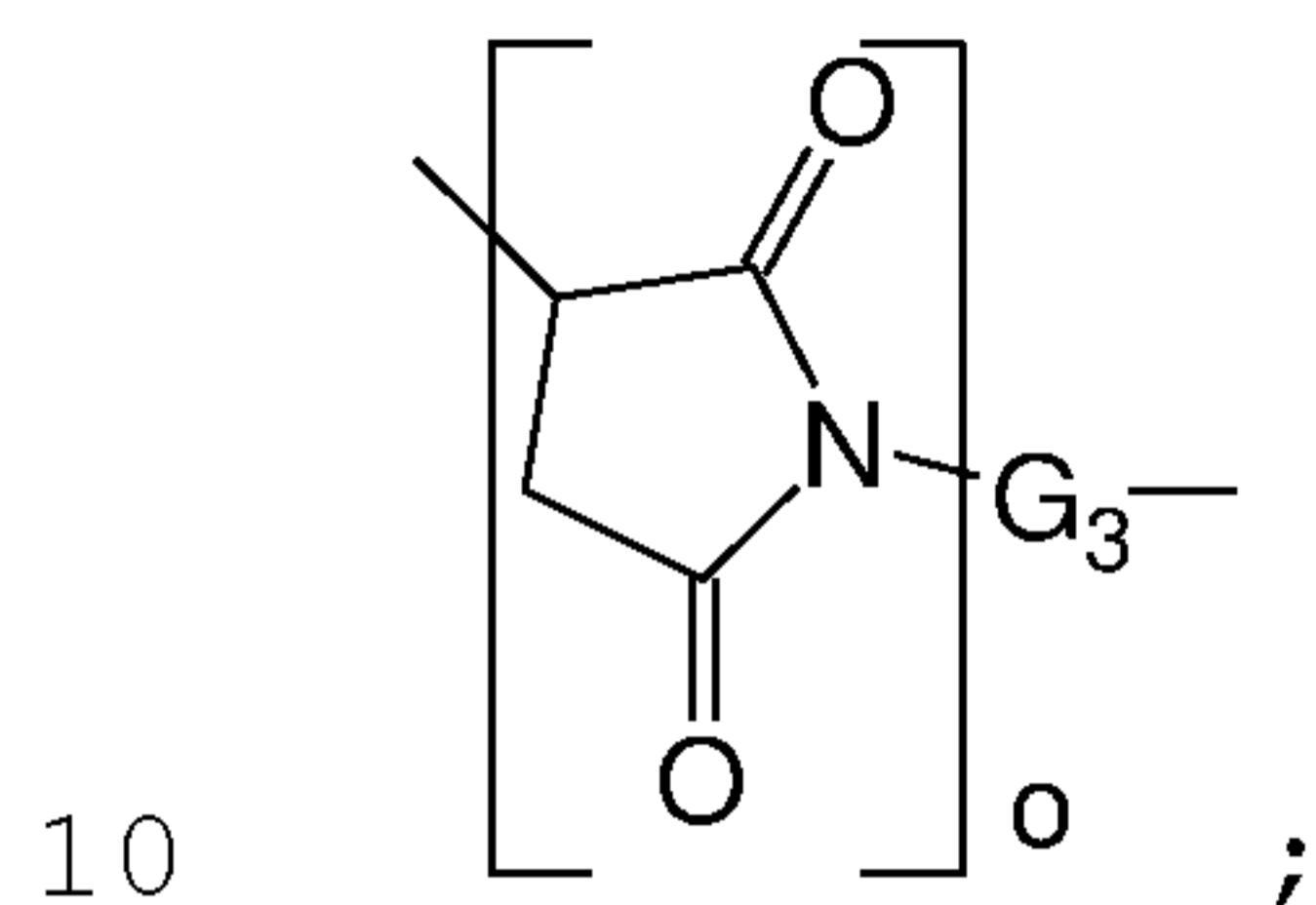
m is 0, 1, 2 or 3;

n is 0, 1 or 2;

5

p is 0 to 20; and

L3 represents



where

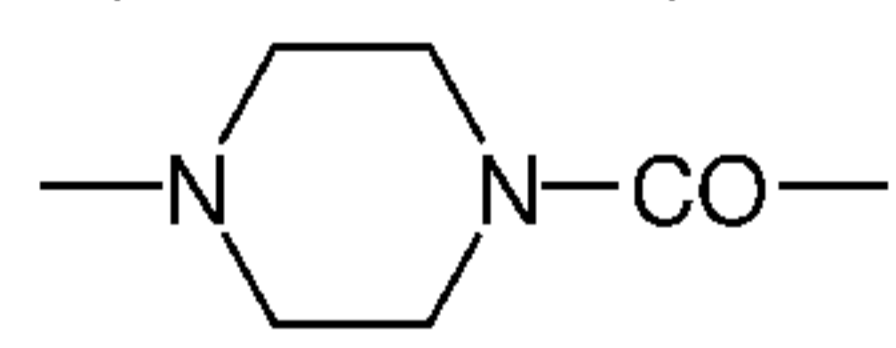
o is 0 or 1;

15

and

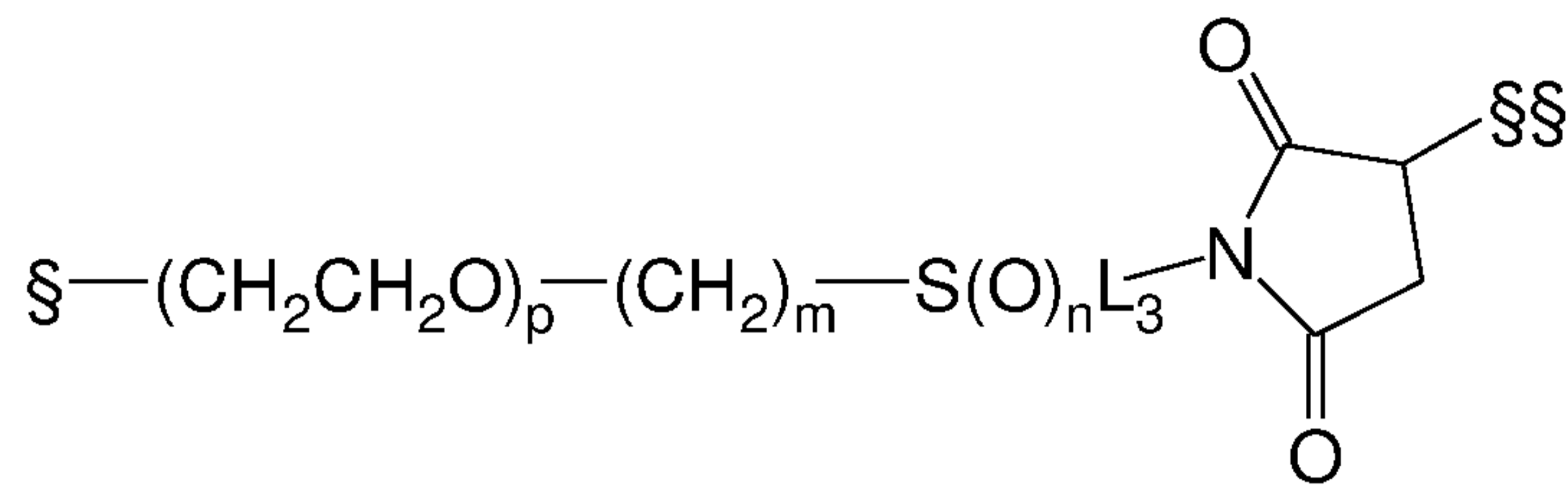
G<sub>3</sub> represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH- and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH-, -SO- or -SO<sub>2</sub>- (preferably

25



), where the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, sulphone, sulphoxide or sulphonic acid.

30 28. Conjugate according to Item 27 where the linker -L- is attached to a cysteine side chain or a cysteine residue and has the formula below:



where

5 § represents the bond to the active compound molecule and

SS represents the bond to the binder peptide or protein,

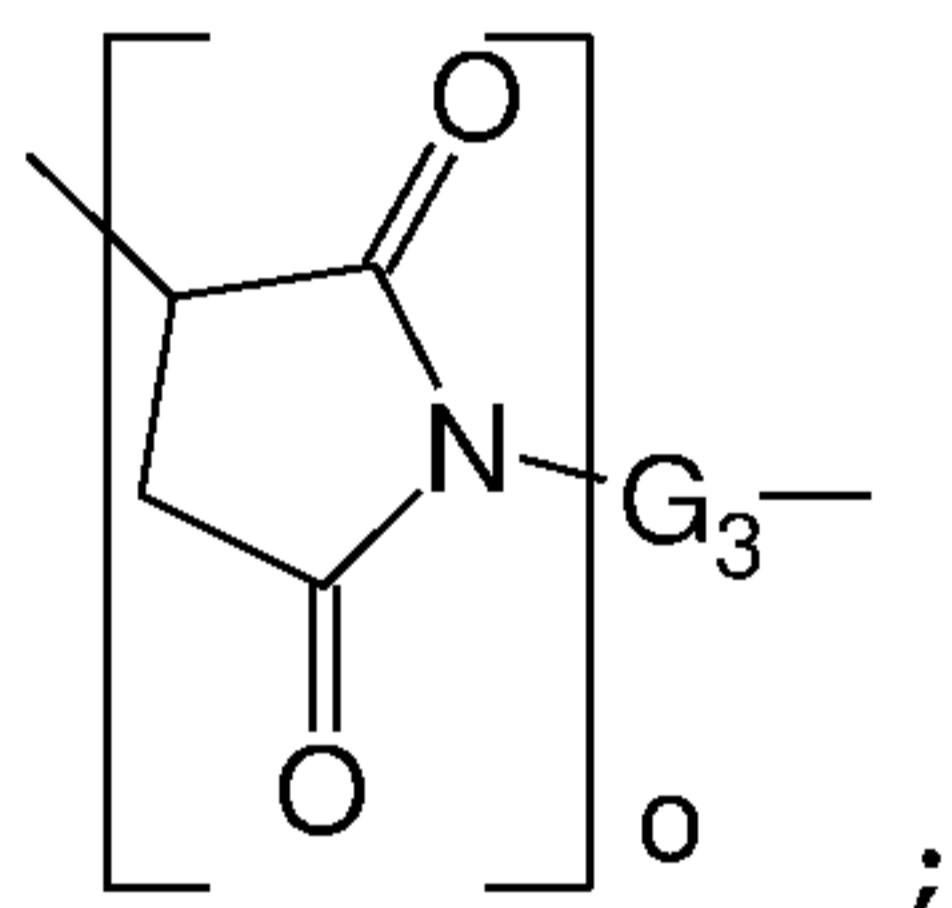
m is 1;

10

p is 0;

n is 0;

15 and L3 represents



where

20

o is 0 or 1; and

G3 represents  $-(\text{CH}_2\text{CH}_2\text{O})_s(\text{CH}_2)_t(\text{CONH})_u(\text{CH}_2\text{CH}_2\text{O})_v(\text{CH}_2)_w-$ , where

25 s, t, v and w each independently of one another are from 0 to 20 and u is 0 or 1.

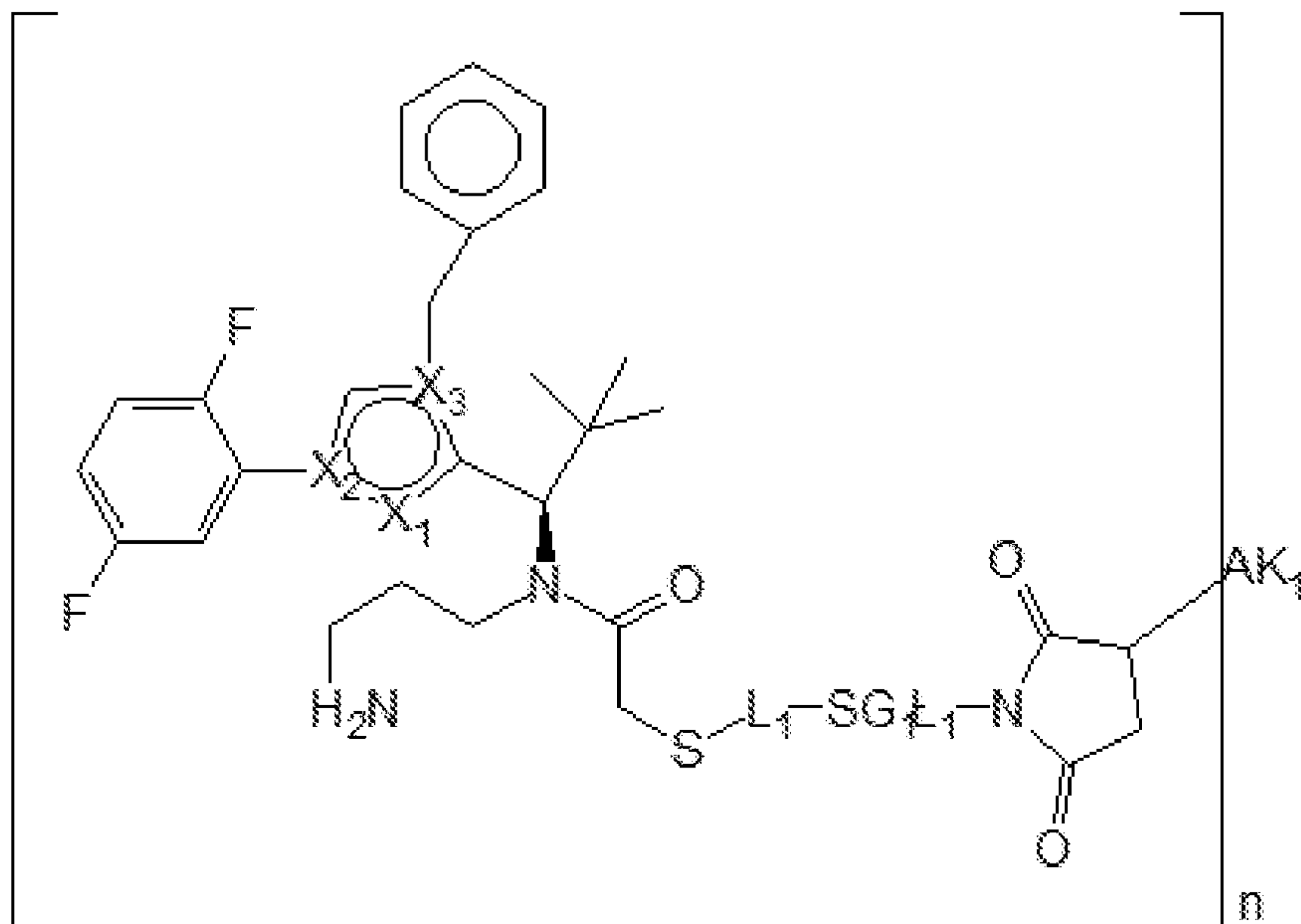
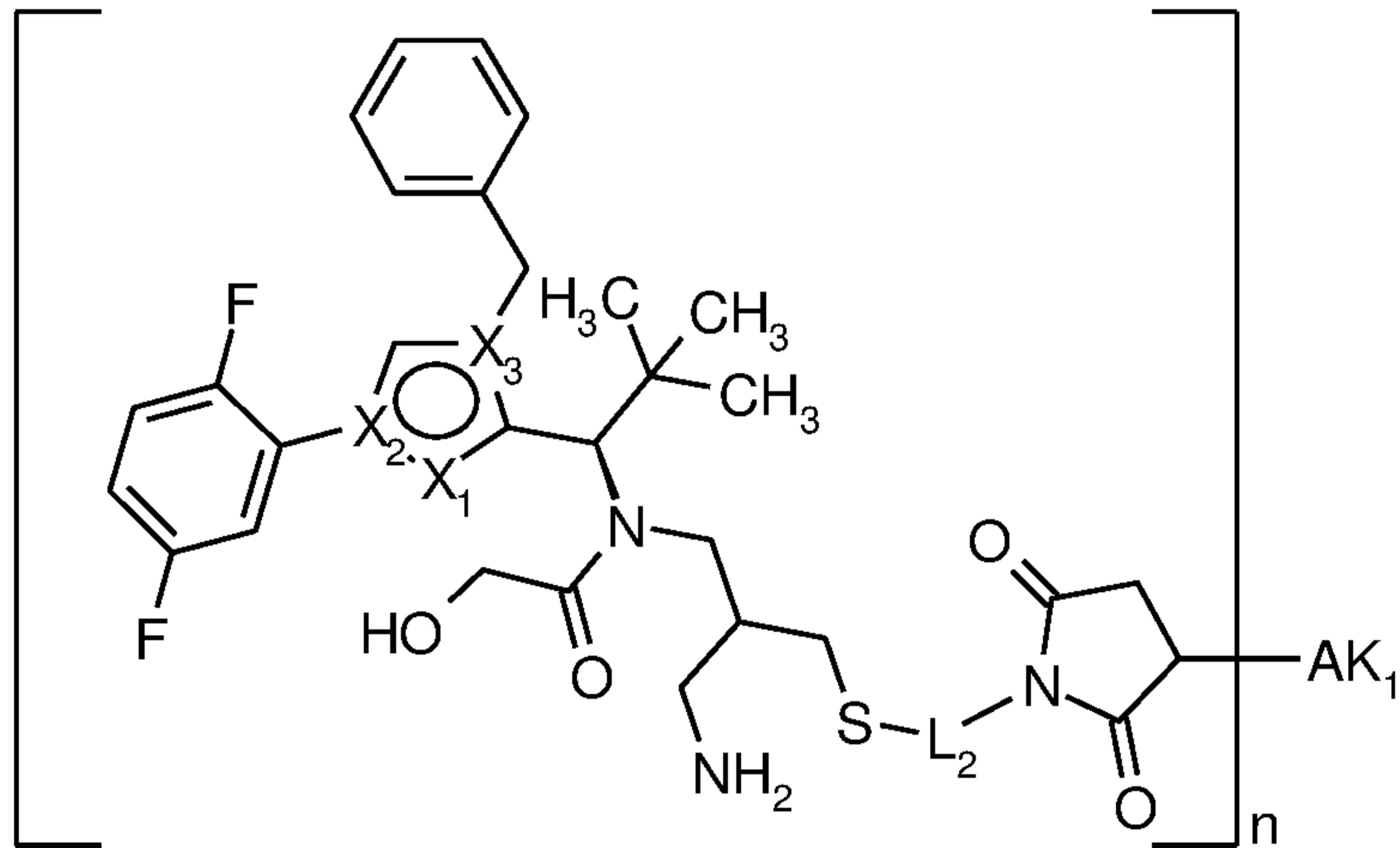
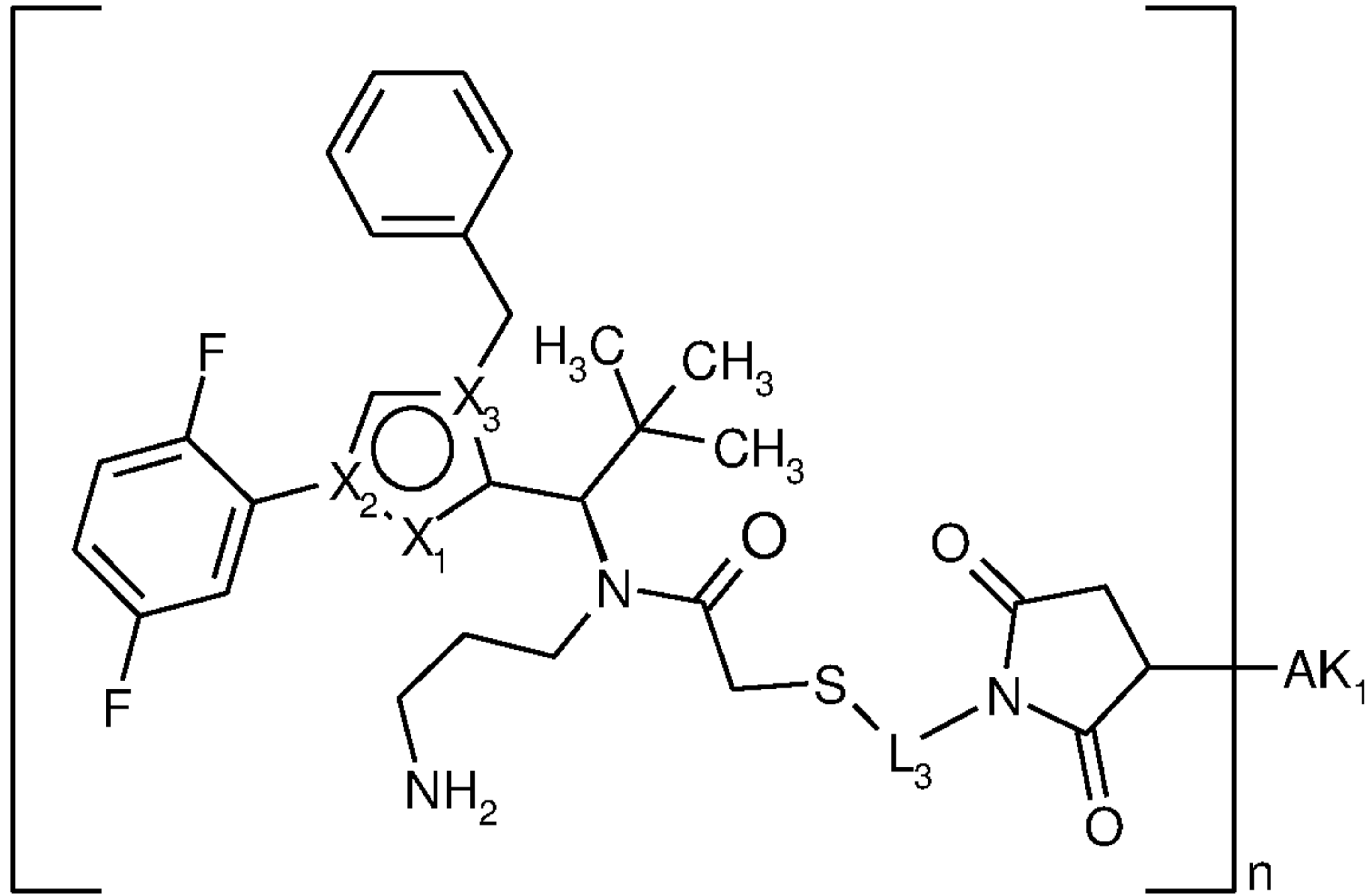
29. Conjugate according to Item 27 or 28 where R<sup>2</sup> or R<sup>3</sup> represents -L-#1.

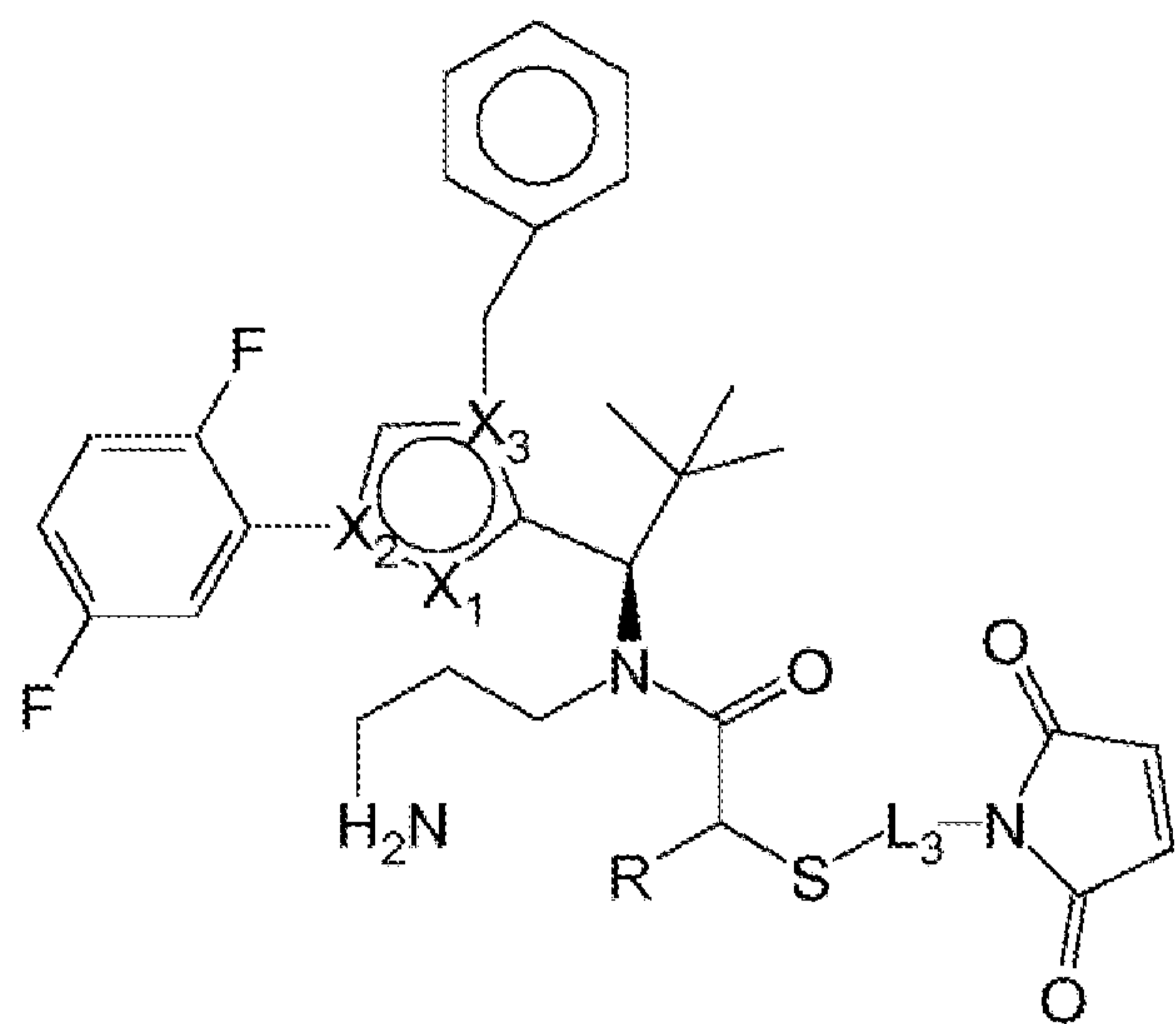
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30. Conjugate according to Item 29 where the conjugate has one



of the formulae below:





TFA

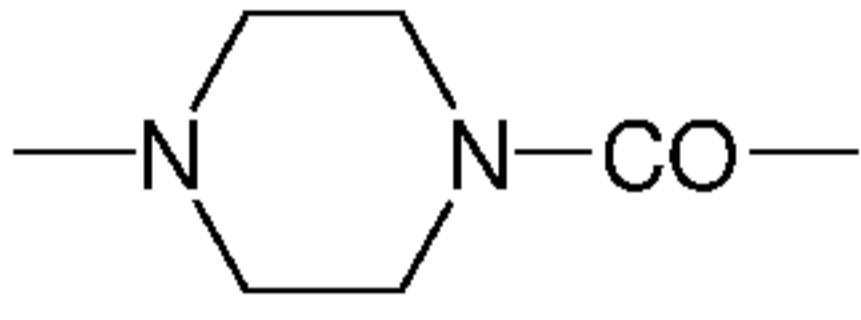


where

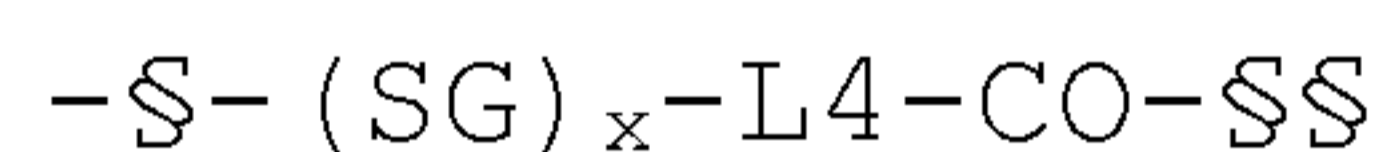
X<sub>1</sub>, X<sub>2</sub> and X<sub>3</sub> have the same meaning as in Item 14,

5

AK<sub>1</sub> represents a binder peptide or protein which is attached via a sulphur atom of the binder; n represents a number from 1 to 20; and L2 and L3 represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -NHCO-, -CONH-, -CONHNH- and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and

15 to 4 heteroatoms selected from the group consisting of N, O and S, -SO- or -SO<sub>2</sub>- (preferably ) , where the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid.

20 31. Conjugate according to one or more of Items 1 to 21 where the linker -L- is attached to a lysine side chain and has the formula below:



25

where

§ represents the bond to the active compound molecule and

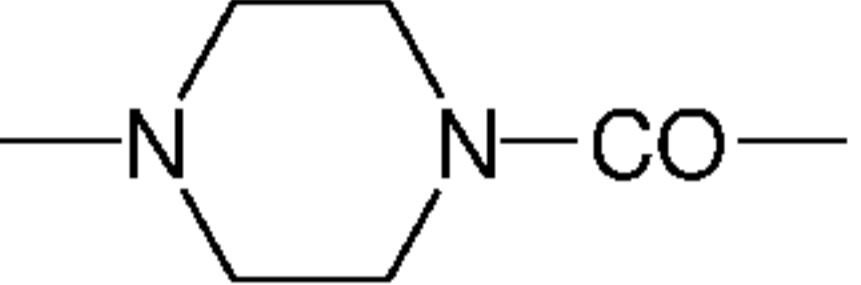
30 §§ represents the bond to the binder peptide or protein,

x represents 0 or 1,

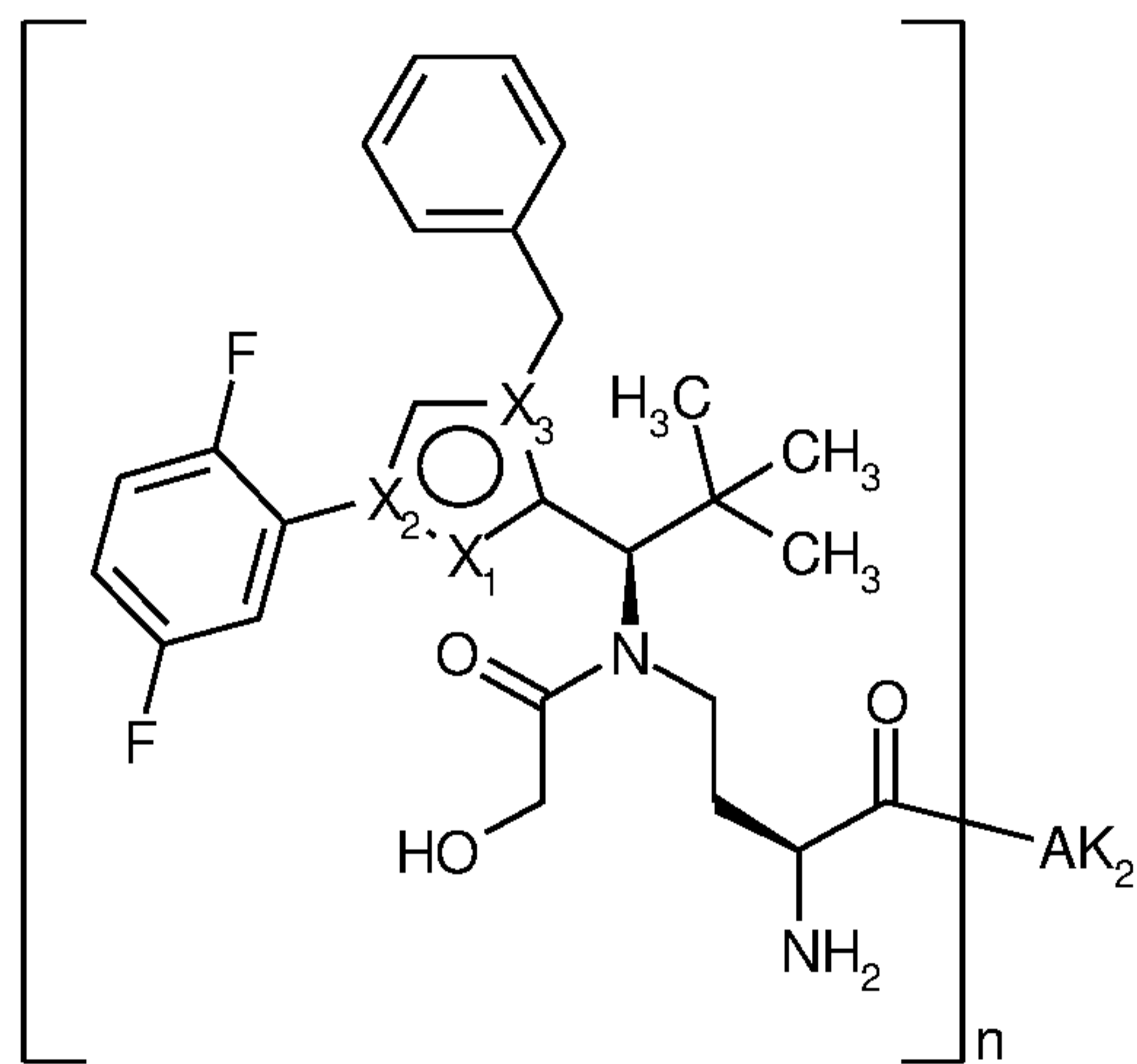
35 SG represents a cleavable group, preferably a 2-8 oligopeptide, particularly preferably a dipeptide,

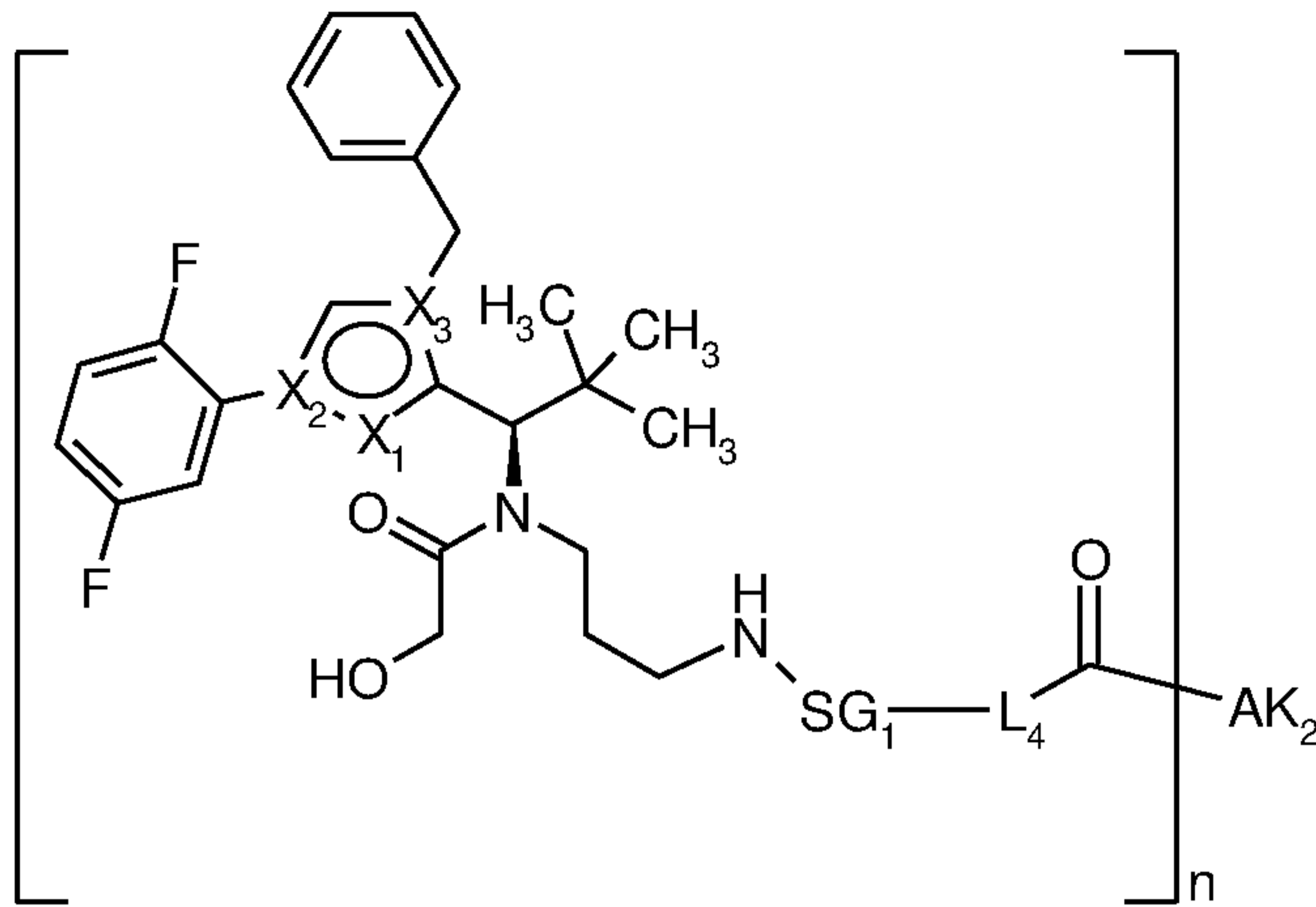
and

L4 represents a single bond or a group  $-(CO)_y-G4-$ , where  $y$  represents 0 or 1, and

- 5 G4 represents a straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups  $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-NHCO-$ ,  $-CONH-$ ,  $-NMe-$ ,  
 10  $-NHNH-$ ,  $-SO_2NHNH-$ ,  $-CONHNH-$  and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and S, or  $-SO-$  (preferably  
 $\text{---N---CO---}$  ) , where the side chains, if present, may be substituted by  $-NHCONH_2$ ,  $-COOH$ ,  $-OH$ ,  $-NH_2$ ,  $NH-CNNH_2$ ,  
 15 sulphonamide, sulphone, sulfoxide or sulphonic acid.

32. Conjugate of a binder peptide or protein according to Item 31 where the conjugate has one of the formulae below:

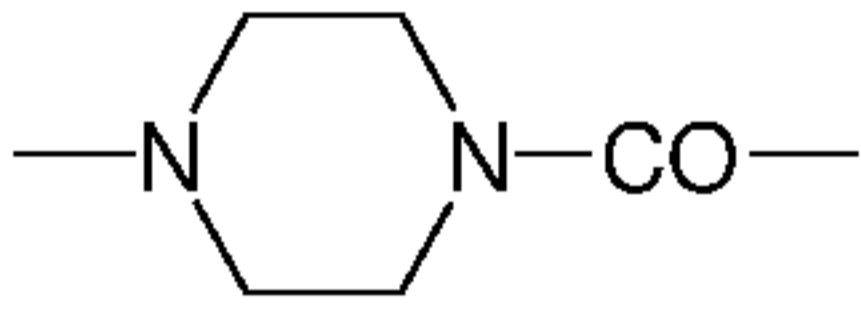




where

5 X<sub>1</sub>, X<sub>2</sub> and X<sub>3</sub> have the same meaning as in Item 14,

AK<sub>2</sub> represents a binder peptide or protein which is attached via a sulphur atom of the binder; n represents a number from 1 to 20; and L<sub>4</sub> represents an optionally straight-chain or branched hydrocarbon chain having 1 to 100 carbon atoms from arylene groups and/or straight-chain and/or branched and/or cyclic alkylene groups and which may be interrupted once or more than once by one or more of the groups -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NMe-, -NHNH-, -SO<sub>2</sub>NHNH-, -NHCO-, -CONH-, -CONHNH- and a 5- to 10-membered aromatic or non-aromatic heterocycle having up to 4 heteroatoms selected from the group consisting of N, O and

15 S, -SO- or -SO<sub>2</sub>- (preferably ) , where the side chains, if present, may be substituted by -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulphonamide, sulphone, sulphoxide or sulphonic acid, and

20 SG<sub>1</sub> represents a cleavable group, preferably a 2-8 oligopeptide, particularly preferably a dipeptide.

25 33. Conjugate according to one or more of Items 10 to 32 where the anti-TWEAKR antibody is an agonistic antibody.

34. Conjugate according to one or more of Items 10 to 33 which

comprises:

a variable heavy chain comprising:

- 5 a. a CDR1 of the heavy chain encoded by an amino acid sequence comprising the formula PYPMX (SEQ ID NO: 171), where X is I or M;
- b. a CDR2 of the heavy chain encoded by an amino acid sequence  
10 comprising the formula YISPSGGXTHYADSVKG (SEQ ID NO: 172), where X is S or K; and
- c. a CDR3 of the heavy chain encoded by an amino acid sequence comprising the formula GGDTYFDYFDY (SEQ ID NO: 173);

15

and a variable light chain comprising:

- d. a CDR1 of the light chain encoded by an amino acid sequence comprising the formula RASQ<sup>S</sup>SISXYLN (SEQ ID NO: 174), where X is  
20 G or S;
- e. a CDR2 of the light chain encoded by an amino acid sequence comprising the formula XASSLQS (SEQ ID NO: 175), where X is Q, A or N; and
- 25 f. a CDR3 of the light chain encoded by an amino acid sequence comprising the formula QQSYXXPXIT (SEQ ID NO: 176), where X at position 5 is T or S, X at position 6 is T or S and X at position 8 is G or F.

30

35. Conjugate according to one or more of Items 10 to 34 which comprises:

- a. a variable sequence of the heavy chain, as shown in SEQ ID  
35 NO:10, and also a variable sequence of the light chain, as shown in SEQ ID NO:9, or
- b. a variable sequence of the heavy chain, as shown in SEQ ID

NO:20, and also a variable sequence of the light chain, as shown in SEQ ID NO:19, or

5 c. a variable sequence of the heavy chain, as shown in SEQ ID NO:30, and also a variable sequence of the light chain, as shown in SEQ ID NO:29, or

10 d. a variable sequence of the heavy chain, as shown in SEQ ID NO:40, and also a variable sequence of the light chain, as shown in SEQ ID NO:39, or

15 e. a variable sequence of the heavy chain, as shown in SEQ ID NO:50, and also a variable sequence of the light chain, as shown in SEQ ID NO:49, or

f. a variable sequence of the heavy chain, as shown in SEQ ID NO:60, and also a variable sequence of the light chain, as shown in SEQ ID NO:59, or

20 g. a variable sequence of the heavy chain, as shown in SEQ ID NO:70, and also a variable sequence of the light chain, as shown in SEQ ID NO:69, or

25 h. a variable sequence of the heavy chain, as shown in SEQ ID NO:80, and also a variable sequence of the light chain, as shown in SEQ ID NO:79, or

30 i. a variable sequence of the heavy chain, as shown in SEQ ID NO:90, and also a variable sequence of the light chain, as shown in SEQ ID NO:89, or

j. a variable sequence of the heavy chain, as shown in SEQ ID NO:100, and also a variable sequence of the light chain, as shown in SEQ ID NO:99, or

35 k. a variable sequence of the heavy chain, as shown in SEQ ID NO:110, and also a variable sequence of the light chain, as shown in SEQ ID NO:109, or



1. a variable sequence of the heavy chain, as shown in SEQ ID NO:120, and also a variable sequence of the light chain, as shown in SEQ ID NO:119.

5

36. Conjugate according to one or more of Items 10 to 35 where the antibody is an IgG antibody.

37. Conjugate according to one or more of Items 10 to 36 which  
10 comprises:

a. a sequence of the heavy chain, as shown in SEQ ID NO:2, and also a sequence of the light chain, as shown in SEQ ID NO:1, or

15 b. a sequence of the heavy chain, as shown in SEQ ID NO:12, and also a sequence of the light chain, as shown in SEQ ID NO:11, or

20 c. a sequence of the heavy chain, as shown in SEQ ID NO:22, and also a sequence of the light chain, as shown in SEQ ID NO:21, or

25 d. a sequence of the heavy chain, as shown in SEQ ID NO:32, and also a sequence of the light chain, as shown in SEQ ID NO:31, or

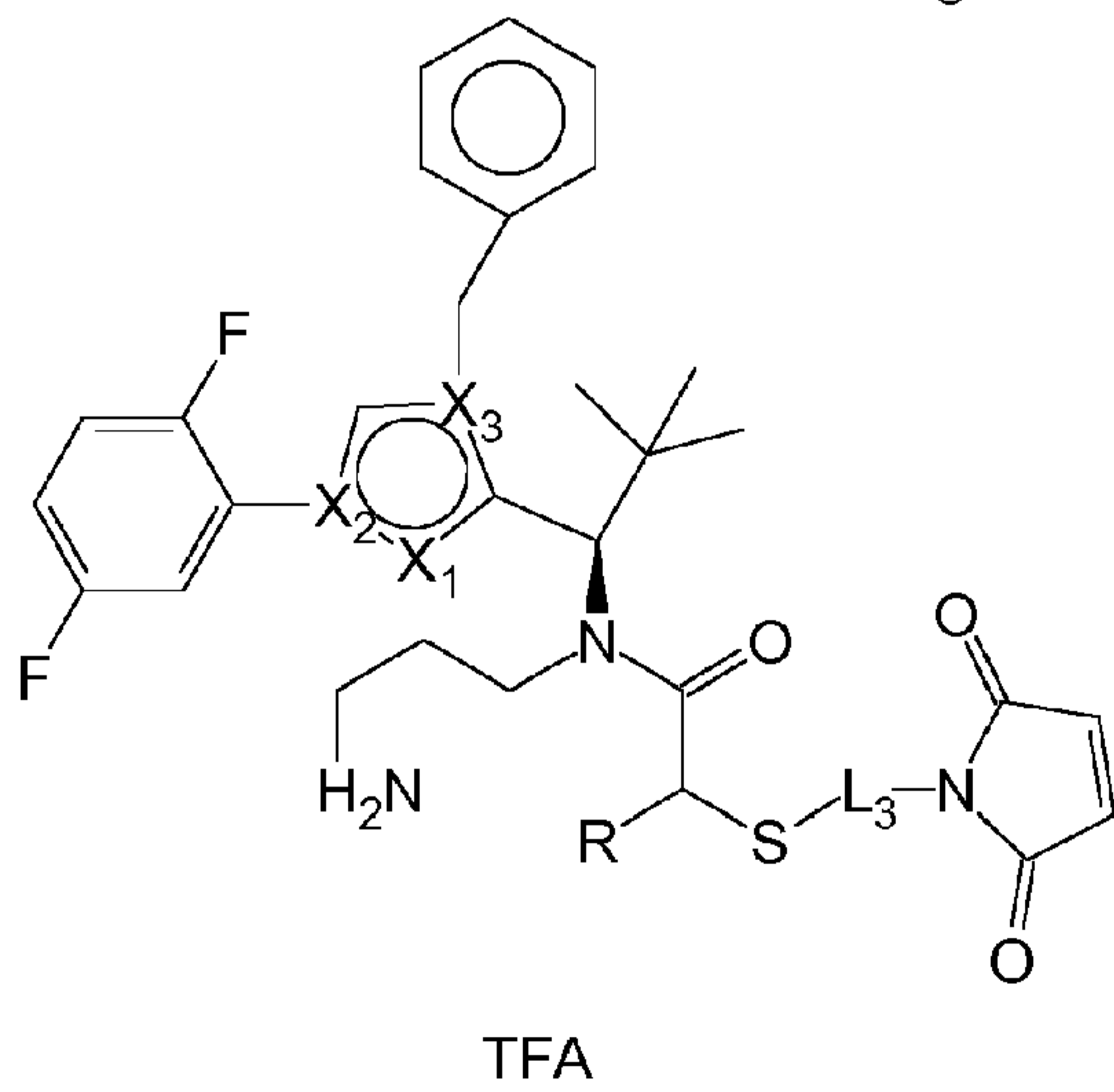
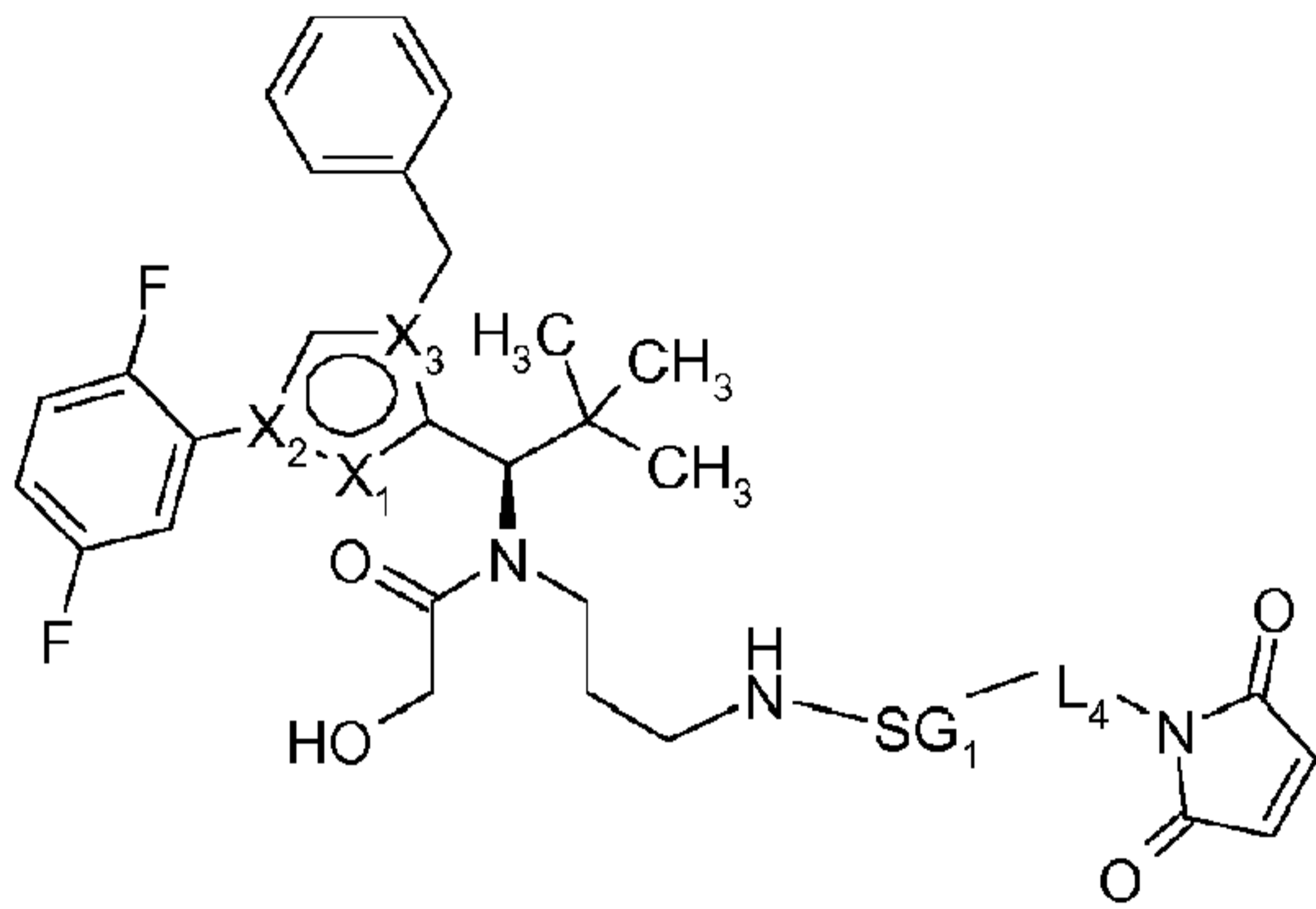
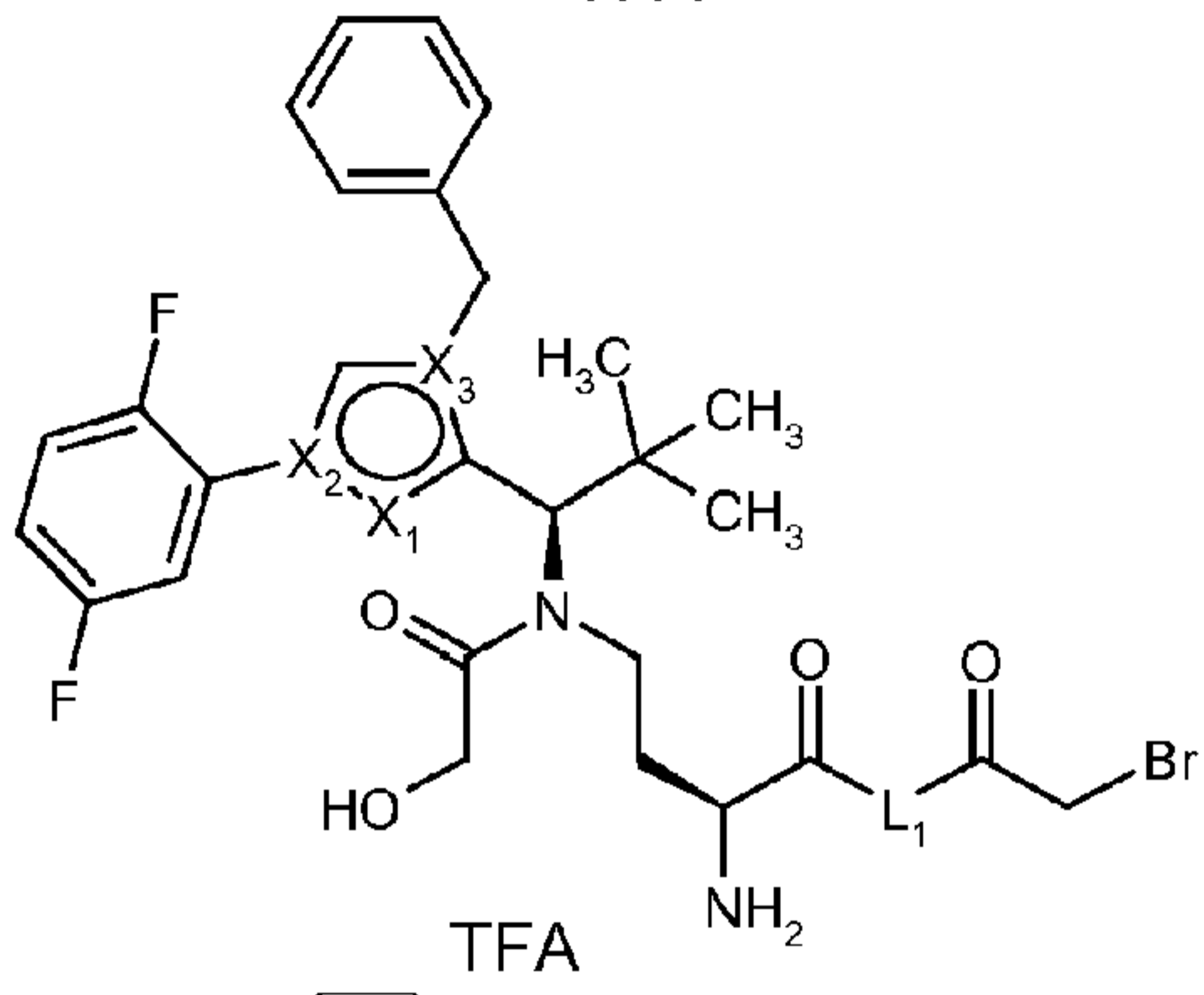
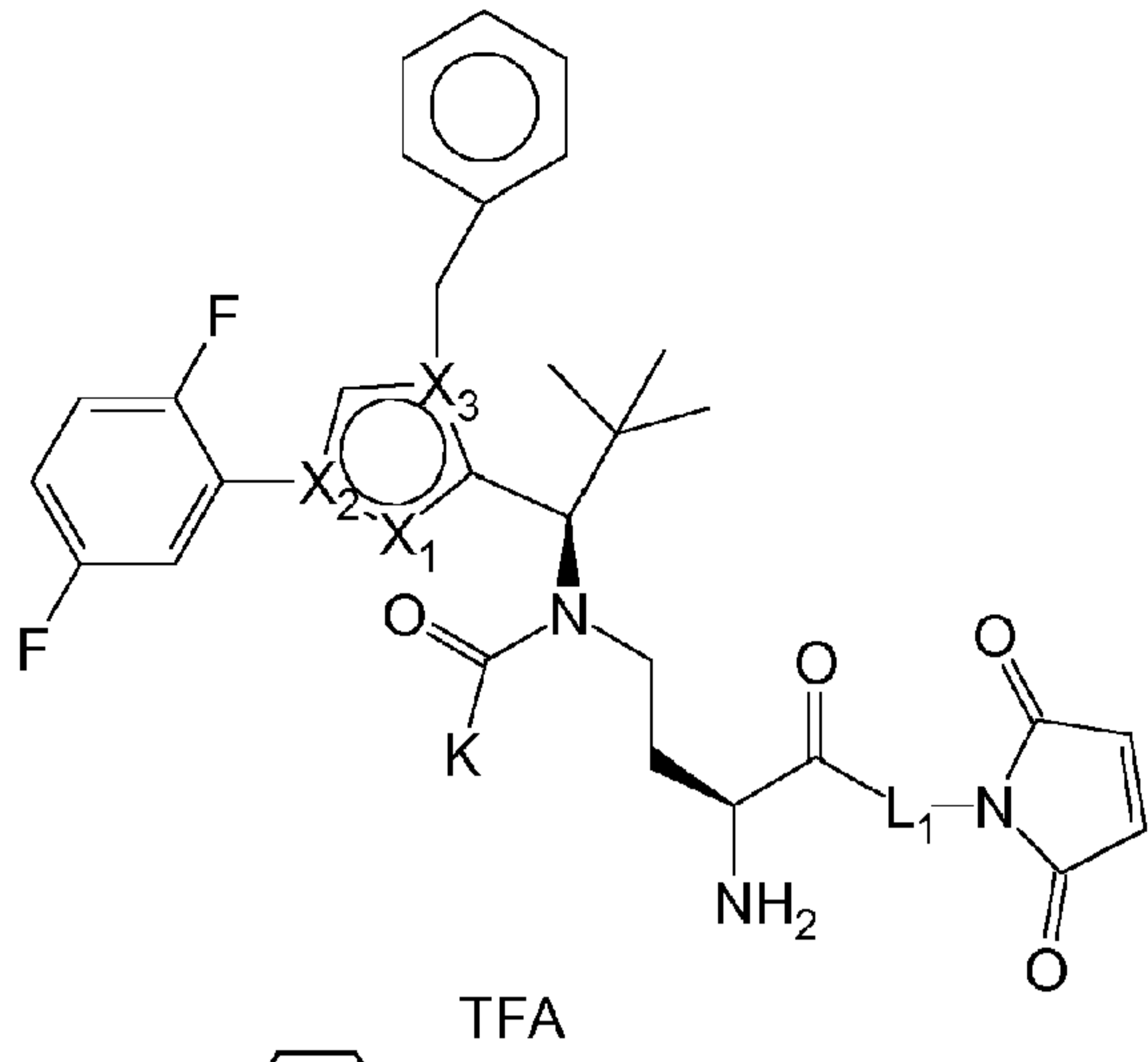
e. a sequence of the heavy chain, as shown in SEQ ID NO:42, and also a sequence of the light chain, as shown in SEQ ID NO:41, or

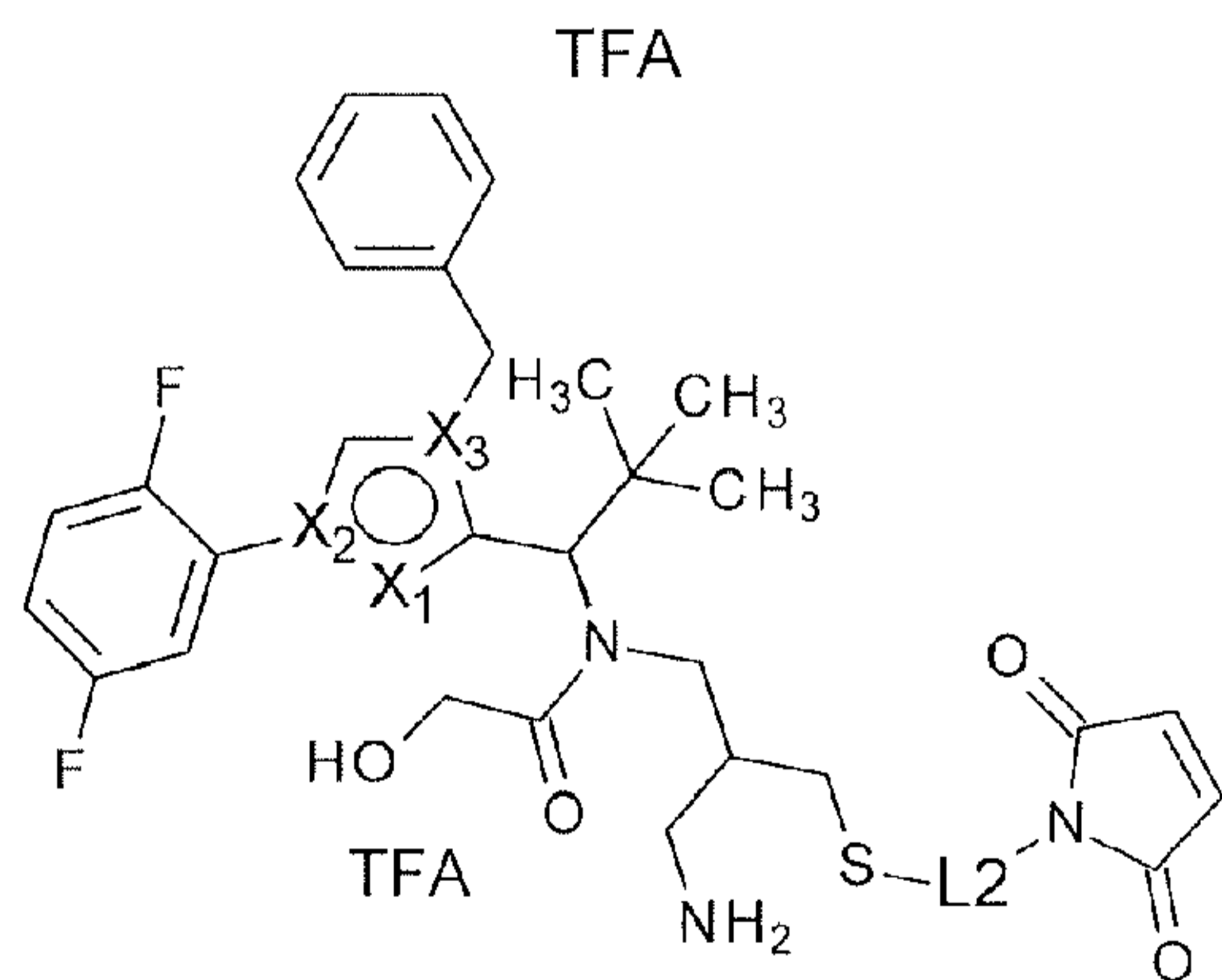
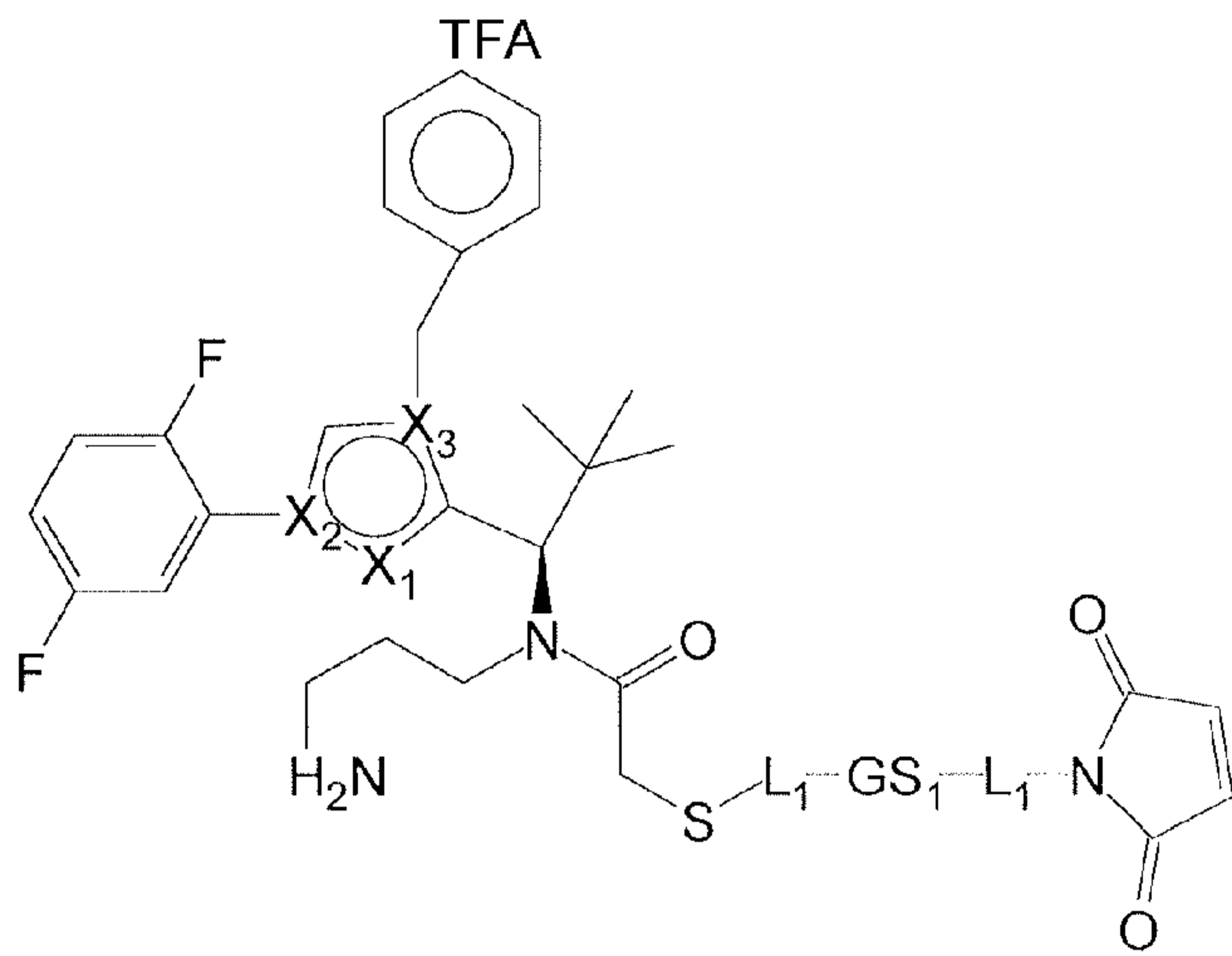
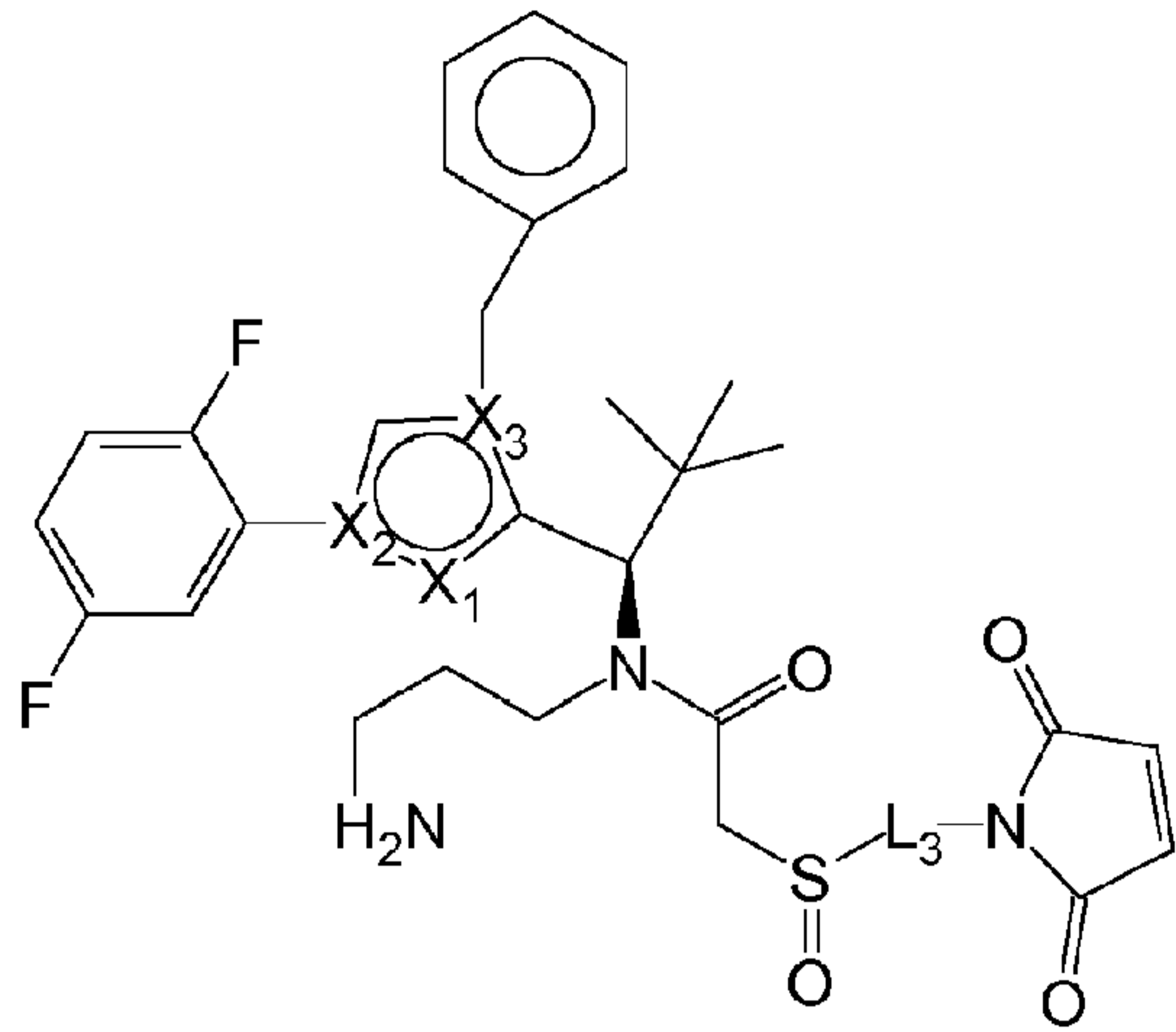
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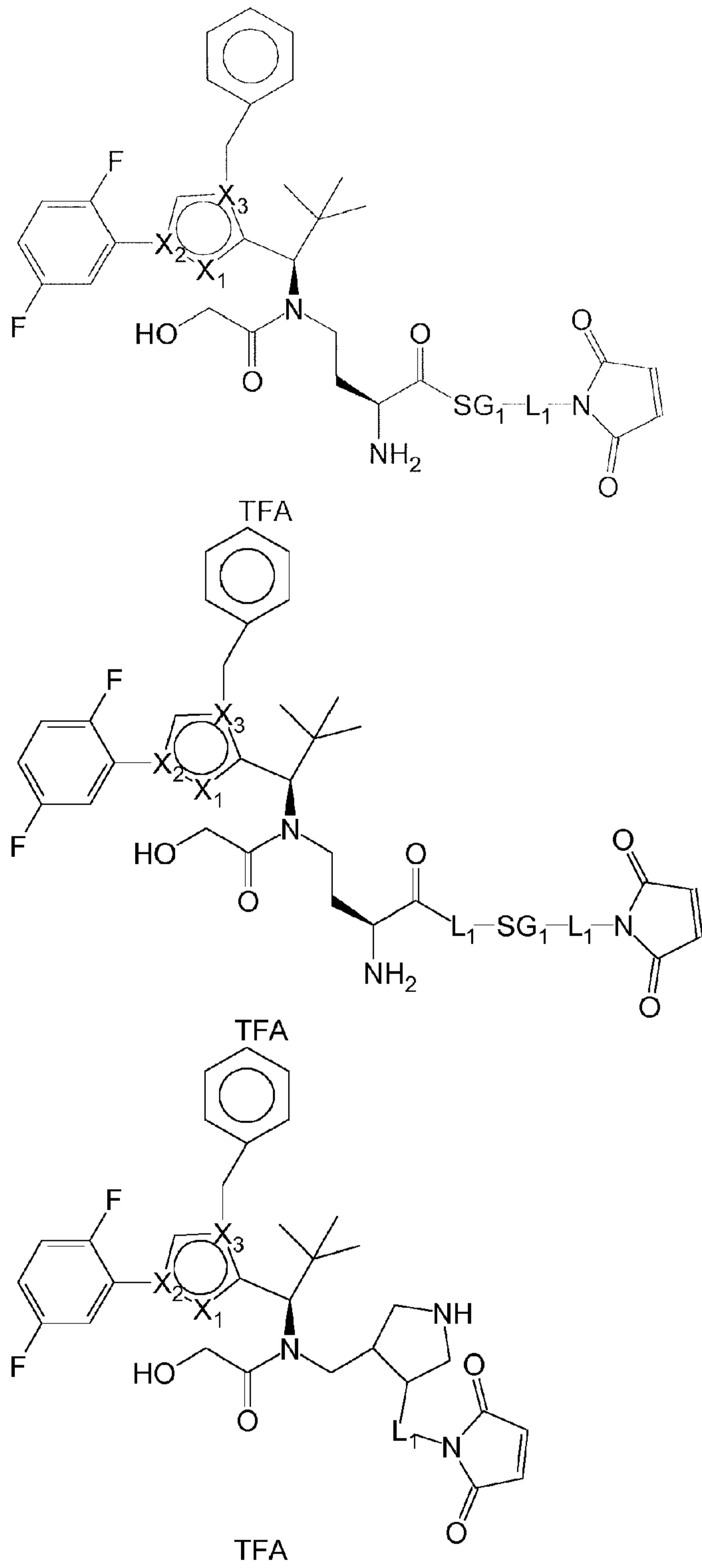
f. a sequence of the heavy chain, as shown in SEQ ID NO:52, and also a sequence of the light chain, as shown in SEQ ID NO:51, or

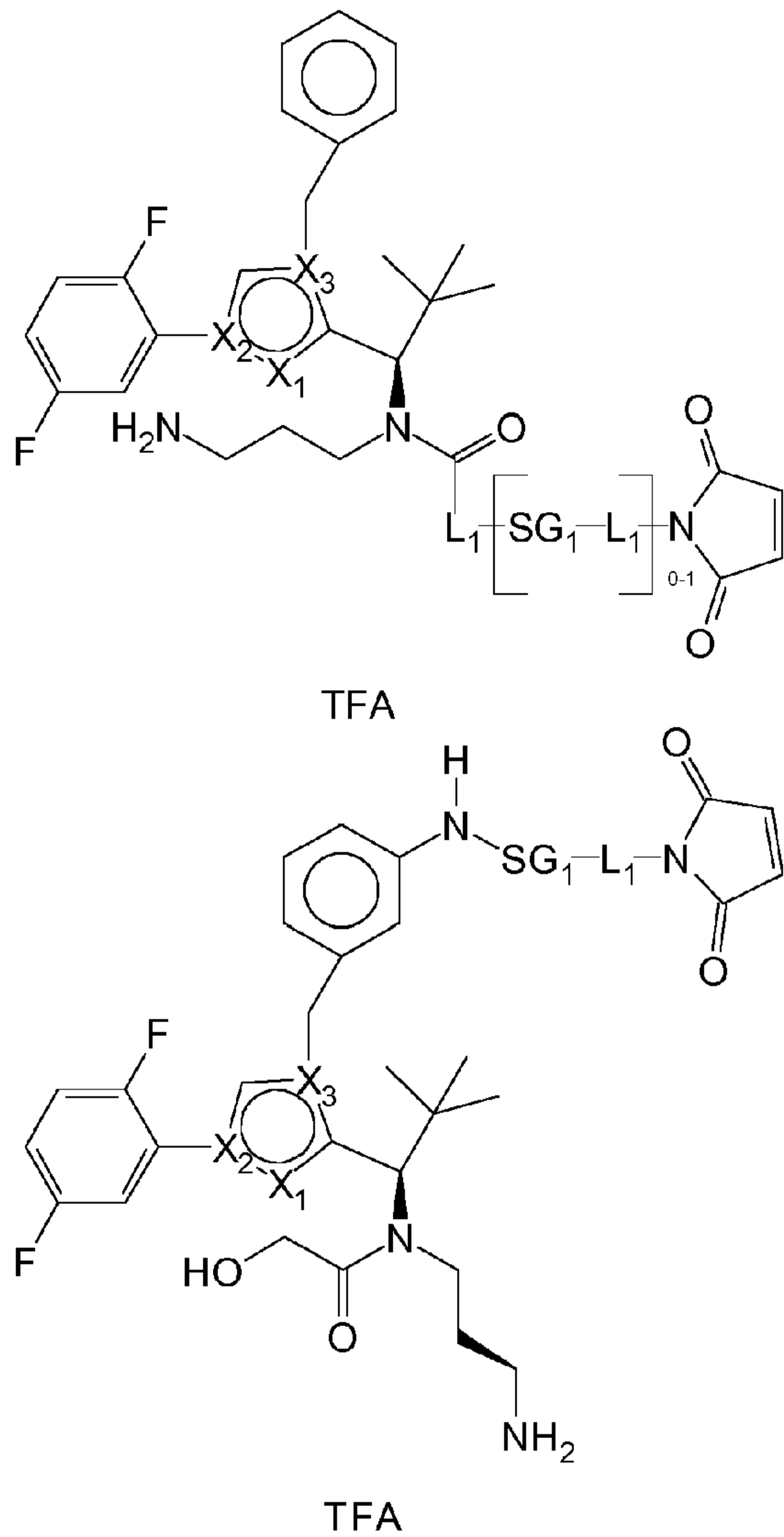
35 g. a sequence of the heavy chain, as shown in SEQ ID NO:62, and also a sequence of the light chain, as shown in SEQ ID NO:61, or

- h. a sequence of the heavy chain, as shown in SEQ ID NO:72,  
and also a sequence of the light chain, as shown in SEQ ID NO:71,  
or
- 5 i. a sequence of the heavy chain, as shown in SEQ ID NO:82,  
and also a sequence of the light chain, as shown in SEQ ID NO:81,  
or
- 10 j. a sequence of the heavy chain, as shown in SEQ ID NO:92,  
and also a sequence of the light chain, as shown in SEQ ID NO:91,  
or
- 15 k. a sequence of the heavy chain, as shown in SEQ ID NO:102,  
and also a sequence of the light chain, as shown in SEQ ID  
NO:101, or
- 20 l. a sequence of the heavy chain, as shown in SEQ ID NO:112,  
and also a sequence of the light chain, as shown in SEQ ID  
NO:111.
38. Spindle conjugate according to one or more of the preceding  
Items where the conjugate has 1 to 10, preferably 2 to 8 active  
compound molecules per binder peptide or protein.
- 25 39. Process for preparing the conjugate according to Item 26  
or 30 where a compound of one of the formulae below, preferably  
in the form of its trifluoroacetic acid salt, is attached to a  
cysteine residue of a binder peptide or protein which is  
optionally partially reduced beforehand, where the compound is  
30 preferably employed in a 2- to 12-fold molar excess with respect  
to the binder peptide or protein:









where R represents -H or -COOH,

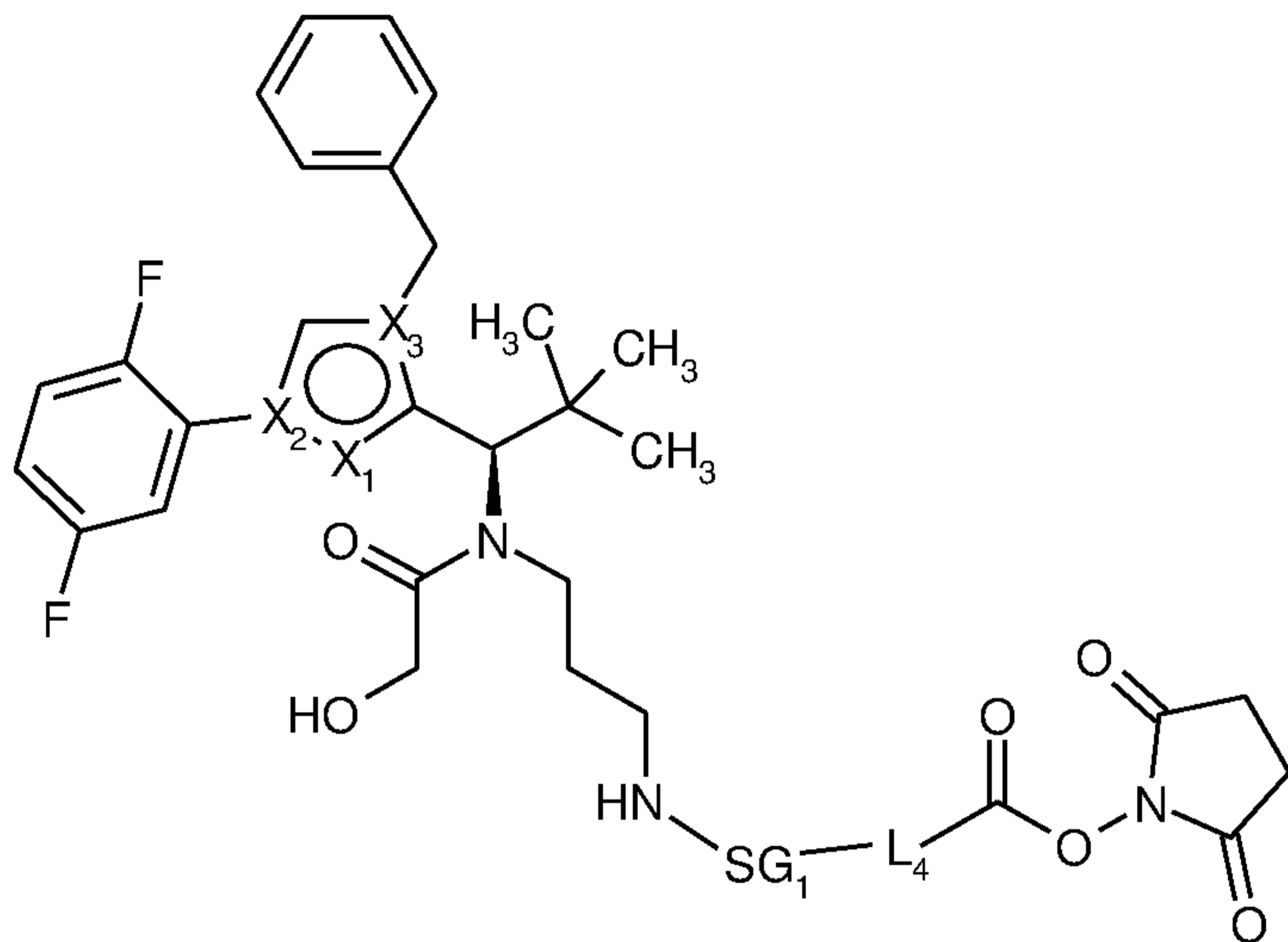
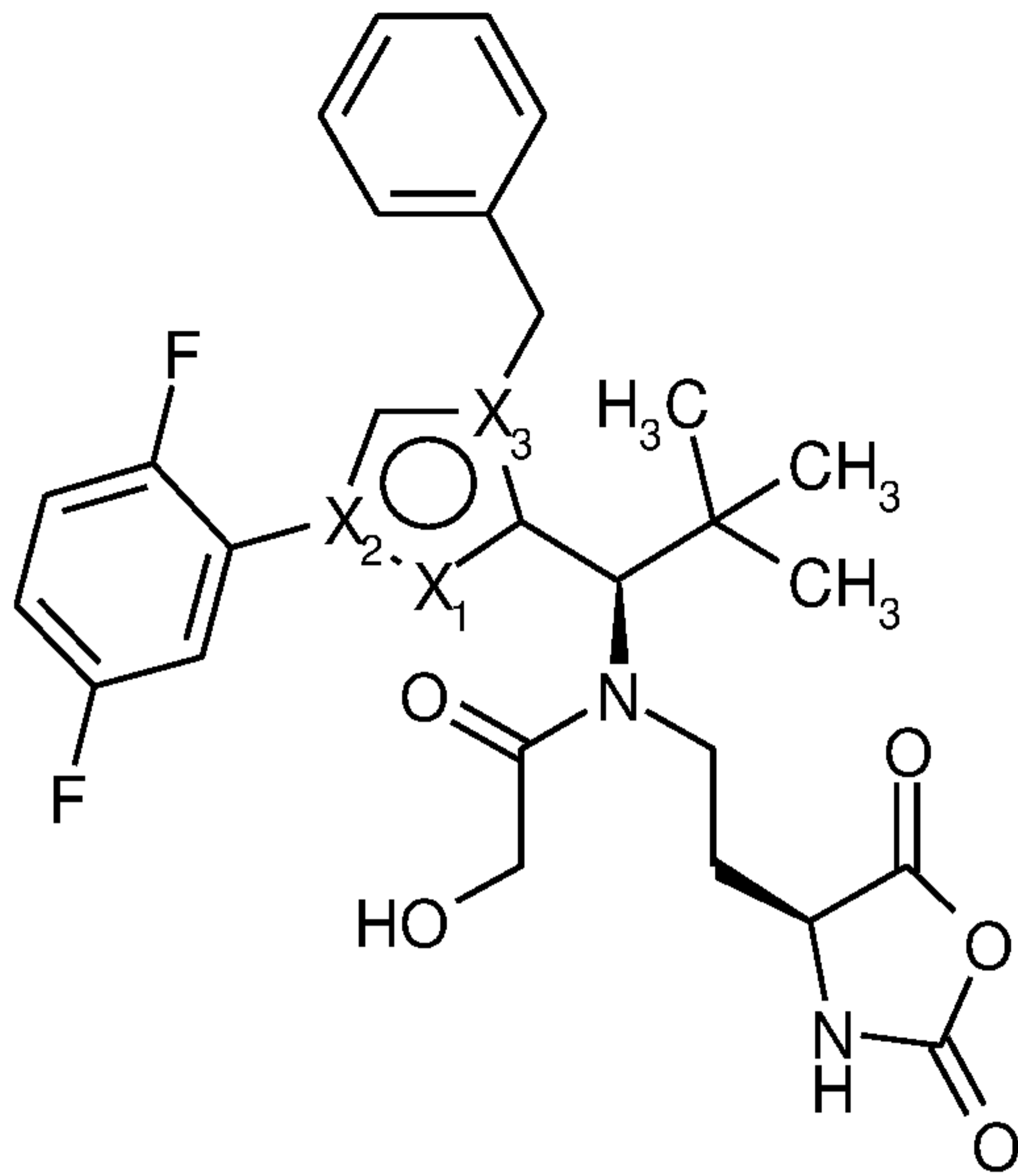
5 where K represents straight-chain or branched optionally substituted by C<sub>1</sub>-C<sub>6</sub>-alkoxy or -OH-C<sub>1</sub>-C<sub>6</sub>-alkyl, and

where X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub>, SG<sub>1</sub>, L<sub>1</sub>, L<sub>2</sub>, L<sub>3</sub> and L<sub>4</sub> have the same meaning as in Item 26 or 30.

10

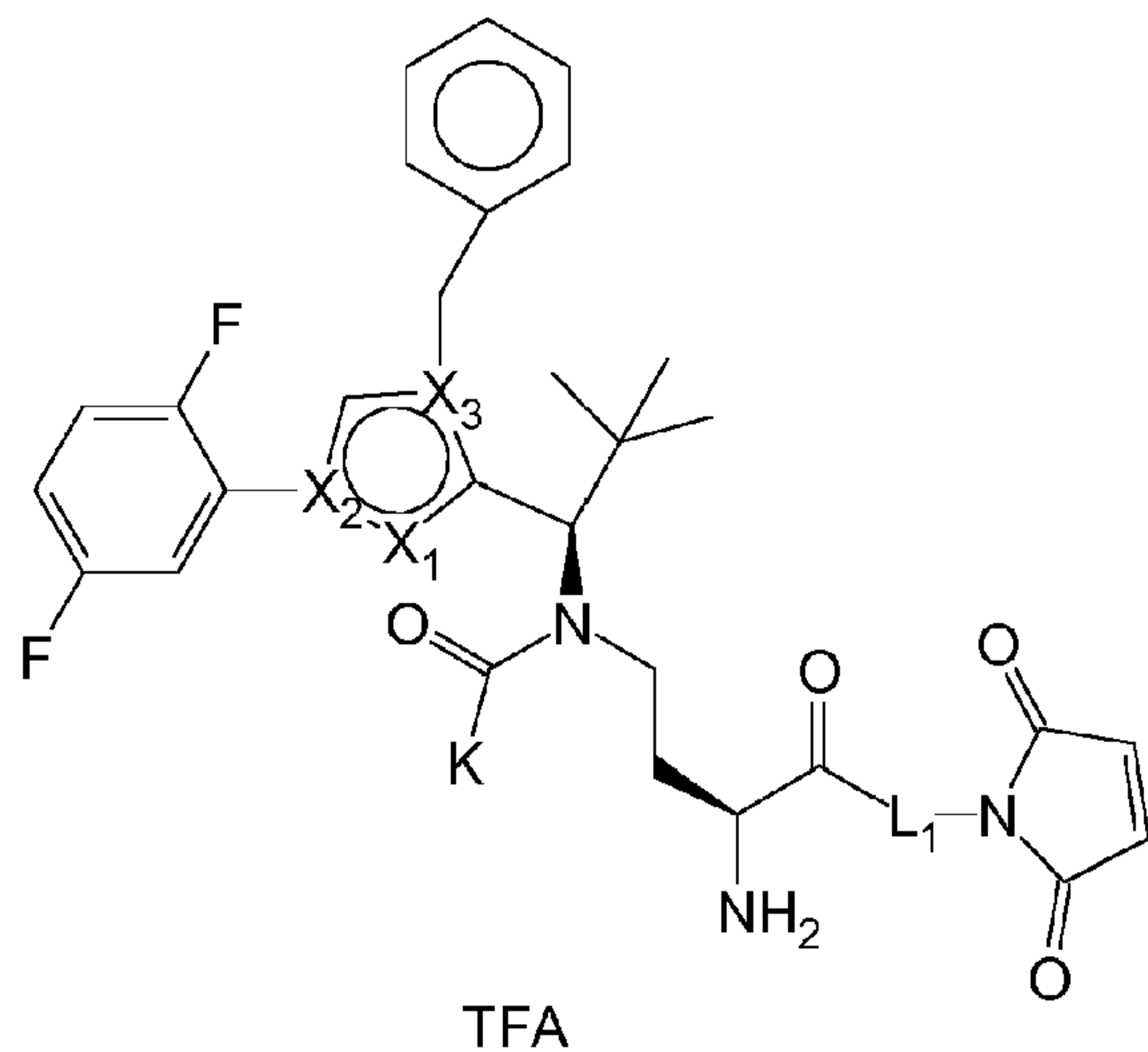
40. Process for preparing the conjugate according to Item 32 where a compound of one of the formulae below, preferably in the form of its trifluoroacetic acid salt, is attached to a lysine residue of a binder peptide or protein, where the compound is preferably employed in a 2- to 12-fold molar excess with respect to the binder peptide or protein:

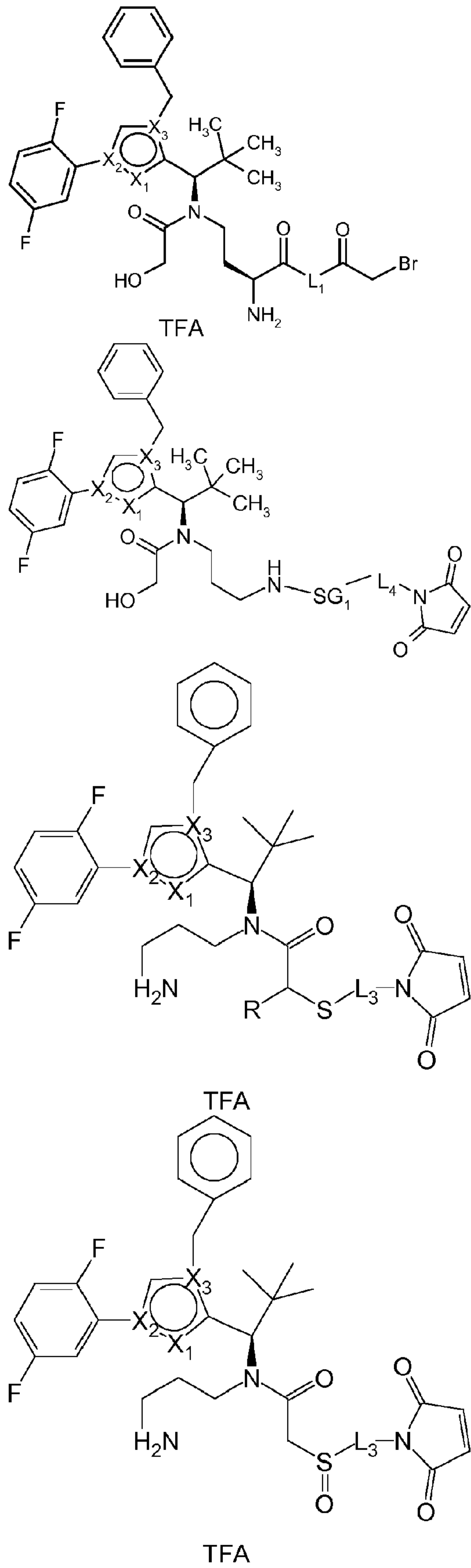
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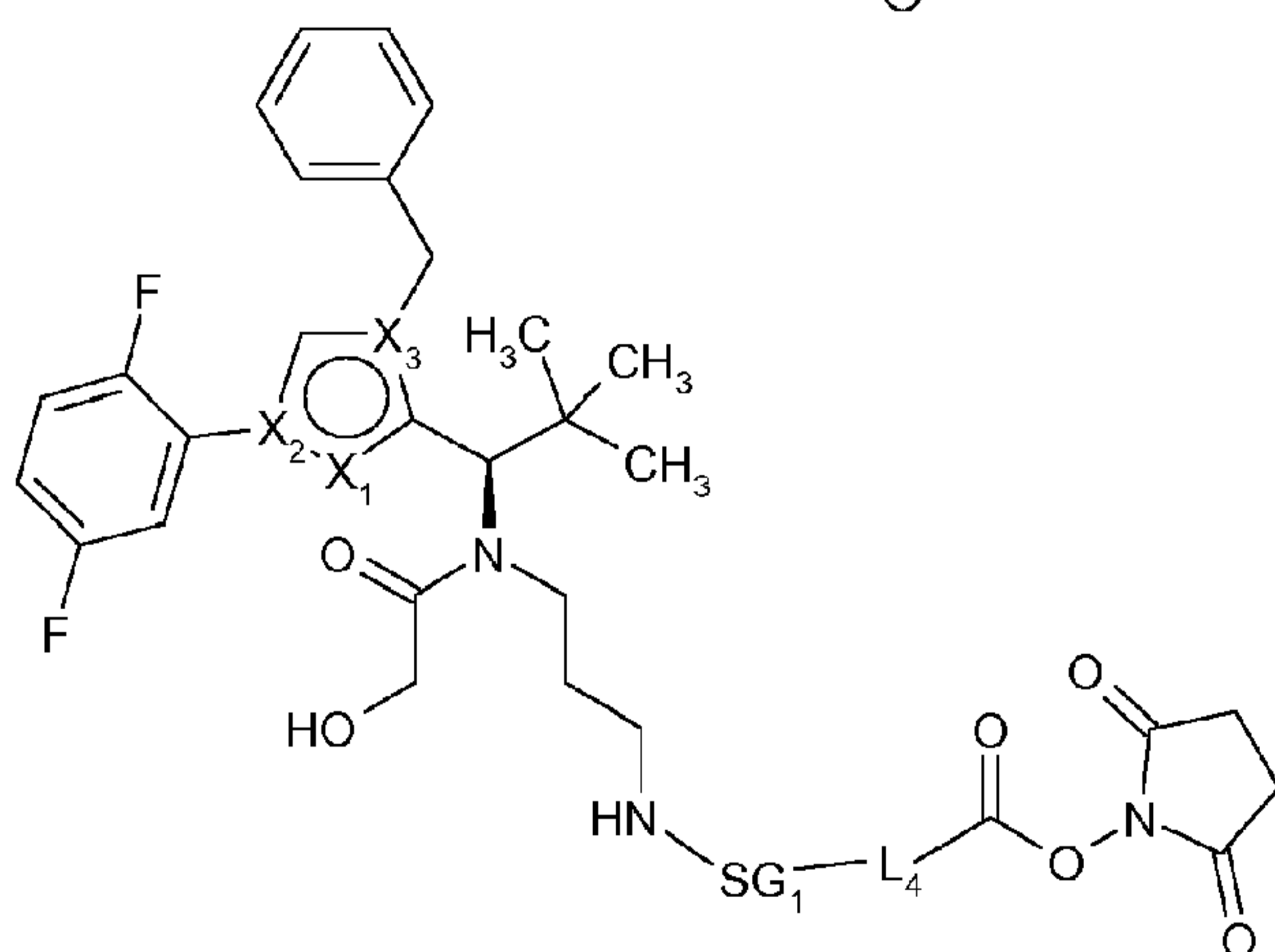
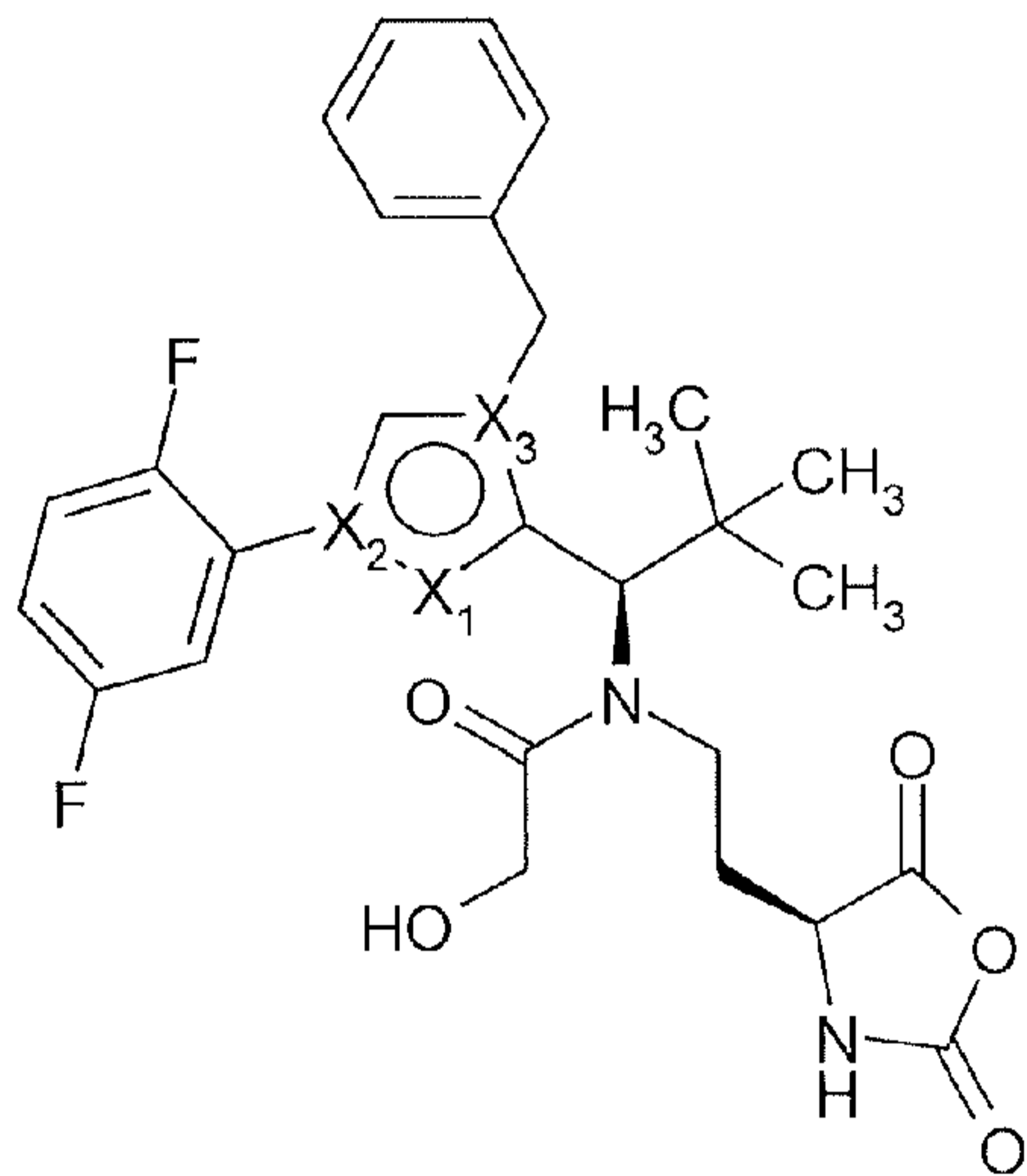
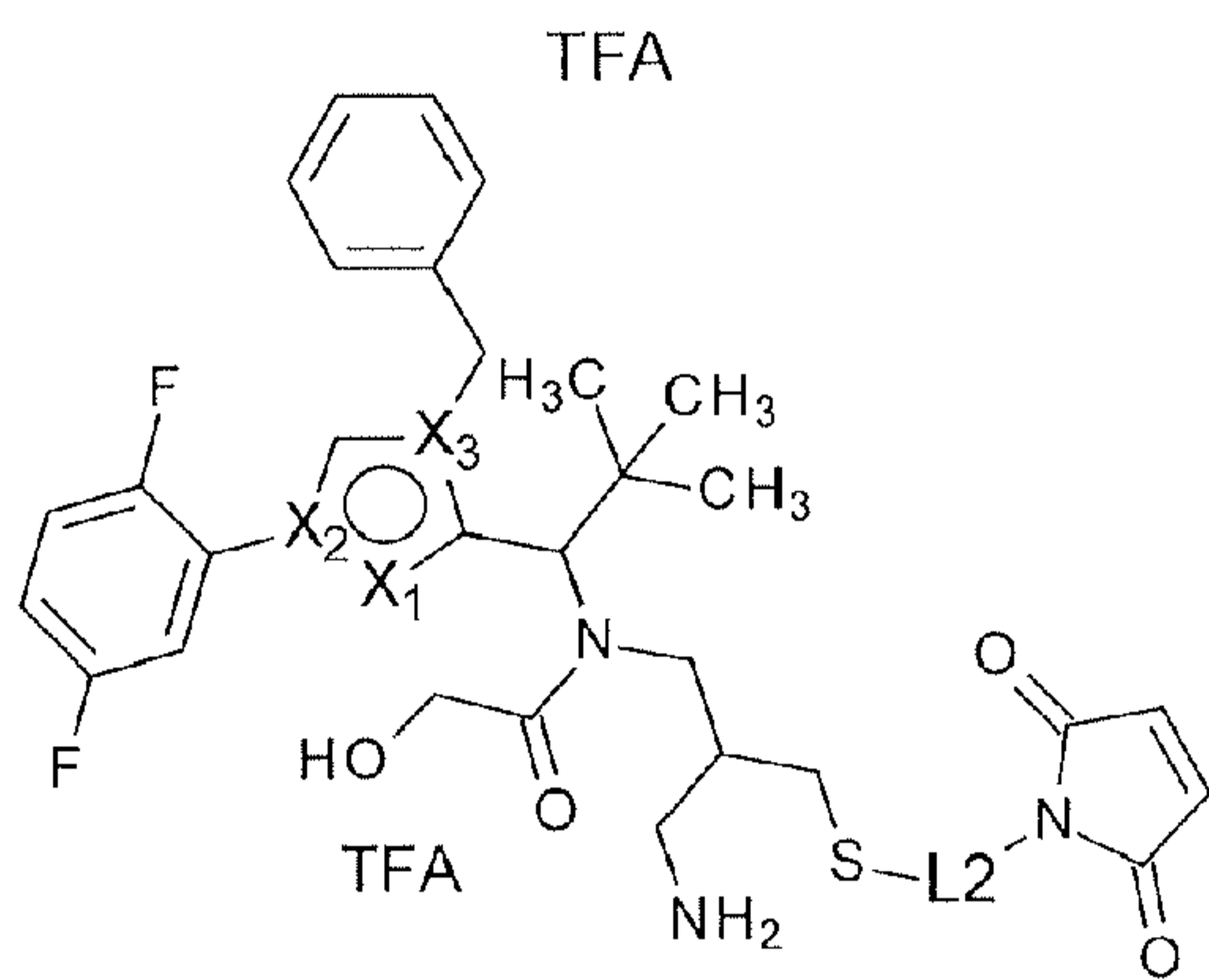
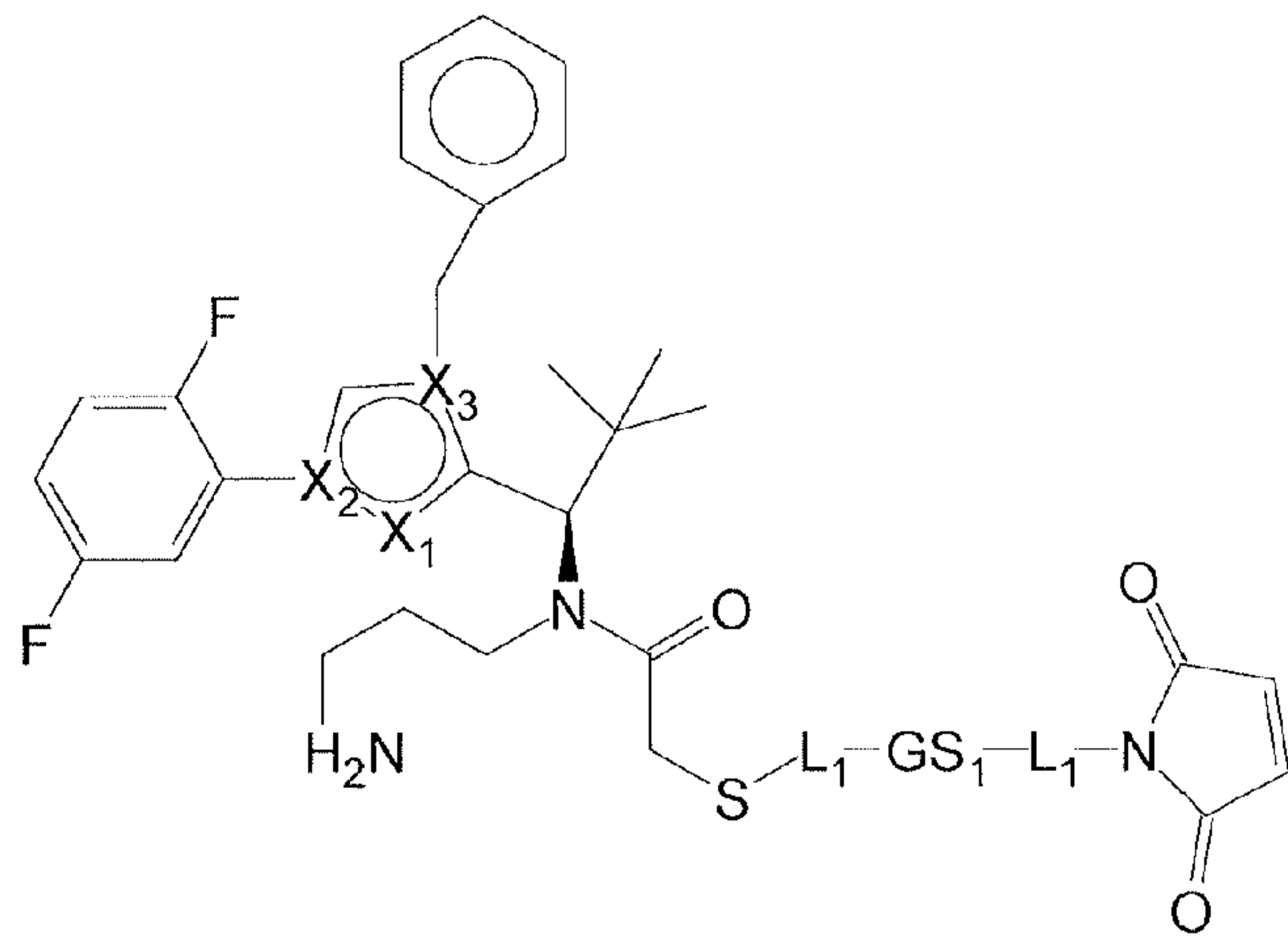
where  $X_1$ ,  $X_2$ ,  $X_3$ ,  $SG_1$  and  $L_4$  have the same meaning as in Item 32.

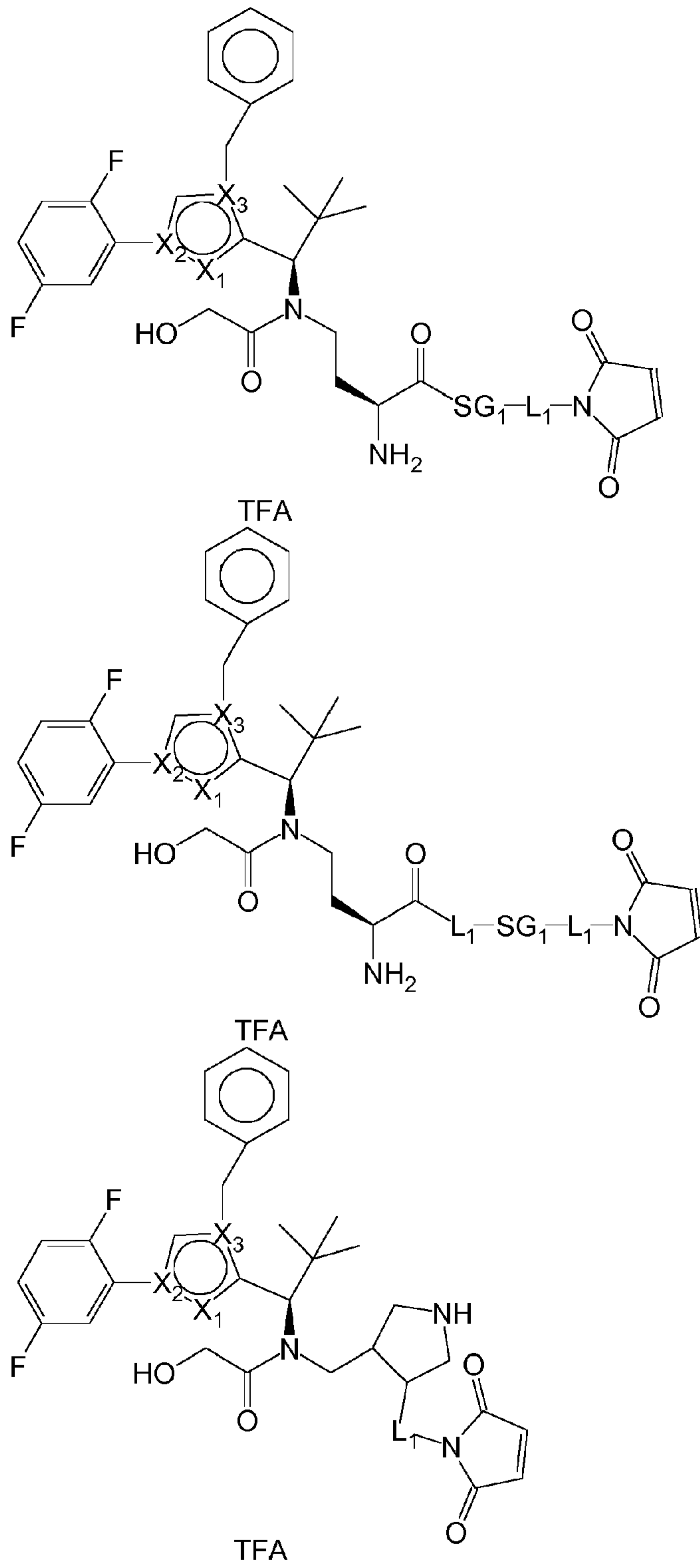
5 41. Compound of one of the formulae below:

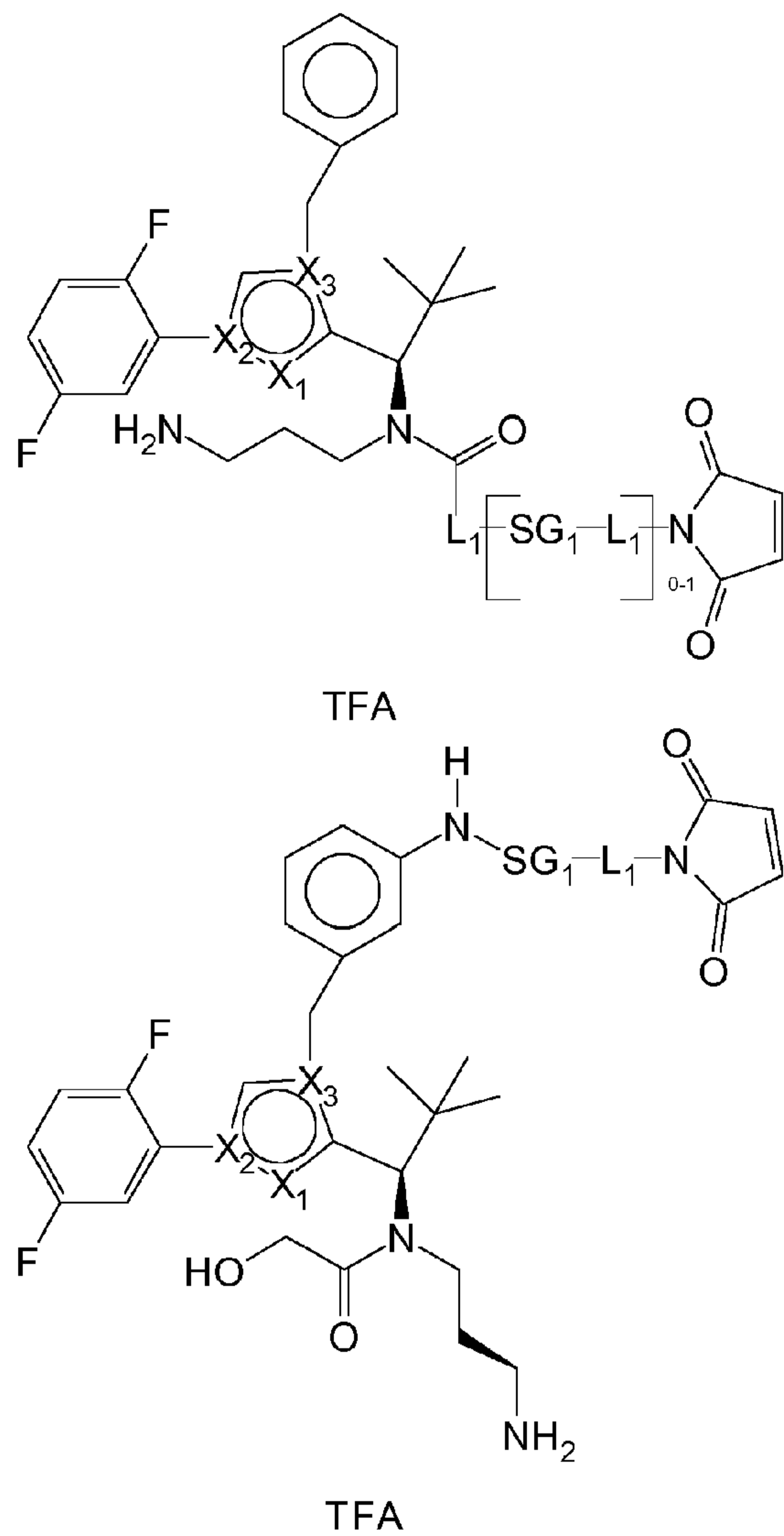












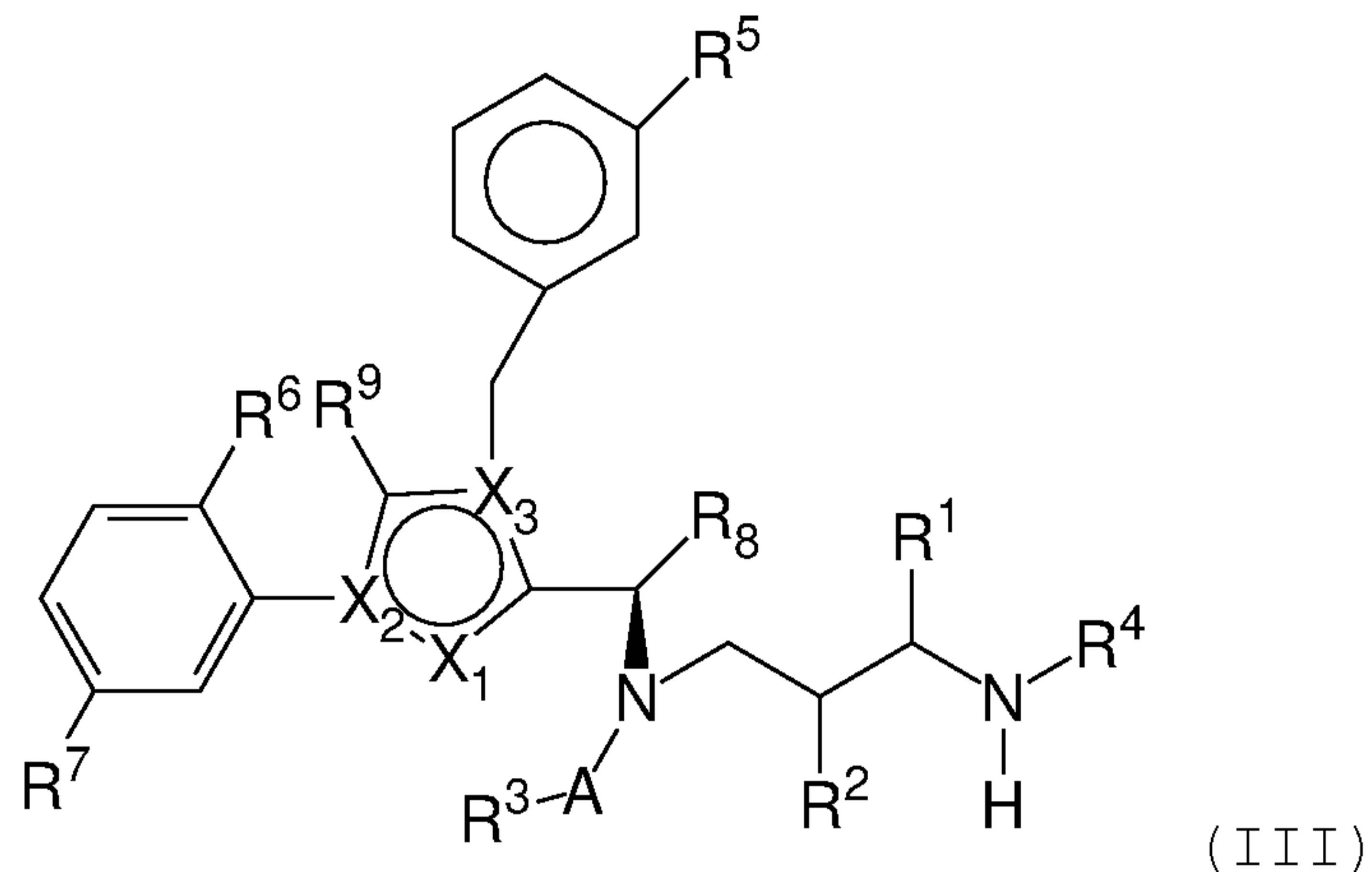
where R represents -H or -COOH,

5 where K represents straight-chain or branched optionally substituted by  $C_1$ - $C_6$ -alkoxy or  $-OH$ - $C_1$ - $C_6$ -alkyl, and

where  $X_1$ ,  $X_2$ ,  $X_3$ ,  $SG_1$ ,  $L_1$ ,  $L_2$ ,  $L_3$  and  $L_4$  have the same meaning as in Item 26, 30 or 32.

10

42. Compounds of the general formula (III):



where

5 X<sub>1</sub> represents N, X<sub>2</sub> represents N and X<sub>3</sub> represents C, or X<sub>1</sub> represents CH, X<sub>2</sub> represents C and X<sub>3</sub> represents N or X<sub>1</sub> represents NH, X<sub>2</sub> represents C and X<sub>3</sub> represents C, or X<sub>1</sub> represents CH, X<sub>2</sub> represents N and X<sub>3</sub> represents C;

10 R<sup>1</sup> represents -L-BINDER, H or -(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>,

where Y<sup>1</sup> and Y<sup>2</sup> independently of one another represent H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' or -CH(CH<sub>2</sub>W)Z', and Y<sup>3</sup> represents H or -  
 15 (CH<sub>2</sub>)<sub>0-3</sub>Z', where Z' represents H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH or -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH; where W represents H or OH;

where Y<sup>4</sup> independently of one another represents straight-chain  
 20 or branched C<sub>1-6</sub>-alkyl which is optionally substituted by -NHCONH<sub>2</sub>, or represents aryl or benzyl which are optionally substituted by -NH<sub>2</sub>;

R<sup>2</sup> and R<sup>4</sup> independently of one another represent -L-BINDER, H, -  
 25 CO-CHY<sup>4</sup>-NHY<sup>5</sup> or -(CH<sub>2</sub>)<sub>0-3</sub>Z, or R<sup>2</sup> and R<sup>4</sup> together (with formation of a pyrrolidine ring) represent -CH<sub>2</sub>-CHR<sup>10</sup>- or -CHR<sup>10</sup>-CH<sub>2</sub>-, where R<sup>10</sup> represents L-#1, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH or OH,

where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> or -  
 30 CO-OY<sup>3</sup>,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  $COOH$ ;

5

where  $Y^4$  independently of one another represents straight-chain or branched  $C_{1-6}$  alkyl which is optionally substituted by  $-NHCONH_2$ , or represents aryl or benzyl which are optionally substituted by  $-NH_2$ , and  $Y^5$  represents H or  $-CO-CHY^6-NH_2$ , where  
 10  $Y^6$  represents straight-chain or branched  $C_{1-6}$ -alkyl;

A represents CO, SO,  $SO_2$ ,  $SO_2NH$  or  $CNNH$ ;

$R^3$  represents  $-L$ -BINDER or an optionally substituted alkyl, aryl,  
 15 heteroaryl, heteroalkyl, heterocycloalkyl group, preferably  $-L$ -  
 #1, or a  $C_{1-10}$ -alkyl,  $C_{6-10}$ -aryl or  $C_{6-10}$ -aralkyl,  $C_{5-10}$ -heteroalkyl,  
 $C_{1-10}$ -alkyl- $O$ - $C_{6-10}$ -aryl or  $C_{5-10}$ -heterocycloalkyl group which may  
 be substituted by 1-3  $-OH$  groups, 1-3 halogen atoms, 1-3  
 halogenated alkyl groups (each having 1-3 halogen atoms), 1-3  
 20  $O$ -alkyl groups, 1-3  $-SH$  groups, 1-3  $-S$ -alkyl groups, 1-3  $-O$ - $CO$ -  
 alkyl groups, 1-3  $-O$ - $CO$ - $NH$ -alkyl groups, 1-3  $-NH$ - $CO$ -alkyl  
 groups, 1-3  $-NH$ - $CO$ - $NH$ -alkyl groups, 1-3  $-S(O)_n$ -alkyl groups, 1-  
 3  $-SO_2$ - $NH$ -alkyl groups, 1-3  $-NH$ -alkyl groups, 1-3  $-N(alkyl)_2$   
 groups, 1-3  $-NH_2$  groups or 1-3  $-(CH_2)_{0-3}Z$  groups, where  $Z$   
 25 represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$  or  $-CO-OY^3$ ,  
 where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  
 $-(CH_2)_{0-3}Z'$  and  $Y^3$  represents H,  $-(CH_2)_{0-3}-CH(NHCOCH_3)Z'$ ,  $-(CH_2)_{0-3}-$   
 $CH(NH_2)Z'$  or  $-(CH_2)_{0-3}Z'$ , where  $Z'$  represents H,  $SO_3H$ ,  $NH_2$  or  
 $COOH$

30

(where "alkyl" preferably represents  $C_{1-10}$ -alkyl);

$R^5$  represents  $-L$ -BINDER, H, F,  $NH_2$ ,  $NO_2$ , halogen, SH or  $-(CH_2)_{0-3}Z$ , where  $Z$  represents  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$   
 35 or  $-CO-OY^3$ ,

where  $Y^1$  and  $Y^2$  independently of one another represent H,  $NH_2$  or  $-(CH_2)_{0-3}Z'$ , and  $Y^3$  represents H or  $-(CH_2)_{0-3}Z'$ , where  $Z'$

represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH;

where L represents a linker and BINDER represents a binder or a derivative thereof, where the binder may optionally be attached  
5 to a plurality of active compound molecules,

R<sup>6</sup> and R<sup>7</sup> independently of one another represent H, cyano, (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>2-10</sub>-alkenyl, (optionally fluorinated) C<sub>2-10</sub>-alkynyl, hydroxy or  
10 halogen,

R<sup>8</sup> represents (optionally fluorinated) C<sub>1-10</sub>-alkyl, (optionally fluorinated) C<sub>4-10</sub>-cycloalkyl or optionally substituted oxetane;  
and  
15

R<sup>9</sup> represents H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F or CHF<sub>2</sub>;

and the salts, solvates and salts of the solvates thereof.

20 43. Pharmaceutical composition comprising a conjugate according to one or more of Items 1 to 38 and 41-42 in combination with an inert non-toxic pharmaceutically suitable auxiliary.

44. Conjugate according to one or more of Items 1 to 38 and 41-  
25 42 for use in a method for the treatment and/or prophylaxis of diseases.

45. Conjugate according to one or more of Items 1 to 38 and 41-  
42 for use in a method for the treatment of hyperproliferative  
30 and/or angiogenic disorders.

46. Conjugate according to one or more of Items 1 to 38 and 41-  
42 where R<sup>3a</sup> or R<sup>3</sup> represents -L-BINDER or a substituted alkyl,  
aryl or heteroaryl group,  
35

preferably -L-#1 or a C<sub>1-10</sub>-alkyl, C<sub>6-10</sub>-aryl or C<sub>6-10</sub>-aralkyl group or C<sub>5-10</sub>-heteroalkyl, which may be substituted by -OH, O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-NH-alkyl, NH-CO-alkyl, NH-CO-NH-

alkyl, S(O)<sub>n</sub>-alkyl, SO<sub>2</sub>-NH-alkyl, NH-alkyl, N(alkyl)<sub>2</sub>, NH<sub>2</sub> or -  
(CH<sub>2</sub>)<sub>0-3</sub>Z, where Z represents -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-  
NY<sup>1</sup>Y<sup>2</sup> or -CO-OY<sup>3</sup>, where Y<sup>1</sup> and Y<sup>2</sup> independently of one another  
represent H, NH<sub>2</sub> or -(CH<sub>2</sub>)<sub>0-3</sub>Z' and Y<sup>3</sup> represents H or -(CH<sub>2</sub>)<sub>0-3</sub>Z',  
5 where Z' represents H, SO<sub>3</sub>H, NH<sub>2</sub> or COOH,

(where "alkyl" preferably represents C<sub>1-10</sub>-alkyl).

47. Method for the treatment and/or prophylaxis of  
10 hyperproliferative and/or angiogenic disorders in humans and  
animals using an effective amount of at least one conjugate  
according to one or more of Items 1 to 38 and 41-42.

## SEQUENCE LISTING

<110> Bayer Pharma AG  
 <120> Drug conjugates (ADCs) with KSP inhibitors  
 <130> 173 506  
 <160> 240  
 <170> BiSSAP 1.3  
 <210> 1  
 <211> 215  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 1

Asp	Ile	Gln	Met	Thr	Gln	Ser	Pro	Ser	Ser	Leu	Ser	Ala	Ser	Val	Gly
1				5				10						15	
Asp	Arg	Val	Thr	Ile	Thr	Cys	Arg	Ala	Ser	Gln	Ser	Ile	Ser	Gly	Tyr
			20					25					30		
Leu	Asn	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Lys	Ala	Pro	Lys	Leu	Leu	Ile
		35				40						45			
Tyr	Gln	Ala	Ser	Ser	Leu	Gln	Ser	Gly	Val	Pro	Ser	Arg	Phe	Ser	Gly
	50					55					60				
Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Leu	Thr	Ile	Ser	Ser	Leu	Gln	Pro
65					70					75					80
Glu	Asp	Phe	Ala	Thr	Tyr	Tyr	Cys	Gln	Gln	Ser	Tyr	Thr	Ser	Pro	Phe
				85						90				95	
Ile	Thr	Phe	Gly	Gln	Gly	Thr	Lys	Val	Glu	Ile	Lys	Arg	Thr	Val	Ala
			100					105					110		
Ala	Pro	Ser	Val	Phe	Ile	Phe	Pro	Pro	Ser	Asp	Glu	Gln	Leu	Lys	Ser
		115				120						125			
Gly	Thr	Ala	Ser	Val	Val	Cys	Leu	Leu	Asn	Asn	Phe	Tyr	Pro	Arg	Glu
	130					135					140				
Ala	Lys	Val	Gln	Trp	Lys	Val	Asp	Asn	Ala	Leu	Gln	Ser	Gly	Asn	Ser
145					150					155					160
Gln	Glu	Ser	Val	Thr	Glu	Gln	Asp	Ser	Lys	Asp	Ser	Thr	Tyr	Ser	Leu
				165					170					175	
Ser	Ser	Thr	Leu	Thr	Leu	Ser	Lys	Ala	Asp	Tyr	Glu	Lys	His	Lys	Val
		180						185					190		
Tyr	Ala	Cys	Glu	Val	Thr	His	Gln	Gly	Leu	Ser	Ser	Pro	Val	Thr	Lys
		195				200						205			
Ser	Phe	Asn	Arg	Gly	Glu	Cys									
	210					215									

<210> 2  
 <211> 449  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 2

Glu	Val	Gln	Leu	Leu	Glu	Ser	Gly	Gly	Gly	Leu	Val	Gln	Pro	Gly	Gly
1				5					10					15	
Ser	Leu	Arg	Leu	Ser	Cys	Ala	Ala	Ser	Gly	Phe	Thr	Phe	Ser	Pro	Tyr
			20					25					30		
Pro	Met	Ile	Trp	Val	Arg	Gln	Ala	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Val
		35				40						45			
Ser	Tyr	Ile	Ser	Pro	Ser	Gly	Gly	Ser	Thr	His	Tyr	Ala	Asp	Ser	Val
	50					55					60				
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr
65					70					75					80



Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 3

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 3

Arg Ala Ser Gln Ser Ile Ser Gly Tyr Leu Asn  
 1 5 10

&lt;210&gt; 4

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 4

Gln Ala Ser Ser Leu Gln Ser  
 1 5

&lt;210&gt; 5

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 5

Gln Gln Ser Tyr Thr Ser Pro Phe Ile Thr  
 1 5 10  
 <210> 6  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 6  
 Pro Tyr Pro Met Ile  
 1 5  
 <210> 7  
 <211> 17  
 <212> PRT  
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 <400> 7  
 Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 8  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 8  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 9  
 <211> 108  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 9  
 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Gly Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Gln Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser Pro Phe  
 85 90 95  
 Ile Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys  
 100 105  
 <210> 10  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 10

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Ile Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 11

&lt;211&gt; 215

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 11

Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Gly Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Gln Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser Pro Phe  
 85 90 95  
 Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala  
 100 105 110  
 Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser  
 115 120 125  
 Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu  
 130 135 140  
 Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser  
 145 150 155 160  
 Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu  
 165 170 175  
 Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Leu  
 180 185 190  
 Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys  
 195 200 205  
 Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 12

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 12

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Ile Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 13

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 13

Arg Ala Ser Gln Ser Ile Ser Gly Tyr Leu Asn  
 1 5 10

&lt;210&gt; 14

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

<400> 14  
 Gln Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 15  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 15  
 Gln Gln Ser Tyr Thr Ser Pro Phe Ile Thr  
 1 5 10  
 <210> 16  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 16  
 Pro Tyr Pro Met Ile  
 1 5  
 <210> 17  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 17  
 Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 18  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 18  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 19  
 <211> 108  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 19  
 Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Gly Tyr  
 20 25 30  
  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Gln Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser Pro Phe  
 85 90 95  
 Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
 100 105  
 <210> 20  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 20

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Ile Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 21

&lt;211&gt; 215

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 21

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Gln Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser Pro Phe  
 85 90 95  
 Ile Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala  
 100 105 110  
 Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser  
 115 120 125  
 Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu  
 130 135 140  
 Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser  
 145 150 155 160  
 Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu  
 165 170 175  
 Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val

Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys  
 180 185 190  
 195 200 205

Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 22

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 22

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp

405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

<210> 23

<211> 11

<212> PRT

<213> Homo Sapiens

<400> 23

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn  
 1 5 10

<210> 24

<211> 7

<212> PRT

<213> Homo Sapiens  
 <400> 24  
 Gln Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 25  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 25  
 Gln Gln Ser Tyr Thr Ser Pro Phe Ile Thr  
 1 5 10  
 <210> 26  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 26  
 Pro Tyr Pro Met Met  
 1 5  
 <210> 27  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 27  
 Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 28  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 28  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 29  
 <211> 108  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 29  
 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Gln Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser Pro Phe  
 85 90 95  
 Ile Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys  
 100 105  
 <210> 30  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 30



Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 31

&lt;211&gt; 215

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 31

Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Gln Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser Pro Phe  
 85 90 95  
  
 Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala  
 100 105 110  
 Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser  
 115 120 125  
 Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu  
 130 135 140  
 Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser  
 145 150 155 160  
 Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu  
 165 170 175  
 Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Leu  
 180 185 190  
 Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys  
 195 200 205  
 Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 32

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 32

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 33

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 33

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn  
 1 5 10

&lt;210&gt; 34

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 34

Gln Ala Ser Ser Leu Gln Ser  
1 5

&lt;210&gt; 35

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 35

Gln Gln Ser Tyr Thr Ser Pro Phe Ile Thr  
1 5 10

&lt;210&gt; 36

&lt;211&gt; 5

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 36

Pro Tyr Pro Met Met  
1 5

&lt;210&gt; 37

&lt;211&gt; 17

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 37

Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

&lt;210&gt; 38

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 38

Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
1 5 10

&lt;210&gt; 39

&lt;211&gt; 108

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 39

Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser Val Gly  
1 5 10 15  
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
20 25 30  
Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
35 40 45  
Tyr Gln Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
50 55 60  
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
65 70 75 80  
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser Pro Phe  
85 90 95  
Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
100 105

&lt;210&gt; 40

&lt;211&gt; 121

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 40

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 41

&lt;211&gt; 215

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 41

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Thr Pro Gly  
 85 90 95  
 Ile Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala  
 100 105 110  
 Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser  
 115 120 125  
 Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu  
 130 135 140  
 Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser  
 145 150 155 160  
 Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu  
 165 170 175  
 Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val  
 180 185 190  
 Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys  
 195 200 205  
 Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 42

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 42

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 43

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 43

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn  
 1 5 10

&lt;210&gt; 44

&lt;211&gt; 7

&lt;212&gt; PRT

<213> Homo Sapiens  
 <400> 44  
 Ala Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 45  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 45  
 Gln Gln Ser Tyr Ser Thr Pro Gly Ile Thr  
 1 5 10  
 <210> 46  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 46  
 Pro Tyr Pro Met Met  
 1 5  
 <210> 47  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 47  
 Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 48  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 4  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 49  
 <211> 108  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 49  
 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Thr Pro Gly  
 85 90 95  
 Ile Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys  
 100 105  
 <210> 50  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 50

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Ser Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln

100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

<210> 51  
 <211> 215  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 51

Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Ser Pro Gly  
 85 90 95  
 Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala  
 100 105 110  
 Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser  
 115 120 125  
 Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu  
 130 135 140  
 Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser  
 145 150 155 160  
 Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu  
 165 170 175  
 Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Leu  
 180 185 190  
 Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys  
 195 200 205  
 Ser Phe Asn Arg Gly Glu Cys  
 210 215

<210> 52  
 <211> 449  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 52

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 53

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 53

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn  
 1 5 10

&lt;210&gt; 54

&lt;211&gt; 7

&lt;212&gt; PRT



<213> Homo Sapiens  
 <400> 54  
 Ala Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 55  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 55  
 Gln Gln Ser Tyr Ser Ser Pro Gly Ile Thr  
 1 5 10  
 <210> 56  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 56  
 Pro Tyr Pro Met Met  
 1 5  
 <210> 57  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 57  
 Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 58  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 58  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 59  
 <211> 108  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 59  
 Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser Ser Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Ser Pro Gly  
 85 90 95  
 Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
 100 105  
 <210> 60  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 60

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 61

&lt;211&gt; 217

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 61

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Ser  
 85 90 95  
 Pro Gly Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr  
 100 105 110  
 Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu  
 115 120 125  
 Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro  
 130 135 140  
 Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly  
 145 150 155 160  
 Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr  
 165 170 175  
 Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His  
 180 185 190  
 Lys Leu Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val  
 195 200 205  
 Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 62

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 62

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val

50						55					60					
Lys	Gly	Arg	Phe	Thr	Ile	Ser	Arg	Asp	Asn	Ser	Lys	Asn	Thr	Leu	Tyr	
65					70					75					80	
Leu	Gln	Met	Asn	Ser	Leu	Arg	Ala	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	
				85					90					95		
Ala	Arg	Gly	Gly	Asp	Thr	Tyr	Phe	Asp	Tyr	Phe	Asp	Tyr	Trp	Gly	Gln	
			100					105					110			
Gly	Thr	Leu	Val	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	
		115						120				125				
Phe	Pro	Leu	Ala	Pro	Ser	Ser	Lys	Ser	Thr	Ser	Gly	Gly	Thr	Ala	Ala	
	130						135				140					
Leu	Gly	Cys	Leu	Val	Lys	Asp	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val	Ser	
145					150					155					160	
Trp	Asn	Ser	Gly	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	
				165					170					175		
Leu	Gln	Ser	Ser	Gly	Leu	Tyr	Ser	Leu	Ser	Ser	Val	Val	Thr	Val	Pro	
			180					185					190			
Ser	Ser	Ser	Leu	Gly	Thr	Gln	Thr	Tyr	Ile	Cys	Asn	Val	Asn	His	Lys	
		195						200				205				
Pro	Ser	Asn	Thr	Lys	Val	Asp	Lys	Lys	Val	Glu	Pro	Lys	Ser	Cys	Asp	
	210					215				220						
Lys	Thr	His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	
225					230					235					240	
Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	
				245					250					255		
Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	
			260					265					270			
Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	
		275						280				285				
Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	
	290					295					300					
Val	Val	Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	
305					310					315					320	
Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	
				325					330					335		
Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	
			340					345					350			
Thr	Leu	Pro	Pro	Ser	Arg	Asp	Glu	Leu	Thr	Lys	Asn	Gln	Val	Ser	Leu	
		355						360					365			
Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	
	370					375					380					
Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	
385					390					395					400	
Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	
				405					410					415		
Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	
			420					425					430			
Glu	Ala	Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	
		435					440					445				

Gly

<210> 63

<211> 11

<212> PRT

<213> Homo Sapiens

<400> 63

Arg	Ala	Ser	Gln	Ser	Ile	Ser	Ser	Tyr	Leu	Asn
1			5						10	

<210> 64

<211> 7

<212> PRT

<213> Homo Sapiens

<400> 64

Ala	Ala	Ser	Ser	Leu	Gln	Ser
1				5		

<210> 65

<211> 10

<212> PRT

<213> Homo Sapiens

<400> 65

Gln Gln Ser Tyr Ser Ser Pro Gly Ile Thr  
1 5 10

<210> 66

<211> 5

<212> PRT

<213> Homo Sapiens

<400> 66

Pro Tyr Pro Met Met  
1 5

<210> 67

<211> 17

<212> PRT

<213> Homo Sapiens

<400> 67

Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 68

<211> 11

<212> PRT

<213> Homo Sapiens

<400> 68

Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
1 5 10

<210> 69

<211> 110

<212> PRT

<213> Homo Sapiens

<400> 69

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
1 5 10 15  
Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
20 25 30  
Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
35 40 45  
Leu Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
50 55 60  
Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
65 70 75 80  
Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Ser  
85 90 95

Pro Gly Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
100 105 110

<210> 70

<211> 121

<212> PRT

<213> Homo Sapiens

<400> 70

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 71

&lt;211&gt; 217

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 71

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Ser Ser  
 85 90 95  
 Pro Gly Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr  
 100 105 110  
 Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu  
 115 120 125  
 Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro  
 130 135 140  
 Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly  
 145 150 155 160  
 Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr  
 165 170 175  
 Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His  
 180 185 190  
 Lys Leu Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val  
 195 200 205  
 Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 72

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 72

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Gly Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 73

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 73

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn  
 1 5 10

&lt;210&gt; 74

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 74

Ala Ala Ser Ser Leu Gln Ser  
1 5

&lt;210&gt; 75

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 75

Gln Gln Ser Tyr Ser Ser Pro Gly Ile Thr  
1 5 10

&lt;210&gt; 76

&lt;211&gt; 5

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 76

Pro Tyr Pro Met Met  
1 5

&lt;210&gt; 77

&lt;211&gt; 17

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 77

Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

&lt;210&gt; 78

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 78

Gly Gly Asp Gly Tyr Phe Asp Tyr Phe Asp Tyr  
1 5 10

&lt;210&gt; 79

&lt;211&gt; 110

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 79

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser

1				5						10					15
Val	Gly	Asp	Arg	Val	Thr	Ile	Thr	Cys	Arg	Ala	Ser	Gln	Ser	Ile	Ser
			20					25					30		
Ser	Tyr	Leu	Asn	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Lys	Ala	Pro	Lys	Leu
		35					40					45			
Leu	Ile	Tyr	Ala	Ala	Ser	Ser	Leu	Gln	Ser	Gly	Val	Pro	Ser	Arg	Phe
		50				55					60				
Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Leu	Thr	Ile	Ser	Ser	Leu
					70					75					80
Gln	Pro	Glu	Asp	Phe	Ala	Thr	Tyr	Tyr	Cys	Gln	Gln	Ser	Tyr	Ser	Ser
				85					90					95	
Pro	Gly	Ile	Thr	Phe	Gly	Pro	Gly	Thr	Lys	Val	Glu	Ile	Lys		
			100					105					110		

&lt;210&gt; 80

&lt;211&gt; 121

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 80

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Gly Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 81

&lt;211&gt; 217

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 81

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Gly Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Phe Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr  
 100 105 110  
 Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu  
 115 120 125  
 Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro  
 130 135 140  
 Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly  
 145 150 155 160  
  
 Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr  
 165 170 175  
 Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His  
 180 185 190  
 Lys Leu Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val  
 195 200 205  
 Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 82

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 82



Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Ile Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380

Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 83

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 83

Arg Ala Ser Gln Ser Ile Ser Gly Tyr Leu Asn  
 1 5 10

&lt;210&gt; 84

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

<400> 84  
 Asn Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 85  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 85  
 Gln Gln Ser Tyr Thr Ser Pro Phe Ile Thr  
 1 5 10  
 <210> 86  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 86  
 Pro Tyr Pro Met Ile  
 1 5  
 <210> 87  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 87  
 Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 88  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 88  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 89  
 <211> 110  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 89  
 Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Gly Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Phe Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
 100 105 110  
 <210> 90  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 90

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Ile Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 91

&lt;211&gt; 217

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 91

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Gly Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr  
 100 105 110  
 Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu  
 115 120 125  
 Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro  
 130 135 140  
 Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly  
 145 150 155 160  
 Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr  
 165 170 175  
 Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His  
 180 185 190  
 Lys Leu Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val  
 195 200 205  
 Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 92

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 92

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 93

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 93

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn  
 1 5 10

&lt;210&gt; 94

&lt;211&gt; 7

&lt;212&gt; PRT

<213> Homo Sapiens  
 <400> 94  
 Asn Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 95  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 95  
 Gln Gln Ser Tyr Thr Ser Pro Gly Ile Thr  
 1 5 10  
 <210> 96  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 96  
 Pro Tyr Pro Met Met  
 1 5  
 <210> 97  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 97  
 Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 98  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 98  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 99  
 <211> 110  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 99  
 Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Gly Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
 100 105 110  
 <210> 100  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 100

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 101

&lt;211&gt; 217

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 101

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Gly Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Gly Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr  
 100 105 110  
 Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu  
 115 120 125  
 Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro  
 130 135 140  
 Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly  
 145 150 155 160  
 Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr  
 165 170 175  
 Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His  
 180 185 190  
 Lys Leu Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val  
 195 200 205  
 Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 102

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 102

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 103

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 103

Arg Ala Ser Gln Ser Ile Ser Gly Tyr Leu Asn  
 1 5 10

&lt;210&gt; 104

&lt;211&gt; 7

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

<400> 104  
 Asn Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 105  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 105  
 Gln Gln Ser Tyr Thr Ser Pro Gly Ile Thr  
 1 5 10  
 <210> 106  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 106  
 Pro Tyr Pro Met Met  
 1 5  
 <210> 107  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 107  
 Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 108  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 108  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 109  
 <211> 110  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 109  
 Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Gly Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Gly Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
 100 105 110  
 <210> 110  
 <211> 121  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 110



Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 111

&lt;211&gt; 217

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 111

Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Phe Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys Arg Thr  
 100 105 110  
 Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu  
 115 120 125  
 Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro  
 130 135 140  
 Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly  
 145 150 155 160  
 Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr  
 165 170 175  
 Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His  
 180 185 190  
 Lys Leu Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val  
 195 200 205  
 Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 112

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 112

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val  
  
 115 120 125  
 Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala  
 130 135 140  
 Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser  
 145 150 155 160  
 Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val  
 165 170 175  
 Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro  
 180 185 190  
 Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
 195 200 205  
 Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp  
 210 215 220  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 225 230 235 240  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 245 250 255  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 260 265 270  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 275 280 285  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 290 295 300  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 305 310 315 320  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 325 330 335  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 340 345 350  
 Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu  
 355 360 365  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 370 375 380  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 385 390 395 400  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 405 410 415  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 420 425 430  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 435 440 445

Gly

&lt;210&gt; 113

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 113

Arg Ala Ser Gln Ser Ile Ser Ser Tyr Leu Asn  
 1 5 10

&lt;210&gt; 114

&lt;211&gt; 7

&lt;212&gt; PRT

<213> Homo Sapiens  
 <400> 114  
 Asn Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 115  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 115  
 Gln Gln Ser Tyr Thr Ser Pro Phe Ile Thr  
 1 5 10  
 <210> 116  
 <211> 5  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 116  
 Pro Tyr Pro Met Met  
 1 5  
 <210> 117  
 <211> 17  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 117  
 Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val Lys  
 1 5 10 15  
 Gly  
 <210> 118  
 <211> 11  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 118  
 Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr  
 1 5 10  
 <210> 119  
 <211> 110  
 <212> PRT  
 <213> Homo Sapiens  
 <400> 119  
 Ala Gln Asp Ile Gln Met Thr Gln Ser Pro Ala Thr Leu Ser Ala Ser  
 1 5 10 15  
 Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Ile Ser  
 20 25 30  
 Ser Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu  
 35 40 45  
 Leu Ile Tyr Asn Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe  
 50 55 60  
 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu  
 65 70 75 80  
 Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Ser Tyr Thr Ser  
 85 90 95  
 Pro Phe Ile Thr Phe Gly Pro Gly Thr Lys Val Glu Ile Lys  
 100 105 110  
 <210> 120  
 <211> 121  
 <212> **PRT**  
 <213> Homo Sapiens  
 <400> 120

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Pro Tyr  
 20 25 30  
 Pro Met Met Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ser Tyr Ile Ser Pro Ser Gly Gly Lys Thr His Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr Trp Gly Gln  
 100 105 110  
 Gly Thr Leu Val Thr Val Ser Ser Ala  
 115 120

&lt;210&gt; 121

&lt;211&gt; 218

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 121

Asp Ile Val Leu Thr Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly  
 1 5 10 15  
 Gln Arg Ala Thr Ile Ser Cys Arg Ala Asn Lys Ser Val Ser Thr Ser  
 20 25 30  
 Ser Tyr Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro  
 35 40 45  
 Lys Leu Leu Ile Lys Tyr Ala Ser Asn Leu Glu Ser Gly Val Pro Ala  
 50 55 60  
 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Ile Leu Asn Ile His  
 65 70 75 80  
 Pro Val Glu Glu Glu Asp Ala Ala Thr Tyr Tyr Cys Gln His Ser Arg  
 85 90 95  
 Glu Leu Pro Phe Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile Lys Arg  
 100 105 110  
 Ala Asp Ala Ala Pro Thr Val Ser Ile Phe Pro Pro Ser Ser Glu Gln  
 115 120 125  
 Leu Thr Ser Gly Gly Ala Ser Val Val Cys Phe Leu Asn Asn Phe Tyr  
 130 135 140  
 Pro Lys Asp Ile Asn Val Lys Trp Lys Ile Asp Gly Ser Glu Arg Gln  
 145 150 155 160  
 Asn Gly Val Leu Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser Thr  
 165 170 175  
 Tyr Ser Met Ser Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu Arg  
 180 185 190  
 His Asn Ser Tyr Thr Cys Glu Ala Thr His Lys Thr Ser Thr Ser Pro  
 195 200 205  
 Ile Val Lys Ser Phe Asn Arg Asn Glu Cys  
 210 215

&lt;210&gt; 122

&lt;211&gt; 450

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 122

Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Val Val Arg Pro Gly Val  
 1 5 10 15  
 Ser Val Lys Ile Ser Cys Lys Gly Ser Gly Tyr Thr Phe Thr Asp Tyr  
 20 25 30



Gln Arg Ala Thr Ile Ser Cys Arg Ala Ser Gln Ser Val Ser Thr Ser  
 20 25 30  
 Ser Tyr Ser Tyr Met Gln Trp Tyr Gln Gln Arg Pro Gly Gln Pro Pro  
 35 40 45  
 Lys Leu Leu Ile Lys Tyr Ala Thr Asn Leu Asp Ser Gly Val Pro Ala  
 50 55 60  
 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His  
 65 70 75 80  
 Pro Val Glu Glu Glu Asp Ala Ala Thr Tyr Tyr Cys Gln His Ser Trp  
 85 90 95  
 Glu Ile Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg  
 100 105 110  
 Ala Asp Ala Ala Pro Thr Val Ser Ile Phe Pro Pro Ser Ser Glu Gln  
 115 120 125  
 Leu Thr Ser Gly Gly Ala Ser Val Val Cys Phe Leu Asn Asn Phe Tyr  
 130 135 140  
 Pro Lys Asp Ile Asn Val Lys Trp Lys Ile Asp Gly Ser Glu Arg Gln  
 145 150 155 160  
 Asn Gly Val Leu Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser Thr  
 165 170 175  
 Tyr Ser Met Ser Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu Arg  
 180 185 190  
 His Asn Ser Tyr Thr Cys Glu Ala Thr His Lys Thr Ser Thr Ser Pro  
 195 200 205  
 Ile Val Lys Ser Phe Asn Arg Asn Glu Cys  
 210 215

&lt;210&gt; 124

&lt;211&gt; 448

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 124

Glu Val Lys Leu Glu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Thr Phe Asn Asn Tyr  
 20 25 30  
 Trp Met Ser Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Leu  
 35 40 45  
 Ala Glu Ile Arg Leu Lys Ser Asp Asn Tyr Ala Thr His Tyr Ala Glu  
 50 55 60  
 Ser Val Lys Gly Lys Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Arg  
 65 70 75 80  
 Leu Tyr Leu Gln Met Asn Asn Leu Arg Ala Glu Asn Thr Gly Ile Tyr  
 85 90 95  
 Tyr Cys Thr Gly Gly Phe Ala Asp Tyr Phe Asp Tyr Trp Gly Gln Gly  
 100 105 110  
 Thr Thr Leu Thr Val Ser Ser Ala Lys Thr Thr Ala Pro Ser Val Tyr  
 115 120 125  
 Pro Leu Ala Pro Val Cys Gly Asp Thr Thr Gly Ser Ser Val Thr Leu  
 130 135 140  
 Gly Cys Leu Val Lys Gly Tyr Phe Pro Glu Pro Val Thr Leu Thr Trp  
 145 150 155 160  
 Asn Ser Gly Ser Leu Ser Ser Gly Val His Thr Phe Pro Ala Val Leu  
 165 170 175  
 Gln Ser Asp Leu Tyr Thr Leu Ser Ser Ser Val Thr Val Thr Ser Ser  
 180 185 190  
 Thr Trp Pro Ser Gln Ser Ile Thr Cys Asn Val Ala His Pro Ala Ser  
 195 200 205  
 Ser Thr Lys Val Asp Lys Lys Ile Glu Pro Arg Gly Pro Thr Ile Lys  
 210 215 220  
 Pro Cys Pro Pro Cys Lys Cys Pro Ala Pro Asn Leu Leu Gly Gly Pro  
 225 230 235 240

Ser Val Phe Ile Phe Pro Pro Lys Ile Lys Asp Val Leu Met Ile Ser  
 245 250 255  
 Leu Ser Pro Ile Val Thr Cys Val Val Val Asp Val Ser Glu Asp Asp  
 260 265 270  
 Pro Asp Val Gln Ile Ser Trp Phe Val Asn Asn Val Glu Val His Thr  
 275 280 285  
 Ala Gln Thr Gln Thr His Arg Glu Asp Tyr Asn Ser Thr Leu Arg Val  
 290 295 300  
 Val Ser Ala Leu Pro Ile Gln His Gln Asp Trp Met Ser Gly Lys Glu  
 305 310 315 320  
 Phe Lys Cys Lys Val Asn Asn Lys Asp Leu Pro Ala Pro Ile Glu Arg  
 325 330 335  
 Thr Ile Ser Lys Pro Lys Gly Ser Val Arg Ala Pro Gln Val Tyr Val  
 340 345 350  
 Leu Pro Pro Pro Glu Glu Glu Met Thr Lys Lys Gln Val Thr Leu Thr  
 355 360 365  
 Cys Met Val Thr Asp Phe Met Pro Glu Asp Ile Tyr Val Glu Trp Thr  
 370 375 380  
 Asn Asn Gly Lys Thr Glu Leu Asn Tyr Lys Asn Thr Glu Pro Val Leu  
 385 390 395 400  
 Asp Ser Asp Gly Ser Tyr Phe Met Tyr Ser Lys Leu Arg Val Glu Lys  
 405 410 415  
 Lys Asn Trp Val Glu Arg Asn Ser Tyr Ser Cys Ser Val Val His Glu  
 420 425 430  
 Gly Leu His Asn His His Thr Thr Lys Ser Phe Ser Arg Thr Pro Gly  
 435 440 445

&lt;210&gt; 125

&lt;211&gt; 218

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 125

Asp Ile Val Leu Thr Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly  
 1 5 10 15  
 Gln Arg Ala Thr Ile Ser Cys Arg Ala Ser Lys Ser Val Ser Thr Ser  
 20 25 30  
 Ser Tyr Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro  
 35 40 45  
 Lys Leu Leu Ile Lys Tyr Ala Ser Asn Leu Glu Ser Gly Val Pro Ala  
 50 55 60  
 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Ser Leu Asn Ile His  
 65 70 75 80  
 Pro Met Glu Glu Asp Asp Thr Ala Met Tyr Phe Cys Gln His Ser Arg  
 85 90 95  
 Glu Leu Pro Phe Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg  
 100 105 110  
 Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln  
 115 120 125  
 Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr  
 130 135 140  
 Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser  
 145 150 155 160  
 Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr  
 165 170 175  
 Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys  
 180 185 190  
 His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro  
 195 200 205  
 Val Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 126

&lt;211&gt; 450

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 126

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala  
 1 5 10 15  
 Ser Val Lys Val Ser Cys Lys Gly Ser Gly Tyr Thr Phe Thr Asp Tyr  
 20 25 30  
 Gly Met His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met  
 35 40 45  
 Gly Val Ile Ser Thr Tyr Asn Gly Tyr Thr Asn Tyr Asn Gln Lys Phe  
 50 55 60  
 Lys Gly Arg Val Thr Met Thr Val Asp Lys Ser Thr Ser Thr Ala Tyr  
 65 70 75 80  
 Met Glu Leu Arg Ser Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys  
 85 90 95  
 Ala Arg Ala Tyr Tyr Gly Asn Leu Tyr Tyr Ala Met Asp Tyr Trp Gly  
 100 105 110  
 Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser  
 115 120 125  
 Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala  
 130 135 140  
 Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val  
 145 150 155 160  
 Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala  
 165 170 175  
 Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val  
 180 185 190  
 Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His  
 195 200 205  
 Lys Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys  
 210 215 220  
 Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly  
 225 230 235 240  
 Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met  
 245 250 255  
 Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His  
 260 265 270  
 Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val  
 275 280 285  
 His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr  
 290 295 300  
 Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly  
 305 310 315 320  
 Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile  
 325 330 335  
 Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val  
 340 345 350  
 Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser  
 355 360 365  
 Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu  
 370 375 380  
 Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro  
 385 390 395 400  
 Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val  
 405 410 415  
 Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met  
 420 425 430  
 His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser  
 435 440 445

Pro Gly  
450

<210> 127

<211> 218

<212> PRT

<213> Homo Sapiens

<400> 127



Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly  
 1 5 10 15  
 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Ser Val Ser Thr Ser  
 20 25 30  
 Ser Tyr Ser Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro  
 35 40 45  
 Lys Leu Leu Ile Lys Tyr Ala Ser Asn Leu Glu Ser Gly Val Pro Ser  
 50 55 60  
 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser  
 65 70 75 80  
 Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys Gln His Ser Trp  
 85 90 95  
 Glu Ile Pro Tyr Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys Arg  
 100 105 110  
 Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln  
 115 120 125  
 Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr  
 130 135 140  
 Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser  
 145 150 155 160  
 Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr  
 165 170 175  
 Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys  
 180 185 190  
 His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro  
 195 200 205  
 Val Thr Lys Ser Phe Asn Arg Gly Glu Cys  
 210 215

&lt;210&gt; 128

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 128

Gln Val Glu Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly  
 1 5 10 15  
 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr  
 20 25 30  
 Trp Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ala Glu Ile Arg Leu Lys Ser Asp Asn Tyr Ala Thr His Tyr Ala Glu  
 50 55 60  
 Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Asn Ser  
 65 70 75 80  
 Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr  
 85 90 95  
 Tyr Cys Thr Gly Tyr Tyr Ala Asp Ala Met Asp Tyr Trp Gly Gln Gly  
 100 105 110  
 Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe  
 115 120 125  
 Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu  
 130 135 140  
 Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp

145					150					155					160
Asn	Ser	Gly	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu
				165					170					175	
Gln	Ser	Ser	Gly	Leu	Tyr	Ser	Leu	Ser	Ser	Val	Val	Thr	Val	Pro	Ser
			180					185					190		
Ser	Ser	Leu	Gly	Thr	Gln	Thr	Tyr	Ile	Cys	Asn	Val	Asn	His	Lys	Pro
		195					200					205			
Ser	Asn	Thr	Lys	Val	Asp	Lys	Lys	Val	Glu	Pro	Lys	Ser	Cys	Asp	Lys
	210					215					220				
Thr	His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro
225					230					235					240
Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser
				245					250					255	
Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp
			260					265					270		
Pro	Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn
		275					280					285			
Ala	Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val
	290					295					300				
Val	Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu
305					310					315					320
Tyr	Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys
				325					330					335	
Thr	Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr
			340					345					350		
Leu	Pro	Pro	Ser	Arg	Asp	Glu	Leu	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr
		355					360					365			
Cys	Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu
	370					375					380				
Ser	Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu
385					390					395					400
Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys
				405					410					415	
Ser	Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu
			420					425					430		
Ala	Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly
		435					440					445			

Lys

&lt;210&gt; 129

&lt;211&gt; 218

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 129

Asp	Ile	Val	Leu	Thr	Gln	Ser	Pro	Ala	Ser	Leu	Ala	Val	Ser	Leu	Gly
1			5					10					15		
Gln	Arg	Ala	Thr	Ile	Ser	Cys	Lys	Ala	Ser	Gln	Ser	Val	Ser	Thr	Ser
			20					25					30		
Thr	Tyr	Ser	Tyr	Met	Gln	Trp	Tyr	Gln	Gln	Arg	Pro	Gly	Gln	Ser	Pro
		35					40					45			
Lys	Leu	Leu	Ile	Lys	Tyr	Ala	Ser	Lys	Leu	Asp	Ser	Gly	Val	Pro	Ala
	50					55				60					
Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Thr	Leu	Asn	Ile	His
65					70				75						80
Pro	Val	Glu	Glu	Glu	Asp	Thr	Ala	Thr	Tyr	Tyr	Cys	Gln	His	Ser	Trp
				85					90					95	
Glu	Leu	Pro	Tyr	Thr	Phe	Gly	Gly	Gly	Thr	Arg	Leu	Glu	Ile	Lys	Arg
			100					105					110		
Ala	Asp	Ala	Ala	Pro	Thr	Val	Ser	Ile	Phe	Pro	Pro	Ser	Ser	Glu	Gln
	115						120					125			
Leu	Thr	Ser	Gly	Gly	Ala	Ser	Val	Val	Cys	Phe	Leu	Asn	Asn	Phe	Tyr

130	135	140
Pro Lys Asp Ile Asn Val Lys Trp Lys Ile Asp Gly Ser Glu Arg Gln		
145	150	155
Asn Gly Val Leu Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser Thr		
	165	170
Tyr Ser Met Ser Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu Arg		
	180	185
His Asn Ser Tyr Thr Cys Glu Ala Thr His Lys Thr Ser Thr Ser Pro		
	195	200
Ile Val Lys Ser Phe Asn Arg Asn Glu Cys		205
	210	215

&lt;210&gt; 130

&lt;211&gt; 448

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 130

Glu Val Lys Leu Gly Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly		
1	5	10
Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Pro Phe Thr Lys Tyr		
	20	25
Trp Met Asn Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Val		
	35	40
Ala Glu Ile Arg Leu Lys Ser Asp Asn Tyr Ala Thr His Tyr Ala Glu		
	50	55
Ser Ala Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Arg Ser Ser		
65	70	75
Val Tyr Leu Gln Met Asn Asn Leu Arg Ala Glu Asp Thr Ala Ile Tyr		
	85	90
Tyr Cys Ser Pro Thr Tyr Ala Asp Thr Met Asp Tyr Trp Gly Gln Gly		
	100	105
Thr Ser Val Thr Val Ser Ser Ala Lys Thr Thr Ala Pro Ser Val Tyr		
	115	120
Pro Leu Ala Pro Val Cys Gly Asp Thr Thr Gly Ser Ser Val Thr Leu		
	130	135
Gly Cys Leu Val Lys Gly Tyr Phe Pro Glu Pro Val Thr Leu Thr Trp		
145	150	155
Asn Ser Gly Ser Leu Ser Ser Gly Val His Thr Phe Pro Ala Val Leu		
	165	170
Gln Ser Asp Leu Tyr Thr Leu Ser Ser Ser Val Thr Val Thr Ser Ser		
	180	185
Thr Trp Pro Ser Gln Ser Ile Thr Cys Asn Val Ala His Pro Ala Ser		
	195	200
Ser Thr Lys Val Asp Lys Lys Ile Glu Pro Arg Gly Pro Thr Ile Lys		
	210	215
Pro Cys Pro Pro Cys Lys Cys Pro Ala Pro Asn Leu Leu Gly Gly Pro		
225	230	235
Ser Val Phe Ile Phe Pro Pro Lys Ile Lys Asp Val Leu Met Ile Ser		
	245	250
Leu Ser Pro Ile Val Thr Cys Val Val Val Asp Val Ser Glu Asp Asp		
	260	265
Pro Asp Val Gln Ile Ser Trp Phe Val Asn Asn Val Glu Val His Thr		
	275	280
Ala Gln Thr Gln Thr His Arg Glu Asp Tyr Asn Ser Thr Leu Arg Val		
	290	295
Val Ser Ala Leu Pro Ile Gln His Gln Asp Trp Met Ser Gly Lys Glu		
305	310	315
Phe Lys Cys Lys Val Asn Asn Lys Asp Leu Pro Ala Pro Ile Glu Arg		
	325	330
Thr Ile Ser Lys Pro Lys Gly Ser Val Arg Ala Pro Gln Val Tyr Val		
	340	345
Leu Pro Pro Pro Glu Glu Glu Met Thr Lys Lys Gln Val Thr Leu Thr		

	355		360		365										
Cys	Met	Val	Thr	Asp	Phe	Met	Pro	Glu	Asp	Ile	Tyr	Val	Glu	Trp	Thr
	370					375					380				
Asn	Asn	Gly	Lys	Thr	Glu	Leu	Asn	Tyr	Lys	Asn	Thr	Glu	Pro	Val	Leu
385					390					395					400
Asp	Ser	Asp	Gly	Ser	Tyr	Phe	Met	Tyr	Ser	Lys	Leu	Arg	Val	Glu	Lys
				405					410					415	
Lys	Asn	Trp	Val	Glu	Arg	Asn	Ser	Tyr	Ser	Cys	Ser	Val	Val	His	Glu
			420					425					430		
Gly	Leu	His	Asn	His	His	Thr	Thr	Lys	Ser	Phe	Ser	Arg	Thr	Pro	Gly
	435					440						445			

&lt;210&gt; 131

&lt;211&gt; 218

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 131

Asp	Ile	Val	Leu	Thr	Gln	Ser	Pro	Ala	Ser	Leu	Ala	Val	Ser	Leu	Gly
1				5					10					15	
Gln	Arg	Ala	Thr	Ile	Ser	Cys	Arg	Ala	Ser	Lys	Ser	Val	Ser	Thr	Ser
			20					25					30		
Ser	Tyr	Ser	Tyr	Met	His	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Gln	Pro	Pro
		35					40					45			
Lys	Leu	Leu	Ile	Lys	Tyr	Thr	Ser	Asn	Leu	Glu	Ser	Gly	Val	Pro	Ala
	50					55					60				
Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	Phe	Ile	Leu	Asn	Ile	His
65					70					75					80
Pro	Val	Glu	Glu	Glu	Asp	Ala	Ala	Thr	Tyr	Tyr	Cys	Gln	His	Ser	Arg
				85					90					95	
Glu	Leu	Pro	Trp	Thr	Phe	Gly	Gly	Gly	Thr	Lys	Leu	Glu	Ile	Lys	Arg
			100					105						110	
Ala	Asp	Ala	Ala	Pro	Thr	Val	Ser	Ile	Phe	Pro	Pro	Ser	Ser	Glu	Gln
	115						120						125		
Leu	Thr	Ser	Gly	Gly	Ala	Ser	Val	Val	Cys	Phe	Leu	Asn	Asn	Phe	Tyr
	130					135					140				
Pro	Lys	Asp	Ile	Asn	Val	Lys	Trp	Lys	Ile	Asp	Gly	Ser	Glu	Arg	Gln
145				150						155					160
Asn	Gly	Val	Leu	Asn	Ser	Trp	Thr	Asp	Gln	Asp	Ser	Lys	Asp	Ser	Thr
				165					170					175	
Tyr	Ser	Met	Ser	Ser	Thr	Leu	Thr	Leu	Thr	Lys	Asp	Glu	Tyr	Glu	Arg
			180					185					190		
His	Asn	Ser	Tyr	Thr	Cys	Glu	Ala	Thr	His	Lys	Thr	Ser	Thr	Ser	Pro
	195						200						205		
Ile	Val	Lys	Ser	Phe	Asn	Arg	Asn	Glu	Cys						
	210					215									

&lt;210&gt; 132

&lt;211&gt; 451

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 132

Gln	Val	Ser	Leu	Lys	Glu	Ser	Gly	Pro	Gly	Ile	Leu	Gln	Pro	Ser	Gln
1				5					10					15	
Thr	Leu	Ser	Leu	Thr	Cys	Ser	Phe	Ser	Gly	Phe	Ser	Leu	Ser	Thr	Ser
			20					25					30		
Gly	Met	Gly	Val	Ser	Trp	Ile	Arg	Gln	Pro	Ser	Gly	Lys	Gly	Leu	Glu
	35						40					45			
Trp	Leu	Ala	His	Ile	Tyr	Trp	Asp	Asp	Asp	Lys	Arg	Tyr	Asn	Pro	Ser
	50					55					60				

Leu Lys Ser Arg Leu Thr Ile Ser Lys Asp Thr Ser Arg Asn Gln Val  
 65 70 75 80  
 Phe Leu Lys Ile Thr Ser Val Asp Thr Ala Asp Thr Ala Thr Tyr Tyr  
 85 90 95  
 Cys Ala Arg Arg Gly Pro Asp Tyr Tyr Gly Tyr Tyr Pro Met Asp Tyr  
 100 105 110  
 Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ala Lys Thr Thr Ala Pro  
 115 120 125  
 Ser Val Tyr Pro Leu Ala Pro Val Cys Gly Asp Thr Thr Gly Ser Ser  
 130 135 140  
 Val Thr Leu Gly Cys Leu Val Lys Gly Tyr Phe Pro Glu Pro Val Thr  
 145 150 155 160  
 Leu Thr Trp Asn Ser Gly Ser Leu Ser Ser Gly Val His Thr Phe Pro  
 165 170 175  
 Ala Val Leu Gln Ser Asp Leu Tyr Thr Leu Ser Ser Ser Val Thr Val  
 180 185 190  
 Thr Ser Ser Thr Trp Pro Ser Gln Ser Ile Thr Cys Asn Val Ala His  
 195 200 205  
 Pro Ala Ser Ser Thr Lys Val Asp Lys Lys Ile Glu Pro Arg Gly Pro  
 210 215 220  
 Thr Ile Lys Pro Cys Pro Pro Cys Lys Cys Pro Ala Pro Asn Leu Leu  
 225 230 235 240  
 Gly Gly Pro Ser Val Phe Ile Phe Pro Pro Lys Ile Lys Asp Val Leu  
 245 250 255  
 Met Ile Ser Leu Ser Pro Ile Val Thr Cys Val Val Val Asp Val Ser  
 260 265 270  
 Glu Asp Asp Pro Asp Val Gln Ile Ser Trp Phe Val Asn Asn Val Glu  
 275 280 285  
 Val His Thr Ala Gln Thr Gln Thr His Arg Glu Asp Tyr Asn Ser Thr  
 290 295 300  
 Leu Arg Val Val Ser Ala Leu Pro Ile Gln His Gln Asp Trp Met Ser  
 305 310 315 320  
 Gly Lys Glu Phe Lys Cys Lys Val Asn Asn Lys Asp Leu Pro Ala Pro  
 325 330 335  
 Ile Glu Arg Thr Ile Ser Lys Pro Lys Gly Ser Val Arg Ala Pro Gln  
 340 345 350  
 Val Tyr Val Leu Pro Pro Pro Glu Glu Glu Met Thr Lys Lys Gln Val  
 355 360 365  
 Thr Leu Thr Cys Met Val Thr Asp Phe Met Pro Glu Asp Ile Tyr Val  
 370 375 380  
 Glu Trp Thr Asn Asn Gly Lys Thr Glu Leu Asn Tyr Lys Asn Thr Glu  
 385 390 395 400  
 Pro Val Leu Asp Ser Asp Gly Ser Tyr Phe Met Tyr Ser Lys Leu Arg  
 405 410 415  
 Val Glu Lys Lys Asn Trp Val Glu Arg Asn Ser Tyr Ser Cys Ser Val  
 420 425 430  
 Val His Glu Gly Leu His Asn His His Thr Thr Lys Ser Phe Ser Arg  
 435 440 445  
 Thr Pro Gly  
 450

&lt;210&gt; 133

&lt;211&gt; 282

&lt;212&gt; PRT

<213> *Macaca fascicularis*

&lt;400&gt; 133

Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser His Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ser Ala Ala Pro Pro Ala Pro Phe  
 35 40 45

Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175  
 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 195 200 205  
 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 210 215 220  
 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 225 230 235 240  
 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 245 250 255  
 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 260 265 270  
 Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 275 280

&lt;210&gt; 134

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Rattus norvegicus

&lt;400&gt; 134

Glu Gln Ala Pro Gly Asn Ala Pro Cys Ser Ser Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Pro Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala His Phe  
 35 40 45  
 Arg Met Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175  
 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 195 200 205

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 210 215 220  
 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 225 230 235 240  
 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 245 250 255  
 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 260 265 270  
 Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 275 280

&lt;210&gt; 135

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Sus scrofa

&lt;400&gt; 135

Glu Arg Val Pro Gly Thr Thr Pro Cys Ser Arg Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Pro Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Ser Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175  
 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 195 200 205  
 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 210 215 220  
 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 225 230 235 240  
 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 245 250 255  
 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 260 265 270  
 Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 275 280

&lt;210&gt; 136

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Canis lupus

&lt;400&gt; 136

Glu Arg Val Pro Gly Thr Thr Pro Cys Pro Arg Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro

			20					25				30				
His	Ser	Asp	Phe	Cys	Leu	Gly	Cys	Thr	Ala	Ala	Pro	Pro	Ala	Pro	Phe	
		35					40					45				
Arg	Leu	Leu	Trp	Pro	Arg	Ser	Asp	Lys	Thr	His	Thr	Cys	Pro	Pro	Cys	
	50					55					60					
Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	
65				70						75					80	
Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	
			85						90					95		
Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp	
			100					105						110		
Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu	
		115						120					125			
Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val	Ser	Val	Leu	Thr	Val	Leu	
		130					135					140				
His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn	
145					150						155				160	
Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly	
				165					170					175		
Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser	Arg	Glu	Glu	
			180					185						190		
Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	
		195					200						205			
Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	
		210					215					220				
Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	
225					230					235					240	
Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn	
				245					250					255		
Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr	Thr	
			260					265					270			
Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys							
		275					280									

&lt;210&gt; 137

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Mus Musculus

&lt;400&gt; 137

Glu	Gln	Ala	Pro	Gly	Thr	Ser	Pro	Cys	Ser	Ser	Gly	Ser	Ser	Trp	Ser	
1				5				10						15		
Ala	Asp	Leu	Asp	Lys	Cys	Met	Asp	Cys	Ala	Ser	Cys	Pro	Ala	Arg	Pro	
			20					25					30			
His	Ser	Asp	Phe	Cys	Leu	Gly	Cys	Ala	Ala	Ala	Pro	Pro	Ala	His	Phe	
		35					40					45				
Arg	Leu	Leu	Trp	Pro	Arg	Ser	Asp	Lys	Thr	His	Thr	Cys	Pro	Pro	Cys	
	50					55					60					
Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	
65				70						75					80	
Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	
			85						90					95		
Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp	
			100					105						110		
Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu	
		115						120					125			
Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val	Ser	Val	Leu	Thr	Val	Leu	
		130					135					140				
His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn	
145					150						155				160	
Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly	
				165					170					175		
Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser	Arg	Glu	Glu	



			180					185				190				
Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	
		195					200					205				
Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	
	210					215					220					
Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	
225					230				235						240	
Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn	
			245					250						255		
Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr	Thr	
			260					265				270				
Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys							
		275					280									

&lt;210&gt; 138

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 138

Glu	Gln	Ala	Pro	Gly	Thr	Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser	
1				5				10						15		
Ala	Asp	Leu	Asp	Lys	Cys	Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	
			20					25					30			
His	Ser	Asp	Phe	Cys	Leu	Gly	Cys	Ala	Ala	Ala	Pro	Pro	Ala	Pro	Phe	
		35					40					45				
Arg	Leu	Leu	Trp	Pro	Arg	Ser	Asp	Lys	Thr	His	Thr	Cys	Pro	Pro	Cys	
	50					55					60					
Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	
65					70					75					80	
Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	
				85					90					95		
Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp	
			100					105					110			
Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu	
		115					120					125				
Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val	Ser	Val	Leu	Thr	Val	Leu	
		130					135					140				
His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn	
145					150					155					160	
Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly	
				165					170					175		
Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser	Arg	Glu	Glu	
			180					185					190			
Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	
		195					200					205				
Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	
	210					215					220					
Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	
225					230				235						240	
Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn	
			245					250						255		
Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr	Thr	
			260					265				270				
Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys							
		275					280									

&lt;210&gt; 139

&lt;211&gt; 296

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 139

Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp  
 50 55 60  
 Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly  
 65 70 75 80  
 Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile  
 85 90 95  
 Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu  
 100 105 110  
 Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His  
 115 120 125  
 Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg  
 130 135 140  
 Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys  
 145 150 155 160  
 Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu  
 165 170 175  
 Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr  
 180 185 190  
 Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu  
 195 200 205  
 Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp  
 210 215 220  
 Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val  
 225 230 235 240  
 Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp  
 245 250 255  
 Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His  
 260 265 270  
 Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro  
 275 280 285  
 Gly Lys His His His His His His  
 290 295

<210> 140

<211> 278

<212> PRT

<213> Homo Sapiens

<400> 140

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140

Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 141

&lt;211&gt; 47

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 141

Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala His His His His His His  
 35 40 45

&lt;210&gt; 142

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 142

Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Gln Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175  
 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr

		195					200					205							
Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn				
	210					215					220								
Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe				
225					230				235						240				
Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn				
			245					250						255					
Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr	Thr				
		260						265				270							
Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys										
	275					280													

&lt;210&gt; 143

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 143

Glu	Gln	Ala	Pro	Gly	Thr	Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser				
1				5				10						15					
Ala	Asp	Leu	Asp	Lys	Cys	Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro				
			20					25					30						
Lys	Ser	Asp	Phe	Cys	Leu	Gly	Cys	Ala	Ala	Ala	Pro	Pro	Ala	Pro	Phe				
		35				40						45							
Arg	Leu	Leu	Trp	Pro	Arg	Ser	Asp	Lys	Thr	His	Thr	Cys	Pro	Pro	Cys				
	50					55					60								
Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro				
65				70						75					80				
Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys				
			85					90						95					
Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp				
			100					105					110						
Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala	Lys	Thr	Lys	Pro	Arg	Glu				
		115					120						125						
Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val	Ser	Val	Leu	Thr	Val	Leu				
		130					135					140							
His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	Lys	Cys	Lys	Val	Ser	Asn				
145				150						155					160				
Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly				
				165					170					175					
Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser	Arg	Glu	Glu				
			180					185					190						
Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr				
		195					200						205						
Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn				
	210					215					220								
Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe				
225					230				235						240				
Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn				
			245					250						255					
Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr	Thr				
		260						265				270							
Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys										
	275					280													

&lt;210&gt; 144

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 144

Glu	Gln	Ala	Pro	Gly	Thr	Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser				
1				5				10						15					

Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Pro Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175  
 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 195 200 205  
 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 210 215 220  
 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 225 230 235 240  
 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 245 250 255  
 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 260 265 270  
 Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 275 280

&lt;210&gt; 145

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 145

Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Ala Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 195 200 205  
 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 210 215 220  
 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 225 230 235 240  
 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 245 250 255  
 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 260 265 270  
 Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 275 280

&lt;210&gt; 146

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 146

Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser Ser Ala Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175  
 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 195 200 205  
 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 210 215 220  
 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 225 230 235 240  
 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 245 250 255  
 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 260 265 270  
 Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 275 280

&lt;210&gt; 147

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 147

Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Arg Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn  
 145 150 155 160  
 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly  
 165 170 175  
 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu  
 180 185 190  
 Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr  
 195 200 205  
 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn  
 210 215 220  
 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe  
 225 230 235 240  
 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn  
 245 250 255  
 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr  
 260 265 270  
 Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 275 280

&lt;210&gt; 148

&lt;211&gt; 282

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 148

Glu Gln Ala Pro Gly Gln Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser  
 1 5 10 15  
 Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro  
 20 25 30  
 His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe  
 35 40 45  
 Arg Leu Leu Trp Pro Arg Ser Asp Lys Thr His Thr Cys Pro Pro Cys  
 50 55 60  
 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro  
 65 70 75 80  
 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys  
 85 90 95  
 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp  
 100 105 110  
 Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu  
 115 120 125  
 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu  
 130 135 140  
 His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn

145					150					155					160
Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly
				165					170					175	
Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	Pro	Pro	Ser	Arg	Glu	Glu
			180					185					190		
Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr
		195					200					205			
Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn
		210				215					220				
Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe
225					230				235						240
Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn
			245					250						255	
Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	Leu	His	Asn	His	Tyr	Thr
			260					265					270		
Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys						
		275					280								

&lt;210&gt; 149

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 149

Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser	Ala	Asp	Leu	Asp	Lys	Cys
1				5					10					15	
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Ala
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr
		35				40						45			
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
		50				55					60				
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg
65					70				75						80
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro
			85						90					95	
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala
			100					105					110		
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val
		115					120					125			
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr
		130				135					140				
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr
145					150					155					160
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu
			165						170					175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys
			180					185					190		
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser
		195					200					205			
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp
		210				215					220				
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser
225					230					235					240
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala
			245					250						255	
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys
		260						265					270		
His	His	His	His	His	His										
		275													

&lt;210&gt; 150

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 150



Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Ala Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His  
 275

&lt;210&gt; 151

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 151

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Ala Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125

Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 152

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 152

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ala Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 153

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 153

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro Ala Ser Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His  
 275

&lt;210&gt; 154

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 154

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Ala His Ser Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala

			100						105				110			
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val	
		115					120					125				
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	
		130				135					140					
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	
145					150					155					160	
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	
				165					170						175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	
			180					185					190			
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	
		195					200					205				
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	
	210					215					220					
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	
225					230					235					240	
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	
				245					250					255		
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys	
			260					265					270			
His	His	His	His	His	His											
			275													

&lt;210&gt; 155

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 155

Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser	Ala	Asp	Leu	Asp	Lys	Cys	
1				5					10					15		
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Ala	Pro	His	Ser	Asp	Phe	Cys	Leu	
			20					25					30			
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr	
		35				40						45				
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser	
		50				55					60					
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg	
65					70					75					80	
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro	
				85					90					95		
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala	
			100					105					110			
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val	
		115					120					125				
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr	
		130				135					140					
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr	
145					150					155					160	
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu	
				165					170						175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys	
			180					185					190			
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser	
		195					200					205				
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp	
	210					215					220					
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser	
225					230					235					240	
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala	
				245					250					255		
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys	
			260					265					270			
His	His	His	His	His	His											
			275													

&lt;210&gt; 156

&lt;211&gt; 240

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 156

Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser	Ala	Asp	Leu	Asp	Lys	Cys
1			5					10						15	
Met	Asp	Cys	Ala	Ser	Cys	Ala	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr
		35				40						45			
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
	50				55						60				
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg
65					70				75						80
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro
			85						90				95		
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala
			100					105					110		
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val
		115					120					125			
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr
		130				135					140				
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr
145					150					155					160
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu
			165						170					175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys
			180					185					190		
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser
		195					200					205			
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp
	210					215					220				
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser
225					230					235					240

&lt;210&gt; 157

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 157

Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser	Ala	Asp	Leu	Asp	Lys	Cys
1			5					10						15	
Met	Asp	Cys	Ala	Ala	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr
		35				40						45			
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
	50				55						60				
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg
65					70				75						80
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro
			85						90				95		
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala
			100					105					110		

Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 158

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 158

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Ala Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 159

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 159

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Ala Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 160

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 160

Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Ala Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro

				85					90					95			
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala		
			100					105					110				
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val		
		115					120					125					
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr		
		130				135					140						
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr		
145					150					155					160		
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu		
			165						170					175			
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys		
			180					185					190				
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser		
		195					200					205					
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp		
	210					215					220						
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser		
225					230					235					240		
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala		
				245					250					255			
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys		
			260					265						270			
His	His	His	His	His	His												
			275														

&lt;210&gt; 161

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 161

Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ser	Ala	Ala	Leu	Asp	Lys	Cys		
1				5				10						15			
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu		
			20					25					30				
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr		
		35				40						45					
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser		
	50					55					60						
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg		
65					70					75					80		
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro		
				85					90					95			
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala		
			100					105					110				
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val		
		115					120					125					
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr		
		130				135					140						
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr		
145					150					155					160		
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu		
			165						170					175			
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys		
			180					185					190				
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser		
		195					200					205					
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp		
	210					215					220						
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser		
225					230					235					240		
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala		
				245					250					255			
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys		
			260					265						270			
His	His	His	His	His	His												
			275														

&lt;210&gt; 162

&lt;211&gt; 278



&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 162

Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Trp	Ala	Ala	Asp	Leu	Asp	Lys	Cys
1			5					10						15	
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr
			35				40					45			
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
			50				55				60				
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg
65						70				75					80
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro
						85			90					95	
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala
			100					105					110		
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val
			115				120					125			
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr
						135					140				
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr
145					150					155					160
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu
					165				170					175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys
			180					185					190		
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser
			195				200					205			
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp
						215					220				
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser
225					230					235					240
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala
					245				250					255	
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys
			260					265					270		
His	His	His	His	His	His										
			275												

&lt;210&gt; 163

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 163

Ala	Pro	Cys	Ser	Arg	Gly	Ser	Ser	Ala	Ser	Ala	Asp	Leu	Asp	Lys	Cys
1			5					10						15	
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr
			35				40					45			
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
			50				55				60				

Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 164

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 164

Ala Pro Cys Ser Arg Gly Ser Ala Trp Ser Ala Asp Leu Asp Lys Cys  
 1 5 10 15  
 Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys Leu  
 20 25 30  
 Gly Cys Ala Ile Glu Gly Arg Met Asp Pro Lys Ser Cys Asp Lys Thr  
 35 40 45  
 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser  
 50 55 60  
 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg  
 65 70 75 80  
 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro  
 85 90 95  
 Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala  
 100 105 110  
 Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val  
 115 120 125  
 Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr  
 130 135 140  
 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr  
 145 150 155 160  
 Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu  
 165 170 175  
 Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys  
 180 185 190  
 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser  
 195 200 205  
 Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp  
 210 215 220  
 Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser  
 225 230 235 240  
 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala  
 245 250 255  
 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys  
 260 265 270  
 His His His His His His  
 275

&lt;210&gt; 165

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 165

Ala	Pro	Cys	Ser	Arg	Gly	Ala	Ser	Trp	Ser	Ala	Asp	Leu	Asp	Lys	Cys
1			5					10						15	
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr
		35					40					45			
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
		50					55					60			
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg
65						70				75					80
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro
				85					90					95	
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala
			100					105					110		
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val
		115					120					125			
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr
		130				135					140				
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr
145					150					155					160
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu
			165						170					175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys
			180					185					190		
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser
		195					200					205			
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp
	210					215						220			
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser
225					230					235					240
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala
				245					250					255	
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys
			260					265					270		
His	His	His	His	His	His										
			275												

&lt;210&gt; 166

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 166

Ala	Pro	Cys	Ser	Ala	Gly	Ser	Ser	Trp	Ser	Ala	Asp	Leu	Asp	Lys	Cys
1			5					10						15	
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr

		35				40					45				
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
	50					55					60				
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg
65					70				75						80
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro
				85					90					95	
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala
			100					105					110		
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val
		115					120						125		
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr
		130				135					140				
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr
145					150					155					160
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu
				165					170					175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys
			180					185					190		
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser
		195				200						205			
Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	Asp
	210					215					220				
Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	Ser
225				230						235					240
Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	Ala
				245					250					255	
Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	Lys
			260					265					270		
His	His	His	His	His	His										
		275													

&lt;210&gt; 167

&lt;211&gt; 278

&lt;212&gt; PRT

&lt;213&gt; Homo Sapiens

&lt;400&gt; 167

Ala	Pro	Cys	Ala	Arg	Gly	Ser	Ser	Trp	Ser	Ala	Asp	Leu	Asp	Lys	Cys
1				5					10					15	
Met	Asp	Cys	Ala	Ser	Cys	Arg	Ala	Arg	Pro	His	Ser	Asp	Phe	Cys	Leu
			20					25					30		
Gly	Cys	Ala	Ile	Glu	Gly	Arg	Met	Asp	Pro	Lys	Ser	Cys	Asp	Lys	Thr
		35					40					45			
His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	Ser
	50					55					60				
Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	Arg
65					70				75						80
Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	Pro
				85					90					95	
Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	Ala
			100					105					110		
Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	Val
		115					120						125		
Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	Tyr
		130				135					140				
Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	Thr
145					150					155					160
Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	Leu
			165						170					175	
Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	Cys
			180					185					190		
Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	Ser

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      195                200                205
Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp
  210                215                220
Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser
225                230                235
Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala
      245                250
Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
      260                265                270
His His His His His His
      275

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<210> 168

<211> 53

<212> PRT

<213> Homo Sapiens

<400> 168

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Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser
 1                5                10                15
Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro
      20                25                30
His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe
      35                40                45
Arg Leu Leu Trp Pro
      50

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<210> 169

<211> 129

<212> PRT

<213> Homo Sapiens

<400> 169

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Met Ala Arg Gly Ser Leu Arg Arg Leu Leu Arg Leu Leu Val Leu Gly
 1                5                10                15
Leu Trp Leu Ala Leu Leu Arg Ser Val Ala Gly Glu Gln Ala Pro Gly
      20                25                30
Thr Ala Pro Cys Ser Arg Gly Ser Ser Trp Ser Ala Asp Leu Asp Lys
      35                40                45
Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys
      50                55                60
Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe Arg Leu Leu Trp Pro
65                70                75                80
Ile Leu Gly Gly Ala Leu Ser Leu Thr Phe Val Leu Gly Leu Leu Ser
      85                90                95
Gly Phe Leu Val Trp Arg Arg Cys Arg Arg Arg Glu Lys Phe Thr Thr
      100                105                110
Pro Ile Glu Glu Thr Gly Gly Glu Gly Cys Pro Ala Val Ala Leu Ile
      115                120                125

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Gln

<210> 170

<211> 959

<212> DNA

<213> Homo Sapiens

<400> 170

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atggctcggg gctcgtgcg cgggttgctg cggctcctcg tgctggggct ctggctggcg      60
ttgctgcgct ccgtggccgg ggagcaagcg ccaggcaccg cccctgctc ccgcggcagc      120
tcctggagcg cggacctgga caagtgcatt gactgcgcgt cttgcagggc gcgaccgcac      180

```

```

agcgacttct gcctgggctg cgctgcagca cctcctgccc ctttccggct gctttggccc 240
atccttgggg gcgctctgag cctgaccttc gtgctggggc tgctttctgg ctttttggtc 300
tggagacgat gccgcaggag agagaagtcc accacccccca tagaggagac cggcggagag 360
ggctgcccag ctgtggcgct gatccagtga caatgtgccc cctgccagcc ggggctcgcc 420
cactcatcat tcattcatcc attctagagc cagtctctgc ctcccagacg cggcgggagc 480
caagctcctc caaccacaag ggggggtgggg ggcgggtgaat cacctctgag gcctgggccc 540
agggttcagg ggaaccttcc aaggtgtctg gttgccctgc ctctggctcc agaacagaaa 600
gggagcctca cgctggctca cacaaaacag ctgacactga ctaaggaact gcagcatttg 660
cacaggggag gggggtgccc tccttcctta ggacctgggg gccaggctga cttggggggc 720
agacttgaca ctaggcccca ctcaactcaga tgtcctgaaa ttccaccacg ggggtcacc 780
tgggggggta gggacctatt ttaaacacta ggggctggcc cactaggagg gctggcccta 840
agatacagac ccccccaact ccccaaagcg gggaggagat atttattttg gggagagttt 900
ggaggggagg gagaatttat taataaaaga atctttaact ttaaaaaaaaa aaaaaaaaa 959

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<210> 171

<211> 5

<212> PRT

<213> Homo Sapiens

<220>

<221> SITE

<222> 5..5

<223> Xaa can be any naturally occurring amino acid

<400> 171

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Pro Tyr Pro Met Xaa
1           5

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<210> 172

<211> 17

<212> PRT

<213> Homo Sapiens

<220>

<221> SITE

<222> 8..8

<223> Xaa can be any naturally occurring amino acid

<400> 172

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Tyr Ile Ser Pro Ser Gly Gly Xaa Thr His Tyr Ala Asp Ser Val Lys
1           5           10           15

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Gly

<210> 173

<211> 11

<212> PRT

<213> Homo Sapiens

<400> 173

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Gly Gly Asp Thr Tyr Phe Asp Tyr Phe Asp Tyr
1           5           10

```

<210> 174

<211> 11

<212> PRT

<213> Homo Sapiens

<220>

<221> SITE  
 <222> 8..8  
 <223> Xaa can be any naturally occurring amino acid  
 <400> 174  
 Arg Ala Ser Gln Ser Ile Ser Xaa Tyr Leu Asn  
 1 5 10  
 <210> 175  
 <211> 7  
 <212> PRT  
 <213> Homo Sapiens  
 <220>  
 <221> SITE  
 <222> 1..1  
 <223> Xaa can be any naturally occurring amino acid  
 <400> 175  
 Xaa Ala Ser Ser Leu Gln Ser  
 1 5  
 <210> 176  
 <211> 10  
 <212> PRT  
 <213> Homo Sapiens  
 <220>  
 <221> SITE  
 <222> 5..6  
 <223> Xaa can be any naturally occurring amino acid  
 <220>  
 <221> SITE  
 <222> 8..8  
 <223> Xaa can be any naturally occurring amino acid  
 <400> 176  
 Gln Gln Ser Tyr Xaa Xaa Pro Xaa Ile Thr  
 1 5 10  
 <210> 177  
 <211> 645  
 <212> DNA  
 <213> Homo Sapiens  
 <400> 177  
 gacatccaga tgaccagag cccagcagc ctgagcgct ccgtgggcga cagagtgacc 60  
 atcacctgtc gggccagcca gagcatcagc ggctacctga actggtatca gcagaagccc 120  
 ggcaaggccc ccaagctgct gatctaccag gccagctccc tgcagagcgg cgtgccaaagc 180  
 agattcagcg gcagcggctc cggcaccgac ttcaccctga ccatcagcag cctgcagccc 240  
 gaggacttcg ccacctacta ctgccagcag agctacacca gcccttcat caccttcggc 300  
 cagggcacca aggtggaat caagcggacc gtggccgctc ccagcgtggt catcttcca 360  
 cccagcgacg agcagctgaa gtccggcaca gccagcgtgg tctgcctgct gaacaacttc 420  
 taccctccgagc aggccaaggt gcagtggaag gtggacaacg ccctgcagtc cggcaactcc 480  
 caggaaagcg tgaccgagca ggacagcaag gactccacct acagcctgag cagcaccctg 540  
 accctgagca aggccgacta cgagaagcac aaggtgtacg cctgcgaagt gaccaccag 600  
 ggctgtcca gcccgtgac caagagcttc aaccggggcg agtgc 645

&lt;210&gt; 178

&lt;211&gt; 1347

&lt;212&gt; DNA

&lt;213&gt; Homo Sapiens

&lt;400&gt; 178

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&lt;213&gt; Mus Musculus

&lt;400&gt; 204

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&lt;212&gt; DNA

&lt;213&gt; Homo Sapiens

&lt;400&gt; 205

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 Gly Asn Lys Asn Leu Cys Tyr Ala Asn Thr Ile Asn Trp Lys Lys Leu  
 465 470 475 480  
 Phe Gly Thr Ser Gly Gln Lys Thr Lys Ile Ile Ser Asn Arg Gly Glu  
 485 490 495  
 Asn Ser Cys Lys Ala Thr Gly Gln Val Cys His Ala Leu Cys Ser Pro  
 500 505 510  
 Glu Gly Cys Trp Gly Pro Glu Pro Arg Asp Cys Val Ser Cys Arg Asn  
 515 520 525  
 Val Ser Arg Gly Arg Glu Cys Val Asp Lys Cys Asn Leu Leu Glu Gly  
 530 535 540  
 Glu Pro Arg Glu Phe Val Glu Asn Ser Glu Cys Ile Gln Cys His Pro  
 545 550 555 560  
 Glu Cys Leu Pro Gln Ala Met Asn Ile Thr Cys Thr Gly Arg Gly Pro  
 565 570 575  
 Asp Asn Cys Ile Gln Cys Ala His Tyr Ile Asp Gly Pro His Cys Val  
 580 585 590  
 Lys Thr Cys Pro Ala Gly Val Met Gly Glu Asn Asn Thr Leu Val Trp  
 595 600 605  
 Lys Tyr Ala Asp Ala Gly His Val Cys His Leu Cys His Pro Asn Cys  
 610 615 620  
 Thr Tyr Gly Cys Thr Gly Pro Gly Leu Glu Gly Cys Pro Thr Asn Gly  
 625 630 635 640  
 Pro Lys Ile Pro Ser Ile Ala Thr Gly Met Val Gly Ala Leu Leu Leu  
 645 650 655  
 Leu Leu Val Val Ala Leu Gly Ile Gly Leu Phe Met Arg Arg Arg His  
 660 665 670  
 Ile Val Arg Lys Arg Thr Leu Arg Arg Leu Leu Gln Glu Arg Glu Leu  
 675 680 685  
 Val Glu Pro Leu Thr Pro Ser Gly Glu Ala Pro Asn Gln Ala Leu Leu  
 690 695 700  
 Arg Ile Leu Lys Glu Thr Glu Phe Lys Lys Ile Lys Val Leu Gly Ser  
 705 710 715 720  
 Gly Ala Phe Gly Thr Val Tyr Lys Gly Leu Trp Ile Pro Glu Gly Glu  
 725 730 735  
 Lys Val Lys Ile Pro Val Ala Ile Lys Glu Leu Arg Glu Ala Thr Ser  
 740 745 750  
 Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Ser  
 755 760 765  
 Val Asp Asn Pro His Val Cys Arg Leu Leu Gly Ile Cys Leu Thr Ser  
 770 775 780  
 Thr Val Gln Leu Ile Thr Gln Leu Met Pro Phe Gly Cys Leu Leu Asp  
 785 790 795 800  
 Tyr Val Arg Glu His Lys Asp Asn Ile Gly Ser Gln Tyr Leu Leu Asn  
 805 810 815  
 Trp Cys Val Gln Ile Ala Lys Gly Met Asn Tyr Leu Glu Asp Arg Arg  
 820 825 830  
 Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Thr Pro



		835					840					845			
Gln	His	Val	Lys	Ile	Thr	Asp	Phe	Gly	Leu	Ala	Lys	Leu	Leu	Gly	Ala
	850					855					860				
Glu	Glu	Lys	Glu	Tyr	His	Ala	Glu	Gly	Gly	Lys	Val	Pro	Ile	Lys	Trp
865					870					875					880
Met	Ala	Leu	Glu	Ser	Ile	Leu	His	Arg	Ile	Tyr	Thr	His	Gln	Ser	Asp
				885					890					895	
Val	Trp	Ser	Tyr	Gly	Val	Thr	Val	Trp	Glu	Leu	Met	Thr	Phe	Gly	Ser
			900					905					910		
Lys	Pro	Tyr	Asp	Gly	Ile	Pro	Ala	Ser	Glu	Ile	Ser	Ser	Ile	Leu	Glu
		915					920						925		
Lys	Gly	Glu	Arg	Leu	Pro	Gln	Pro	Pro	Ile	Cys	Thr	Ile	Asp	Val	Tyr
	930					935					940				
Met	Ile	Met	Val	Lys	Cys	Trp	Met	Ile	Asp	Ala	Asp	Ser	Arg	Pro	Lys
945					950					955					960
Phe	Arg	Glu	Leu	Ile	Ile	Glu	Phe	Ser	Lys	Met	Ala	Arg	Asp	Pro	Gln
				965					970					975	
Arg	Tyr	Leu	Val	Ile	Gln	Gly	Asp	Glu	Arg	Met	His	Leu	Pro	Ser	Pro
			980					985					990		
Thr	Asp	Ser	Asn	Phe	Tyr	Arg	Ala	Leu	Met	Asp	Glu	Glu	Asp	Met	Asp
		995					1000						1005		
Asp	Val	Val	Asp	Ala	Asp	Glu	Tyr	Leu	Ile	Pro	Gln	Gln	Gly	Phe	Phe
	1010					1015					1020				
Ser	Ser	Pro	Ser	Thr	Ser	Arg	Thr	Pro	Leu	Leu	Ser	Ser	Leu	Ser	Ala
1025					1030					1035					1040
Thr	Ser	Asn	Asn	Ser	Thr	Val	Ala	Cys	Ile	Asp	Arg	Asn	Gly	Leu	Gln
			1045						1050					1055	
Ser	Cys	Pro	Ile	Lys	Glu	Asp	Ser	Phe	Leu	Gln	Arg	Tyr	Ser	Ser	Asp
			1060					1065					1070		
Pro	Thr	Gly	Ala	Leu	Thr	Glu	Asp	Ser	Ile	Asp	Asp	Thr	Phe	Leu	Pro
	1075					1080						1085			
Val	Pro	Glu	Tyr	Ile	Asn	Gln	Ser	Val	Pro	Lys	Arg	Pro	Ala	Gly	Ser
	1090					1095					1100				
Val	Gln	Asn	Pro	Val	Tyr	His	Asn	Gln	Pro	Leu	Asn	Pro	Ala	Pro	Ser
1105					1110					1115					1120
Arg	Asp	Pro	His	Tyr	Gln	Asp	Pro	His	Ser	Thr	Ala	Val	Gly	Asn	Pro
			1125						1130					1135	
Glu	Tyr	Leu	Asn	Thr	Val	Gln	Pro	Thr	Cys	Val	Asn	Ser	Thr	Phe	Asp
			1140					1145					1150		
Ser	Pro	Ala	His	Trp	Ala	Gln	Lys	Gly	Ser	His	Gln	Ile	Ser	Leu	Asp
		1155					1160						1165		
Asn	Pro	Asp	Tyr	Gln	Gln	Asp	Phe	Phe	Pro	Lys	Glu	Ala	Lys	Pro	Asn
	1170					1175					1180				
Gly	Ile	Phe	Lys	Gly	Ser	Thr	Ala	Glu	Asn	Ala	Glu	Tyr	Leu	Arg	Val
1185					1190					1195					1200
Ala	Pro	Gln	Ser	Ser	Glu	Phe	Ile	Gly	Ala						
				1205				1210							

&lt;210&gt; 214

&lt;211&gt; 622

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 214

Met	Ala	Leu	Pro	Thr	Ala	Arg	Pro	Leu	Leu	Gly	Ser	Cys	Gly	Thr	Pro
1				5					10					15	
Ala	Leu	Gly	Ser	Leu	Leu	Phe	Leu	Leu	Phe	Ser	Leu	Gly	Trp	Val	Gln
			20					25					30		
Pro	Ser	Arg	Thr	Leu	Ala	Gly	Glu	Thr	Gly	Gln	Glu	Ala	Ala	Pro	Leu
		35					40					45			
Asp	Gly	Val	Leu	Ala	Asn	Pro	Pro	Asn	Ile	Ser	Ser	Leu	Ser	Pro	Arg
	50					55					60				
Gln	Leu	Leu	Gly	Phe	Pro	Cys	Ala	Glu	Val	Ser	Gly	Leu	Ser	Thr	Glu



&lt;400&gt; 215

Met Ala Pro Leu Cys Pro Ser Pro Trp Leu Pro Leu Leu Ile Pro Ala  
 1 5 10 15  
 Pro Ala Pro Gly Leu Thr Val Gln Leu Leu Leu Ser Leu Leu Leu  
 20 25 30  
 Val Pro Val His Pro Gln Arg Leu Pro Arg Met Gln Glu Asp Ser Pro  
 35 40 45  
 Leu Gly Gly Gly Ser Ser Gly Glu Asp Asp Pro Leu Gly Glu Glu Asp  
 50 55 60  
 Leu Pro Ser Glu Glu Asp Ser Pro Arg Glu Glu Asp Pro Pro Gly Glu  
 65 70 75 80  
 Glu Asp Leu Pro Gly Glu Glu Asp Leu Pro Gly Glu Glu Asp Leu Pro  
 85 90 95  
 Glu Val Lys Pro Lys Ser Glu Glu Glu Gly Ser Leu Lys Leu Glu Asp  
 100 105 110  
 Leu Pro Thr Val Glu Ala Pro Gly Asp Pro Gln Glu Pro Gln Asn Asn  
 115 120 125  
 Ala His Arg Asp Lys Glu Gly Asp Asp Gln Ser His Trp Arg Tyr Gly  
 130 135 140  
 Gly Asp Pro Pro Trp Pro Arg Val Ser Pro Ala Cys Ala Gly Arg Phe  
 145 150 155 160  
 Gln Ser Pro Val Asp Ile Arg Pro Gln Leu Ala Ala Phe Cys Pro Ala  
 165 170 175  
 Leu Arg Pro Leu Glu Leu Leu Gly Phe Gln Leu Pro Pro Leu Pro Glu  
 180 185 190  
 Leu Arg Leu Arg Asn Asn Gly His Ser Val Gln Leu Thr Leu Pro Pro  
 195 200 205  
 Gly Leu Glu Met Ala Leu Gly Pro Gly Arg Glu Tyr Arg Ala Leu Gln  
 210 215 220  
 Leu His Leu His Trp Gly Ala Ala Gly Arg Pro Gly Ser Glu His Thr  
 225 230 235 240  
 Val Glu Gly His Arg Phe Pro Ala Glu Ile His Val Val His Leu Ser  
 245 250 255  
 Thr Ala Phe Ala Arg Val Asp Glu Ala Leu Gly Arg Pro Gly Gly Leu  
 260 265 270  
 Ala Val Leu Ala Ala Phe Leu Glu Glu Gly Pro Glu Glu Asn Ser Ala  
 275 280 285  
 Tyr Glu Gln Leu Leu Ser Arg Leu Glu Glu Ile Ala Glu Glu Gly Ser  
 290 295 300  
 Glu Thr Gln Val Pro Gly Leu Asp Ile Ser Ala Leu Leu Pro Ser Asp  
 305 310 315 320  
 Phe Ser Arg Tyr Phe Gln Tyr Glu Gly Ser Leu Thr Thr Pro Pro Cys  
 325 330 335  
 Ala Gln Gly Val Ile Trp Thr Val Phe Asn Gln Thr Val Met Leu Ser  
 340 345 350  
 Ala Lys Gln Leu His Thr Leu Ser Asp Thr Leu Trp Gly Pro Gly Asp  
 355 360 365  
 Ser Arg Leu Gln Leu Asn Phe Arg Ala Thr Gln Pro Leu Asn Gly Arg  
 370 375 380  
 Val Ile Glu Ala Ser Phe Pro Ala Gly Val Asp Ser Ser Pro Arg Ala  
 385 390 395 400  
  
 Ala Glu Pro Val Gln Leu Asn Ser Cys Leu Ala Ala Gly Asp Ile Leu  
 405 410 415  
 Ala Leu Val Phe Gly Leu Leu Phe Ala Val Thr Ser Val Ala Phe Leu  
 420 425 430  
 Val Gln Met Arg Arg Gln His Arg Arg Gly Thr Lys Gly Gly Val Ser  
 435 440 445  
 Tyr Arg Pro Ala Glu Val Ala Glu Thr Gly Ala  
 450 455

&lt;210&gt; 216

&lt;211&gt; 346

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 216

Met Asp Pro Ala Arg Lys Ala Gly Ala Gln Ala Met Ile Trp Thr Ala  
 1 5 10 15  
 Gly Trp Leu Leu Leu Leu Leu Leu Arg Gly Gly Ala Gln Ala Leu Glu  
 20 25 30  
 Cys Tyr Ser Cys Val Gln Lys Ala Asp Asp Gly Cys Ser Pro Asn Lys  
 35 40 45  
 Met Lys Thr Val Lys Cys Ala Pro Gly Val Asp Val Cys Thr Glu Ala  
 50 55 60  
 Val Gly Ala Val Glu Thr Ile His Gly Gln Phe Ser Leu Ala Val Arg  
 65 70 75 80  
 Gly Cys Gly Ser Gly Leu Pro Gly Lys Asn Asp Arg Gly Leu Asp Leu  
 85 90 95  
 His Gly Leu Leu Ala Phe Ile Gln Leu Gln Gln Cys Ala Gln Asp Arg  
 100 105 110  
 Cys Asn Ala Lys Leu Asn Leu Thr Ser Arg Ala Leu Asp Pro Ala Gly  
 115 120 125  
 Asn Glu Ser Ala Tyr Pro Pro Asn Gly Val Glu Cys Tyr Ser Cys Val  
 130 135 140  
 Gly Leu Ser Arg Glu Ala Cys Gln Gly Thr Ser Pro Pro Val Val Ser  
 145 150 155 160  
 Cys Tyr Asn Ala Ser Asp His Val Tyr Lys Gly Cys Phe Asp Gly Asn  
 165 170 175  
 Val Thr Leu Thr Ala Ala Asn Val Thr Val Ser Leu Pro Val Arg Gly  
 180 185 190  
 Cys Val Gln Asp Glu Phe Cys Thr Arg Asp Gly Val Thr Gly Pro Gly  
 195 200 205  
 Phe Thr Leu Ser Gly Ser Cys Cys Gln Gly Ser Arg Cys Asn Ser Asp  
 210 215 220  
 Leu Arg Asn Lys Thr Tyr Phe Ser Pro Arg Ile Pro Pro Leu Val Arg  
 225 230 235 240  
 Leu Pro Pro Pro Glu Pro Thr Thr Val Ala Ser Thr Thr Ser Val Thr  
 245 250 255  
 Thr Ser Thr Ser Ala Pro Val Arg Pro Thr Ser Thr Thr Lys Pro Met  
 260 265 270  
 Pro Ala Pro Thr Ser Gln Thr Pro Arg Gln Gly Val Glu His Glu Ala  
 275 280 285  
 Ser Arg Asp Glu Glu Pro Arg Leu Thr Gly Gly Ala Ala Gly His Gln  
 290 295 300  
 Asp Arg Ser Asn Ser Gly Gln Tyr Pro Ala Lys Gly Gly Pro Gln Gln  
 305 310 315 320  
 Pro His Asn Lys Gly Cys Val Ala Pro Thr Ala Gly Leu Ala Ala Leu  
 325 330 335  
 Leu Leu Ala Val Ala Ala Gly Val Leu Leu  
 340 345

&lt;210&gt; 217

&lt;211&gt; 61

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 217

Met Lys Arg Phe Leu Phe Leu Leu Leu Thr Ile Ser Leu Leu Val Met  
 1 5 10 15  
 Val Gln Ile Gln Thr Gly Leu Ser Gly Gln Asn Asp Thr Ser Gln Thr  
 20 25 30  
 Ser Ser Pro Ser Ala Ser Ser Asn Ile Ser Gly Gly Ile Phe Leu Phe  
 35 40 45  
 Phe Val Ala Asn Ala Ile Ile His Leu Phe Cys Phe Ser  
 50 55 60

&lt;210&gt; 218

&lt;211&gt; 1255

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 218

Met Glu Leu Ala Ala Leu Cys Arg Trp Gly Leu Leu Leu Ala Leu Leu  
 1 5 10 15  
 Pro Pro Gly Ala Ala Ser Thr Gln Val Cys Thr Gly Thr Asp Met Lys  
 20 25 30  
 Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His  
 35 40 45  
 Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr  
 50 55 60  
 Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val  
 65 70 75 80  
 Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln Val Pro Leu  
 85 90 95  
 Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Asn Tyr  
 100 105 110  
 Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn Thr Thr Pro  
 115 120 125  
 Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln Leu Arg Ser  
 130 135 140  
 Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg Asn Pro Gln  
 145 150 155 160  
 Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe His Lys Asn  
 165 170 175  
 Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser Arg Ala Cys  
 180 185 190  
 His Pro Cys Ser Pro Met Cys Lys Gly Ser Arg Cys Trp Gly Glu Ser  
 195 200 205  
 Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala Gly Gly Cys  
 210 215 220  
 Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His Glu Gln Cys  
 225 230 235 240  
 Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu Ala Cys Leu  
 245 250 255  
 His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro Ala Leu Val  
 260 265 270  
 Thr Tyr Asn Thr Asp Thr Phe Glu Ser Met Pro Asn Pro Glu Gly Arg  
 275 280 285  
 Tyr Thr Phe Gly Ala Ser Cys Val Thr Ala Cys Pro Tyr Asn Tyr Leu  
 290 295 300  
 Ser Thr Asp Val Gly Ser Cys Thr Leu Val Cys Pro Leu His Asn Gln  
 305 310 315 320  
 Glu Val Thr Ala Glu Asp Gly Thr Gln Arg Cys Glu Lys Cys Ser Lys  
 325 330 335  
 Pro Cys Ala Arg Val Cys Tyr Gly Leu Gly Met Glu His Leu Arg Glu  
 340 345 350  
 Val Arg Ala Val Thr Ser Ala Asn Ile Gln Glu Phe Ala Gly Cys Lys

		355				360					365				
Lys	Ile	Phe	Gly	Ser	Leu	Ala	Phe	Leu	Pro	Glu	Ser	Phe	Asp	Gly	Asp
	370					375					380				
Pro	Ala	Ser	Asn	Thr	Ala	Pro	Leu	Gln	Pro	Glu	Gln	Leu	Gln	Val	Phe
385					390					395					400
Glu	Thr	Leu	Glu	Glu	Ile	Thr	Gly	Tyr	Leu	Tyr	Ile	Ser	Ala	Trp	Pro
			405						410					415	
Asp	Ser	Leu	Pro	Asp	Leu	Ser	Val	Phe	Gln	Asn	Leu	Gln	Val	Ile	Arg
			420					425					430		
Gly	Arg	Ile	Leu	His	Asn	Gly	Ala	Tyr	Ser	Leu	Thr	Leu	Gln	Gly	Leu
	435					440						445			
Gly	Ile	Ser	Trp	Leu	Gly	Leu	Arg	Ser	Leu	Arg	Glu	Leu	Gly	Ser	Gly
	450					455					460				
Leu	Ala	Leu	Ile	His	His	Asn	Thr	His	Leu	Cys	Phe	Val	His	Thr	Val
465					470					475					480
Pro	Trp	Asp	Gln	Leu	Phe	Arg	Asn	Pro	His	Gln	Ala	Leu	Leu	His	Thr
			485					490						495	
Ala	Asn	Arg	Pro	Glu	Asp	Glu	Cys	Val	Gly	Glu	Gly	Leu	Ala	Cys	His
			500					505					510		
Gln	Leu	Cys	Ala	Arg	Gly	His	Cys	Trp	Gly	Pro	Gly	Pro	Thr	Gln	Cys
	515					520						525			
Val	Asn	Cys	Ser	Gln	Phe	Leu	Arg	Gly	Gln	Glu	Cys	Val	Glu	Glu	Cys
	530					535					540				
Arg	Val	Leu	Gln	Gly	Leu	Pro	Arg	Glu	Tyr	Val	Asn	Ala	Arg	His	Cys
545					550					555					560
Leu	Pro	Cys	His	Pro	Glu	Cys	Gln	Pro	Gln	Asn	Gly	Ser	Val	Thr	Cys
			565					570						575	
Phe	Gly	Pro	Glu	Ala	Asp	Gln	Cys	Val	Ala	Cys	Ala	His	Tyr	Lys	Asp
			580					585					590		
Pro	Pro	Phe	Cys	Val	Ala	Arg	Cys	Pro	Ser	Gly	Val	Lys	Pro	Asp	Leu
		595					600					605			
Ser	Tyr	Met	Pro	Ile	Trp	Lys	Phe	Pro	Asp	Glu	Glu	Gly	Ala	Cys	Gln
	610					615						620			
Pro	Cys	Pro	Ile	Asn	Cys	Thr	His	Ser	Cys	Val	Asp	Leu	Asp	Asp	Lys
625					630					635					640
Gly	Cys	Pro	Ala	Glu	Gln	Arg	Ala	Ser	Pro	Leu	Thr	Ser	Ile	Ile	Ser
			645					650						655	
Ala	Val	Val	Gly	Ile	Leu	Leu	Val	Val	Val	Leu	Gly	Val	Val	Phe	Gly
			660					665						670	
Ile	Leu	Ile	Lys	Arg	Arg	Gln	Gln	Lys	Ile	Arg	Lys	Tyr	Thr	Met	Arg
	675					680						685			
Arg	Leu	Leu	Gln	Glu	Thr	Glu	Leu	Val	Glu	Pro	Leu	Thr	Pro	Ser	Gly
	690					695					700				
Ala	Met	Pro	Asn	Gln	Ala	Gln	Met	Arg	Ile	Leu	Lys	Glu	Thr	Glu	Leu
705					710					715					720
Arg	Lys	Val	Lys	Val	Leu	Gly	Ser	Gly	Ala	Phe	Gly	Thr	Val	Tyr	Lys
			725					730						735	
Gly	Ile	Trp	Ile	Pro	Asp	Gly	Glu	Asn	Val	Lys	Ile	Pro	Val	Ala	Ile
			740					745					750		
Lys	Val	Leu	Arg	Glu	Asn	Thr	Ser	Pro	Lys	Ala	Asn	Lys	Glu	Ile	Leu
	755					760						765			
Asp	Glu	Ala	Tyr	Val	Met	Ala	Gly	Val	Gly	Ser	Pro	Tyr	Val	Ser	Arg
	770				775						780				
Leu	Leu	Gly	Ile	Cys	Leu	Thr	Ser	Thr	Val	Gln	Leu	Val	Thr	Gln	Leu
785					790					795					800
Met	Pro	Tyr	Gly	Cys	Leu	Leu	Asp	His	Val	Arg	Glu	Asn	Arg	Gly	Arg
			805					810						815	
Leu	Gly	Ser	Gln	Asp	Leu	Leu	Asn	Trp	Cys	Met	Gln	Ile	Ala	Lys	Gly
			820					825					830		
Met	Ser	Tyr	Leu	Glu	Asp	Val	Arg	Leu	Val	His	Arg	Asp	Leu	Ala	Ala
	835					840						845			
Arg	Asn	Val	Leu	Val	Lys	Ser	Pro	Asn	His	Val	Lys	Ile	Thr	Asp	Phe
	850					855						860			

Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr Glu Tyr His Ala Asp  
 865 870 875 880  
 Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Glu Ser Ile Leu Arg  
 885 890 895  
 Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr Val  
 900 905 910  
 Trp Glu Leu Met Thr Phe Gly Ala Lys Pro Tyr Asp Gly Ile Pro Ala  
 915 920 925  
 Arg Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu Pro Gln Pro  
 930 935 940  
 Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met Val Lys Cys Trp Met  
 945 950 955 960  
 Ile Asp Ser Glu Cys Arg Pro Arg Phe Arg Glu Leu Val Ser Glu Phe  
 965 970 975  
 Ser Arg Met Ala Arg Asp Pro Gln Arg Phe Val Val Ile Gln Asn Glu  
 980 985 990  
 Asp Leu Gly Pro Ala Ser Pro Leu Asp Ser Thr Phe Tyr Arg Ser Leu  
 995 1000 1005  
 Leu Glu Asp Asp Asp Met Gly Asp Leu Val Asp Ala Glu Glu Tyr Leu  
 1010 1015 1020  
 Val Pro Gln Gln Gly Phe Phe Cys Pro Asp Pro Ala Pro Gly Ala Gly  
 1025 1030 1035 1040  
 Gly Met Val His His Arg His Arg Ser Ser Ser Thr Arg Ser Gly Gly  
 1045 1050 1055  
 Gly Asp Leu Thr Leu Gly Leu Glu Pro Ser Glu Glu Glu Ala Pro Arg  
 1060 1065 1070  
 Ser Pro Leu Ala Pro Ser Glu Gly Ala Gly Ser Asp Val Phe Asp Gly  
 1075 1080 1085  
 Asp Leu Gly Met Gly Ala Ala Lys Gly Leu Gln Ser Leu Pro Thr His  
 1090 1095 1100  
 Asp Pro Ser Pro Leu Gln Arg Tyr Ser Glu Asp Pro Thr Val Pro Leu  
 1105 1110 1115 1120  
 Pro Ser Glu Thr Asp Gly Tyr Val Ala Pro Leu Thr Cys Ser Pro Gln  
 1125 1130 1135  
 Pro Glu Tyr Val Asn Gln Pro Asp Val Arg Pro Gln Pro Pro Ser Pro  
 1140 1145 1150  
 Arg Glu Gly Pro Leu Pro Ala Ala Arg Pro Ala Gly Ala Thr Leu Glu  
 1155 1160 1165  
 Arg Pro Lys Thr Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val  
 1170 1175 1180  
 Phe Ala Phe Gly Gly Ala Val Glu Asn Pro Glu Tyr Leu Thr Pro Gln  
 1185 1190 1195 1200  
 Gly Gly Ala Ala Pro Gln Pro His Pro Pro Pro Ala Phe Ser Pro Ala  
 1205 1210 1215  
 Phe Asp Asn Leu Tyr Tyr Trp Asp Gln Asp Pro Pro Glu Arg Gly Ala  
 1220 1225 1230  
 Pro Pro Ser Thr Phe Lys Gly Thr Pro Thr Ala Glu Asn Pro Glu Tyr  
 1235 1240 1245  
 Leu Gly Leu Asp Val Pro Val  
 1250 1255

&lt;210&gt; 219

&lt;211&gt; 297

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 219

Met Thr Thr Pro Arg Asn Ser Val Asn Gly Thr Phe Pro Ala Glu Pro  
 1 5 10 15  
 Met Lys Gly Pro Ile Ala Met Gln Ser Gly Pro Lys Pro Leu Phe Arg  
 20 25 30  
 Arg Met Ser Ser Leu Val Gly Pro Thr Gln Ser Phe Phe Met Arg Glu  
 35 40 45

Ser Lys Thr Leu Gly Ala Val Gln Ile Met Asn Gly Leu Phe His Ile  
 50 55 60  
 Ala Leu Gly Gly Leu Leu Met Ile Pro Ala Gly Ile Tyr Ala Pro Ile  
 65 70 75 80  
 Cys Val Thr Val Trp Tyr Pro Leu Trp Gly Gly Ile Met Tyr Ile Ile  
 85 90 95  
 Ser Gly Ser Leu Leu Ala Ala Thr Glu Lys Asn Ser Arg Lys Cys Leu  
 100 105 110  
 Val Lys Gly Lys Met Ile Met Asn Ser Leu Ser Leu Phe Ala Ala Ile  
 115 120 125  
 Ser Gly Met Ile Leu Ser Ile Met Asp Ile Leu Asn Ile Lys Ile Ser  
 130 135 140  
 His Phe Leu Lys Met Glu Ser Leu Asn Phe Ile Arg Ala His Thr Pro  
 145 150 155 160  
 Tyr Ile Asn Ile Tyr Asn Cys Glu Pro Ala Asn Pro Ser Glu Lys Asn  
 165 170 175  
 Ser Pro Ser Thr Gln Tyr Cys Tyr Ser Ile Gln Ser Leu Phe Leu Gly  
 180 185 190  
 Ile Leu Ser Val Met Leu Ile Phe Ala Phe Phe Gln Glu Leu Val Ile  
 195 200 205  
 Ala Gly Ile Val Glu Asn Glu Trp Lys Arg Thr Cys Ser Arg Pro Lys  
 210 215 220  
 Ser Asn Ile Val Leu Leu Ser Ala Glu Glu Lys Lys Glu Gln Thr Ile  
 225 230 235 240  
 Glu Ile Lys Glu Glu Val Val Gly Leu Thr Glu Thr Ser Ser Gln Pro  
 245 250 255  
 Lys Asn Glu Glu Asp Ile Glu Ile Ile Pro Ile Gln Glu Glu Glu Glu  
 260 265 270  
 Glu Glu Thr Glu Thr Asn Phe Pro Glu Pro Pro Gln Asp Gln Glu Ser  
 275 280 285  
 Ser Pro Ile Glu Asn Asp Ser Ser Pro  
 290 295

&lt;210&gt; 220

&lt;211&gt; 595

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 220

Met Arg Val Leu Leu Ala Ala Leu Gly Leu Leu Phe Leu Gly Ala Leu  
 1 5 10 15  
 Arg Ala Phe Pro Gln Asp Arg Pro Phe Glu Asp Thr Cys His Gly Asn  
 20 25 30  
 Pro Ser His Tyr Tyr Asp Lys Ala Val Arg Arg Cys Cys Tyr Arg Cys  
 35 40 45  
 Pro Met Gly Leu Phe Pro Thr Gln Gln Cys Pro Gln Arg Pro Thr Asp  
 50 55 60  
 Cys Arg Lys Gln Cys Glu Pro Asp Tyr Tyr Leu Asp Glu Ala Asp Arg  
 65 70 75 80  
 Cys Thr Ala Cys Val Thr Cys Ser Arg Asp Asp Leu Val Glu Lys Thr  
 85 90 95  
 Pro Cys Ala Trp Asn Ser Ser Arg Val Cys Glu Cys Arg Pro Gly Met  
 100 105 110  
 Phe Cys Ser Thr Ser Ala Val Asn Ser Cys Ala Arg Cys Phe Phe His  
 115 120 125  
 Ser Val Cys Pro Ala Gly Met Ile Val Lys Phe Pro Gly Thr Ala Gln  
 130 135 140  
 Lys Asn Thr Val Cys Glu Pro Ala Ser Pro Gly Val Ser Pro Ala Cys  
 145 150 155 160  
 Ala Ser Pro Glu Asn Cys Lys Glu Pro Ser Ser Gly Thr Ile Pro Gln  
 165 170 175  
 Ala Lys Pro Thr Pro Val Ser Pro Ala Thr Ser Ser Ala Ser Thr Met  
 180 185 190



Pro Val Arg Gly Gly Thr Arg Leu Ala Gln Glu Ala Ala Ser Lys Leu  
 195 200 205  
 Thr Arg Ala Pro Asp Ser Pro Ser Ser Val Gly Arg Pro Ser Ser Asp  
 210 215 220  
 Pro Gly Leu Ser Pro Thr Gln Pro Cys Pro Glu Gly Ser Gly Asp Cys  
 225 230 235 240  
 Arg Lys Gln Cys Glu Pro Asp Tyr Tyr Leu Asp Glu Ala Gly Arg Cys  
 245 250 255  
 Thr Ala Cys Val Ser Cys Ser Arg Asp Asp Leu Val Glu Lys Thr Pro  
 260 265 270  
 Cys Ala Trp Asn Ser Ser Arg Thr Cys Glu Cys Arg Pro Gly Met Ile  
 275 280 285  
 Cys Ala Thr Ser Ala Thr Asn Ser Arg Ala Arg Cys Val Pro Tyr Pro  
 290 295 300  
 Ile Cys Ala Ala Glu Thr Val Thr Lys Pro Gln Asp Met Ala Glu Lys  
 305 310 315 320  
 Asp Thr Thr Phe Glu Ala Pro Pro Leu Gly Thr Gln Pro Asp Cys Asn  
 325 330 335  
 Pro Thr Pro Glu Asn Gly Glu Ala Pro Ala Ser Thr Ser Pro Thr Gln  
 340 345 350  
 Ser Leu Leu Val Asp Ser Gln Ala Ser Lys Thr Leu Pro Ile Pro Thr  
 355 360 365  
 Ser Ala Pro Val Ala Leu Ser Ser Thr Gly Lys Pro Val Leu Asp Ala  
 370 375 380  
 Gly Pro Val Leu Phe Trp Val Ile Leu Val Leu Val Val Val Gly  
 385 390 395 400  
 Ser Ser Ala Phe Leu Leu Cys His Arg Arg Ala Cys Arg Lys Arg Ile  
 405 410 415  
 Arg Gln Lys Leu His Leu Cys Tyr Pro Val Gln Thr Ser Gln Pro Lys  
 420 425 430  
 Leu Glu Leu Val Asp Ser Arg Pro Arg Arg Ser Ser Thr Gln Leu Arg  
 435 440 445  
 Ser Gly Ala Ser Val Thr Glu Pro Val Ala Glu Glu Arg Gly Leu Met  
 450 455 460  
 Ser Gln Pro Leu Met Glu Thr Cys His Ser Val Gly Ala Ala Tyr Leu  
 465 470 475 480  
 Glu Ser Leu Pro Leu Gln Asp Ala Ser Pro Ala Gly Gly Pro Ser Ser  
 485 490 495  
 Pro Arg Asp Leu Pro Glu Pro Arg Val Ser Thr Glu His Thr Asn Asn  
 500 505 510  
 Lys Ile Glu Lys Ile Tyr Ile Met Lys Ala Asp Thr Val Ile Val Gly  
 515 520 525  
 Thr Val Lys Ala Glu Leu Pro Glu Gly Arg Gly Leu Ala Gly Pro Ala  
 530 535 540  
 Glu Pro Glu Leu Glu Glu Glu Leu Glu Ala Asp His Thr Pro His Tyr  
 545 550 555 560  
 Pro Glu Gln Glu Thr Glu Pro Pro Leu Gly Ser Cys Ser Asp Val Met  
 565 570 575  
 Leu Ser Val Glu Glu Glu Gly Lys Glu Asp Pro Leu Pro Thr Ala Ala  
 580 585 590  
 Ser Gly Lys  
 595

&lt;210&gt; 221

&lt;211&gt; 847

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 221

Met His Leu Leu Gly Pro Trp Leu Leu Leu Leu Val Leu Glu Tyr Leu  
 1 5 10 15  
 Ala Phe Ser Asp Ser Ser Lys Trp Val Phe Glu His Pro Glu Thr Leu  
 20 25 30

Tyr Ala Trp Glu Gly Ala Cys Val Trp Ile Pro Cys Thr Tyr Arg Ala  
 35 40 45  
 Leu Asp Gly Asp Leu Glu Ser Phe Ile Leu Phe His Asn Pro Glu Tyr  
 50 55 60  
 Asn Lys Asn Thr Ser Lys Phe Asp Gly Thr Arg Leu Tyr Glu Ser Thr  
 65 70 75 80  
 Lys Asp Gly Lys Val Pro Ser Glu Gln Lys Arg Val Gln Phe Leu Gly  
 85 90 95  
 Asp Lys Asn Lys Asn Cys Thr Leu Ser Ile His Pro Val His Leu Asn  
 100 105 110  
 Asp Ser Gly Gln Leu Gly Leu Arg Met Glu Ser Lys Thr Glu Lys Trp  
 115 120 125  
 Met Glu Arg Ile His Leu Asn Val Ser Glu Arg Pro Phe Pro Pro His  
 130 135 140  
 Ile Gln Leu Pro Pro Glu Ile Gln Glu Ser Gln Glu Val Thr Leu Thr  
 145 150 155 160  
 Cys Leu Leu Asn Phe Ser Cys Tyr Gly Tyr Pro Ile Gln Leu Gln Trp  
 165 170 175  
 Leu Leu Glu Gly Val Pro Met Arg Gln Ala Ala Val Thr Ser Thr Ser  
 180 185 190  
 Leu Thr Ile Lys Ser Val Phe Thr Arg Ser Glu Leu Lys Phe Ser Pro  
 195 200 205  
 Gln Trp Ser His His Gly Lys Ile Val Thr Cys Gln Leu Gln Asp Ala  
 210 215 220  
 Asp Gly Lys Phe Leu Ser Asn Asp Thr Val Gln Leu Asn Val Lys His  
 225 230 235 240  
 Thr Pro Lys Leu Glu Ile Lys Val Thr Pro Ser Asp Ala Ile Val Arg  
 245 250 255  
 Glu Gly Asp Ser Val Thr Met Thr Cys Glu Val Ser Ser Ser Asn Pro  
 260 265 270  
 Glu Tyr Thr Thr Val Ser Trp Leu Lys Asp Gly Thr Ser Leu Lys Lys  
 275 280 285  
 Gln Asn Thr Phe Thr Leu Asn Leu Arg Glu Val Thr Lys Asp Gln Ser  
 290 295 300  
 Gly Lys Tyr Cys Cys Gln Val Ser Asn Asp Val Gly Pro Gly Arg Ser  
 305 310 315 320  
 Glu Glu Val Phe Leu Gln Val Gln Tyr Ala Pro Glu Pro Ser Thr Val  
 325 330 335  
 Gln Ile Leu His Ser Pro Ala Val Glu Gly Ser Gln Val Glu Phe Leu  
 340 345 350  
 Cys Met Ser Leu Ala Asn Pro Leu Pro Thr Asn Tyr Thr Trp Tyr His  
 355 360 365  
 Asn Gly Lys Glu Met Gln Gly Arg Thr Glu Glu Lys Val His Ile Pro  
 370 375 380  
 Lys Ile Leu Pro Trp His Ala Gly Thr Tyr Ser Cys Val Ala Glu Asn  
 385 390 395 400  
 Ile Leu Gly Thr Gly Gln Arg Gly Pro Gly Ala Glu Leu Asp Val Gln  
 405 410 415  
 Tyr Pro Pro Lys Lys Val Thr Thr Val Ile Gln Asn Pro Met Pro Ile  
 420 425 430  
 Arg Glu Gly Asp Thr Val Thr Leu Ser Cys Asn Tyr Asn Ser Ser Asn  
 435 440 445  
 Pro Ser Val Thr Arg Tyr Glu Trp Lys Pro His Gly Ala Trp Glu Glu  
 450 455 460  
 Pro Ser Leu Gly Val Leu Lys Ile Gln Asn Val Gly Trp Asp Asn Thr  
 465 470 475 480  
 Thr Ile Ala Cys Ala Ala Cys Asn Ser Trp Cys Ser Trp Ala Ser Pro  
 485 490 495  
 Val Ala Leu Asn Val Gln Tyr Ala Pro Arg Asp Val Arg Val Arg Lys  
 500 505 510  
 Ile Lys Pro Leu Ser Glu Ile His Ser Gly Asn Ser Val Ser Leu Gln  
 515 520 525  
 Cys Asp Phe Ser Ser Ser His Pro Lys Glu Val Gln Phe Phe Trp Glu

530						535					540				
Lys	Asn	Gly	Arg	Leu	Leu	Gly	Lys	Glu	Ser	Gln	Leu	Asn	Phe	Asp	Ser
545					550					555					560
Ile	Ser	Pro	Glu	Asp	Ala	Gly	Ser	Tyr	Ser	Cys	Trp	Val	Asn	Asn	Ser
				565					570					575	
Ile	Gly	Gln	Thr	Ala	Ser	Lys	Ala	Trp	Thr	Leu	Glu	Val	Leu	Tyr	Ala
			580					585					590		
Pro	Arg	Arg	Leu	Arg	Val	Ser	Met	Ser	Pro	Gly	Asp	Gln	Val	Met	Glu
		595					600					605			
Gly	Lys	Ser	Ala	Thr	Leu	Thr	Cys	Glu	Ser	Asp	Ala	Asn	Pro	Pro	Val
	610					615					620				
Ser	His	Tyr	Thr	Trp	Phe	Asp	Trp	Asn	Asn	Gln	Ser	Leu	Pro	Tyr	His
625					630					635					640
Ser	Gln	Lys	Leu	Arg	Leu	Glu	Pro	Val	Lys	Val	Gln	His	Ser	Gly	Ala
				645					650					655	
Tyr	Trp	Cys	Gln	Gly	Thr	Asn	Ser	Val	Gly	Lys	Gly	Arg	Ser	Pro	Leu
			660					665					670		
Ser	Thr	Leu	Thr	Val	Tyr	Tyr	Ser	Pro	Glu	Thr	Ile	Gly	Arg	Arg	Val
		675					680					685			
Ala	Val	Gly	Leu	Gly	Ser	Cys	Leu	Ala	Ile	Leu	Ile	Leu	Ala	Ile	Cys
	690					695					700				
Gly	Leu	Lys	Leu	Gln	Arg	Arg	Trp	Lys	Arg	Thr	Gln	Ser	Gln	Gln	Gly
705					710					715					720
Leu	Gln	Glu	Asn	Ser	Ser	Gly	Gln	Ser	Phe	Phe	Val	Arg	Asn	Lys	Lys
				725					730					735	
Val	Arg	Arg	Ala	Pro	Leu	Ser	Glu	Gly	Pro	His	Ser	Leu	Gly	Cys	Tyr
			740					745					750		
Asn	Pro	Met	Met	Glu	Asp	Gly	Ile	Ser	Tyr	Thr	Thr	Leu	Arg	Phe	Pro
		755					760					765			
Glu	Met	Asn	Ile	Pro	Arg	Thr	Gly	Asp	Ala	Glu	Ser	Ser	Glu	Met	Gln
	770					775						780			
Arg	Pro	Pro	Pro	Asp	Cys	Asp	Asp	Thr	Val	Thr	Tyr	Ser	Ala	Leu	His
785					790					795					800
Lys	Arg	Gln	Val	Gly	Asp	Tyr	Glu	Asn	Val	Ile	Pro	Asp	Phe	Pro	Glu
				805					810					815	
Asp	Glu	Gly	Ile	His	Tyr	Ser	Glu	Leu	Ile	Gln	Phe	Gly	Val	Gly	Glu
			820					825					830		
Arg	Pro	Gln	Ala	Gln	Glu	Asn	Val	Asp	Tyr	Val	Ile	Leu	Lys	His	
		835					840					845			

&lt;210&gt; 222

&lt;211&gt; 364

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 222

Met	Pro	Leu	Leu	Leu	Leu	Pro	Leu	Leu	Trp	Ala	Gly	Ala	Leu	Ala	
1			5					10					15		
Met	Asp	Pro	Asn	Phe	Trp	Leu	Gln	Val	Gln	Glu	Ser	Val	Thr	Val	Gln
			20					25					30		
Glu	Gly	Leu	Cys	Val	Leu	Val	Pro	Cys	Thr	Phe	Phe	His	Pro	Ile	Pro
		35					40					45			
Tyr	Tyr	Asp	Lys	Asn	Ser	Pro	Val	His	Gly	Tyr	Trp	Phe	Arg	Glu	Gly
	50					55					60				
Ala	Ile	Ile	Ser	Arg	Asp	Ser	Pro	Val	Ala	Thr	Asn	Lys	Leu	Asp	Gln
65					70					75					80
Glu	Val	Gln	Glu	Glu	Thr	Gln	Gly	Arg	Phe	Arg	Leu	Leu	Gly	Asp	Pro
				85					90					95	
Ser	Arg	Asn	Asn	Cys	Ser	Leu	Ser	Ile	Val	Asp	Ala	Arg	Arg	Arg	Asp
			100					105					110		
Asn	Gly	Ser	Tyr	Phe	Phe	Arg	Met	Glu	Arg	Gly	Ser	Thr	Lys	Tyr	Ser
		115					120						125		
Tyr	Lys	Ser	Pro	Gln	Leu	Ser	Val	His	Val	Thr	Asp	Leu	Thr	His	Arg

130	Pro	Lys	Ile	Leu	Ile	Pro	Gly	Thr	Leu	Glu	Pro	Gly	His	Ser	Lys	Asn
145	Leu	Thr	Cys	Ser	Val	Ser	Trp	Ala	Cys	Glu	Gln	Gly	Thr	Pro	Pro	Ile
				165	Ala	Ala	Pro	Thr	Ser	Leu	Gly	Pro	Arg	Thr	Thr	
				180				185						190		
				195				200						205		
				210				215						220		
				225				230						235		
				245				250						255		
				260				265						270		
				275				280						285		
				290				295						300		
				305				310						315		
				325				330						335		
				340				345						350		
				355				360								

&lt;210&gt; 223

&lt;211&gt; 572

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 223

1	Met	Glu	Cys	Leu	Tyr	Tyr	Phe	Leu	Gly	Phe	Leu	Leu	Leu	Ala	Ala	Arg
				5						10					15	
				20						25					30	
				35						40					45	
				50						55					60	
				65						70					75	
				85						90					95	
				100						105					110	
				115						120					125	
				130						135					140	
				145						150					155	
				165						170					175	
				180						185					190	
				195						200					205	
				355						360						

210		215		220
Gln Val Lys Asp Val Tyr Val Val Thr Asp Gln Ile Pro Val Phe Val				
225		230		240
Thr Met Phe Gln Lys Asn Asp Arg Asn Ser Ser Asp Glu Thr Phe Leu				
	245		250	255
Lys Asp Leu Pro Ile Met Phe Asp Val Leu Ile His Asp Pro Ser His				
	260		265	270
Phe Leu Asn Tyr Ser Thr Ile Asn Tyr Lys Trp Ser Phe Gly Asp Asn				
	275		280	285
Thr Gly Leu Phe Val Ser Thr Asn His Thr Val Asn His Thr Tyr Val				
	290		295	300
Leu Asn Gly Thr Phe Ser Leu Asn Leu Thr Val Lys Ala Ala Ala Pro				
305		310		320
Gly Pro Cys Pro Pro Pro Pro Pro Pro Arg Pro Ser Lys Pro Thr				
	325		330	335
Pro Ser Leu Ala Thr Thr Leu Lys Ser Tyr Asp Ser Asn Thr Pro Gly				
	340		345	350
Pro Ala Gly Asp Asn Pro Leu Glu Leu Ser Arg Ile Pro Asp Glu Asn				
	355		360	365
Cys Gln Ile Asn Arg Tyr Gly His Phe Gln Ala Thr Ile Thr Ile Val				
	370		375	380
Glu Gly Ile Leu Glu Val Asn Ile Ile Gln Met Thr Asp Val Leu Met				
385		390		400
Pro Val Pro Trp Pro Glu Ser Ser Leu Ile Asp Phe Val Val Thr Cys				
	405		410	415
Gln Gly Ser Ile Pro Thr Glu Val Cys Thr Ile Ile Ser Asp Pro Thr				
	420		425	430
Cys Glu Ile Thr Gln Asn Thr Val Cys Ser Pro Val Asp Val Asp Glu				
	435		440	445
Met Cys Leu Leu Thr Val Arg Arg Thr Phe Asn Gly Ser Gly Thr Tyr				
	450		455	460
Cys Val Asn Leu Thr Leu Gly Asp Asp Thr Ser Leu Ala Leu Thr Ser				
465		470		480
Thr Leu Ile Ser Val Pro Asp Arg Asp Pro Ala Ser Pro Leu Arg Met				
	485		490	495
Ala Asn Ser Ala Leu Ile Ser Val Gly Cys Leu Ala Ile Phe Val Thr				
	500		505	510
Val Ile Ser Leu Leu Val Tyr Lys Lys His Lys Glu Tyr Asn Pro Ile				
	515		520	525
Glu Asn Ser Pro Gly Asn Val Val Arg Ser Lys Gly Leu Ser Val Phe				
	530		535	540
Leu Asn Arg Ala Lys Ala Val Phe Phe Pro Gly Asn Gln Glu Lys Asp				
545		550		560
Pro Leu Leu Lys Asn Gln Glu Phe Lys Gly Val Ser				
	565		570	

&lt;210&gt; 224

&lt;211&gt; 848

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 224

Met Leu Gln Thr Lys Asp Leu Ile Trp Thr Leu Phe Phe Leu Gly Thr				
1	5		10	15
Ala Val Ser Leu Gln Val Asp Ile Val Pro Ser Gln Gly Glu Ile Ser				
	20		25	30
Val Gly Glu Ser Lys Phe Phe Leu Cys Gln Val Ala Gly Asp Ala Lys				
	35		40	45
Asp Lys Asp Ile Ser Trp Phe Ser Pro Asn Gly Glu Lys Leu Thr Pro				
	50		55	60
Asn Gln Gln Arg Ile Ser Val Val Trp Asn Asp Asp Ser Ser Ser Thr				
65	70		75	80
Leu Thr Ile Tyr Asn Ala Asn Ile Asp Asp Ala Gly Ile Tyr Lys Cys				



Phe Lys Thr Gln Pro Val Gln Gly Glu Pro Ser Ala Pro Lys Leu Glu  
                   595                                  600                                  605  
 Gly Gln Met Gly Glu Asp Gly Asn Ser Ile Lys Val Asn Leu Ile Lys  
           610                                  615                                  620  
 Gln Asp Asp Gly Gly Ser Pro Ile Arg His Tyr Leu Val Arg Tyr Arg  
 625                                  630                                  635                                  640  
 Ala Leu Ser Ser Glu Trp Lys Pro Glu Ile Arg Leu Pro Ser Gly Ser  
                   645                                  650                                  655  
 Asp His Val Met Leu Lys Ser Leu Asp Trp Asn Ala Glu Tyr Glu Val  
                   660                                  665                                  670  
 Tyr Val Val Ala Glu Asn Gln Gln Gly Lys Ser Lys Ala Ala His Phe  
                   675                                  680                                  685  
 Val Phe Arg Thr Ser Ala Gln Pro Thr Ala Ile Pro Ala Asn Gly Ser  
           690                                  695                                  700  
 Pro Thr Ser Gly Leu Ser Thr Gly Ala Ile Val Gly Ile Leu Ile Val  
 705                                  710                                  715                                  720  
 Ile Phe Val Leu Leu Leu Val Val Val Asp Ile Thr Cys Tyr Phe Leu  
                   725                                  730                                  735  
 Asn Lys Cys Gly Leu Phe Met Cys Ile Ala Val Asn Leu Cys Gly Lys  
                   740                                  745                                  750  
 Ala Gly Pro Gly Ala Lys Gly Lys Asp Met Glu Glu Gly Lys Ala Ala  
           755                                  760                                  765  
 Phe Ser Lys Asp Glu Ser Lys Glu Pro Ile Val Glu Val Arg Thr Glu  
           770                                  775                                  780  
 Glu Glu Arg Thr Pro Asn His Asp Gly Gly Lys His Thr Glu Pro Asn  
 785                                  790                                  795                                  800  
 Glu Thr Thr Pro Leu Thr Glu Pro Glu Lys Gly Pro Val Glu Ala Lys  
                   805                                  810                                  815  
 Pro Glu Cys Gln Glu Thr Glu Thr Lys Pro Ala Pro Ala Glu Val Lys  
                   820                                  825                                  830  
 Thr Val Pro Asn Asp Ala Thr Gln Thr Lys Glu Asn Glu Ser Lys Ala  
                   835                                  840                                  845

&lt;210&gt; 225

&lt;211&gt; 193

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 225

Met Pro Glu Glu Gly Ser Gly Cys Ser Val Arg Arg Arg Pro Tyr Gly  
 1                  5                                  10                                  15  
 Cys Val Leu Arg Ala Ala Leu Val Pro Leu Val Ala Gly Leu Val Ile  
                   20                                  25                                  30  
 Cys Leu Val Val Cys Ile Gln Arg Phe Ala Gln Ala Gln Gln Leu  
           35                                  40                                  45  
 Pro Leu Glu Ser Leu Gly Trp Asp Val Ala Glu Leu Gln Leu Asn His  
           50                                  55                                  60  
 Thr Gly Pro Gln Gln Asp Pro Arg Leu Tyr Trp Gln Gly Gly Pro Ala  
 65                                  70                                  75                                  80  
 Leu Gly Arg Ser Phe Leu His Gly Pro Glu Leu Asp Lys Gly Gln Leu  
                   85                                  90                                  95  
 Arg Ile His Arg Asp Gly Ile Tyr Met Val His Ile Gln Val Thr Leu  
           100                                  105                                  110  
 Ala Ile Cys Ser Ser Thr Thr Ala Ser Arg His His Pro Thr Thr Leu  
           115                                  120                                  125  
 Ala Val Gly Ile Cys Ser Pro Ala Ser Arg Ser Ile Ser Leu Leu Arg  
           130                                  135                                  140  
 Leu Ser Phe His Gln Gly Cys Thr Ile Ala Ser Gln Arg Leu Thr Pro  
 145                                  150                                  155                                  160  
 Leu Ala Arg Gly Asp Thr Leu Cys Thr Asn Leu Thr Gly Thr Leu Leu  
                   165                                  170                                  175  
  
 Pro Ser Arg Asn Thr Asp Glu Thr Phe Phe Gly Val Gln Trp Val Arg  
                   180                                  185                                  190

Pro

&lt;210&gt; 226

&lt;211&gt; 232

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 226

Met His Arg Arg Arg Ser Arg Ser Cys Arg Glu Asp Gln Lys Pro Val  
 1 5 10 15  
 Met Asp Asp Gln Arg Asp Leu Ile Ser Asn Asn Glu Gln Leu Pro Met  
 20 25 30  
 Leu Gly Arg Arg Pro Gly Ala Pro Glu Ser Lys Cys Ser Arg Gly Ala  
 35 40 45  
 Leu Tyr Thr Gly Phe Ser Ile Leu Val Thr Leu Leu Leu Ala Gly Gln  
 50 55 60  
 Ala Thr Thr Ala Tyr Phe Leu Tyr Gln Gln Gln Gly Arg Leu Asp Lys  
 65 70 75 80  
 Leu Thr Val Thr Ser Gln Asn Leu Gln Leu Glu Asn Leu Arg Met Lys  
 85 90 95  
 Leu Pro Lys Pro Pro Lys Pro Val Ser Lys Met Arg Met Ala Thr Pro  
 100 105 110  
 Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro Gln Gly Pro Met  
 115 120 125  
 Gln Asn Ala Thr Lys Tyr Gly Asn Met Thr Glu Asp His Val Met His  
 130 135 140  
 Leu Leu Gln Asn Ala Asp Pro Leu Lys Val Tyr Pro Pro Leu Lys Gly  
 145 150 155 160  
 Ser Phe Pro Glu Asn Leu Arg His Leu Lys Asn Thr Met Glu Thr Ile  
 165 170 175  
 Asp Trp Lys Val Phe Glu Ser Trp Met His His Trp Leu Leu Phe Glu  
 180 185 190  
 Met Ser Arg His Ser Leu Glu Gln Lys Pro Thr Asp Ala Pro Pro Lys  
 195 200 205  
 Glu Ser Leu Glu Leu Glu Asp Pro Ser Ser Gly Leu Gly Val Thr Lys  
 210 215 220  
 Gln Asp Leu Gly Pro Val Pro Met  
 225 230

<210> 227

<211> 557

<212> PRT

<213> Homo sapiens

<400> 227

Met Pro Pro Pro Arg Leu Leu Phe Phe Leu Leu Phe Leu Thr Pro Met  
 1 5 10 15  
 Glu Val Arg Pro Glu Glu Pro Leu Val Val Lys Val Glu Glu Gly Asp  
 20 25 30  
 Asn Ala Val Leu Gln Cys Leu Lys Gly Thr Ser Asp Gly Pro Thr Gln  
 35 40 45  
 Gln Leu Thr Trp Ser Arg Glu Ser Pro Leu Lys Pro Phe Leu Lys Leu  
 50 55 60  
 Ser Leu Gly Leu Pro Gly Leu Gly Ile His Met Arg Pro Leu Ala Ile  
 65 70 75 80  
 Trp Leu Phe Ile Phe Asn Val Ser Gln Gln Met Gly Gly Phe Tyr Leu  
 85 90 95  
 Cys Gln Pro Gly Pro Pro Ser Glu Lys Ala Trp Gln Pro Gly Trp Thr  
 100 105 110  
 Val Asn Val Glu Gly Ser Gly Glu Leu Phe Arg Trp Asn Val Ser Asp





1				5					10					15		
Val	Leu	Thr	Val	Val	Thr	Gly	Ser	Gly	His	Ala	Ser	Ser	Thr	Pro	Gly	
			20					25					30			
Gly	Glu	Lys	Glu	Thr	Ser	Ala	Thr	Gln	Arg	Ser	Ser	Val	Pro	Ser	Ser	
		35					40					45				
Thr	Glu	Lys	Asn	Ala	Leu	Ser	Thr	Gly	Val	Ser	Phe	Phe	Phe	Leu	Ser	
	50					55					60					
Phe	His	Ile	Ser	Asn	Leu	Gln	Phe	Asn	Ser	Ser	Leu	Glu	Asp	Pro	Ser	
65				70						75					80	
Thr	Asp	Tyr	Tyr	Gln	Glu	Leu	Gln	Arg	Asp	Ile	Ser	Glu	Met	Phe	Leu	
			85						90					95		
Gln	Ile	Tyr	Lys	Gln	Gly	Gly	Phe	Leu	Gly	Leu	Ser	Asn	Ile	Lys	Phe	
			100					105					110			
Arg	Pro	Gly	Ser	Val	Val	Val	Gln	Leu	Thr	Leu	Ala	Phe	Arg	Glu	Gly	
		115					120					125				
Thr	Ile	Asn	Val	His	Asp	Val	Glu	Thr	Gln	Phe	Asn	Gln	Tyr	Lys	Thr	
	130					135					140					
Glu	Ala	Ala	Ser	Arg	Tyr	Asn	Leu	Thr	Ile	Ser	Asp	Val	Ser	Val	Ser	
145				150						155					160	
Asp	Val	Pro	Phe	Pro	Phe	Ser	Ala	Gln	Ser	Gly	Ala	Gly	Val	Pro	Gly	
			165						170					175		
Trp	Gly	Ile	Ala	Leu	Leu	Val	Leu	Val	Cys	Val	Leu	Val	Ala	Leu	Ala	
			180					185					190			
Ile	Val	Tyr	Leu	Ile	Ala	Leu	Ala	Val	Cys	Gln	Cys	Arg	Arg	Lys	Asn	
		195					200					205				
Tyr	Gly	Gln	Leu	Asp	Ile	Phe	Pro	Ala	Arg	Asp	Thr	Tyr	His	Pro	Met	
	210					215					220					
Ser	Glu	Tyr	Pro	Thr	Tyr	His	Thr	His	Gly	Arg	Tyr	Val	Pro	Pro	Ser	
225				230						235					240	
Ser	Thr	Asp	Arg	Ser	Pro	Tyr	Glu	Lys	Val	Ser	Ala	Gly	Asn	Gly	Gly	
			245						250					255		
Ser	Ser	Leu	Ser	Tyr	Thr	Asn	Pro	Ala	Val	Ala	Ala	Thr	Ser	Ala	Asn	
			260					265					270			

Leu

&lt;210&gt; 229

&lt;211&gt; 310

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 229

Met	Arg	Arg	Ala	Ala	Leu	Trp	Leu	Trp	Leu	Cys	Ala	Leu	Ala	Leu	Ser	
1				5					10					15		
Leu	Gln	Pro	Ala	Leu	Pro	Gln	Ile	Val	Ala	Thr	Asn	Leu	Pro	Pro	Glu	
			20					25					30			
Asp	Gln	Asp	Gly	Ser	Gly	Asp	Asp	Ser	Asp	Asn	Phe	Ser	Gly	Ser	Gly	
		35				40						45				
Ala	Gly	Ala	Leu	Gln	Asp	Ile	Thr	Leu	Ser	Gln	Gln	Thr	Pro	Ser	Thr	
	50					55					60					
Trp	Lys	Asp	Thr	Gln	Leu	Leu	Thr	Ala	Ile	Pro	Thr	Ser	Pro	Glu	Pro	
65				70						75					80	
Thr	Gly	Leu	Glu	Ala	Thr	Ala	Ala	Ser	Thr	Ser	Thr	Leu	Pro	Ala	Gly	
				85					90					95		
Glu	Gly	Pro	Lys	Glu	Gly	Glu	Ala	Val	Val	Leu	Pro	Glu	Val	Glu	Pro	
			100					105					110			
Gly	Leu	Thr	Ala	Arg	Glu	Gln	Glu	Ala	Thr	Pro	Arg	Pro	Arg	Glu	Thr	
			115					120					125			
Thr	Gln	Leu	Pro	Thr	Thr	His	Gln	Ala	Ser	Thr	Thr	Thr	Ala	Thr	Thr	
	130					135						140				
Ala	Gln	Glu	Pro	Ala	Thr	Ser	His	Pro	His	Arg	Asp	Met	Gln	Pro	Gly	
145				150						155					160	
His	His	Glu	Thr	Ser	Thr	Pro	Ala	Gly	Pro	Ser	Gln	Ala	Asp	Leu	His	

				165					170					175		
Thr	Pro	His	Thr	Glu	Asp	Gly	Gly	Pro	Ser	Ala	Thr	Glu	Arg	Ala	Ala	
			180					185					190			
Glu	Asp	Gly	Ala	Ser	Ser	Gln	Leu	Pro	Ala	Ala	Glu	Gly	Ser	Gly	Glu	
		195					200					205				
Gln	Asp	Phe	Thr	Phe	Glu	Thr	Ser	Gly	Glu	Asn	Thr	Ala	Val	Val	Ala	
	210					215					220					
Val	Glu	Pro	Asp	Arg	Arg	Asn	Gln	Ser	Pro	Val	Asp	Gln	Gly	Ala	Thr	
225				230						235					240	
Gly	Ala	Ser	Gln	Gly	Leu	Leu	Asp	Arg	Lys	Glu	Val	Leu	Gly	Gly	Val	
			245					250						255		
Ile	Ala	Gly	Gly	Leu	Val	Gly	Leu	Ile	Phe	Ala	Val	Cys	Leu	Val	Gly	
			260					265					270			
Phe	Met	Leu	Tyr	Arg	Met	Lys	Lys	Lys	Asp	Glu	Gly	Ser	Tyr	Ser	Leu	
		275				280						285				
Glu	Glu	Pro	Lys	Gln	Ala	Asn	Gly	Gly	Ala	Tyr	Gln	Lys	Pro	Thr	Lys	
		290				295					300					
Gln	Glu	Glu	Phe	Tyr	Ala											
305					310											

&lt;210&gt; 230

&lt;211&gt; 1048

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 230

Met	Ala	Phe	Pro	Pro	Arg	Arg	Arg	Leu	Arg	Leu	Gly	Pro	Arg	Gly	Leu	
1			5					10					15			
Pro	Leu	Leu	Leu	Ser	Gly	Leu	Leu	Leu	Pro	Leu	Cys	Arg	Ala	Phe	Asn	
		20						25					30			
Leu	Asp	Val	Asp	Ser	Pro	Ala	Glu	Tyr	Ser	Gly	Pro	Glu	Gly	Ser	Tyr	
		35				40						45				
Phe	Gly	Phe	Ala	Val	Asp	Phe	Phe	Val	Pro	Ser	Ala	Ser	Ser	Arg	Met	
	50				55						60					
Phe	Leu	Leu	Val	Gly	Ala	Pro	Lys	Ala	Asn	Thr	Thr	Gln	Pro	Gly	Ile	
65				70						75					80	
Val	Glu	Gly	Gly	Gln	Val	Leu	Lys	Cys	Asp	Trp	Ser	Ser	Thr	Arg	Arg	
			85					90						95		
Cys	Gln	Pro	Ile	Glu	Phe	Asp	Ala	Thr	Gly	Asn	Arg	Asp	Tyr	Ala	Lys	
		100						105					110			
Asp	Asp	Pro	Leu	Glu	Phe	Lys	Ser	His	Gln	Trp	Phe	Gly	Ala	Ser	Val	
		115				120						125				
Arg	Ser	Lys	Gln	Asp	Lys	Ile	Leu	Ala	Cys	Ala	Pro	Leu	Tyr	His	Trp	
	130				135						140					
Arg	Thr	Glu	Met	Lys	Gln	Glu	Arg	Glu	Pro	Val	Gly	Thr	Cys	Phe	Leu	
145				150						155					160	
Gln	Asp	Gly	Thr	Lys	Thr	Val	Glu	Tyr	Ala	Pro	Cys	Arg	Ser	Gln	Asp	
			165						170					175		
Ile	Asp	Ala	Asp	Gly	Gln	Gly	Phe	Cys	Gln	Gly	Gly	Phe	Ser	Ile	Asp	
		180						185					190			
Phe	Thr	Lys	Ala	Asp	Arg	Val	Leu	Leu	Gly	Gly	Pro	Gly	Ser	Phe	Tyr	
		195				200						205				
Trp	Gln	Gly	Gln	Leu	Ile	Ser	Asp	Gln	Val	Ala	Glu	Ile	Val	Ser	Lys	
	210				215						220					
Tyr	Asp	Pro	Asn	Val	Tyr	Ser	Ile	Lys	Tyr	Asn	Asn	Gln	Leu	Ala	Thr	
225				230						235					240	
Arg	Thr	Ala	Gln	Ala	Ile	Phe	Asp	Asp	Ser	Tyr	Leu	Gly	Tyr	Ser	Val	
			245					250						255		
Ala	Val	Gly	Asp	Phe	Asn	Gly	Asp	Gly	Ile	Asp	Asp	Phe	Val	Ser	Gly	
		260						265					270			
Val	Pro	Arg	Ala	Ala	Arg	Thr	Leu	Gly	Met	Val	Tyr	Ile	Tyr	Asp	Gly	
		275					280					285				
Lys	Asn	Met	Ser	Ser	Leu	Tyr	Asn	Phe	Thr	Gly	Glu	Gln	Met	Ala	Ala	

290						295						300					
Tyr	Phe	Gly	Phe	Ser	Val	Ala	Ala	Thr	Asp	Ile	Asn	Gly	Asp	Asp	Tyr		
305					310					315					320		
Ala	Asp	Val	Phe	Ile	Gly	Ala	Pro	Leu	Phe	Met	Asp	Arg	Gly	Ser	Asp		
				325					330					335			
Gly	Lys	Leu	Gln	Glu	Val	Gly	Gln	Val	Ser	Val	Ser	Leu	Gln	Arg	Ala		
			340					345					350				
Ser	Gly	Asp	Phe	Gln	Thr	Thr	Lys	Leu	Asn	Gly	Phe	Glu	Val	Phe	Ala		
		355					360					365					
Arg	Phe	Gly	Ser	Ala	Ile	Ala	Pro	Leu	Gly	Asp	Leu	Asp	Gln	Asp	Gly		
	370					375					380						
Phe	Asn	Asp	Ile	Ala	Ile	Ala	Ala	Pro	Tyr	Gly	Gly	Glu	Asp	Lys	Lys		
385					390					395					400		
Gly	Ile	Val	Tyr	Ile	Phe	Asn	Gly	Arg	Ser	Thr	Gly	Leu	Asn	Ala	Val		
				405					410					415			
Pro	Ser	Gln	Ile	Leu	Glu	Gly	Gln	Trp	Ala	Ala	Arg	Ser	Met	Pro	Pro		
			420					425					430				
Ser	Phe	Gly	Tyr	Ser	Met	Lys	Gly	Ala	Thr	Asp	Ile	Asp	Lys	Asn	Gly		
		435					440					445					
Tyr	Pro	Asp	Leu	Ile	Val	Gly	Ala	Phe	Gly	Val	Asp	Arg	Ala	Ile	Leu		
	450					455					460						
Tyr	Arg	Ala	Arg	Pro	Val	Ile	Thr	Val	Asn	Ala	Gly	Leu	Glu	Val	Tyr		
465					470					475					480		
Pro	Ser	Ile	Leu	Asn	Gln	Asp	Asn	Lys	Thr	Cys	Ser	Leu	Pro	Gly	Thr		
				485					490					495			
Ala	Leu	Lys	Val	Ser	Cys	Phe	Asn	Val	Arg	Phe	Cys	Leu	Lys	Ala	Asp		
			500					505					510				
Gly	Lys	Gly	Val	Leu	Pro	Arg	Lys	Leu	Asn	Phe	Gln	Val	Glu	Leu	Leu		
		515						520				525					
Leu	Asp	Lys	Leu	Lys	Gln	Lys	Gly	Ala	Ile	Arg	Arg	Ala	Leu	Phe	Leu		
	530					535					540						
Tyr	Ser	Arg	Ser	Pro	Ser	His	Ser	Lys	Asn	Met	Thr	Ile	Ser	Arg	Gly		
545					550					555					560		
Gly	Leu	Met	Gln	Cys	Glu	Glu	Leu	Ile	Ala	Tyr	Leu	Arg	Asp	Glu	Ser		
				565					570					575			
Glu	Phe	Arg	Asp	Lys	Leu	Thr	Pro	Ile	Thr	Ile	Phe	Met	Glu	Tyr	Arg		
			580					585					590				
Leu	Asp	Tyr	Arg	Thr	Ala	Ala	Asp	Thr	Thr	Gly	Leu	Gln	Pro	Ile	Leu		
		595					600					605					
Asn	Gln	Phe	Thr	Pro	Ala	Asn	Ile	Ser	Arg	Gln	Ala	His	Ile	Leu	Leu		
						615				620							
Asp	Cys	Gly	Glu	Asp	Asn	Val	Cys	Lys	Pro	Lys	Leu	Glu	Val	Ser	Val		
625					630					635					640		
Asp	Ser	Asp	Gln	Lys	Lys	Ile	Tyr	Ile	Gly	Asp	Asp	Asn	Pro	Leu	Thr		
				645					650					655			
Leu	Ile	Val	Lys	Ala	Gln	Asn	Gln	Gly	Glu	Gly	Ala	Tyr	Glu	Ala	Glu		
			660					665					670				
Leu	Ile	Val	Ser	Ile	Pro	Leu	Gln	Ala	Asp	Phe	Ile	Gly	Val	Val	Arg		
		675					680					685					
Asn	Asn	Glu	Ala	Leu	Ala	Arg	Leu	Ser	Cys	Ala	Phe	Lys	Thr	Glu	Asn		
						695					700						
Gln	Thr	Arg	Gln	Val	Val	Cys	Asp	Leu	Gly	Asn	Pro	Met	Lys	Ala	Gly		
705					710					715					720		
Thr	Gln	Leu	Leu	Ala	Gly	Leu	Arg	Phe	Ser	Val	His	Gln	Gln	Ser	Glu		
				725					730					735			
Met	Asp	Thr	Ser	Val	Lys	Phe	Asp	Leu	Gln	Ile	Gln	Ser	Ser	Asn	Leu		
			740					745						750			
Phe	Asp	Lys	Val	Ser	Pro	Val	Val	Ser	His	Lys	Val	Asp	Leu	Ala	Val		
		755					760						765				
Leu	Ala	Ala	Val	Glu	Ile	Arg	Gly	Val	Ser	Ser	Pro	Asp	His	Ile	Phe		
	770					775					780						
Leu	Pro	Ile	Pro	Asn	Trp	Glu	His	Lys	Glu	Asn	Pro	Glu	Thr	Glu	Glu		
785					790					795					800		

Asp Val Gly Pro Val Val Gln His Ile Tyr Glu Leu Arg Asn Asn Gly  
 805 810 815  
 Pro Ser Ser Phe Ser Lys Ala Met Leu His Leu Gln Trp Pro Tyr Lys  
 820 825 830  
 Tyr Asn Asn Asn Thr Leu Leu Tyr Ile Leu His Tyr Asp Ile Asp Gly  
 835 840 845  
 Pro Met Asn Cys Thr Ser Asp Met Glu Ile Asn Pro Leu Arg Ile Lys  
 850 855 860  
 Ile Ser Ser Leu Gln Thr Thr Glu Lys Asn Asp Thr Val Ala Gly Gln  
 865 870 875  
 Gly Glu Arg Asp His Leu Ile Thr Lys Arg Asp Leu Ala Leu Ser Glu  
 885 890 895  
 Gly Asp Ile His Thr Leu Gly Cys Gly Val Ala Gln Cys Leu Lys Ile  
 900 905 910  
 Val Cys Gln Val Gly Arg Leu Asp Arg Gly Lys Ser Ala Ile Leu Tyr  
 915 920 925  
 Val Lys Ser Leu Leu Trp Thr Glu Thr Phe Met Asn Lys Glu Asn Gln  
 930 935 940  
 Asn His Ser Tyr Ser Leu Lys Ser Ser Ala Ser Phe Asn Val Ile Glu  
 945 950 955  
 Phe Pro Tyr Lys Asn Leu Pro Ile Glu Asp Ile Thr Asn Ser Thr Leu  
 965 970 975  
 Val Thr Thr Asn Val Thr Trp Gly Ile Gln Pro Ala Pro Met Pro Val  
 980 985 990  
 Pro Val Trp Val Ile Ile Leu Ala Val Leu Ala Gly Leu Leu Leu Leu  
 995 1000 1005  
 Ala Val Leu Val Phe Val Met Tyr Arg Met Gly Phe Phe Lys Arg Val  
 1010 1015 1020  
 Arg Pro Pro Gln Glu Glu Gln Glu Arg Glu Gln Leu Gln Pro His Glu  
 1025 1030 1035 1040  
 Asn Gly Glu Gly Asn Ser Glu Thr  
 1045

&lt;210&gt; 231

&lt;211&gt; 188

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 231

Met Asp Cys Arg Lys Met Ala Arg Phe Ser Tyr Ser Val Ile Trp Ile  
 1 5 10 15  
 Met Ala Ile Ser Lys Val Phe Glu Leu Gly Leu Val Ala Gly Leu Gly  
 20 25 30  
 His Gln Glu Phe Ala Arg Pro Ser Arg Gly Tyr Leu Ala Phe Arg Asp  
 35 40 45  
 Asp Ser Ile Trp Pro Gln Glu Glu Pro Ala Ile Arg Pro Arg Ser Ser  
 50 55 60  
 Gln Arg Val Pro Pro Met Gly Ile Gln His Ser Lys Glu Leu Asn Arg  
 65 70 75 80  
 Thr Cys Cys Leu Asn Gly Gly Thr Cys Met Leu Gly Ser Phe Cys Ala  
 85 90 95  
 Cys Pro Pro Ser Phe Tyr Gly Arg Asn Cys Glu His Asp Val Arg Lys  
 100 105 110  
 Glu Asn Cys Gly Ser Val Pro His Asp Thr Trp Leu Pro Lys Lys Cys  
 115 120 125  
 Ser Leu Cys Lys Cys Trp His Gly Gln Leu Arg Cys Phe Pro Gln Ala  
 130 135 140  
 Phe Leu Pro Gly Cys Asp Gly Leu Val Met Asp Glu His Leu Val Ala  
 145 150 155 160  
 Ser Arg Thr Pro Glu Leu Pro Pro Ser Ala Arg Thr Thr Thr Phe Met  
 165 170 175  
 Leu Val Gly Ile Cys Leu Ser Ile Gln Ser Tyr Tyr  
 180 185

&lt;210&gt; 232

&lt;211&gt; 750

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 232

Met	Trp	Asn	Leu	Leu	His	Glu	Thr	Asp	Ser	Ala	Val	Ala	Thr	Ala	Arg
1			5						10					15	
Arg	Pro	Arg	Trp	Leu	Cys	Ala	Gly	Ala	Leu	Val	Leu	Ala	Gly	Gly	Phe
			20					25					30		
Phe	Leu	Leu	Gly	Phe	Leu	Phe	Gly	Trp	Phe	Ile	Lys	Ser	Ser	Asn	Glu
		35					40					45			
Ala	Thr	Asn	Ile	Thr	Pro	Lys	His	Asn	Met	Lys	Ala	Phe	Leu	Asp	Glu
	50					55					60				
Leu	Lys	Ala	Glu	Asn	Ile	Lys	Lys	Phe	Leu	Tyr	Asn	Phe	Thr	Gln	Ile
65				70						75					80
Pro	His	Leu	Ala	Gly	Thr	Glu	Gln	Asn	Phe	Gln	Leu	Ala	Lys	Gln	Ile
				85					90					95	
Gln	Ser	Gln	Trp	Lys	Glu	Phe	Gly	Leu	Asp	Ser	Val	Glu	Leu	Ala	His
			100					105					110		
Tyr	Asp	Val	Leu	Leu	Ser	Tyr	Pro	Asn	Lys	Thr	His	Pro	Asn	Tyr	Ile
	115						120					125			
Ser	Ile	Ile	Asn	Glu	Asp	Gly	Asn	Glu	Ile	Phe	Asn	Thr	Ser	Leu	Phe
	130					135					140				
Glu	Pro	Pro	Pro	Pro	Gly	Tyr	Glu	Asn	Val	Ser	Asp	Ile	Val	Pro	Pro
145					150					155					160
Phe	Ser	Ala	Phe	Ser	Pro	Gln	Gly	Met	Pro	Glu	Gly	Asp	Leu	Val	Tyr
				165					170					175	
Val	Asn	Tyr	Ala	Arg	Thr	Glu	Asp	Phe	Phe	Lys	Leu	Glu	Arg	Asp	Met
			180					185						190	
Lys	Ile	Asn	Cys	Ser	Gly	Lys	Ile	Val	Ile	Ala	Arg	Tyr	Gly	Lys	Val
		195					200					205			
Phe	Arg	Gly	Asn	Lys	Val	Lys	Asn	Ala	Gln	Leu	Ala	Gly	Ala	Lys	Gly
	210					215					220				
Val	Ile	Leu	Tyr	Ser	Asp	Pro	Ala	Asp	Tyr	Phe	Ala	Pro	Gly	Val	Lys
225					230					235					240
Ser	Tyr	Pro	Asp	Gly	Trp	Asn	Leu	Pro	Gly	Gly	Gly	Val	Gln	Arg	Gly
			245						250					255	
Asn	Ile	Leu	Asn	Leu	Asn	Gly	Ala	Gly	Asp	Pro	Leu	Thr	Pro	Gly	Tyr
			260					265					270		
Pro	Ala	Asn	Glu	Tyr	Ala	Tyr	Arg	Arg	Gly	Ile	Ala	Glu	Ala	Val	Gly
		275					280					285			
Leu	Pro	Ser	Ile	Pro	Val	His	Pro	Ile	Gly	Tyr	Tyr	Asp	Ala	Gln	Lys
	290					295						300			
Leu	Leu	Glu	Lys	Met	Gly	Gly	Ser	Ala	Pro	Pro	Asp	Ser	Ser	Trp	Arg
305					310					315					320
Gly	Ser	Leu	Lys	Val	Pro	Tyr	Asn	Val	Gly	Pro	Gly	Phe	Thr	Gly	Asn
				325					330					335	
Phe	Ser	Thr	Gln	Lys	Val	Lys	Met	His	Ile	His	Ser	Thr	Asn	Glu	Val
			340					345					350		
Thr	Arg	Ile	Tyr	Asn	Val	Ile	Gly	Thr	Leu	Arg	Gly	Ala	Val	Glu	Pro
		355					360					365			
Asp	Arg	Tyr	Val	Ile	Leu	Gly	Gly	His	Arg	Asp	Ser	Trp	Val	Phe	Gly
	370					375					380				
Gly	Ile	Asp	Pro	Gln	Ser	Gly	Ala	Ala	Val	Val	His	Glu	Ile	Val	Arg
385					390					395					400
Ser	Phe	Gly	Thr	Leu	Lys	Lys	Glu	Gly	Trp	Arg	Pro	Arg	Arg	Thr	Ile
				405					410					415	
Leu	Phe	Ala	Ser	Trp	Asp	Ala	Glu	Glu	Phe	Gly	Leu	Leu	Gly	Ser	Thr
			420					425					430		
Glu	Trp	Ala	Glu	Glu	Asn	Ser	Arg	Leu	Leu	Gln	Glu	Arg	Gly	Val	Ala
		435					440					445			

Tyr Ile Asn Ala Asp Ser Ser Ile Glu Gly Asn Tyr Thr Leu Arg Val  
 450 455 460  
 Asp Cys Thr Pro Leu Met Tyr Ser Leu Val His Asn Leu Thr Lys Glu  
 465 470 475 480  
 Leu Lys Ser Pro Asp Glu Gly Phe Glu Gly Lys Ser Leu Tyr Glu Ser  
 485 490 495  
 Trp Thr Lys Lys Ser Pro Ser Pro Glu Phe Ser Gly Met Pro Arg Ile  
 500 505 510  
 Ser Lys Leu Gly Ser Gly Asn Asp Phe Glu Val Phe Phe Gln Arg Leu  
 515 520 525  
 Gly Ile Ala Ser Gly Arg Ala Arg Tyr Thr Lys Asn Trp Glu Thr Asn  
 530 535 540  
 Lys Phe Ser Gly Tyr Pro Leu Tyr His Ser Val Tyr Glu Thr Tyr Glu  
 545 550 555 560  
 Leu Val Glu Lys Phe Tyr Asp Pro Met Phe Lys Tyr His Leu Thr Val  
 565 570 575  
 Ala Gln Val Arg Gly Gly Met Val Phe Glu Leu Ala Asn Ser Ile Val  
 580 585 590  
 Leu Pro Phe Asp Cys Arg Asp Tyr Ala Val Val Leu Arg Lys Tyr Ala  
 595 600 605  
 Asp Lys Ile Tyr Ser Ile Ser Met Lys His Pro Gln Glu Met Lys Thr  
 610 615 620  
 Tyr Ser Val Ser Phe Asp Ser Leu Phe Ser Ala Val Lys Asn Phe Thr  
 625 630 635 640  
 Glu Ile Ala Ser Lys Phe Ser Glu Arg Leu Gln Asp Phe Asp Lys Ser  
 645 650 655  
 Asn Pro Ile Val Leu Arg Met Met Asn Asp Gln Leu Met Phe Leu Glu  
 660 665 670  
 Arg Ala Phe Ile Asp Pro Leu Gly Leu Pro Asp Arg Pro Phe Tyr Arg  
 675 680 685  
 His Val Ile Tyr Ala Pro Ser Ser His Asn Lys Tyr Ala Gly Glu Ser  
 690 695 700  
 Phe Pro Gly Ile Tyr Asp Ala Leu Phe Asp Ile Glu Ser Lys Val Asp  
 705 710 715 720  
 Pro Ser Lys Ala Trp Gly Glu Val Lys Arg Gln Ile Tyr Val Ala Ala  
 725 730 735  
 Phe Thr Val Gln Ala Ala Ala Glu Thr Leu Ser Glu Val Ala  
 740 745 750

&lt;210&gt; 233

&lt;211&gt; 976

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 233

Met Glu Leu Gln Ala Ala Arg Ala Cys Phe Ala Leu Leu Trp Gly Cys  
 1 5 10 15  
 Ala Leu Ala Ala Ala Ala Ala Ala Gln Gly Lys Glu Val Val Leu Leu  
 20 25 30  
 Asp Phe Ala Ala Ala Gly Gly Glu Leu Gly Trp Leu Thr His Pro Tyr  
 35 40 45  
 Gly Lys Gly Trp Asp Leu Met Gln Asn Ile Met Asn Asp Met Pro Ile  
 50 55 60  
 Tyr Met Tyr Ser Val Cys Asn Val Met Ser Gly Asp Gln Asp Asn Trp  
 65 70 75 80  
 Leu Arg Thr Asn Trp Val Tyr Arg Gly Glu Ala Glu Arg Ile Phe Ile  
 85 90 95  
 Glu Leu Lys Phe Thr Val Arg Asp Cys Asn Ser Phe Pro Gly Gly Ala  
 100 105 110  
 Ser Ser Cys Lys Glu Thr Phe Asn Leu Tyr Tyr Ala Glu Ser Asp Leu  
 115 120 125  
 Asp Tyr Gly Thr Asn Phe Gln Lys Arg Leu Phe Thr Lys Ile Asp Thr  
 130 135 140

Ile	Ala	Pro	Asp	Glu	Ile	Thr	Val	Ser	Ser	Asp	Phe	Glu	Ala	Arg	His
145					150					155					160
Val	Lys	Leu	Asn	Val	Glu	Glu	Arg	Ser	Val	Gly	Pro	Leu	Thr	Arg	Lys
			165					170							175
Gly	Phe	Tyr	Leu	Ala	Phe	Gln	Asp	Ile	Gly	Ala	Cys	Val	Ala	Leu	Leu
			180					185							190
Ser	Val	Arg	Val	Tyr	Tyr	Lys	Lys	Cys	Pro	Glu	Leu	Leu	Gln	Gly	Leu
		195					200					205			
Ala	His	Phe	Pro	Glu	Thr	Ile	Ala	Gly	Ser	Asp	Ala	Pro	Ser	Leu	Ala
	210					215				220					
Thr	Val	Ala	Gly	Thr	Cys	Val	Asp	His	Ala	Val	Val	Pro	Pro	Gly	Gly
225					230					235					240
Glu	Glu	Pro	Arg	Met	His	Cys	Ala	Val	Asp	Gly	Glu	Trp	Leu	Val	Pro
				245					250						255
Ile	Gly	Gln	Cys	Leu	Cys	Gln	Ala	Gly	Tyr	Glu	Lys	Val	Glu	Asp	Ala
			260					265						270	
Cys	Gln	Ala	Cys	Ser	Pro	Gly	Phe	Phe	Lys	Phe	Glu	Ala	Ser	Glu	Ser
		275					280					285			
Pro	Cys	Leu	Glu	Cys	Pro	Glu	His	Thr	Leu	Pro	Ser	Pro	Glu	Gly	Ala
		290					295					300			
Thr	Ser	Cys	Glu	Cys	Glu	Glu	Gly	Phe	Phe	Arg	Ala	Pro	Gln	Asp	Pro
305					310					315					320
Ala	Ser	Met	Pro	Cys	Thr	Arg	Pro	Pro	Ser	Ala	Pro	His	Tyr	Leu	Thr
				325					330						335
Ala	Val	Gly	Met	Gly	Ala	Lys	Val	Glu	Leu	Arg	Trp	Thr	Pro	Pro	Gln
			340					345						350	
Asp	Ser	Gly	Gly	Arg	Glu	Asp	Ile	Val	Tyr	Ser	Val	Thr	Cys	Glu	Gln
		355					360						365		
Cys	Trp	Pro	Glu	Ser	Gly	Glu	Cys	Gly	Pro	Cys	Glu	Ala	Ser	Val	Arg
		370				375					380				
Tyr	Ser	Glu	Pro	Pro	His	Gly	Leu	Thr	Arg	Thr	Ser	Val	Thr	Val	Ser
385					390					395					400
Asp	Leu	Glu	Pro	His	Met	Asn	Tyr	Thr	Phe	Thr	Val	Glu	Ala	Arg	Asn
				405					410						415
Gly	Val	Ser	Gly	Leu	Val	Thr	Ser	Arg	Ser	Phe	Arg	Thr	Ala	Ser	Val
			420					425						430	
Ser	Ile	Asn	Gln	Thr	Glu	Pro	Pro	Lys	Val	Arg	Leu	Glu	Gly	Arg	Ser
		435						440					445		
Thr	Thr	Ser	Leu	Ser	Val	Ser	Trp	Ser	Ile	Pro	Pro	Pro	Gln	Gln	Ser
	450					455						460			
Arg	Val	Trp	Lys	Tyr	Glu	Val	Thr	Tyr	Arg	Lys	Lys	Gly	Asp	Ser	Asn
465					470					475					480
Ser	Tyr	Asn	Val	Arg	Arg	Thr	Glu	Gly	Phe	Ser	Val	Thr	Leu	Asp	Asp
				485					490						495
Leu	Ala	Pro	Asp	Thr	Thr	Tyr	Leu	Val	Gln	Val	Gln	Ala	Leu	Thr	Gln
			500					505						510	
Glu	Gly	Gln	Gly	Ala	Gly	Ser	Lys	Val	His	Glu	Phe	Gln	Thr	Leu	Ser
		515					520					525			
Pro	Glu	Gly	Ser	Gly	Asn	Leu	Ala	Val	Ile	Gly	Gly	Val	Ala	Val	Gly
		530				535						540			
Val	Val	Leu	Leu	Leu	Val	Leu	Ala	Gly	Val	Gly	Phe	Phe	Ile	His	Arg
545					550					555					560
Arg	Arg	Lys	Asn	Gln	Arg	Ala	Arg	Gln	Ser	Pro	Glu	Asp	Val	Tyr	Phe
				565					570						575
Ser	Lys	Ser	Glu	Gln	Leu	Lys	Pro	Leu	Lys	Thr	Tyr	Val	Asp	Pro	His
			580					585					590		
Thr	Tyr	Glu	Asp	Pro	Asn	Gln	Ala	Val	Leu	Lys	Phe	Thr	Thr	Glu	Ile
		595					600						605		
His	Pro	Ser	Cys	Val	Thr	Arg	Gln	Lys	Val	Ile	Gly	Ala	Gly	Glu	Phe
	610					615						620			
Gly	Glu	Val	Tyr	Lys	Gly	Met	Leu	Lys	Thr	Ser	Ser	Gly	Lys	Lys	Glu
625					630					635					640
Val	Pro	Val	Ala	Ile	Lys	Thr	Leu	Lys	Ala	Gly	Tyr	Thr	Glu	Lys	Gln



				645					650					655		
Arg	Val	Asp	Phe	Leu	Gly	Glu	Ala	Gly	Ile	Met	Gly	Gln	Phe	Ser	His	
			660					665					670			
His	Asn	Ile	Ile	Arg	Leu	Glu	Gly	Val	Ile	Ser	Lys	Tyr	Lys	Pro	Met	
		675					680					685				
Met	Ile	Ile	Thr	Glu	Tyr	Met	Glu	Asn	Gly	Ala	Leu	Asp	Lys	Phe	Leu	
	690					695					700					
Arg	Glu	Lys	Asp	Gly	Glu	Phe	Ser	Val	Leu	Gln	Leu	Val	Gly	Met	Leu	
705					710					715					720	
Arg	Gly	Ile	Ala	Ala	Gly	Met	Lys	Tyr	Leu	Ala	Asn	Met	Asn	Tyr	Val	
			725					730						735		
His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Ile	Leu	Val	Asn	Ser	Asn	Leu	Val	
		740					745						750			
Cys	Lys	Val	Ser	Asp	Phe	Gly	Leu	Ser	Arg	Val	Leu	Glu	Asp	Asp	Pro	
		755				760						765				
Glu	Ala	Thr	Tyr	Thr	Thr	Ser	Gly	Gly	Lys	Ile	Pro	Ile	Arg	Trp	Thr	
	770					775					780					
Ala	Pro	Glu	Ala	Ile	Ser	Tyr	Arg	Lys	Phe	Thr	Ser	Ala	Ser	Asp	Val	
785					790					795					800	
Trp	Ser	Phe	Gly	Ile	Val	Met	Trp	Glu	Val	Met	Thr	Tyr	Gly	Glu	Arg	
			805					810						815		
Pro	Tyr	Trp	Glu	Leu	Ser	Asn	His	Glu	Val	Met	Lys	Ala	Ile	Asn	Asp	
			820					825						830		
Gly	Phe	Arg	Leu	Pro	Thr	Pro	Met	Asp	Cys	Pro	Ser	Ala	Ile	Tyr	Gln	
		835					840					845				
Leu	Met	Met	Gln	Cys	Trp	Gln	Gln	Glu	Arg	Ala	Arg	Arg	Pro	Lys	Phe	
	850					855					860					
Ala	Asp	Ile	Val	Ser	Ile	Leu	Asp	Lys	Leu	Ile	Arg	Ala	Pro	Asp	Ser	
865					870					875					880	
Leu	Lys	Thr	Leu	Ala	Asp	Phe	Asp	Pro	Arg	Val	Ser	Ile	Arg	Leu	Pro	
			885					890						895		
Ser	Thr	Ser	Gly	Ser	Glu	Gly	Val	Pro	Phe	Arg	Thr	Val	Ser	Glu	Trp	
			900					905						910		
Leu	Glu	Ser	Ile	Lys	Met	Gln	Gln	Tyr	Thr	Glu	His	Phe	Met	Ala	Ala	
		915					920					925				
Gly	Tyr	Thr	Ala	Ile	Glu	Lys	Val	Val	Gln	Met	Thr	Asn	Asp	Asp	Ile	
	930					935					940					
Lys	Arg	Ile	Gly	Val	Arg	Leu	Pro	Gly	His	Gln	Lys	Arg	Ile	Ala	Tyr	
945					950					955					960	
Ser	Leu	Leu	Gly	Leu	Lys	Asp	Gln	Val	Asn	Thr	Val	Gly	Ile	Pro	Ile	
			965					970						975		

&lt;210&gt; 234

&lt;211&gt; 668

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 234

Met	Gly	Gly	Lys	Gln	Arg	Asp	Glu	Asp	Asp	Glu	Ala	Tyr	Gly	Lys	Pro	
1				5					10					15		
Val	Lys	Tyr	Asp	Pro	Ser	Phe	Arg	Gly	Pro	Ile	Lys	Asn	Arg	Ser	Cys	
			20					25					30			
Thr	Asp	Val	Ile	Cys	Cys	Val	Leu	Phe	Leu	Leu	Phe	Ile	Leu	Gly	Tyr	
		35					40					45				
Ile	Val	Val	Gly	Ile	Val	Ala	Trp	Leu	Tyr	Gly	Asp	Pro	Arg	Gln	Val	
	50					55					60					
Leu	Tyr	Pro	Arg	Asn	Ser	Thr	Gly	Ala	Tyr	Cys	Gly	Met	Gly	Glu	Asn	
65					70					75					80	
Lys	Asp	Lys	Pro	Tyr	Leu	Leu	Tyr	Phe	Asn	Ile	Phe	Ser	Cys	Ile	Leu	
				85					90					95		
Ser	Ser	Asn	Ile	Ile	Ser	Val	Ala	Glu	Asn	Gly	Leu	Gln	Cys	Pro	Thr	

100 105 110  
 Pro Gln Thr Val Ile Thr Ser Leu Gln Gln Glu Leu Cys Pro Ser Phe  
 115 120 125  
 Leu Leu Pro Ser Ala Pro Ala Leu Gly Arg Cys Phe Pro Trp Thr Asn  
 130 135 140  
 Val Thr Pro Pro Ala Leu Pro Gly Ile Thr Asn Asp Thr Thr Ile Gln  
 145 150 155 160  
 Gln Gly Ile Ser Gly Leu Ile Asp Ser Leu Asn Ala Arg Asp Ile Ser  
 165 170 175  
 Val Lys Ile Phe Glu Asp Phe Ala Gln Ser Trp Tyr Trp Ile Leu Val  
 180 185 190  
 Ala Leu Gly Val Ala Leu Val Leu Ser Leu Leu Phe Ile Leu Leu Leu  
 195 200 205  
 Arg Leu Val Ala Gly Pro Leu Val Leu Val Leu Ile Leu Gly Val Leu  
 210 215 220  
 Gly Val Leu Ala Tyr Gly Ile Tyr Tyr Cys Trp Glu Glu Tyr Arg Val  
 225 230 235 240  
 Leu Arg Asp Lys Gly Ala Ser Ile Ser Gln Leu Gly Phe Thr Thr Asn  
 245 250 255  
 Leu Ser Ala Tyr Gln Ser Val Gln Glu Thr Trp Leu Ala Ala Leu Ile  
 260 265 270  
 Val Leu Ala Val Leu Glu Ala Ile Leu Leu Leu Met Leu Ile Phe Leu  
 275 280 285  
 Arg Gln Arg Ile Arg Ile Ala Ile Ala Leu Leu Lys Glu Ala Ser Lys  
 290 295 300  
 Ala Val Gly Gln Met Met Ser Thr Met Phe Tyr Pro Leu Val Thr Phe  
 305 310 315 320  
 Val Leu Leu Leu Ile Cys Ile Ala Tyr Trp Ala Met Thr Ala Leu Tyr  
 325 330 335  
 Leu Ala Thr Ser Gly Gln Pro Gln Tyr Val Leu Trp Ala Ser Asn Ile  
 340 345 350  
 Ser Ser Pro Gly Cys Glu Lys Val Pro Ile Asn Thr Ser Cys Asn Pro  
 355 360 365  
 Thr Ala His Leu Val Asn Ser Ser Cys Pro Gly Leu Met Cys Val Phe  
 370 375 380  
 Gln Gly Tyr Ser Ser Lys Gly Leu Ile Gln Arg Ser Val Phe Asn Leu  
 385 390 395 400  
 Gln Ile Tyr Gly Val Leu Gly Leu Phe Trp Thr Leu Asn Trp Val Leu  
 405 410 415  
 Ala Leu Gly Gln Cys Val Leu Ala Gly Ala Phe Ala Ser Phe Tyr Trp  
 420 425 430  
 Ala Phe His Lys Pro Gln Asp Ile Pro Thr Phe Pro Leu Ile Ser Ala  
 435 440 445  
 Phe Ile Arg Thr Leu Arg Tyr His Thr Gly Ser Leu Ala Phe Gly Ala  
 450 455 460  
 Leu Ile Leu Thr Leu Val Gln Ile Ala Arg Val Ile Leu Glu Tyr Ile  
 465 470 475 480  
 Asp His Lys Leu Arg Gly Val Gln Asn Pro Val Ala Arg Cys Ile Met  
 485 490 495  
 Cys Cys Phe Lys Cys Cys Leu Trp Cys Leu Glu Lys Phe Ile Lys Phe  
 500 505 510  
 Leu Asn Arg Asn Ala Tyr Ile Met Ile Ala Ile Tyr Gly Lys Asn Phe  
 515 520 525  
 Cys Val Ser Ala Lys Asn Ala Phe Met Leu Leu Met Arg Asn Ile Val  
 530 535 540  
 Arg Val Val Val Leu Asp Lys Val Thr Asp Leu Leu Leu Phe Phe Gly  
 545 550 555 560  
 Lys Leu Leu Val Val Gly Gly Val Gly Val Leu Ser Phe Phe Phe Phe  
 565 570 575  
 Ser Gly Arg Ile Pro Gly Leu Gly Lys Asp Phe Lys Ser Pro His Leu  
 580 585 590  
 Asn Tyr Tyr Trp Leu Pro Ile Met Thr Ser Ile Leu Gly Ala Tyr Val  
 595 600 605  
  
 Ile Ala Ser Gly Phe Phe Ser Val Phe Gly Met Cys Val Asp Thr Leu  
 610 615 620  
 Phe Leu Cys Phe Leu Glu Asp Leu Glu Arg Asn Asn Gly Ser Leu Asp  
 625 630 635 640  
 Arg Pro Tyr Tyr Met Ser Lys Ser Leu Leu Lys Ile Leu Gly Lys Lys  
 645 650 655  
 Asn Glu Ala Pro Pro Asp Asn Lys Lys Arg Lys Lys  
 660 665

&lt;210&gt; 235

&lt;211&gt; 214

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; antibody constituent

&lt;400&gt; 235

```

Asp Ile Leu Leu Thr Gln Ser Pro Val Ile Leu Ser Val Ser Pro Gly
1      5      10      15
Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Ser Ile Gly Thr Asn
20     25     30
Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
35     40     45
Lys Tyr Ala Ser Glu Ser Ile Ser Gly Ile Pro Ser Arg Phe Ser Gly
50     55     60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Ser Val Glu Ser
65     70     75     80
Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Asn Asn Asn Trp Pro Thr
85     90     95
Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg Thr Val Ala Ala
100    105    110
Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly
115    120    125
Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala
130    135    140
Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln
145    150    155    160
Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser
165    170    175
Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr
180    185    190
Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser
195    200    205
Phe Asn Arg Gly Glu Cys
210

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&lt;210&gt; 236

&lt;211&gt; 449

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; antibody constituent

&lt;400&gt; 236

```

Gln Val Gln Leu Lys Gln Ser Gly Pro Gly Leu Val Gln Pro Ser Gln
1      5      10      15
Ser Leu Ser Ile Thr Cys Thr Val Ser Gly Phe Ser Leu Thr Asn Tyr
20     25     30
Gly Val His Trp Val Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Leu
35     40     45
Gly Val Ile Trp Ser Gly Gly Asn Thr Asp Tyr Asn Thr Pro Phe Thr

```

50						55					60					
Ser	Arg	Leu	Ser	Ile	Asn	Lys	Asp	Asn	Ser	Lys	Ser	Gln	Val	Phe	Phe	
65					70					75					80	
Lys	Met	Asn	Ser	Leu	Gln	Ser	Asn	Asp	Thr	Ala	Ile	Tyr	Tyr	Cys	Ala	
				85					90					95		
Arg	Ala	Leu	Thr	Tyr	Tyr	Asp	Tyr	Glu	Phe	Ala	Tyr	Trp	Gly	Gln	Gly	
			100					105					110			
Thr	Leu	Val	Thr	Val	Ser	Ala	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	
		115						120					125			
Pro	Leu	Ala	Pro	Ser	Ser	Lys	Ser	Thr	Ser	Gly	Gly	Thr	Ala	Ala	Leu	
130						135					140					
Gly	Cys	Leu	Val	Lys	Asp	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val	Ser	Trp	
145					150				155						160	
Asn	Ser	Gly	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu	
				165					170					175		
Gln	Ser	Ser	Gly	Leu	Tyr	Ser	Leu	Ser	Ser	Val	Val	Thr	Val	Pro	Ser	
			180					185					190			
Ser	Ser	Leu	Gly	Thr	Gln	Thr	Tyr	Ile	Cys	Asn	Val	Asn	His	Lys	Pro	
		195					200					205				
Ser	Asn	Thr	Lys	Val	Asp	Lys	Arg	Val	Glu	Pro	Lys	Ser	Cys	Asp	Lys	
210						215					220					
Thr	His	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu	Leu	Leu	Gly	Gly	Pro	
225					230					235					240	
Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile	Ser	
				245					250					255		
Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	Ser	His	Glu	Asp	
			260					265					270			
Pro	Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His	Asn	
		275					280						285			
Ala	Lys	Thr	Lys	Pro	Arg	Glu	Glu	Gln	Tyr	Asn	Ser	Thr	Tyr	Arg	Val	
290						295					300					
Val	Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp	Trp	Leu	Asn	Gly	Lys	Glu	
305					310					315					320	
Tyr	Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro	Ala	Pro	Ile	Glu	Lys	
				325					330					335		
Thr	Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr	Thr	
		340						345					350			
Leu	Pro	Pro	Ser	Arg	Glu	Glu	Met	Thr	Lys	Asn	Gln	Val	Ser	Leu	Thr	
		355					360						365			
Cys	Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp	Glu	
370					375					380						
Ser	Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val	Leu	
385					390					395					400	
Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp	Lys	
				405					410					415		
Ser	Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His	Glu	
		420						425					430			
Ala	Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro	Gly	
		435					440					445				

Lys

&lt;210&gt; 237

&lt;211&gt; 214

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; antibody constituent

&lt;400&gt; 237

Asp	Ile	Gln	Met	Thr	Gln	Ser	Pro	Ser	Ser	Leu	Ser	Ala	Ser	Val	Gly
1				5				10						15	

Asp Arg Val Thr Ile Thr Cys Gln Ala Ser Gln Asp Ile Ser Asn Tyr  
 20 25 30  
 Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile  
 35 40 45  
 Tyr Asp Ala Ser Asn Leu Glu Thr Gly Val Pro Ser Arg Phe Ser Gly  
 50 55 60  
 Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Leu Gln Pro  
 65 70 75 80  
 Glu Asp Ile Ala Thr Tyr Phe Cys Gln His Phe Asp His Leu Pro Leu  
 85 90 95  
 Ala Phe Gly Gly Thr Lys Val Glu Ile Lys Arg Thr Val Ala Ala  
 100 105 110  
 Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly  
 115 120 125  
 Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala  
 130 135 140  
 Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln  
 145 150 155 160  
 Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser  
 165 170 175  
 Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr  
 180 185 190  
 Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser  
 195 200 205  
 Phe Asn Arg Gly Glu Cys  
 210

<210> 238

<211> 444

<212> PRT

<213> Artificial Sequence

<220>

<223> antibody constituent

<400> 238

Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Glu  
 1 5 10 15  
 Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Gly Ser Val Ser Ser Gly  
 20 25 30  
 Asp Tyr Tyr Trp Thr Trp Ile Arg Gln Ser Pro Gly Lys Gly Leu Glu  
 35 40 45  
 Trp Ile Gly His Ile Tyr Tyr Ser Gly Asn Thr Asn Tyr Asn Pro Ser  
 50 55 60  
 Leu Lys Ser Arg Leu Thr Ile Ser Ile Asp Thr Ser Lys Thr Gln Phe  
 65 70 75 80  
 Ser Leu Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Ile Tyr Tyr  
 85 90 95  
 Cys Val Arg Asp Arg Val Thr Gly Ala Phe Asp Ile Trp Gly Gln Gly  
 100 105 110  
 Thr Met Val Thr Val Ser Ser Ala Ser Thr Lys Gly Pro Ser Val Phe  
 115 120 125  
 Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu  
 130 135 140  
 Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp  
 145 150 155 160  
 Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu  
 165 170 175  
 Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser  
 180 185 190  
 Ser Asn Phe Gly Thr Gln Thr Tyr Thr Cys Asn Val Asp His Lys Pro  
 195 200 205  
 Ser Asn Thr Lys Val Asp Lys Thr Val Glu Arg Lys Cys Cys Val Glu

210		215		220
Cys Pro Pro Cys Pro	Ala Pro Pro Val Ala Gly Pro Ser Val Phe Leu			
225	230		235	240
Phe Pro Pro Lys Pro	Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu			
	245		250	255
Val Thr Cys Val Val	Val Asp Val Ser His Glu Asp Pro Glu Val Gln			
	260		265	270
Phe Asn Trp Tyr Val	Asp Gly Val Glu Val His Asn Ala Lys Thr Lys			
	275		280	285
Pro Arg Glu Glu Gln	Phe Asn Ser Thr Phe Arg Val Val Ser Val Leu			
	290		295	300
Thr Val Val His Gln	Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys			
	310		315	320
Val Ser Asn Lys Gly	Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys			
	325		330	335
Thr Lys Gly Gln Pro	Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser			
	340		345	350
Arg Glu Glu Met Thr	Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys			
	355		360	365
Gly Phe Tyr Pro Ser	Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln			
	370		375	380
Pro Glu Asn Asn Tyr	Lys Thr Thr Pro Pro Met Leu Asp Ser Asp Gly			
	385		390	400
Ser Phe Phe Leu Tyr	Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln			
	405		410	415
Gln Gly Asn Val Phe	Ser Cys Ser Val Met His Glu Ala Leu His Asn			
	420		425	430
His Tyr Thr Gln Lys	Ser Leu Ser Leu Ser Pro Gly			
	435		440	

&lt;210&gt; 239

&lt;211&gt; 219

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; antibody constituent

&lt;400&gt; 239

Asp Ile Gln Met Thr	Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1	5 10 15
Asp Arg Val Thr Ile	Thr Cys Arg Ser Ser Gln Asn Ile Val His Ser
	20 25 30
Asn Gly Asn Thr Tyr	Leu Asp Trp Tyr Gln Gln Thr Pro Gly Lys Ala
	35 40 45
Pro Lys Leu Leu Ile	Tyr Lys Val Ser Asn Arg Phe Ser Gly Val Pro
	50 55 60
Ser Arg Phe Ser Gly	Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile
	65 70 75 80
Ser Ser Leu Gln Pro	Glu Asp Ile Ala Thr Tyr Tyr Cys Phe Gln Tyr
	85 90 95
Ser His Val Pro Trp	Thr Phe Gly Gln Gly Thr Lys Leu Gln Ile Thr
	100 105 110
Arg Thr Val Ala Ala	Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
	115 120 125
Gln Leu Lys Ser Gly	Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
	130 135 140
Tyr Pro Arg Glu Ala	Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
	145 150 155 160
Ser Gly Asn Ser Gln	Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
	165 170 175
Thr Tyr Ser Leu Ser	Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
	180 185 190
Lys His Lys Val Tyr	Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
	195 200 205
Pro Val Thr Lys Ser	Phe Asn Arg Gly Glu Cys
	210 215

&lt;210&gt; 240

&lt;211&gt; 453

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; antibody constituent

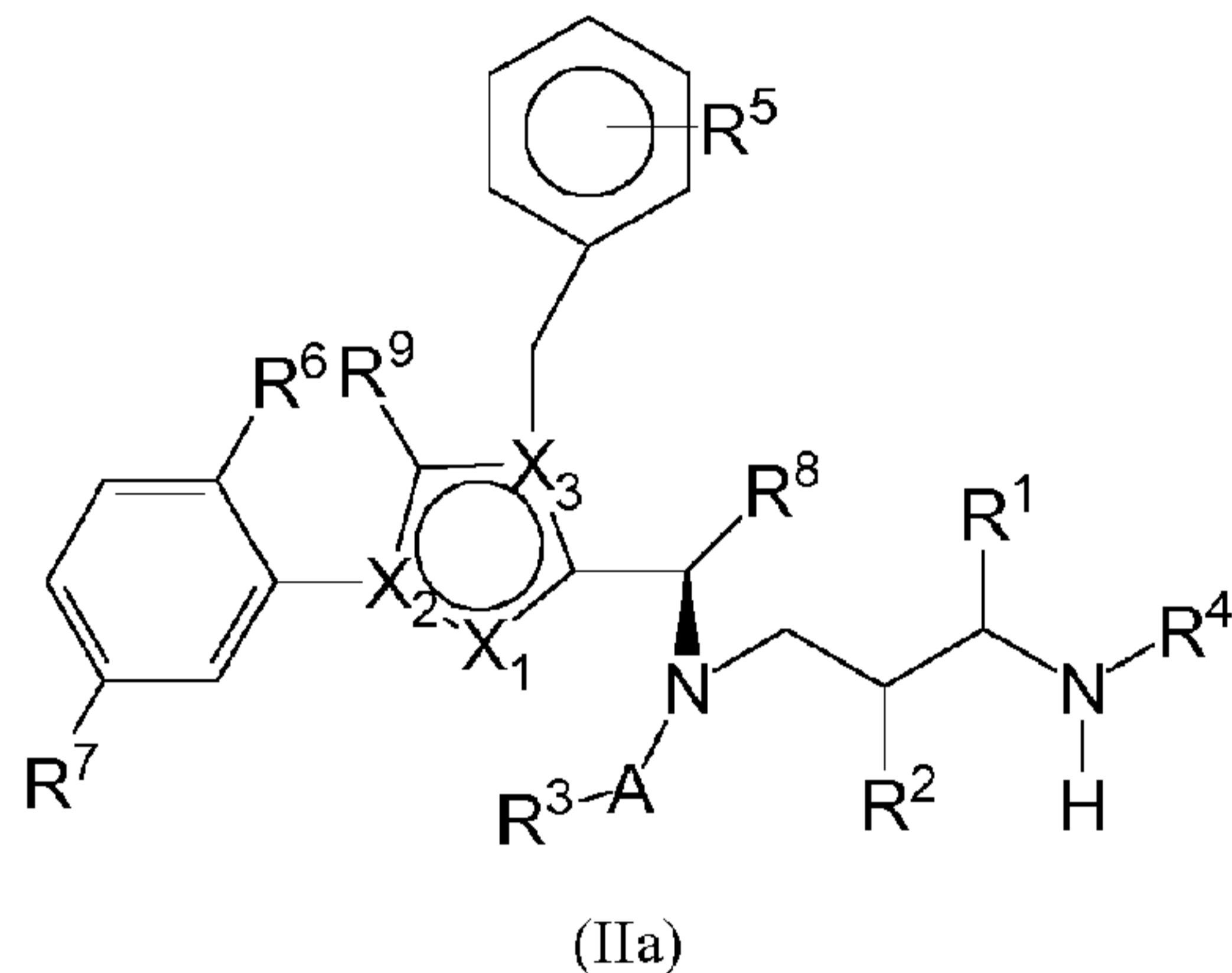
&lt;400&gt; 240

Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser  
 1 5 10 15  
 Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asn Tyr  
 20 25 30  
 Tyr Ile Tyr Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Ile  
 35 40 45  
 Gly Gly Ile Asn Pro Thr Ser Gly Gly Ser Asn Phe Asn Glu Lys Phe  
 50 55 60  
 Lys Thr Arg Val Thr Ile Thr Ala Asp Glu Ser Ser Thr Thr Ala Tyr  
 65 70 75 80  
 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Phe Tyr Phe Cys  
 85 90 95  
 Thr Arg Gln Gly Leu Trp Phe Asp Ser Asp Gly Arg Gly Phe Asp Phe  
 100 105 110  
 Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys Gly  
 115 120 125  
 Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys Ser Thr Ser Gly Gly  
 130 135 140  
 Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val  
 145 150 155 160  
 Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe  
 165 170 175  
 Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val  
 180 185 190  
 Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val  
 195 200 205  
 Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Lys Val Glu Pro Lys  
 210 215 220  
 Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu  
 225 230 235 240  
 Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr  
 245 250 255  
 Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val  
 260 265 270  
 Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val  
 275 280 285  
 Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser  
 290 295 300  
 Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu  
 305 310 315 320  
 Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala  
 325 330 335  
 Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro  
 340 345 350  
 Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln  
 355 360 365  
 Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala  
 370 375 380  
 Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr  
 385 390 395 400  
 Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu  
 405 410 415  
 Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser  
 420 425 430  
 Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser  
 435 440 445  
 Leu Ser Pro Gly Lys  
 450

## Patentkrav

1. Konjugat af en binder eller et derivat heraf med et eller flere aktivstofmolekyler,

- 5 idet aktivstofmolekylet er en kinesin spindel proteininhibitor, som er forbundet med binderen via en linker L, idet kinesin spindel proteininhibitoren har følgende formel (IIa):

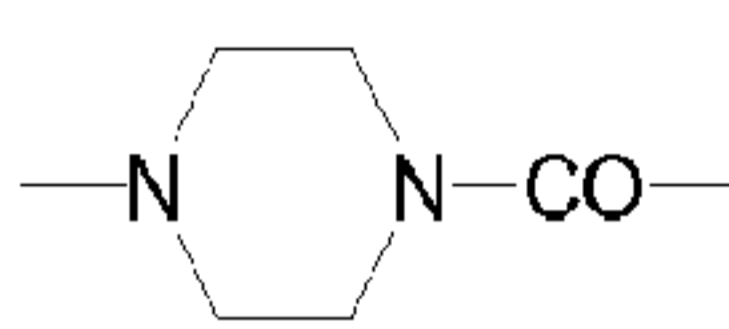
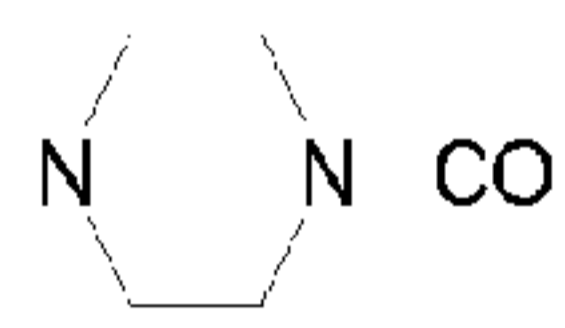


idet

- 10  $X_1$  N,  $X_2$  N og  $X_3$  er C; eller  
 $X_1$  er CH eller CF,  $X_2$ , C og  $X_3$  er N; eller  
 $X_1$  er NH,  $X_2$  C og  $X_3$  er C; eller  
 $X_1$  er CH,  $X_2$  N, og  $X_3$  er C;  
 (idet  $X_1$  er CH,  $X_2$  C, og  $X_3$  er N er foretrukket);
- 15  $R^1$  er H, -L-#1, -MOD eller  $-(CH_2)_{0-3}$  er Z, idet Z er -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup>, eller -CO-OY<sup>3</sup>,  
 idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden er H, NH<sub>2</sub>,  $-(CH_2CH_2O)_{0-3}$ -  
 $(CH_2)_{0-3}Z'$  (f.eks.  $-(CH_2)_{0-3}Z'$ ),  
 eller -CH(CH<sub>2</sub>W)Z', og Y<sup>3</sup> er H eller  $-(CH_2)_{0-3}$  er Z', idet Z' er H,  
 20 NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH eller  $-(CO-NH-CHY^4)_{1-3}$ COOH; idet W er H eller OH,  
 idet Y<sup>4</sup> uafhængigt af hinanden er lineært eller forgrenet, i givet fald med -NHCONH<sub>2</sub> substitueret C<sub>1-6</sub>-alkyl eller i givet fald med -NH<sub>2</sub> substitueret aryl eller benzyl;
- 25  $R^2$  er -L-#1, H, -MOD, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> eller  $-(CH_2)_{0-3}Z$ , eller  $R^2$  og  $R^4$  i fællesskab (under dannelse af en pyrrolidinring) er -CH<sub>2</sub>-CHR<sup>10</sup>- eller -CHR<sup>10</sup>-CH<sub>2</sub>-, idet R<sup>10</sup> er L-#1, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH, eller OH;  
 idet Z er -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup>, eller -CO-
- 30 OY<sup>3</sup>,  
 idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden er H, NH<sub>2</sub>, eller  $-(CH_2)_{0-3}Z'$ ,



og  $Y^3$  er H eller  $-(CH_2)_{0-3}Z'$ , idet  $Z'$  er H,  $SO_3H$ ,  $NH_2$  eller  $COOH$ ;  
 idet  $Y^4$  uafhængigt af hinanden er lineært eller forgrenet, i  
 givet fald med  $-NHCONH_2$  substitueret  $C_{1-6}$  alkyl eller i givet  
 fald med  $-NH_2$  substitueret aryl eller benzyl, og  $Y^5$  er H eller -  
 5  $CO-CHY^6-NH_2$ , idet  $Y^6$  lineært eller forgrenet  $C_{1-6}$  alkyl;  
 $R^4$  er  $-L-#1$ , H,  $-CO-CHY^4-NHY^5$  eller  $-(CH_2)_{0-3}Z$ ,  
 idet  $Z$  er  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $NHY^3$ ,  $-CO-NY^1Y^2$ , eller  $-CO-$   
 $OY^3$ ,  
 idet  $Y^1$  og  $Y^2$  uafhængigt af hinanden er H,  $NH_2$ , eller  $-(CH_2)_{0-3}Z'$ ,  
 10 og  $Y^3$  er H eller  $-(CH_2)_{0-3}Z'$ , idet  $Z'$  er H,  $SO_3H$ ,  $NH_2$  eller  $COOH$ ;  
 idet  $Y^4$  uafhængigt af hinanden er lineært eller forgrenet, i  
 givet fald med  $-NHCONH_2$  substitueret  $C_{1-6}$  alkyl eller i givet  
 fald med  $-NH_2$  substitueret aryl eller benzyl, og  $Y^5$  er H eller -  
 $CO-CHY^6-NH_2$ , idet  $Y^6$  er lineært eller forgrenet  $C_{1-6}$  alkyl;  
 15 eller  $R^2$  og  $R^4$  i fællesskab (under dannelse af en pyrrolidinring)  
 er  $-CH_2-CHR^{10}-$  eller  $-CHR^{10}-CH_2-$ , idet  $R^{10}$  er  $L-#1$ , H,  $NH_2$ ,  $SO_3H$ ,  
 $COOH$ , SH eller OH;  
 $A$  er CO, SO,  $SO_2$ ,  $SO_2NH$  eller CNNH;  
 $R^3$  er  $-L-#1$ ,  $-MOD$  eller en i givet fald substitueret alkyl-,  
 20 cycloalkyl-, aryl-, heteroaryl-heteroalkyl-,  
 heterocycloalkylgruppe, fortrinsvis  $-L-#1$  eller en  $C_{1-10}$ -alkyl-,  
 $C_{6-10}$ -aryl- eller  $C_{6-10}$ -aralkyl-,  $C_{5-10}$ -heteroalkyl-,  $C_{1-10}$ -alkyl-O-  
 $C_{6-10}$ -aryl- eller  $C_{5-10}$ -heterocycloalkylgruppe, som kan være  
 substitueret med 1-3  $-OH$ -grupper, 1-3 halogenatomer, 1-3  
 25 halogenerede alkylgrupper (som hver især har 1-3 halogenatomer),  
 1-3 O-alkylgrupper, 1-3  $-SH$ -grupper, 1-3  $-S$ -alkylgrupper, 1-3  $-$   
 $O-CO$ -alkylgrupper, 1-3  $-O-CO-NH$ -alkylgrupper, 1-3  $-NH-CO-$   
 alkylgrupper, 1-3  $-NH-CO-NH$ -alkylgrupper, 1-3  $-S(O)_n-$   
 alkylgrupper, 1-3  $-SO_2-NH$ -alkylgrupper, 1-3  $-NH$ -alkylgrupper, 1-  
 30 3  $-N(alkyl)_2$ -grupper, 1-3  $-NH_2$ -grupper eller 1-3  $-(CH_2)_{0-3}Z-$   
 grupper, idet  $Z$  er  $-H$ , halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$   
 eller  $-CO-OY^3$ , idet  $Y^1$  og  $Y^2$  uafhængigt af hinanden er H,  $NH_2$  eller  
 $-(CH_2)_{0-3}Z'$ , og  $Y^3$  er H,  $-(CH_2)_{0-3}-CH(NHCOCH_3)Z'$ ,  $-(CH_2)_{0-3}-$   
 $CH(NH_2)Z'$ , eller  $-(CH_2)_{0-3}Z'$ , idet  $Z'$  er H,  $SO_3H$ ,  $NH_2$  eller  $COOH$ ,  
 35 (idet "alkyl" fortrinsvis betegner  $C_{1-10}$ -alkyl);  
 $R^5$  er  $-L-#1$ , H,  $NH_2$ ,  $NO_2$ , halogen (navnlig F, Cl, Br),  $-CN$ ,  $CF_3$ ,  
 $-OCF_3$ ,  $-CH_2F$ ,  $-CH_2F$ , SH eller  $-(CH_2)_{0-3}Z$ , idet  $Z$  er  $-H$ ,  $-OY^3$ ,  $-$   
 $SY^3$ , halogen,  $NHY^3$ ,  $-CO-NY^1Y^2$  eller  $-CO-OY^3$ ,

idet  $Y^1$  og  $Y^2$  uafhængigt af hinanden er H,  $NH_2$ , eller  $-(CH_2)_{0-3}Z'$ ,  
og  $Y^3$  er H eller  $-(CH_2)_{0-3}Z'$ , idet  $Z'$  er H,  $SO_3H$ ,  $NH_2$  eller  $COOH$ ;  
 $R^6$  og  $R^7$  uafhængigt af hinanden er H, cyano, (i givet fald  
fluoreret)  $C_{1-10}$ -alkyl, (i givet fald fluoreret)  $C_{2-10}$ -alkenyl, (i  
5 givet fald fluoreret)  $C_{2-10}$ -alkynyl, hydroxy,  $NO_2$ ,  $NH_2$ ,  $COOH$  eller  
halogen (navnlig F, Cl, Br),  
 $R^8$  er (i givet fald fluoreret)  $C_{1-10}$ -alkyl, (i givet fald  
fluoreret)  $C_{2-10}$ -alkenyl, (i givet fald fluoreret)  $C_{2-10}$ -alkynyl,  
(i givet fald fluoreret)  $C_{4-10}$ -cycloalkyl eller  $-(CH_2)_{0-2}-(HZ^2)$ ,  
10 idet  $HZ^2$  er en 4- til 7-leddet heterocyclus med op til to  
heteroatomer udvalgt af N, O og S, idet hver af disse grupper  
kan være substitueret med  $-OH$ ,  $CO_2H$  or  $NH_2$  eller  $-L-#1$ ;  
idet en eller ingen af substituenterne  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$ ,  $R^5$ ,  $R^8$  og  
 $R^{10}$  er eller (i tilfælde af  $R^8$ ) har  $-L-#1$ , eller  
15 L står for linkerens, og #1 står for bindingen til binderen eller  
derivatet heraf,  
 $R^9$  er H, F,  $CH_3$ ,  $CF_3$ ,  $CH_2F$  eller  $CHF_2$ ;  
idet  $-MOD$  er  $-(NR^{10})_n-(G1)_o-G2-H$ , idet  
 $R^{10}$  er H eller  $C_1-C_3$ -alkyl;  
20 G1 er  $-NHCO-$ ,  $-CONH-$  eller  

  
(idet, hvis G1 er  $-NHCO-$  eller  

  
 $R^{10}$  ikke er  $NH_2$ );  
25 n er 0 eller 1;  
o er 0 eller 1; og  
G2 er en ligekædet og/eller forgrenet carbonhydridgruppe med 1  
til 10 carbonatomer, som kan være afbrudt en gang eller flere  
gange med en eller flere af grupperne  $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NRY-$   
30 ,  $-NRYCO-$ ,  $CONRY-$ ,  $-NRYNRY-$ ,  $-SO_2NR^yNR^y-$ ,  $-CONR^yNR^y-$ , (idet  $R^y$  er  
H, phenyl,  $C_1-C_{10}$ -alkyl,  $C_2-C_{10}$ -alkenyl eller  $C_2-C_{10}$ -alkynyl, som  
hver især kan være substitueret med  $NHCONH_2$ ,  $-COOH$ ,  $-OH$ ,  $-NH_2$ ,  
 $NH-CNNH_2$ , sulfonamid, sulfon, sulfoxid, eller sulfonsyre),  $-CO-$   
,  $-CR^x=N-O-$ , idet  $R^x$  er H,  $C_1-C_3$ -alkyl eller phenyl), idet  
35 carbonhydridkæden inklusiv sidekæderne, såfremt sådanne  
eksisterer, kan være substitueret med  $-NHCONH_2$ ,  $-COOH$ ,  $-OH$ ,  $-$   
 $NH_2$ ,  $NH-CNNH_2$ , sulfonamid, sulfon, sulfoxid, eller sulfonsyre,

idet gruppen -MOD fortrinsvis har i det mindste en gruppe -COOH; samt deres salte, solvater samt salte af solvaterne.

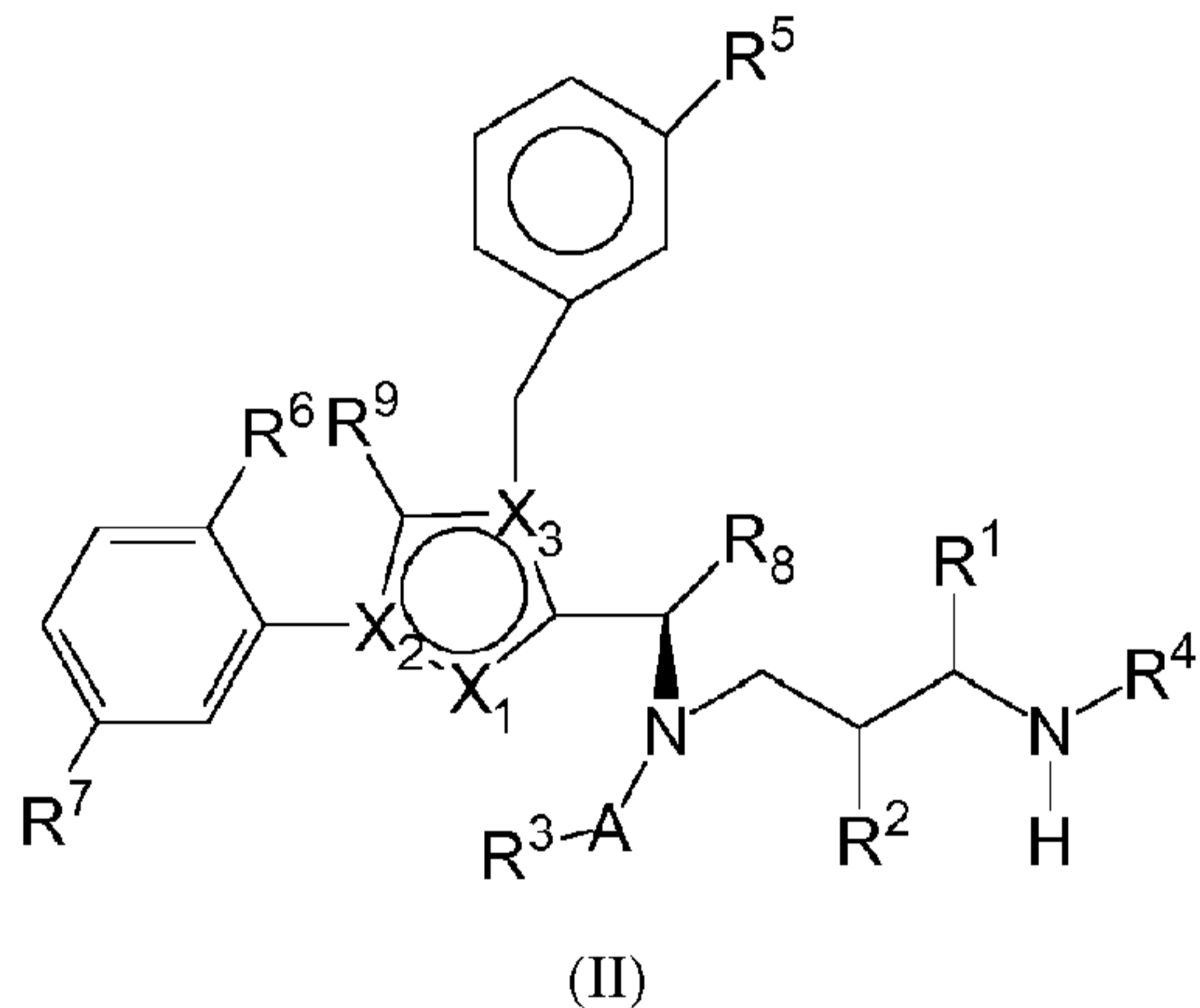
2. Konjugat ifølge krav 1, idet  $X_1$  er CH,  $X_2$  C, og  $X_3$  er N.

5

3. Konjugat ifølge krav 1 eller 2, idet substituenten  $R^1$  er -L-#1.

4. Konjugat ifølge et eller flere af de foregående krav, idet aktivstofmolekyle-linkeren vises i den almene formel (II):

10



idet

$X_1$  er N,  $X_2$  N, og  $X_3$  er C; eller

$X_1$  er CH,  $X_2$  C, og  $X_3$  er N; eller

15  $X_1$  er NH,  $X_2$  C, og  $X_3$  er C; eller

$X_1$  er CH,  $X_2$  N, og  $X_3$  er C;

$R^1$  er H, -L-#1 eller  $-(CH_2)_{0-3}Z$ , idet Z er -H, -NH $Y^3$ , -O $Y^3$ , -S $Y^3$ , halogen, -CO-N $Y^1Y^2$ , eller -CO-O $Y^3$ ,

idet  $Y^1$  og  $Y^2$  uafhængigt af hinanden er H, NH $_2$ ,  $-(CH_2CH_2O)_{0-3}$ -

20  $(CH_2)_{0-3}Z'$  eller -CH(CH $_2W$ )Z', og  $Y^3$  er H eller  $-(CH_2)_{0-3}Z'$ , idet Z'

er H, NH $_2$ , SO $_3H$ , COOH, -NH-CO-CH $_2$ -CH $_2$ -CH(NH $_2$ )COOH eller -(CO-NH-

CH $Y^4$ ) $_{1-3}$ COOH, idet W er H eller OH; idet  $Y^4$  uafhængigt af hinanden

er lineært eller forgrenet, i givet fald med -NHCONH $_2$

substitueret C $_{1-6}$  alkyl eller i givet fald med -NH $_2$  substitueret

25 aryl eller benzyl;

$R^2$  og  $R^4$  uafhængigt af hinanden er -L-#1, H, -CO-CH $Y^4$ -NH $Y^5$  eller

$-(CH_2)_{0-3}Z$ , eller  $R^2$  og  $R^4$  i fællesskab (under dannelsen af en

pyrrolidinring) er -CH $_2$ -CH $R^{10}$ - eller -CH $R^{10}$ -CH $_2$ -, idet  $R^{10}$  er L-

#1, H, NH $_2$ , SO $_3H$ , COOH, SH eller OH;

30 idet Z er -H, O $Y^3$ , -S $Y^3$ , halogen, NH $Y^3$ , -CO-N $Y^1Y^2$  eller -CO-O $Y^3$ ;

idet  $Y^1$  og  $Y^2$  uafhængigt af hinanden er H, NH $_2$ , eller  $-(CH_2)_{0-3}Z'$ ,

og  $Y^3$  er H eller  $-(CH_2)_{0-3}Z'$ , idet  $Z'$  er H,  $SO_3H$ ,  $NH_2$  eller  $COOH$ ;  
idet  $Y^4$  uafhængigt af hinanden er lineært eller forgrenet, i  
givet fald med  $-NHCONH_2$  substitueret  $C_{1-6}$  alkyl eller i givet  
fald med  $-NH_2$  substitueret aryl eller benzyl, og  $Y^5$  er H eller -  
5  $CO-CHY^6-NH_2$ , idet  $Y^6$  er lineært eller forgrenet  $C_{1-6}$  alkyl;  
A er CO, SO,  $SO_2$ ,  $SO_2NH$  eller CNNH;  
 $R^3$  er -L-#1 eller en i givet fald substitueret alkyl-, aryl-,  
heteroaryl- heteroalkyl-, heterocycloalkylgruppe,  
fortrinsvis -L-#1 eller en  $C_{1-10}$ -alkyl-,  $C_{6-10}$ -aryl-,  $C_{6-10}$ -aralkyl-  
10 ,  $C_{5-10}$ -heteroalkyl-,  $C_{1-10}$ -alkyl-O- $C_{6-10}$ -aryl- eller  $C_{5-10}$ -  
heterocycloalkylgruppe,  
som kan være substitueret med 1-3 -OH-grupper, 1-3  
halogenatomer, 1-3 halogenerede alkylgrupper (som hver især kan  
have 1-3 halogenatomer), 1-3 O-alkylgrupper, 1-3 -SH-grupper,  
15 1-3 -S-alkylgrupper, 1-3 -O-CO-alkylgrupper, 1-3 -O-CO-NH-  
alkylgrupper, 1-3 -NH-CO-alkylgrupper, 1-3 -NH-CO-NH-  
alkylgrupper, 1-3 -S(O)<sub>n</sub>-alkylgrupper, 1-3 -SO<sub>2</sub>-NH-alkylgrupper,  
1-3 -NH-alkylgrupper, 1-3 -N(alkyl)<sub>2</sub>-grupper, 1-3 -NH<sub>2</sub>-grupper  
eller 1-3-(CH<sub>2</sub>)<sub>0-3</sub>Z-grupper, idet Z er -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -  
20 NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> eller -CO-OY<sup>3</sup>, idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden  
er H, NH<sub>2</sub> eller  $-(CH_2)_{0-3}Z'$ , og Y<sup>3</sup> er H,  $-(CH_2)_{0-3}-CH(NHCOCH_3)Z\%$ -  
 $(CH_2)_{0-3}-CH(NH_2)Z'$  eller  $-(CH_2)_{0-3}Z'$ , idet Z' er H,  $SO_3H$ ,  $NH_2$  eller  
COOH,  
(idet "alkyl" fortrinsvis betegner  $C_{1-10}$ -alkyl);  
25  $R^5$  er -L-#1, H, F,  $NH_2$ ,  $NO_2$ , halogen, SH eller  $-(CH_2)_{0-3}Z$ , idet Z  
er -H, OY<sup>3</sup>, -SY<sup>3</sup>, halogen, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> eller -CO-OY<sup>3</sup>,  
idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden er H,  $NH_2$ , eller  $-(CH_2)_{0-3}Z'$ ,  
og Y<sup>3</sup> er H eller  $-(CH_2)_{0-3}Z'$ , idet Z' er H,  $SO_3H$ ,  $NH_2$  eller  $COOH$ ;  
idet en af substituenterne  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$  og  $R^5$  er -L-#1,  
30 L står for linkeren, og #1 står for bindingen til binderen eller  
derivatet heraf,  
 $R^6$  og  $R^7$  uafhængigt af hinanden er H, cyano, (i givet fald  
fluoreret)  $C_{1-10}$ -alkyl, (i givet fald fluoreret)  $C_{2-10}$ -alkenyl, (i  
givet fald fluoreret)  $C_{2-10}$ -alkynyl, hydroxy eller halogen,  
35  $R^8$  er (i givet fald fluoreret)  $C_{1-10}$ -alkyl, (i givet fald  
fluoreret)  $C_{4-10}$ -cycloalkyl eller optionalt substitueret oxetan;  
og  
 $R^9$  er H, F,  $CH_3$ ,  $CF_3$ ,  $CH_2F$  eller  $CHF_2$ ;

samt deres salte, solvater samt salte af solvaterne.

5. Konjugat ifølge krav 4,  
 idet  $R^1$  er  $-L-#1$ , H,  $-(CH_2)_{0-3}Z$ , idet Z er  $-H$ ,  $-CO-NY^1Y^2$ ,  $NHY^3$ ,  
 5  $OY^3$ ,  $-SY^3$  eller  $-CO-OY^3$ ,  
 idet  $Y^1$  og  $Y^2$  uafhængigt af hinanden er H,  $NH_2$ ,  $-(CH_2CH_2O)_{0-3}-$   
 $(CH_2)_{0-3}Z'$  eller  $-CH(CH_2W)Z'$ , og  $Y^3$  er H eller  $-(CH_2)_{0-3}Z'$ , idet  $Z'$   
 er H,  $NH_2$ ,  $COOH$ ,  $-NH-CO-CH_2-CH_2-CH(NH_2)COOH$  eller  $-(CO-NH-CHY^4)_{1-}$   
 10  $_3COOH$ , idet W er H eller OH, idet  $Y^4$  uafhængigt af hinanden er  
 lineært eller forgrenet, i givet fald med  $-NHCONH_2$  substitueret  
 $C_{1-6}$  alkyl eller i givet fald med  $-NH_2$  substitueret aryl eller  
 benzyl;  
 $R^2$  og  $R^4$  uafhængigt af hinanden er  $-L-#1$ ,  $-CO-CHY^4-NHY^5$  eller H,  
 eller  $R^2$  og  $R^4$  i fællesskab (under dannelse af en pyrrolidinring)  
 15 er  $-CH_2-CHR^{10}-$ , idet  $R^{10}$  er H,  $-L-#1$ ,  $NH_2$ ,  $COOH$ , SH, OH eller  $SO_3H$ ;  
 idet  $Y^4$  uafhængigt af hinanden er lineært eller forgrenet, i  
 givet fald med  $-NHCONH_2$  substitueret  $C_{1-6}$  alkyl eller i givet  
 fald med  $-NH_2$  substitueret aryl eller benzyl, og  $Y^5$  er H eller  $-$   
 $CO-CHY^6-NH_2$ , idet  $Y^6$  er lineært eller forgrenet  $C_{1-6}$  alkyl;  
 20 A er CO,  
 $R^3$  er  $-(CH_2)OH$  eller  $-L-#1$ , og  
 $R^5$  er  $-L-#1$  eller H,  
 idet en af substituenterne  $R^1$ ,  $R^2$ ,  $R^3$ ,  $R^4$  og  $R^5$  er  $-L-#1$ .
- 25 6. Konjugat ifølge krav 4 eller 5, idet  $R^6$  og  $R^7$  uafhængigt af  
 hinanden er H,  $C_{1-3}$ -alkyl eller halogen.
7. Konjugat ifølge et eller flere af kravene 4 til 6, idet  $R^8$   
 er  $C_{1-4}$ -alkyl (fortrinsvis tert-butyl).
- 30 8. Konjugat ifølge et eller flere af kravene 4 til 7, idet  $R^9$   
 er H.
9. Konjugat ifølge et eller flere af kravene 4 til 8, idet  $R^6$   
 35 og  $R^7$  er F.
10. Konjugat ifølge et eller flere af de foregående krav, idet  
 binderen eller derivatet heraf er et binderpeptid eller -protein

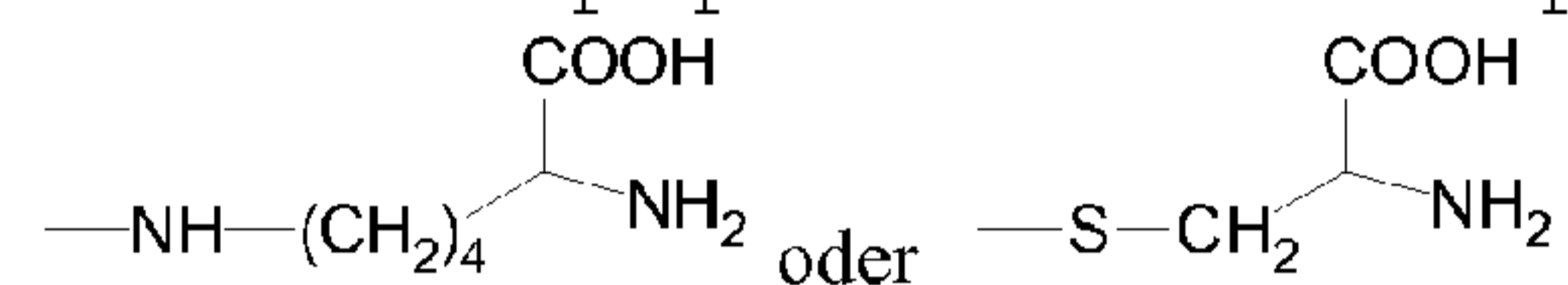
eller et derivat af et binderpeptid eller -protein.

11. Konjugat ifølge krav 10, idet hvert aktivstofmolekyle er bundet til forskellige aminosyrer af binderpeptidet eller -proteinet eller derivatet heraf via linkeren.

12. Konjugat ifølge et eller flere af de foregående krav, idet konjugatet i gennemsnit har 1,2 til 20 aktivstofmolekyler pr. binder.

10

13. Konjugat ifølge et eller flere af kravene 10 til 12, idet binderpeptidet eller -proteinet er et antistof eller derivatet af binderpeptidet eller -proteinet



15

14. Konjugat ifølge et eller flere af de foregående krav, idet binderen binder sig til et cancer-målmolekyle.

15. Konjugat ifølge krav 14, idet binderen binder sig til et ekstracellulært målmolekyle.

20

16. Konjugat ifølge krav 15, idet binderen efter binding til det ekstracellulære målmolekyle, hvorfra målmolekylet internaliserer eksprimerende celle og processeres intracellulært (fortrinsvis lysosomalt).

25

17. Konjugat ifølge et eller flere af kravene 10 til 16, idet binderpeptidet eller -proteinet er et humant, humaniseret eller kimært monoklonalt antistof eller et antigen-bindende fragment heraf.

30

18. Konjugat ifølge krav 17, idet binderpeptidet eller -proteinet er et anti-HER2-antistof, et anti-EGFR-antistof, et anti-TWEAKR-antistof eller et antigen-bindende fragment heraf.

35

19. Konjugat ifølge krav 18, idet anti-TWEAKR-antistoffet binder sig specifikt til aminosyren D i position 47 (D47) på TWEAKR (SEQ ID NO:169), fortrinsvis anti-TWEAKR-antistoffet TPP-

2090.

20. Konjugat ifølge krav 18, idet binderpeptidet eller -proteinet er et anti-EGFR-antistof, og R3 er -L-#1.

5

21. Konjugat ifølge et eller flere af de foregående krav, idet linkerens -L- har en af de følgende grundstrukturer (i) til (iv):

1.  $-(CO)_m-SG1-L1-L2-$

(ii)  $-(CO)_m-L1-SG-L1-L2-$

10 (iii)  $-(CO)_m-L1-L2-$

(iv)  $-(CO)_m-L1-SG-L2$

idet m er 0 eller 1, SG og SG1 er en in vivo spaltbar gruppe, L1 uafhængigt af hinanden står for in vivo ikke spaltbare organiske grupper, og L2 står for en koblingsgruppe til  
15 binderen.

22. Konjugat ifølge krav 21, idet den in vivo spaltbare gruppe SG er en 2-8 oligopeptidgruppe, fortrinsvis en dipeptidgruppe eller et disulfid, en hydrazon, en acetal eller en aminor, og  
20 SG1 er en 2-8 oligopeptidgruppe, fortrinsvis en dipeptidgruppe.

23. Konjugat ifølge et eller flere af de foregående krav, idet linkerens er bundet til en cystein-sidekæde eller cysteingrouppe og har den følgende formel:

25  $\S-(CO)_m-L1-L2-\S\S$

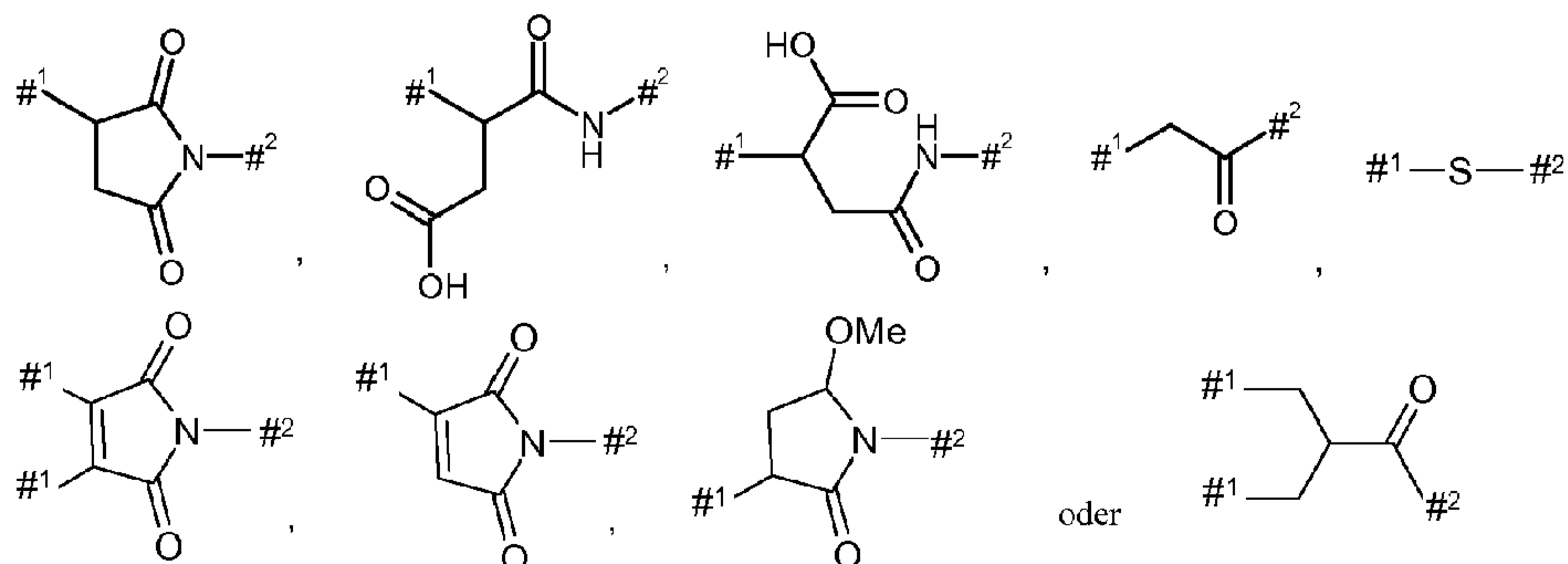
idet

m er 0 eller 1;

§ er bindingen til aktivstofmolekylet, og

§§ er bindingen til binderpeptidet eller -proteinet, og

30 -L2- står for



idet

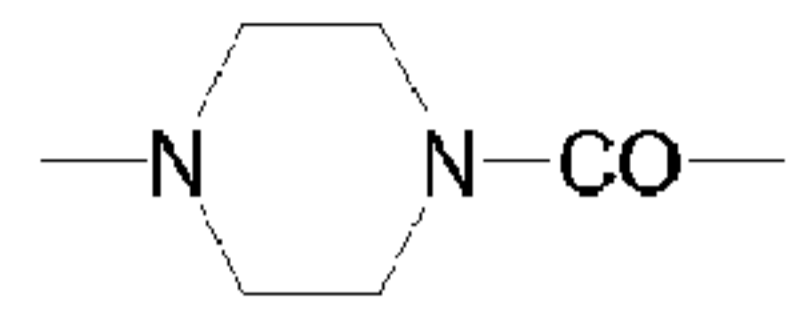
#<sup>1</sup> kendetegner bindingsstedet med svovlatomet i binderen,

#<sup>2</sup> kendetegner bindingsstedet med gruppen L<sup>1</sup>,

L<sup>1</sup> er  $-(NR^{10})_n-(G1)_o-G2-$ ,

idet R<sup>10</sup> er H, NH<sub>2</sub> eller C<sub>1</sub>-C<sub>3</sub>-;

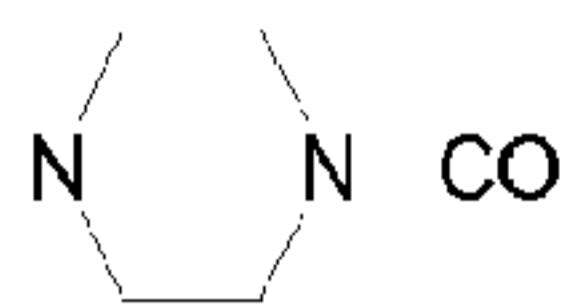
5 G<sub>1</sub> er -NHCO- eller



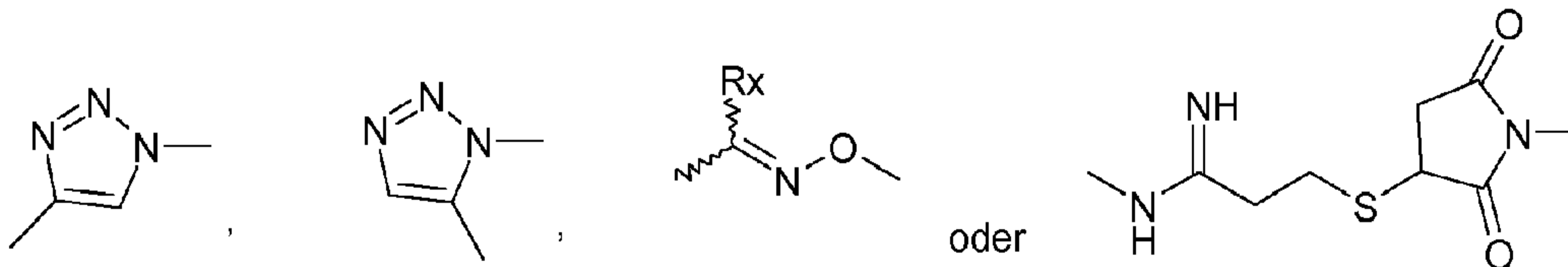
n er 0 eller 1;

o er 0 eller 1; og

G<sub>2</sub> er en ligekædet eller forgrenet carbonhydriddkæde med 1 til  
 10 100 carbonatomer fra arylengrupper og/eller ligekædede og/eller  
 forgrenede og/eller cykliske alkylengrupper, som kan være  
 afbrudt en eller flere gange af en eller flere af grupperne -O-  
 , -S-, , -SO-, SO<sub>2</sub>, -NH-, -CO-, -Nme-, -NHNH-, -SO<sub>2</sub>NHNH-, -NHCO-  
 , -CONH-, -CONHNH- og en 5- til 10-leddet, aromatisk eller ikke  
 15 aromatisk heterocyclus med op til 4 heteroatomer udvalgt af N,  
 O og S, -SO- eller -SO<sub>2</sub>- (fortrinsvis

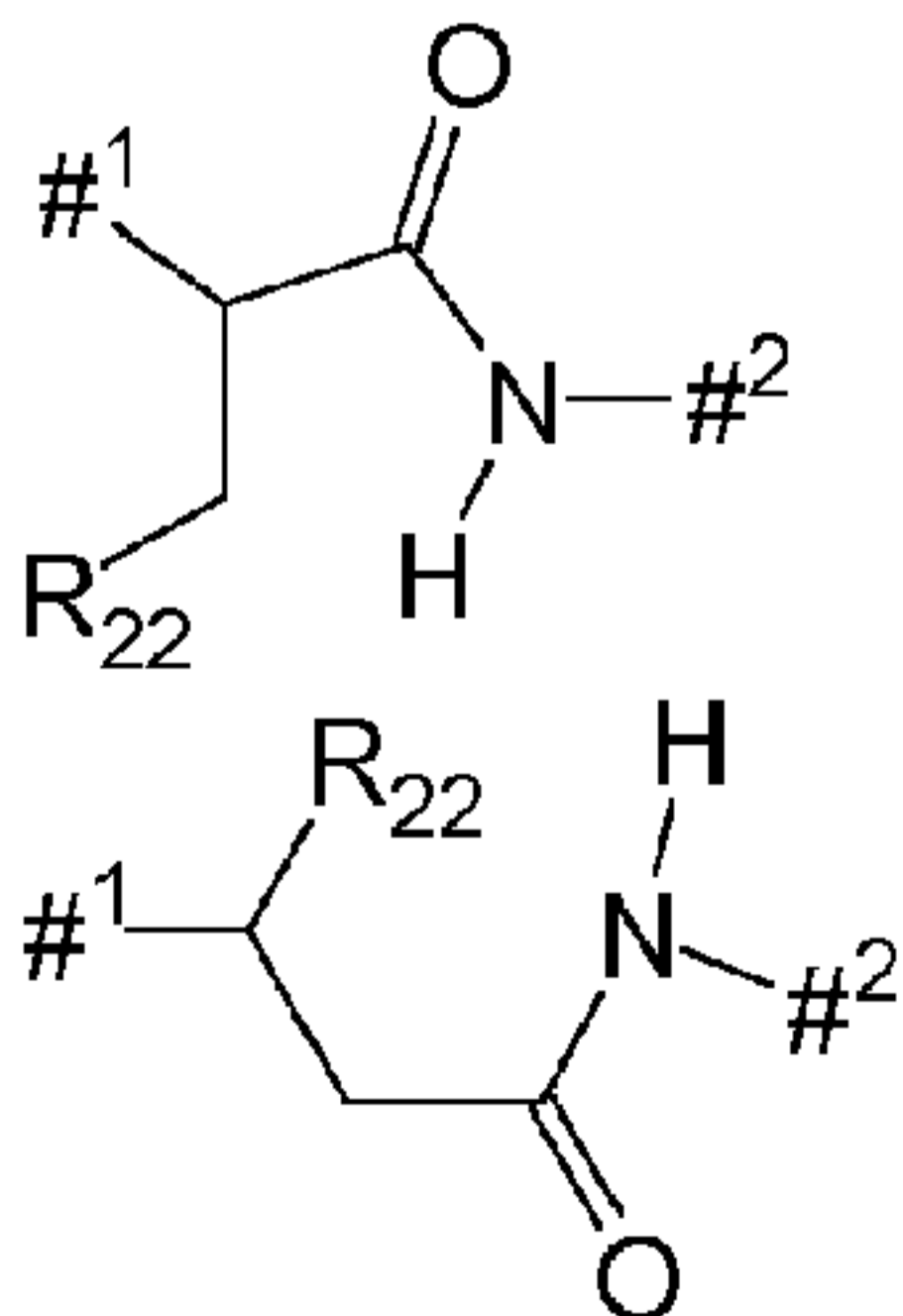


), idet sidekæderne, såfremt sådanne eksisterer, kan være  
 substitueret med -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>,  
 20 sulfonamid, sulfon, sulfoxid eller sulfonsyre,  
 eller er en af de følgende grupper:



idet Rx er H, C<sub>1</sub>-C<sub>3</sub>-alkyl eller phenyl.

25 24. Konjugat ifølge krav 23, idet L<sub>2</sub> vises ved en eller begge  
 af de følgende formler:

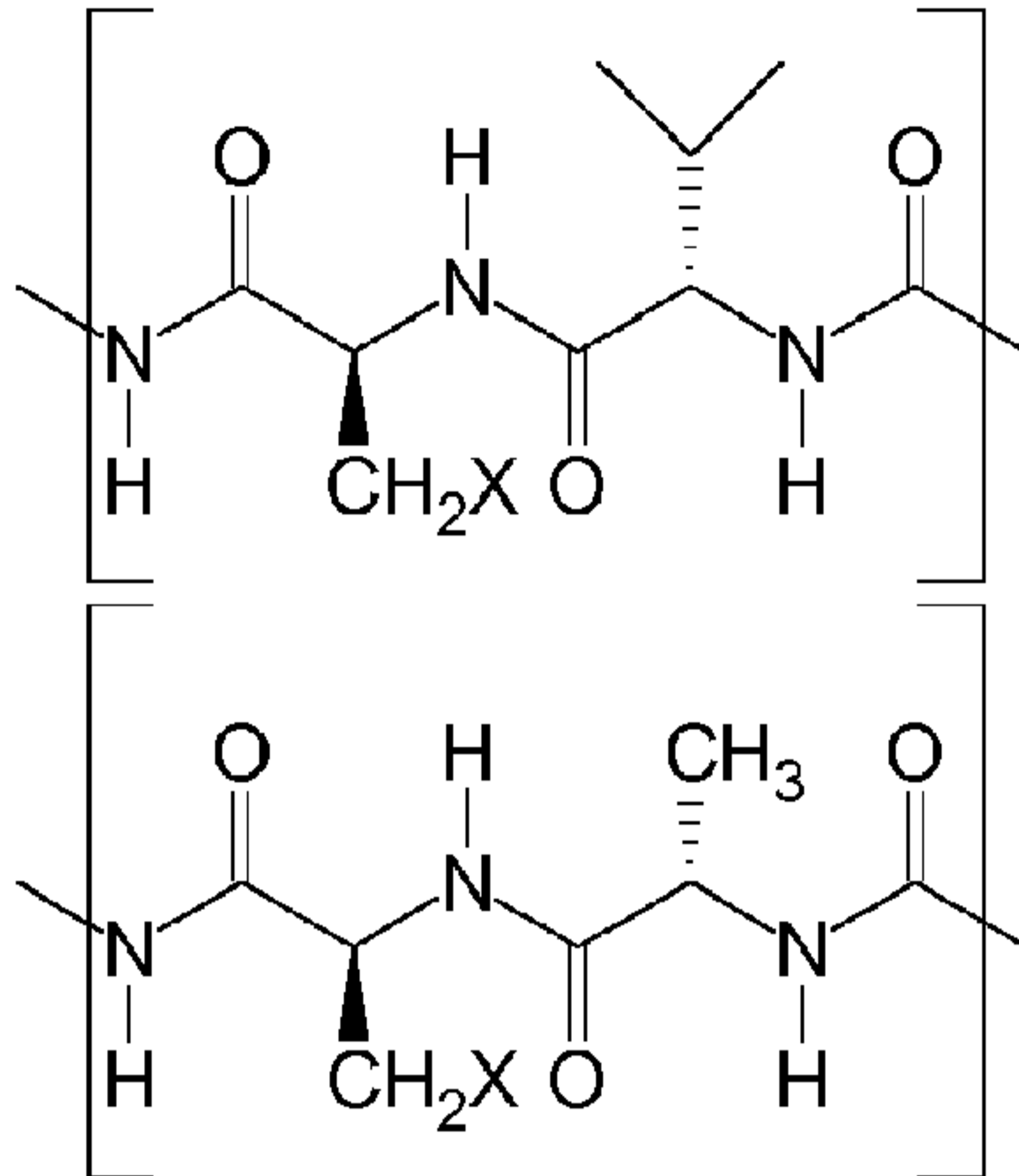


idet #<sup>1</sup> kendetegner bindingsstedet med svovlatomet i binderen,  
 #<sup>2</sup> kendetegner bindingsstedet med gruppen L<sup>1</sup>, R<sup>22</sup> er COOH, og



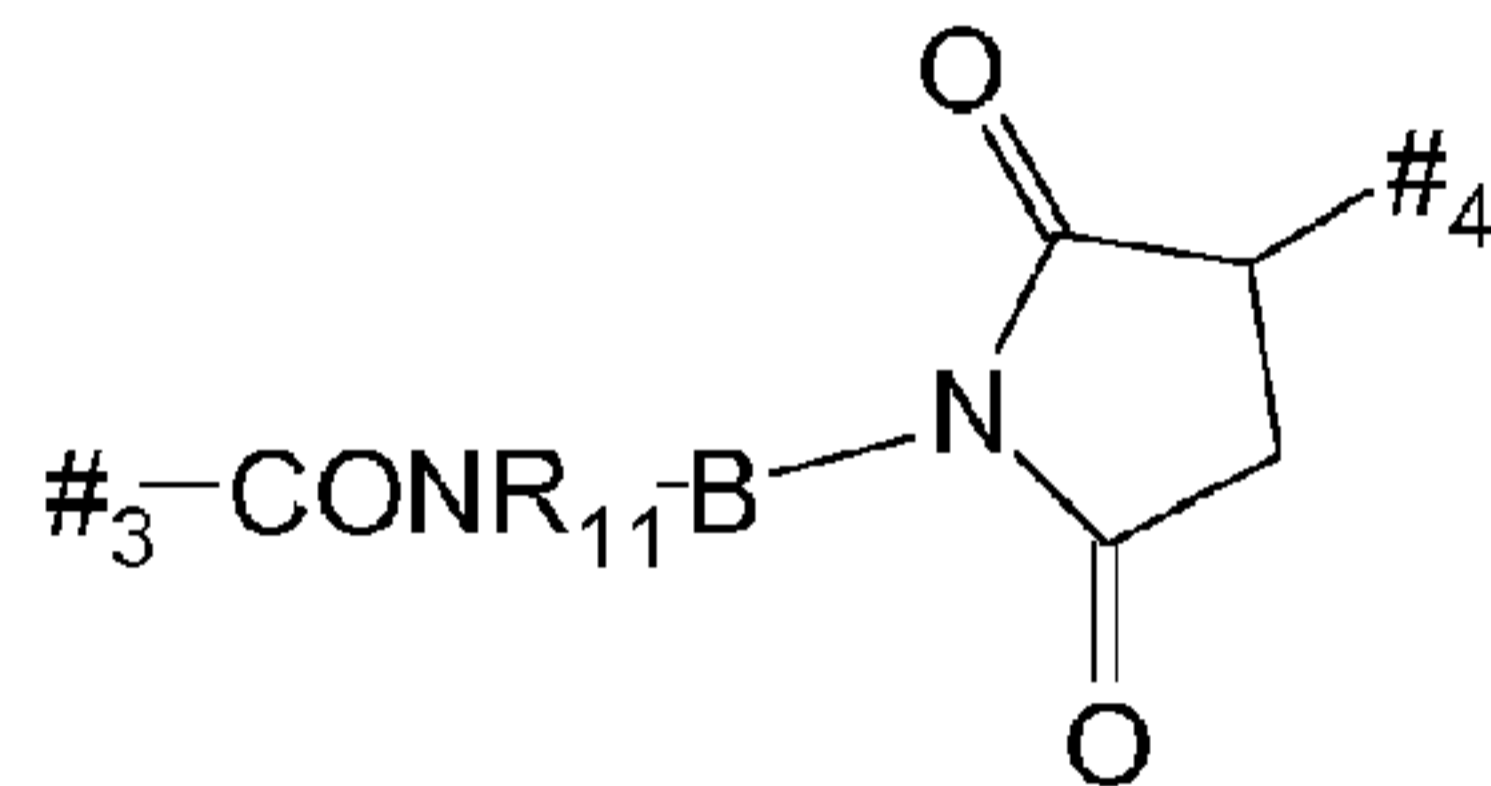
bindingerne til svovlatomet i binderen foreligger med over 80 % (baseret på det samlede antal bindinger i linkerens til binderen) i en af disse to strukturer.

- 5 25. Konjugat ifølge krav 23 eller 24, idet carbonhydriddkæden er afbrudt af en af de følgende grupper:



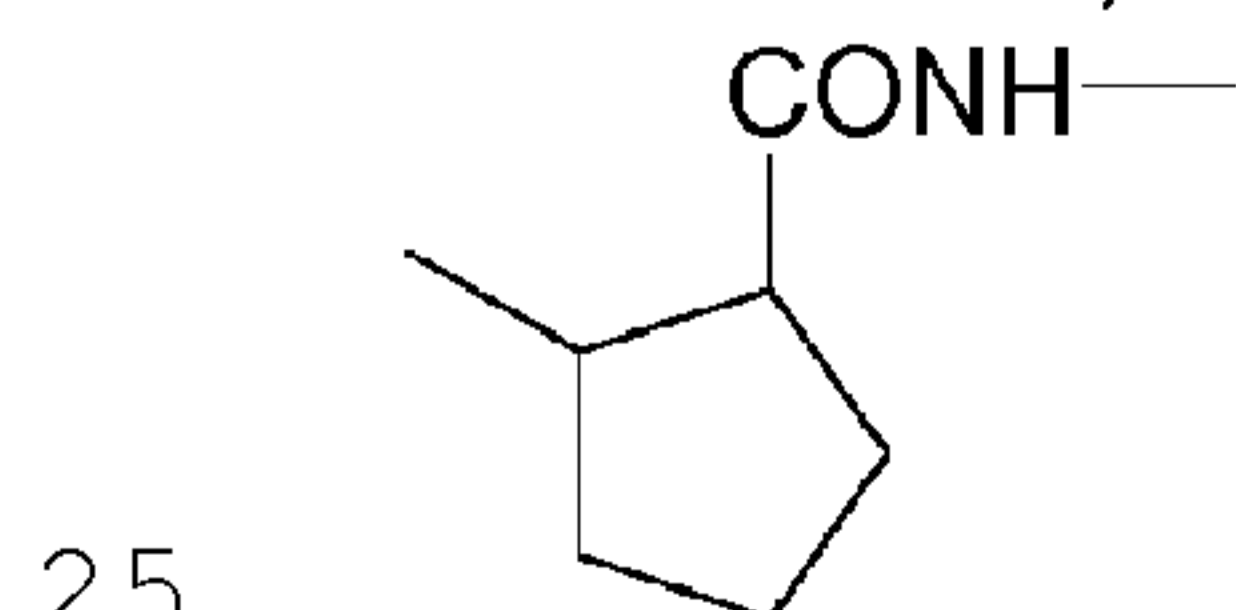
- 10 idet X er H, en C<sub>1-10</sub>-alkylgruppe, som i givet fald kan være substitueret med -NHCONH<sub>2</sub>, -COOH, -OH, NH<sub>2</sub>, -NH-CNNH<sub>2</sub>, sulfon, sulfoxid eller sulfonsyre.

26. Konjugat ifølge krav 23, idet linkerens har følgende formel:



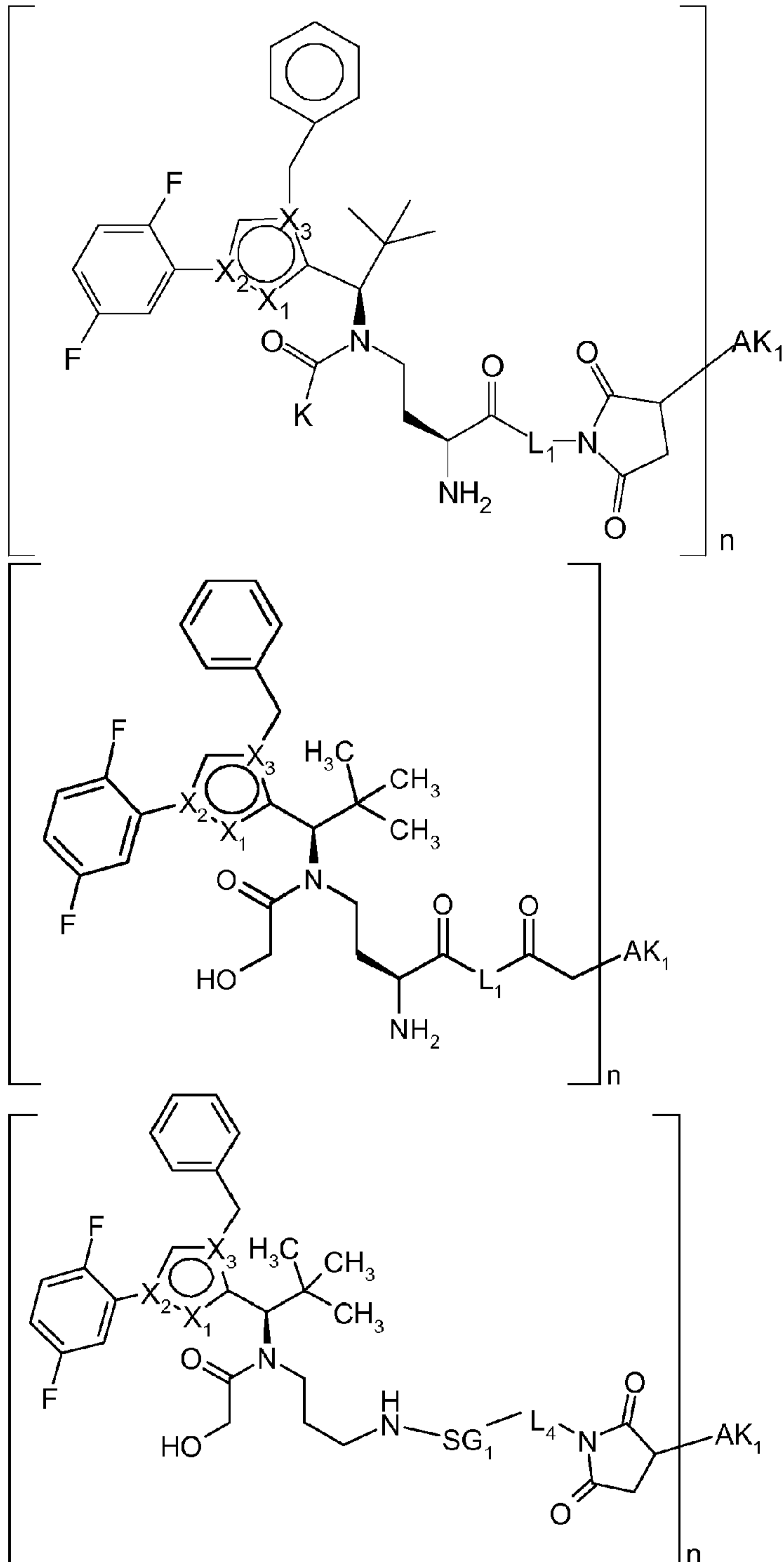
idet

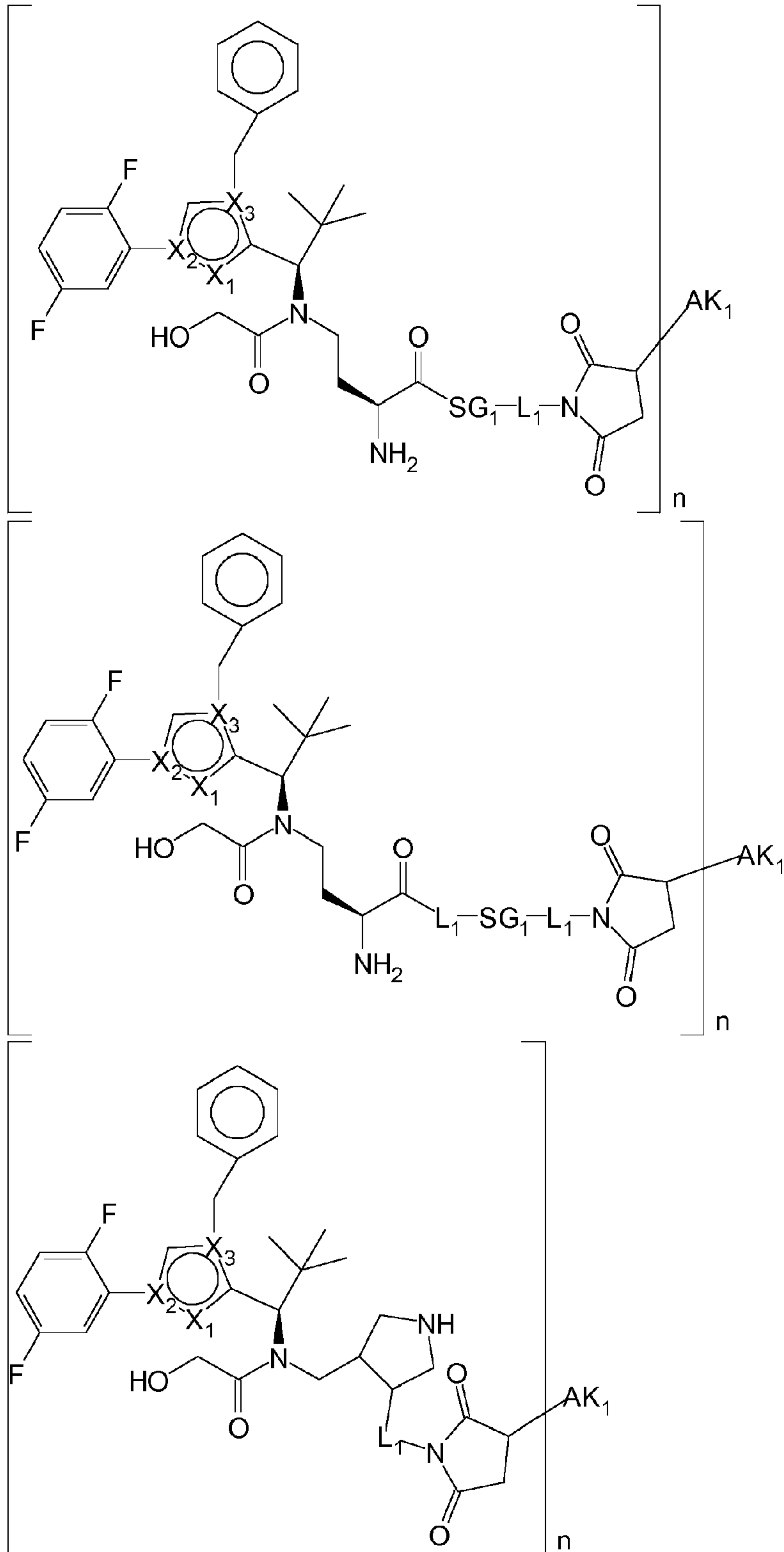
- 15 #3 er bindingen til aktivstofmolekylet,  
 #4 er bindingen til binderpeptidet eller -proteinet,  
 R<sup>11</sup> er H eller NH<sub>2</sub>;  
 B er -[(CH<sub>2</sub>)<sub>x</sub>-(X<sup>4</sup>)<sub>y</sub>]<sub>w</sub>-(CH<sub>2</sub>)<sub>z</sub>-,  
 w = 0 til 20;  
 20 x = 0 til 5;  
 x = 0 til 5;  
 y = 0 eller 1;  
 z = 0 til 5; og  
 X<sup>4</sup> er -O-, -CONH- eller -NHCO-

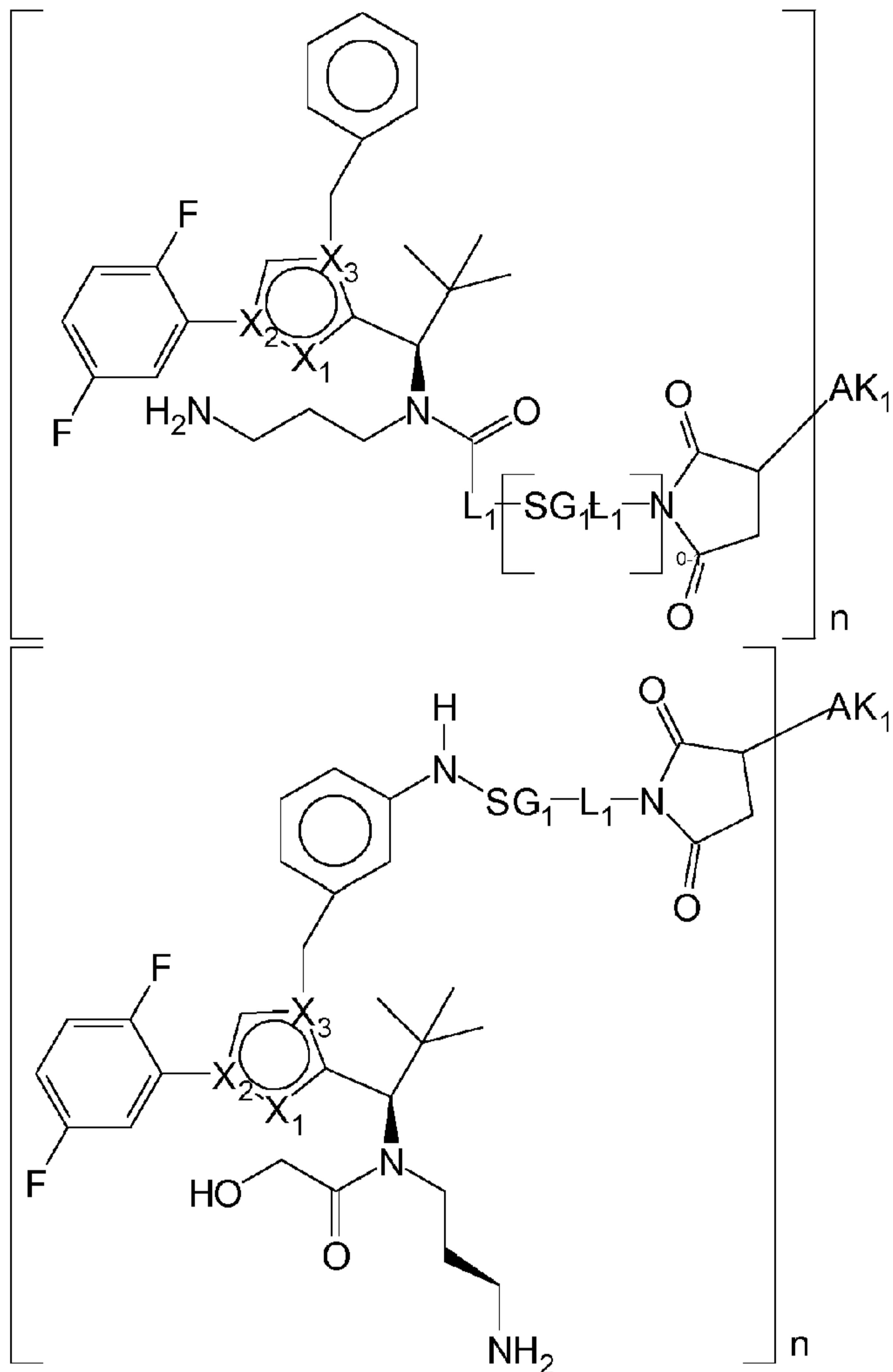


27. Konjugat ifølge krav 23, idet  $R^1$  eller  $R^4$  er  $-L-#1$ .

28. Konjugat ifølge et eller flere af kravene 23 til 27, idet 5 konjugatet har en af de følgende formler:



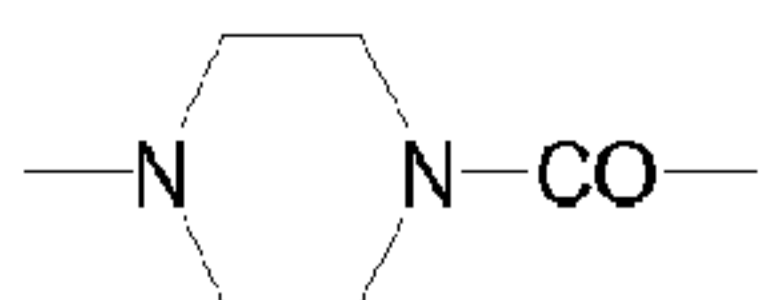




idet

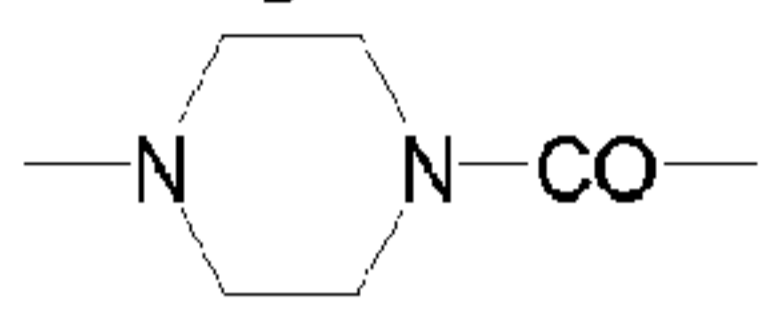
$X_1$ ,  $X_2$  og  $X_3$  har samme betegnelse som i krav 4,

$AK_1$  er et binderpeptid eller -protein, som er bundet via et  
 5 svovlatom i binderen;  $n$  er et tal fra 1 til 20; og  $L_1$  er en i  
 givet fald forgrenet carbonhydridgruppe med 1 til 70  
 carbonatomer, som er en ligekædet eller forgrenet  
 carbonhydridkæde med 1 til 100 carbonatomer fra arylengrupper  
 og/eller ligekædede og/eller forgrenede og/eller cykliske  
 10 alkylengrupper, som kan være afbrudt en eller flere gange af en  
 eller flere af grupperne  $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-$   
 $CONH-$ ,  $-NHCO-$ ,  $-Nme-$ ,  $-NHNH-$ ,  $-SO_2NHNH-$ ,  $-CONHNH-$  og en 5- til  
 10-leddet, aromatisk eller ikke aromatisk heterocyclus med op  
 til 4 heteroatomer udvalgt af N, O og S,  $-SO-$  eller  $-SO_2-$   
 15 (fortrinsvis



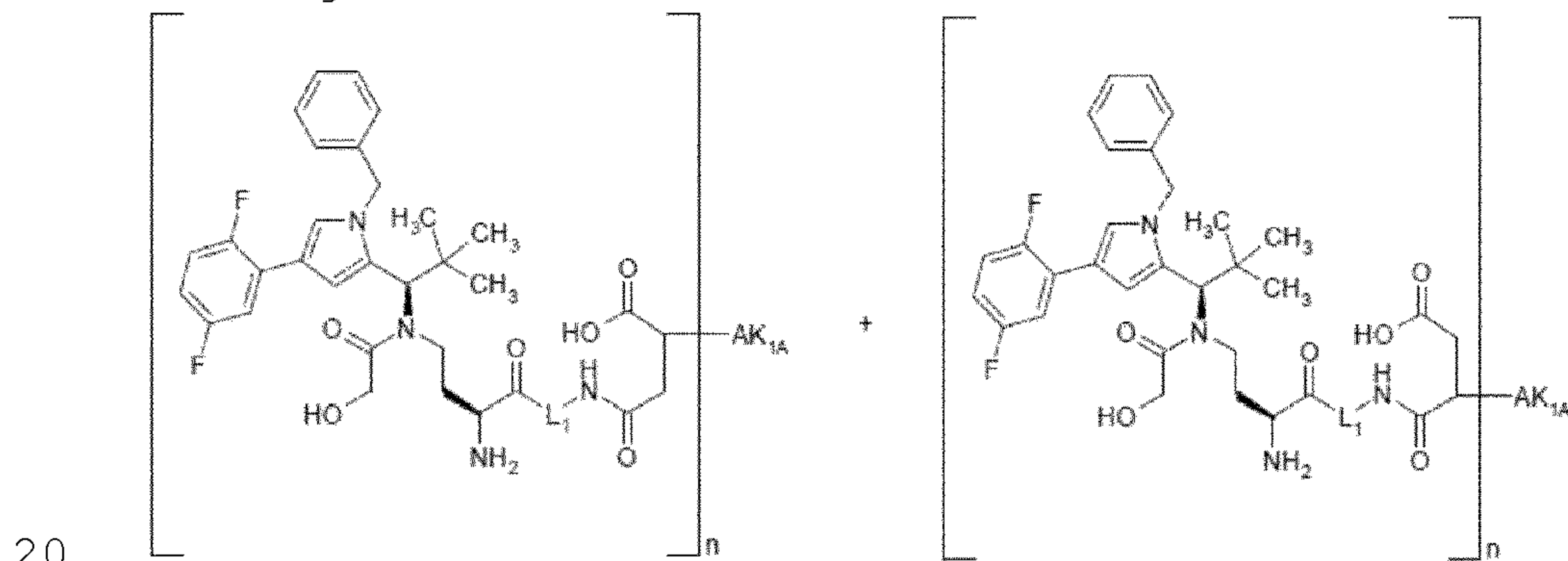
), idet sidekæderne, såfremt sådanne eksisterer, kan være  
 substitueret med  $-NHCONH_2$ ,  $-COOH$ ,  $-OH$ ,  $-NH_2$ ,  $NH-CNNH_2$ ,

sulfonamid, sulfon, sulfoxid eller sulfonsyre,  
 og SG1 er et 2-8 oligopeptid, fortrinsvis et dipeptid;  
 L4 er en enkeltbinding eller en gruppe  $-(CO)_y-G4-$ , idet y er 0  
 eller 1, og G4 er en ligekædet eller forgrenet carbonhydriddkæde  
 5 med 1 til 100 carbonatomer fra arylengrupper og/eller ligekædede  
 og/eller forgrenede og/eller cykliske alkylengrupper, som kan  
 være afbrudt en eller flere gange af en eller flere af grupperne  
 $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-NHCO-$ ,  $-CONH-$ ,  $-Nme-$ ,  $-NHNH-$ ,  
 $-SO_2NHNH-$ ,  $-CONHNNH-$  og en 5- til 10-leddet, aromatisk eller ikke  
 10 aromatisk heterocyclus med op til 4 heteroatomer udvalgt fra N,  
 O og S,  $-SO-$  eller  $-SO_2-$  (fortrinsvis



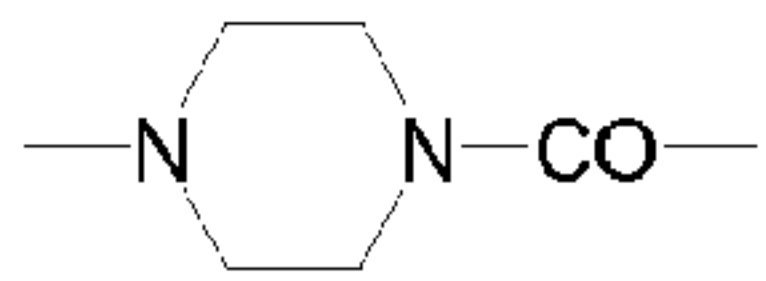
), idet sidekæderne, såfremt sådanne eksisterer, kan være  
 substitueret med  $-NHCONH_2$ ,  $-COOH$ ,  $-OH$ ,  $-NH_2$ ,  $NH-CNNH_2$ ,  
 15 sulfonamid, sulfon, sulfoxid eller sulfonsyre.

29. Konjugat af en binder eller et derivat heraf med et eller  
 flere aktivstofmolekyler, idet konjugatet har en eller begge af  
 de følgende formler:



idet

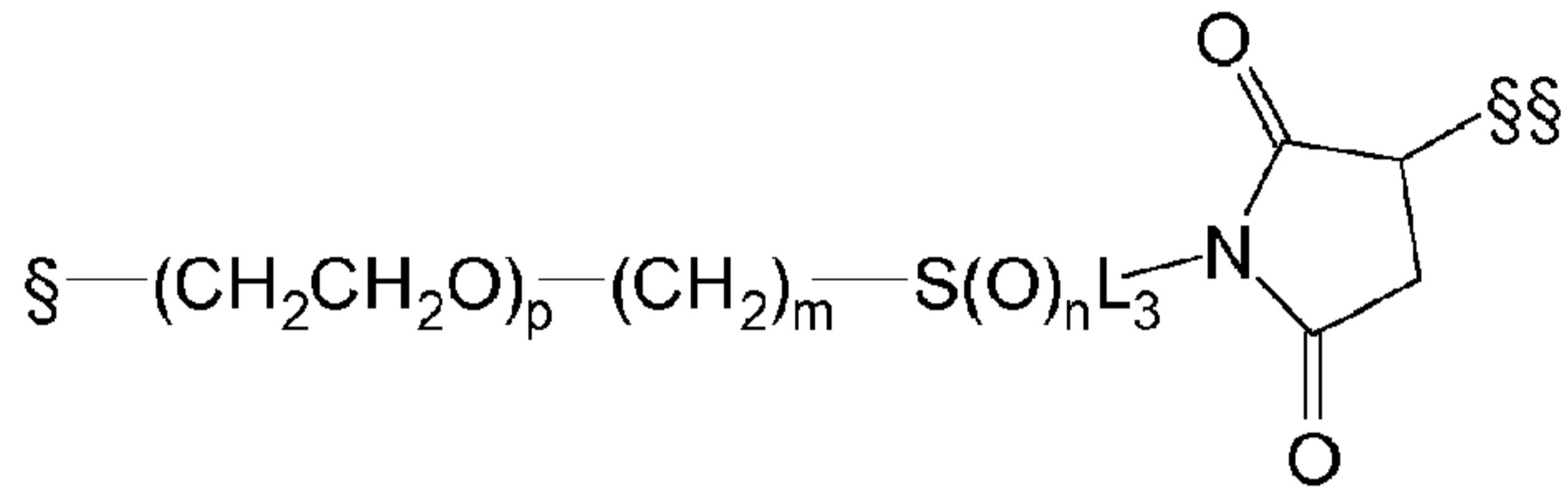
AK<sub>1A</sub> er et binderpeptid eller -protein, som er bundet via et  
 svovlatom i binderen; n er et tal fra 1 til 20; og L1 er en  
 ligekædet eller forgrenet carbonhydriddkæde med 1 til 100  
 25 carbonatomer fra arylengrupper og/eller ligekædede og/eller  
 forgrenede og/eller cykliske alkylengrupper, som kan være  
 afbrudt en eller flere gange af en eller flere af grupperne  $-$   
 $O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-CONH-$ ,  $-NHCO-$ ,  $-NMe-$ ,  $-NHNH-$ ,  
 $-SO_2NHNH-$ ,  $-CONHNNH-$  og en 5- til 10-leddet, aromatisk eller  
 30 ikke aromatisk heterocyclus med op til 4 heteroatomer udvalgt  
 fra N, O og S,  $-SO-$  eller  $-SO_2-$  (fortrinsvis



), idet sidekæderne, såfremt sådanne eksisterer, kan være substitueret med  $-NHCONH_2$ ,  $-COOH$ ,  $-OH$ ,  $-NH_2$ ,  $NH-CNNH_2$ , sulfonamid, sulfon, sulfoxid eller sulfonsyre.

5

30. Konjugat ifølge et eller flere af de foregående krav, idet linkeren  $-L-$  er bundet til en cystein-sidekæde eller cysteingruppe og har den følgende formel:



10 idet

§ er bindingen til aktivstofmolekylet, og

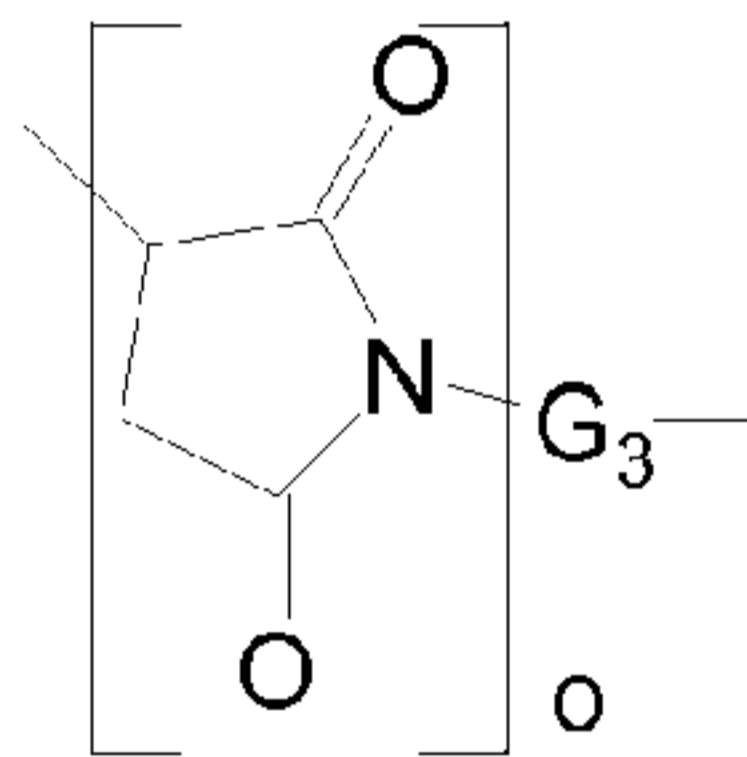
§§ er bindingen til binderpeptidet eller -proteinet,

m er 0, 1, 2, eller 3;

n er 0, 1 eller 2;

15 p er 0 til 20; og

L3 er

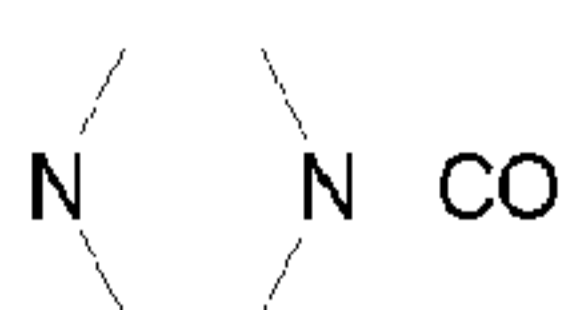


idet

o er 0 eller 1;

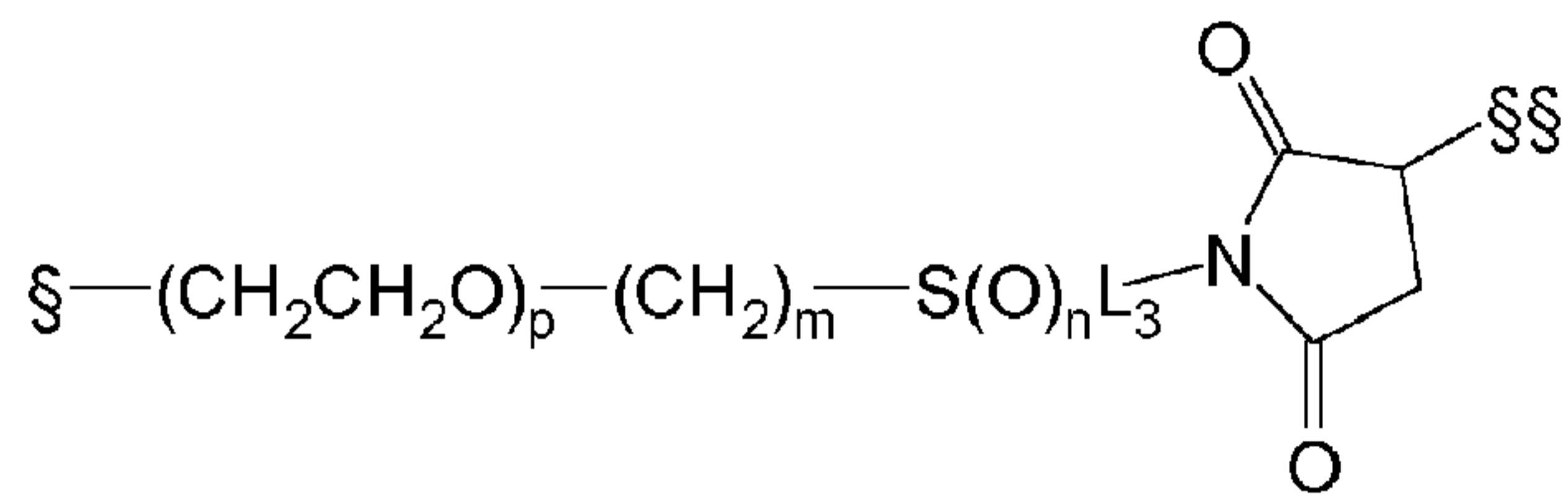
20 og

G3 er en ligekædet eller forgrenet carbonhydriddkæde med 1 til 100 carbonatomer fra arylengrupper og/eller ligekædede og/eller forgrenede og/eller cykliske alkylengrupper, som kan være afbrudt en eller flere gange af en eller flere af grupperne  $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-NHCO$ ,  $-CONH-$  og en 5- til 10-leddet aromatisk eller ikke aromatisk heterocyclus med op til 4 heteroatomer udvalgt fra N, O og S,  $-$ ,  $-Nme-$ ,  $-NHNH-$ ,  $-SO_2NHNH-$ ,  $-CONHNH-$   $-SO-$  eller  $-SO_2-$  (fortrinsvis



30 ), idet sidekæderne, såfremt sådanne eksisterer, kan være substitueret med  $-NHCONH_2$ ,  $-COOH$ ,  $-OH$ , sulfon, sulfoxid eller sulfonsyre.

31. Konjugat ifølge krav 30, idet linkeren -L- er bundet til en cystein-sidekæde eller cysteingruppe og har den følgende formel:



5

idet

$\S$  er bindingen til aktivstofmolekylet, og

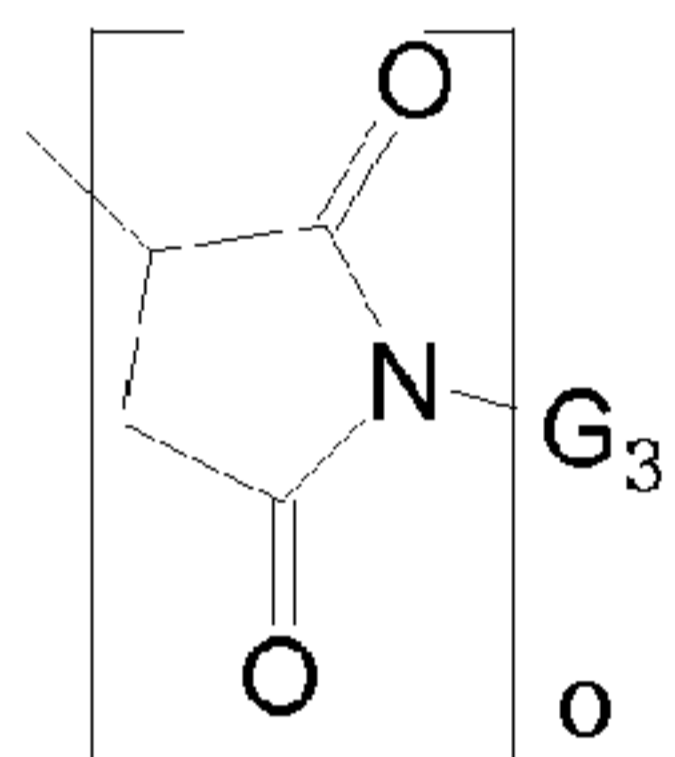
$\S \S$  er bindingen til binderpeptidet eller -proteinet,

$m$  er 1;

10  $p$  er 0;

$n$  er 0;

og  $\text{L}_3$  er



idet

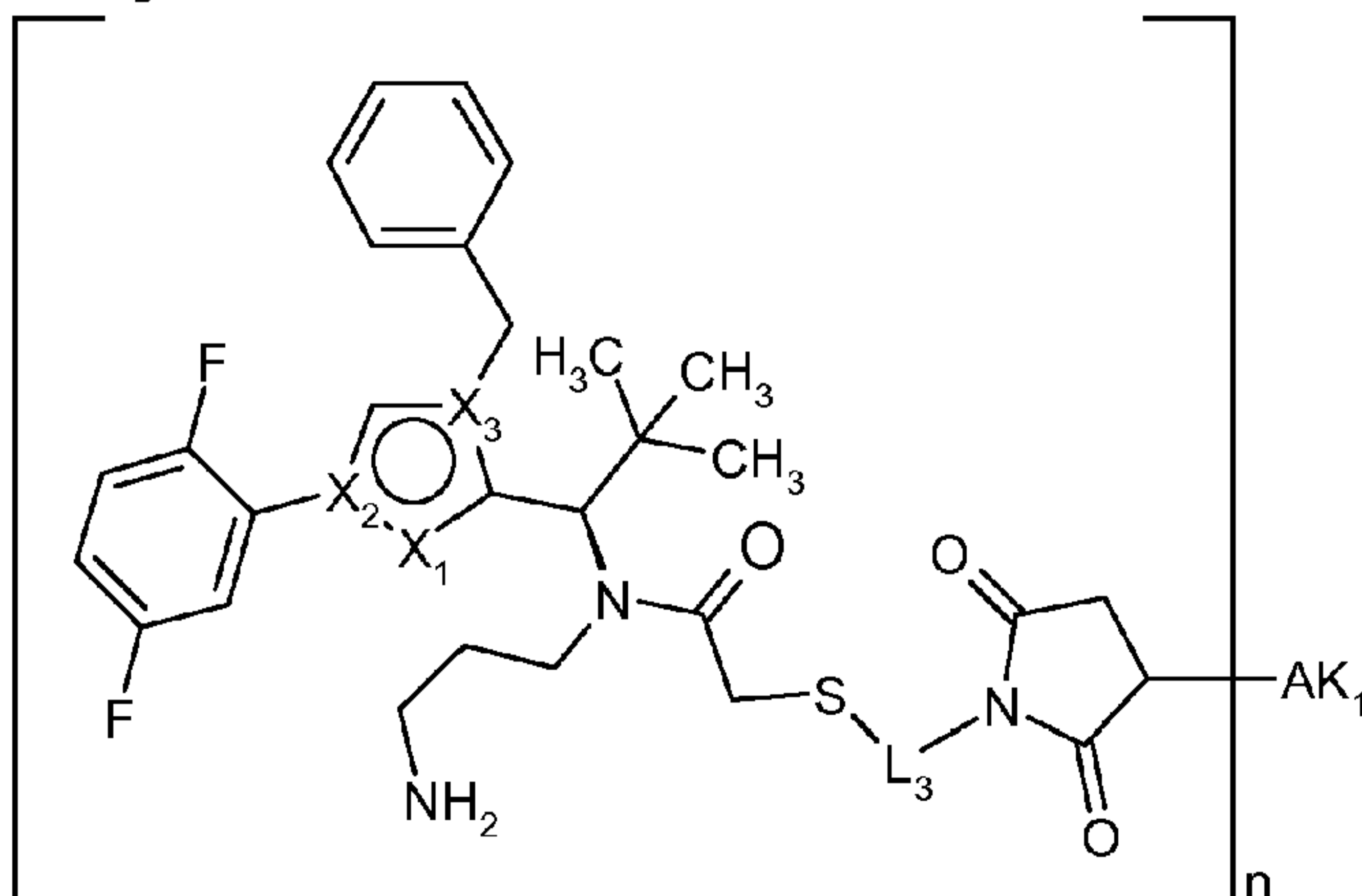
15  $o$  er 0 eller 1; og

$\text{G}_3$  er  $-(\text{CH}_2\text{CH}_2\text{O})_s(\text{CH}_2)_t(\text{CONH})_u(\text{CH}_2\text{CH}_2\text{O})_v(\text{CH}_2)_w-$ , idet

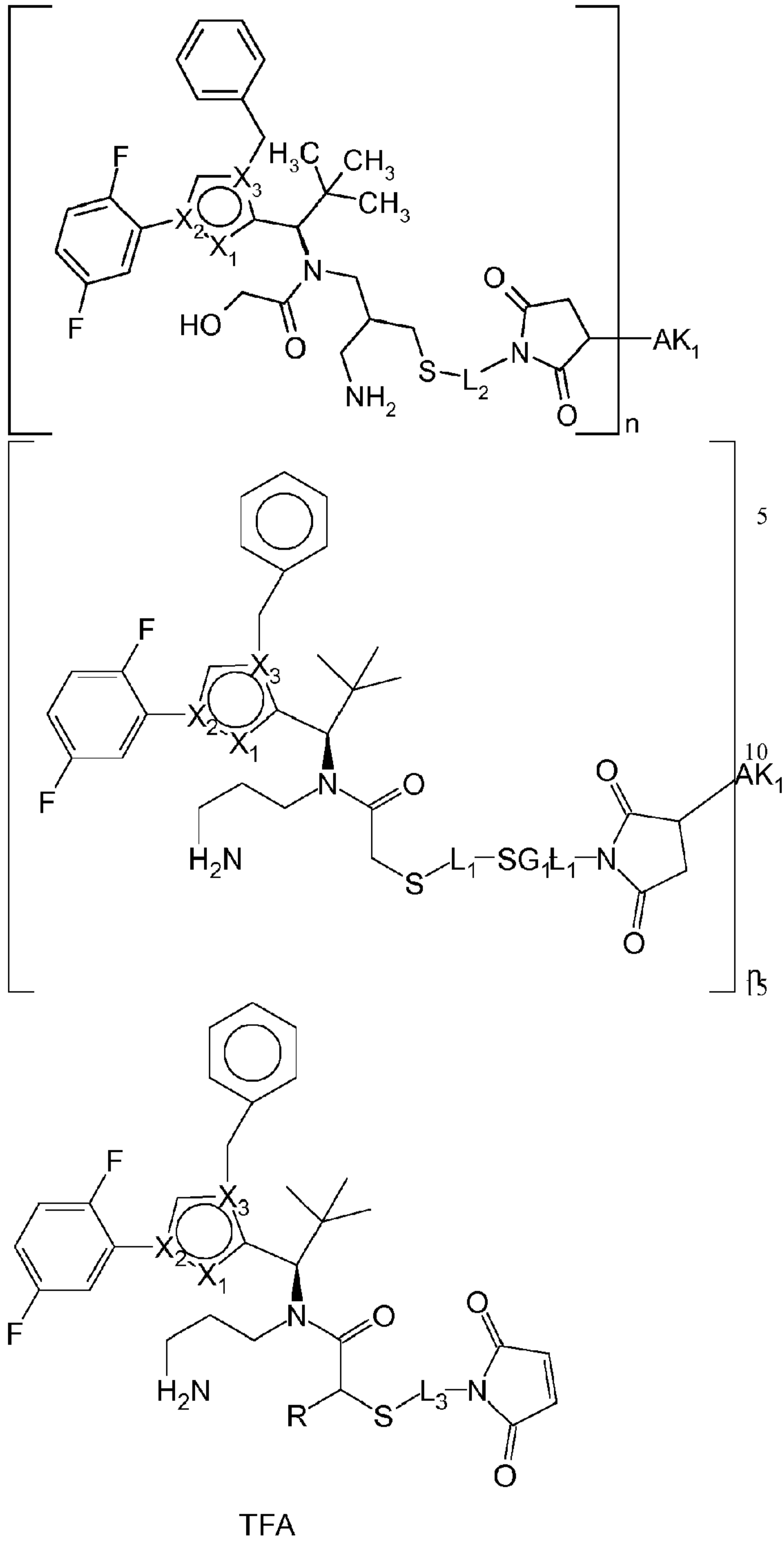
$s$ ,  $t$ ,  $v$  og  $w$  hver især uafhængigt af hinanden er 0 til 20, og  $u$  er 0 eller 1.

20 32. Konjugat ifølge krav 30 eller 31, idet  $\text{R}^2$  eller  $\text{R}^3$  er -L-  
#1.

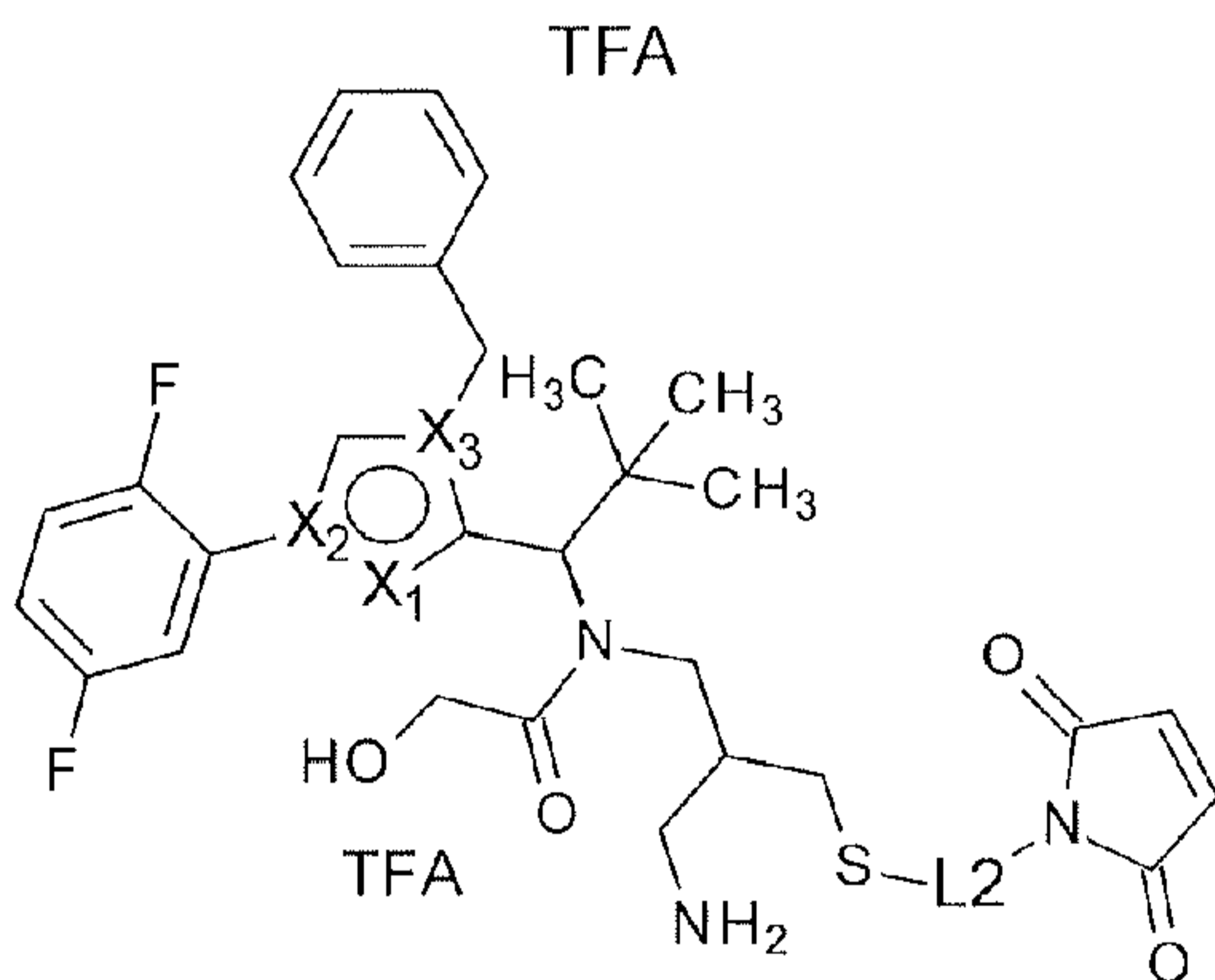
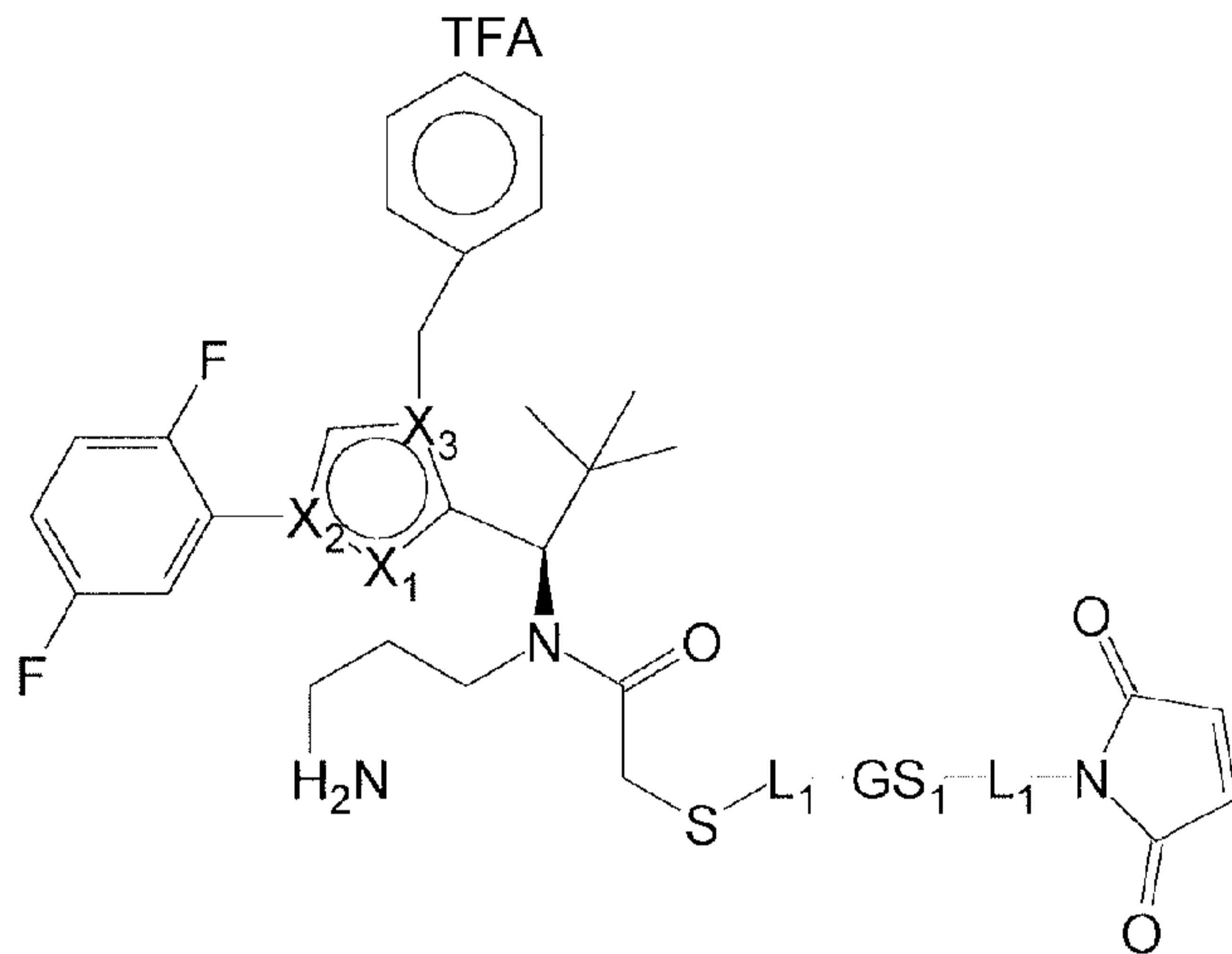
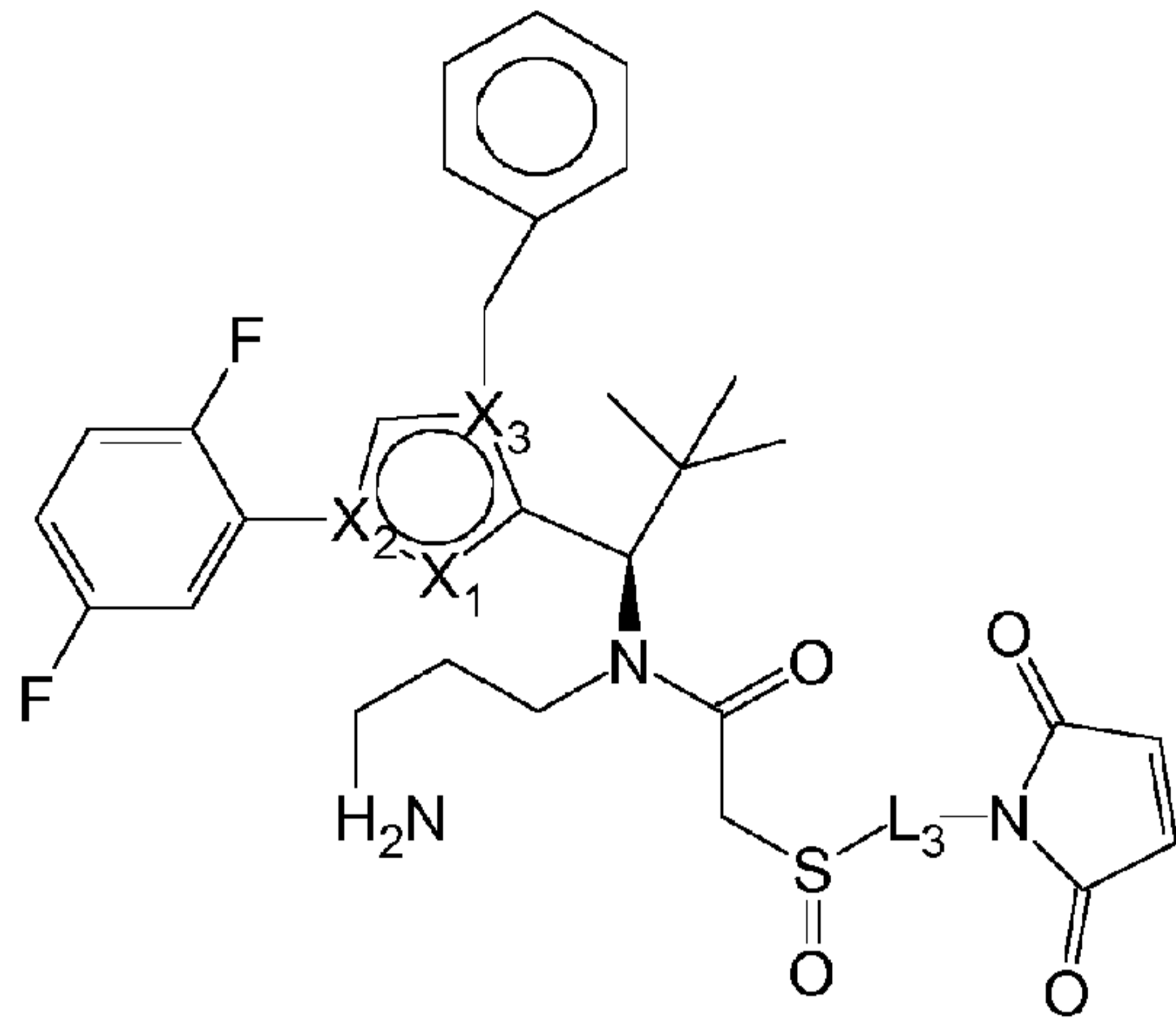
33. Konjugat ifølge krav 32, idet konjugatet har en af de følgende formler:



25



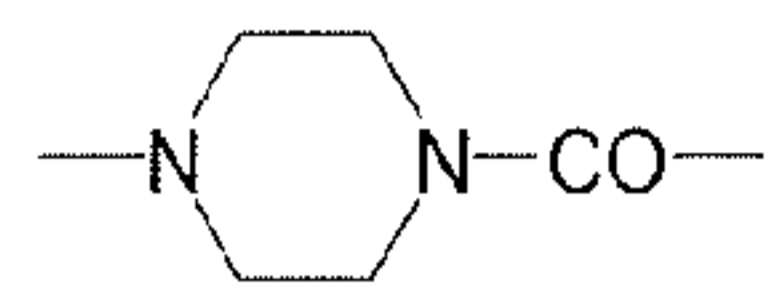




idet

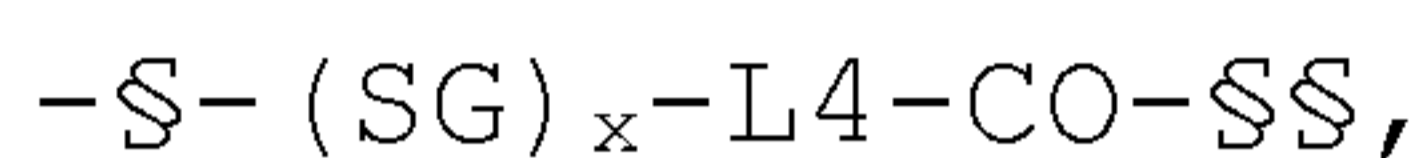
$X_1$ ,  $X_2$  og  $X_3$  har den samme betegnelse som i krav 4,  
 $AK_1$  er et binderpeptid eller -protein, som er bundet via et  
 5 svovlatom i binderen;  $n$  er et tal fra 1 til 20; og  $L_2$  og  $L_3$  er  
 en i givet fald forgrenet, ligekædet eller forgrenet  
 carbonhydridkæde med 1 til 100 carbonatomer fra arylengrupper  
 og/eller ligekædede og/eller forgrenede og/eller cykliske  
 alkylengrupper, som kan være afbrudt en eller flere gange af en  
 10 eller flere af grupperne  $-O-$ ,  $-S-$ ,  $-SO-$ ,  $SO_2$ ,  $-NH-$ ,  $-CO-$ ,  $-$   
 $Nme-$ ,  $-NHNH-$ ,  $-SO_2NHNH-$ ,  $-NHCO-$ ,  $-CONH-$ ,  $-CONHNH-$  og en 5- til

10-leddet aromatisk eller ikke aromatisk heterocyclus med op til 4 heteroatomer udvalgt fra N, O og S, -SO- eller -SO<sub>2</sub>- (fortrinsvis



5 ), idet sidekæderne, såfremt sådanne eksisterer, kan være substitueret med -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulfonamid, sulfon, sulfoxid eller sulfonsyre.

34. Konjugat ifølge et eller flere af kravene 1 til 29, idet linkerens -L- er bundet til en lysin-sidekæde og har følgende formel:



idet

§ er bindingen til aktivstofmolekylet, og

15 §§ er bindingen til binderpeptidet eller -proteinet,

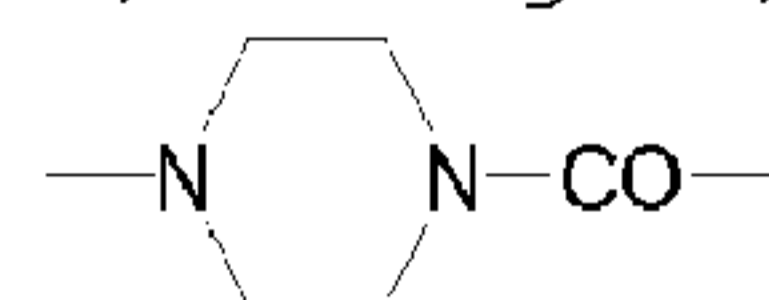
x er 0 eller 1,

SG er en spaltbar gruppe, fortrinsvis et 2-8 oligopeptid, særligt foretrukket et dipeptid,

og

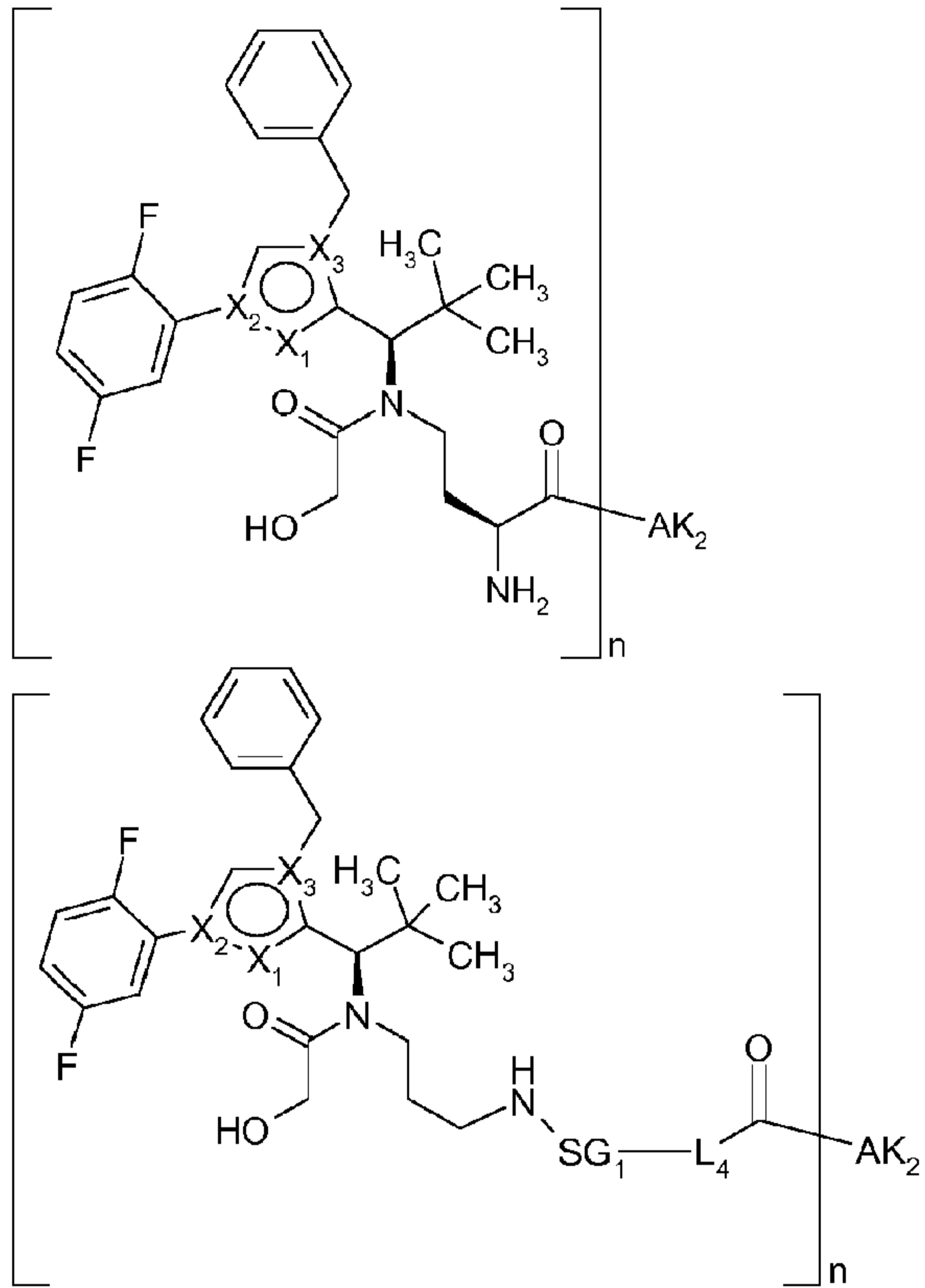
20 L4 er en enkeltbinding eller en gruppe -(CO)<sub>y</sub>-G4-, idet y er 0 eller 1, og G4 er en ligekædet eller forgrenet carbonhydriddkæde med 1 til 100 carbonatomer af arylengrupper og/eller ligekædede og/eller forgrenede og/eller cykliske alkylengrupper, som kan være afbrudt e eller flere gange af en eller flere af grupperne

25 -O-, -S-, -SO-, SO<sub>2</sub>, -NH-, -CO-, -NHCO-, -CONH-, -Nme-, -NHNH-, -SO<sub>2</sub>NHNH-, -CONHNH- og en 5- til 10-leddet aromatisk eller ikke aromatisk heterocyclus med op til 4 heteroatomer udvalgt fra N, O og S, eller -SO- (fortrinsvis



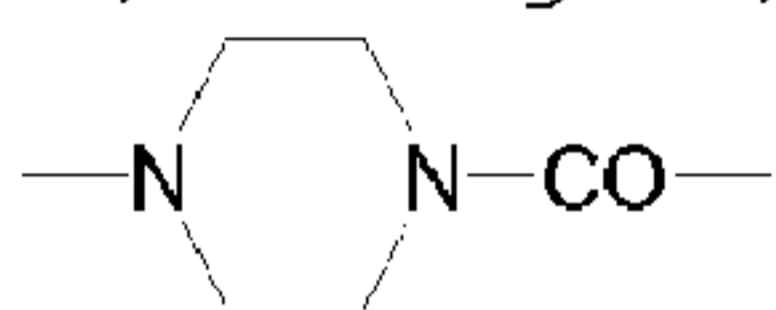
30 ), idet sidekæderne, såfremt sådanne eksisterer, kan være substitueret med -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>, sulfonamid, sulfon, sulfoxid eller sulfonsyre.

35 35. Konjugat af et binderpeptid eller -protein ifølge krav 34, idet konjugatet har en af de følgende formler:



idet

X<sub>1</sub>, X<sub>2</sub> og X<sub>3</sub> har den samme betegnelse som i krav 4,  
 AK<sub>2</sub> er et binderpeptid eller -protein, som er bundet via et  
 5 svovlatom i binderen; n er et tal fra 1 til 20; og L<sub>4</sub> er en i  
 givet fald ligekædet eller forgrenet carbonhydriddkæde med 1 til  
 100 carbonatomer af arylengrupper og/eller ligekædede og/eller  
 forgrenede og/eller cykliske alkylengrupper, som kan være  
 afbrudt en eller flere gange af en eller flere af grupperne -O-  
 10 , -S-, , -SO-, SO<sub>2</sub>, -NH-, -CO-, -Nme-, -NHNH-, -SO<sub>2</sub>NHNH-, -NHCO-,  
 -CONH-, -CONHNH- og en 5- til 10-leddet aromatisk eller ikke  
 aromatisk heterocyclus med op til 4 heteroatomer udvalgt fra  
 N, O og S, -SO- eller -SO<sub>2</sub>- (fortrinsvis



15 ), idet sidekæderne, såfremt sådanne eksisterer, kan være  
 substitueret med -NHCONH<sub>2</sub>, -COOH, -OH, -NH<sub>2</sub>, NH-CNNH<sub>2</sub>,  
 sulfonamid, sulfon, sulfoxid eller sulfonsyre, og SG<sub>1</sub> er en  
 spaltbar gruppe, fortrinsvis et 2-8 oligopeptid, særligt  
 foretrukket et dipeptid.

20

36. Konjugat ifølge et eller flere af kravene 18 til 35, idet  
 anti-TWEAKR-antistoffet er et agonistisk antistof.

37. Konjugat ifølge et eller flere af kravene 18 til 36, som omfatter:

en variabel tung kæde omfattende:

- 5 a. en CDR1 af den tunge kæde, som kodes af en aminosyresekvens omfattende formlen PYPMX (SEQ ID NO: 171), idet X er I eller M;
- b. en CDR2 af den tunge kæde, som kodes af en aminosyresekvens omfattende formlen YISPSGGXTHYADSVKG (SEQ ID NO: 172), idet X er S eller K; og
- 10 c. en CDR3 af den tunge kæde, som kodes af en aminosyresekvens omfattende formlen GGDTYFDYFDY (SEQ ID NO: 173);
- og en variabel let kæde omfattende:
- d. en CDR1 af den lette kæde, som kodes af en aminosyresekvens omfattende formlen RASQSISXYLN (SEQ ID NO: 174), idet X er G
- 15 eller S;
- e. en CDR2 af den lette kæde, som kodes af en aminosyresekvens omfattende formlen XASSLQS (SEQ ID NO: 175), idet X er Q, A eller N; og
- f. en CDR3 af den lette kæde, som kodes af en aminosyresekvens
- 20 omfattende formlen QQSYXXPXIT (SEQ ID NO: 176), idet X på position 5 er T eller S, X på position 6 er T eller S, og X på position 8 er G eller F.

38. Konjugat ifølge et eller flere af kravene 18 til 37

25 omfattende:

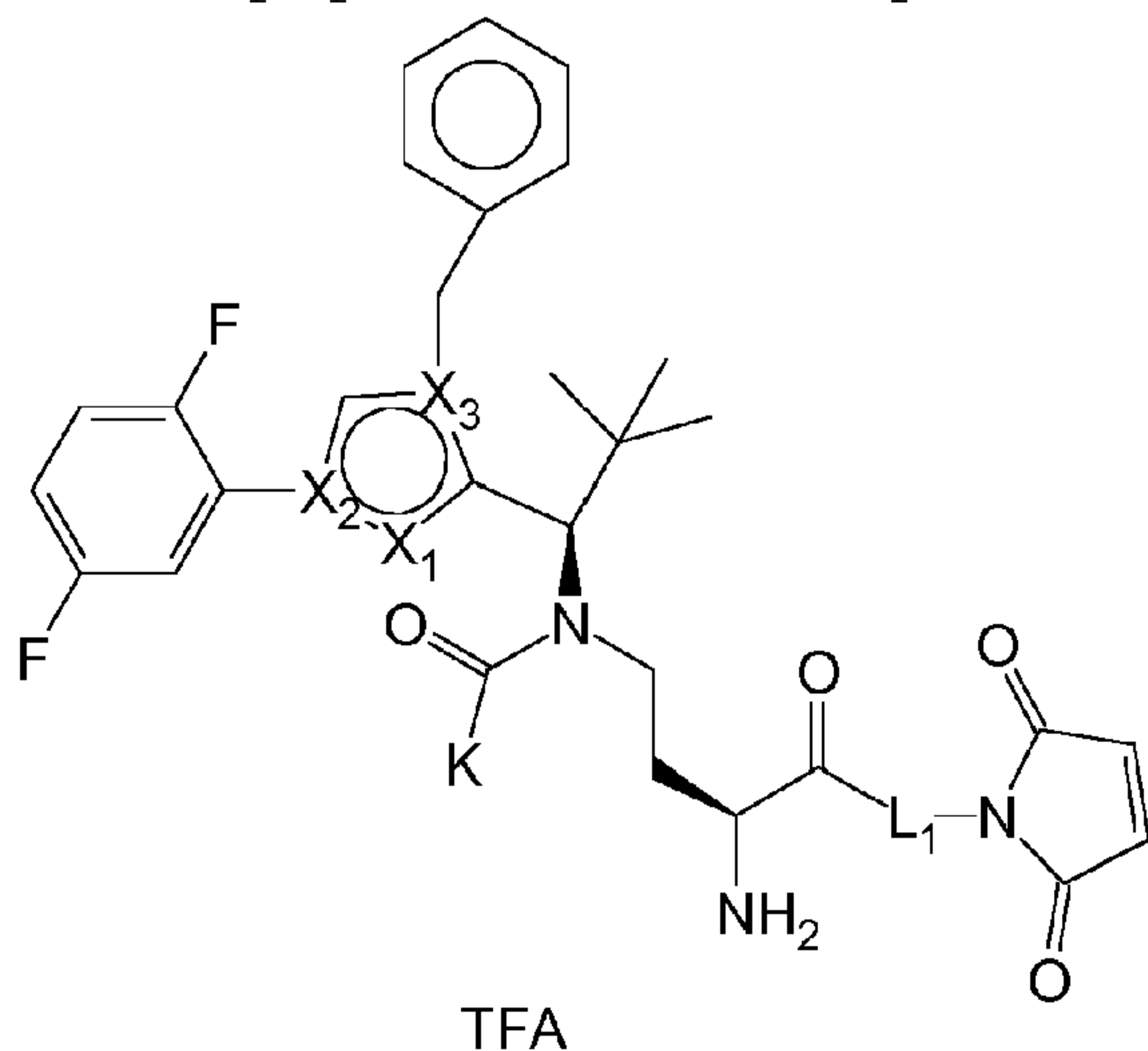
- a. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:10, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:9 eller
- b. en variabel sekvens af den tunge kæde, som vist ved SEQ ID
- 30 NO:20, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:19 eller
- c. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:30, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:29 eller
- 35 d. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:40, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:39 eller
- e. en variabel sekvens af den tunge kæde, som vist ved SEQ ID

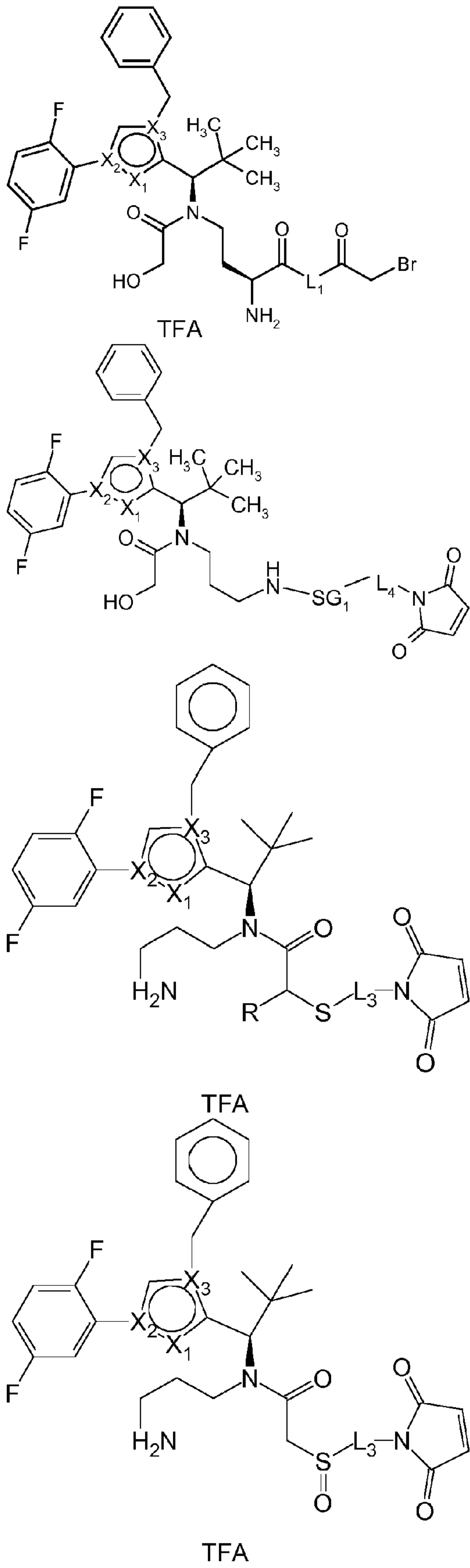
- NO:50, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:49 eller
- f. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:60, samt en variabel sekvens af den lette kæde, som vist ved
- 5 SEQ ID NO:59 eller
- g. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:70, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:69 eller
- h. en variabel sekvens af den tunge kæde, som vist ved SEQ ID
- 10 NO:80, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:79 eller
- i. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:90, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:89 eller
- 15 j. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:100, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:99 eller
- k. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:110, samt en variabel sekvens af den lette kæde, som vist ved
- 20 SEQ ID NO:109 eller
- l. en variabel sekvens af den tunge kæde, som vist ved SEQ ID NO:120, samt en variabel sekvens af den lette kæde, som vist ved SEQ ID NO:119.
- 25 39. Konjugat ifølge et eller flere af kravene 18 til 38, idet antistoffet er et IgG antistof.
40. Konjugat ifølge et eller flere af kravene 18 til 39 omfattende:
- 30 a. en sekvens af den tunge kæde, som vist ved SEQ ID NO:2, samt en sekvens af den lette kæde, som vist ved SEQ ID NO:1, eller
- b. en sekvens af den tunge kæde, som vist ved SEQ ID NO:12, samt en sekvens af den lette kæde, som vist ved SEQ ID NO:11, eller
- c. en sekvens af den tunge kæde, som vist ved SEQ ID NO:22, samt
- 35 en sekvens af den lette kæde, som vist ved SEQ ID NO:21, eller
- d. en sekvens af den tunge kæde, som vist ved SEQ ID NO:32, samt en sekvens af den lette kæde, som vist ved SEQ ID NO:31, eller
- e. en sekvens af den tunge kæde, som vist ved SEQ ID NO:42, samt

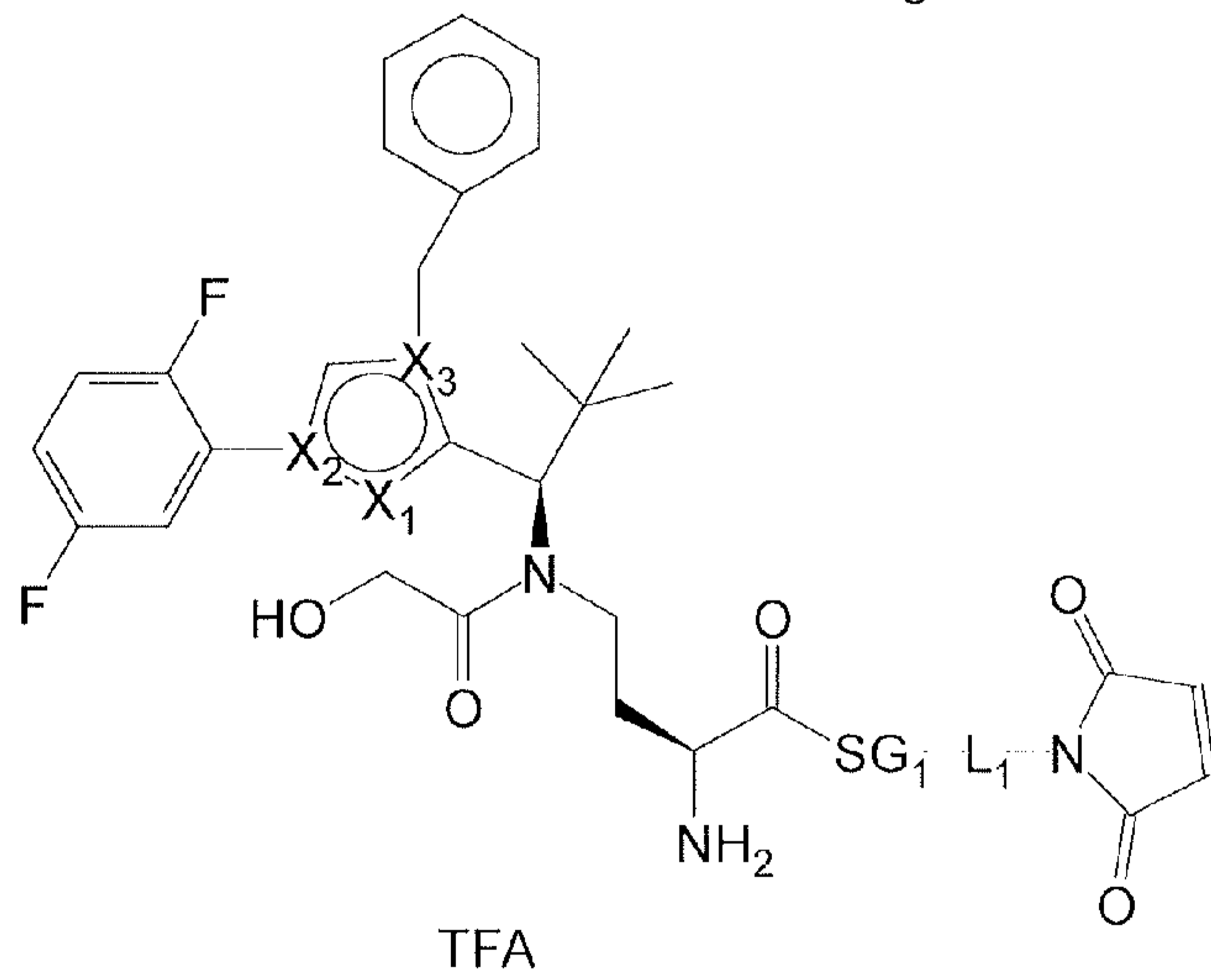
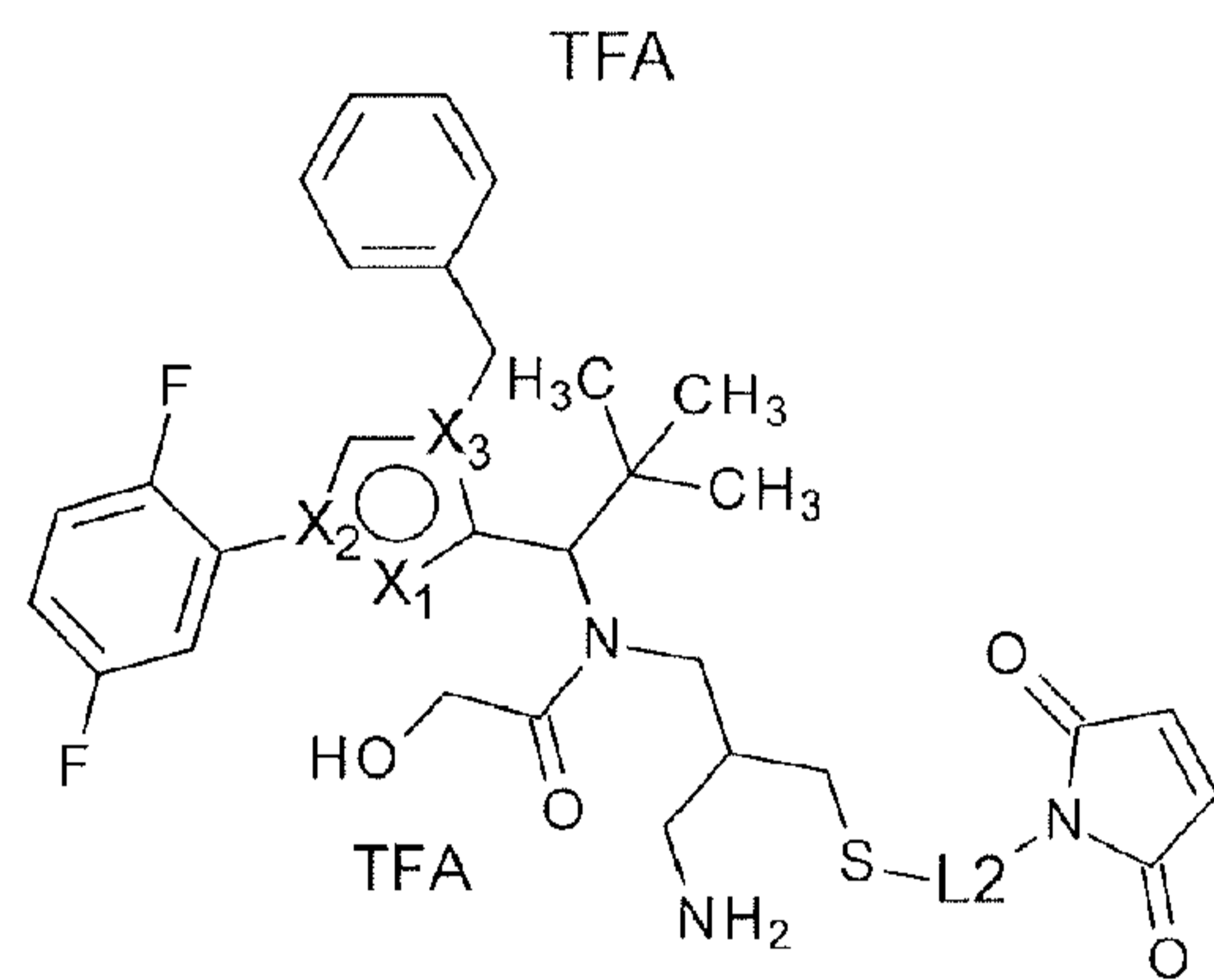
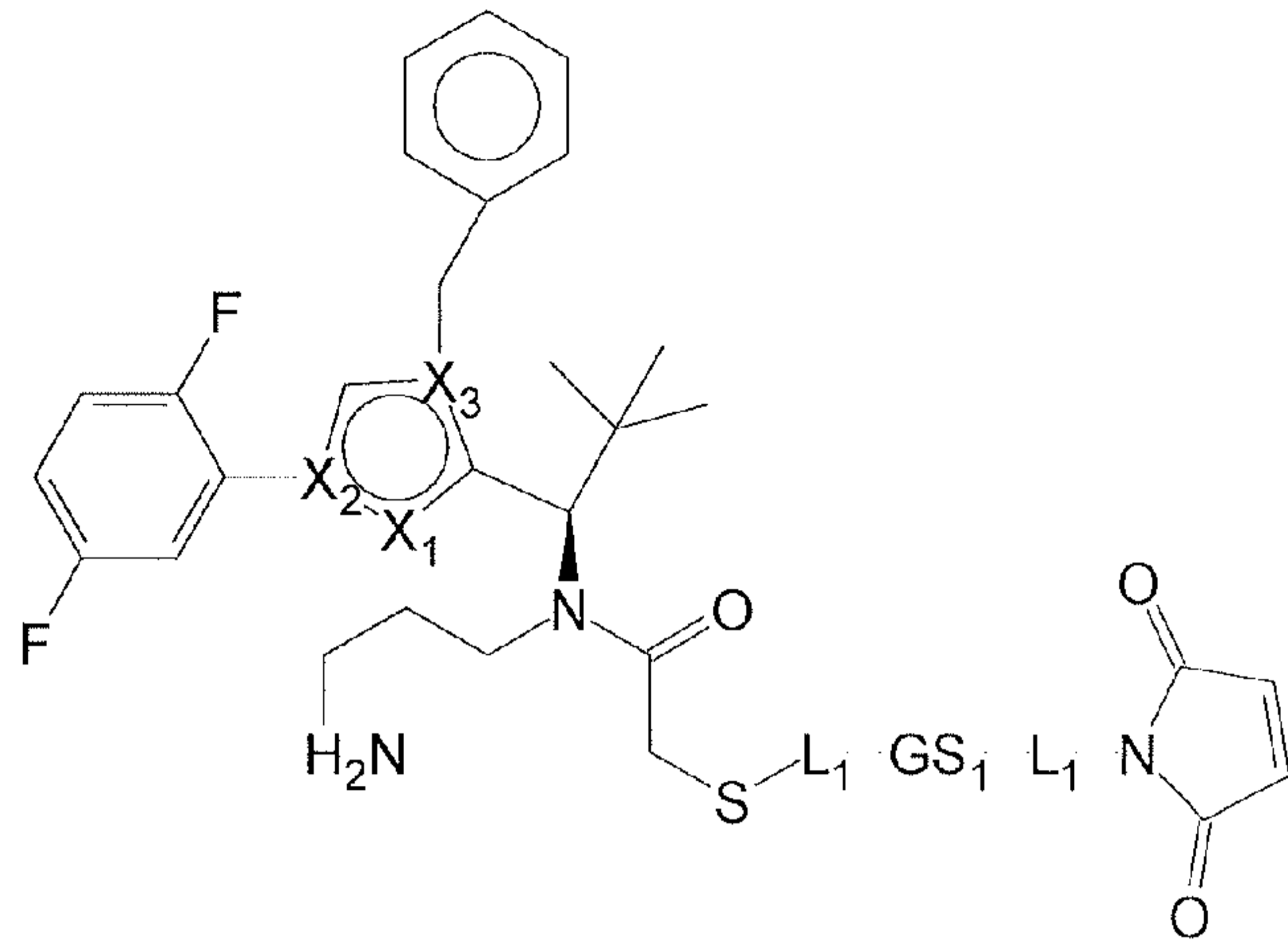
- en sekvens af den lette kæde, som vist ved SEQ ID NO:41, eller  
 f. en sekvens af den tunge kæde, som vist ved SEQ ID NO:52, samt  
 en sekvens af den lette kæde, som vist ved SEQ ID NO:51, eller  
 g. en sekvens af den tunge kæde, som vist ved SEQ ID NO:62, samt  
 5 en sekvens af den lette kæde, som vist ved SEQ ID NO:61, eller  
 h. en sekvens af den tunge kæde, som vist ved SEQ ID NO:72, samt  
 en sekvens af den lette kæde, som vist ved SEQ ID NO:71, eller  
 i. en sekvens af den tunge kæde, som vist ved SEQ ID NO:82, samt  
 en sekvens af den lette kæde, som vist ved SEQ ID NO:81, eller  
 10 j. en sekvens af den tunge kæde, som vist ved SEQ ID NO:92, samt  
 en sekvens af den lette kæde, som vist ved SEQ ID NO:91, eller  
 k. en sekvens af den tunge kæde, som vist ved SEQ ID NO:102,  
 samt en sekvens af den lette kæde, som vist ved SEQ ID NO:101,  
 eller  
 15 l. en sekvens af den tunge kæde, som vist ved SEQ ID NO:112,  
 samt en sekvens af den lette kæde, som vist ved SEQ ID NO:111.

41. Spindel-konjugat ifølge et eller flere af de foregående  
 krav, idet konjugatet har 1 til 10, fortrinsvis 2 til 8,  
 20 aktivstofmolekyler pr. binderpeptid eller -protein.

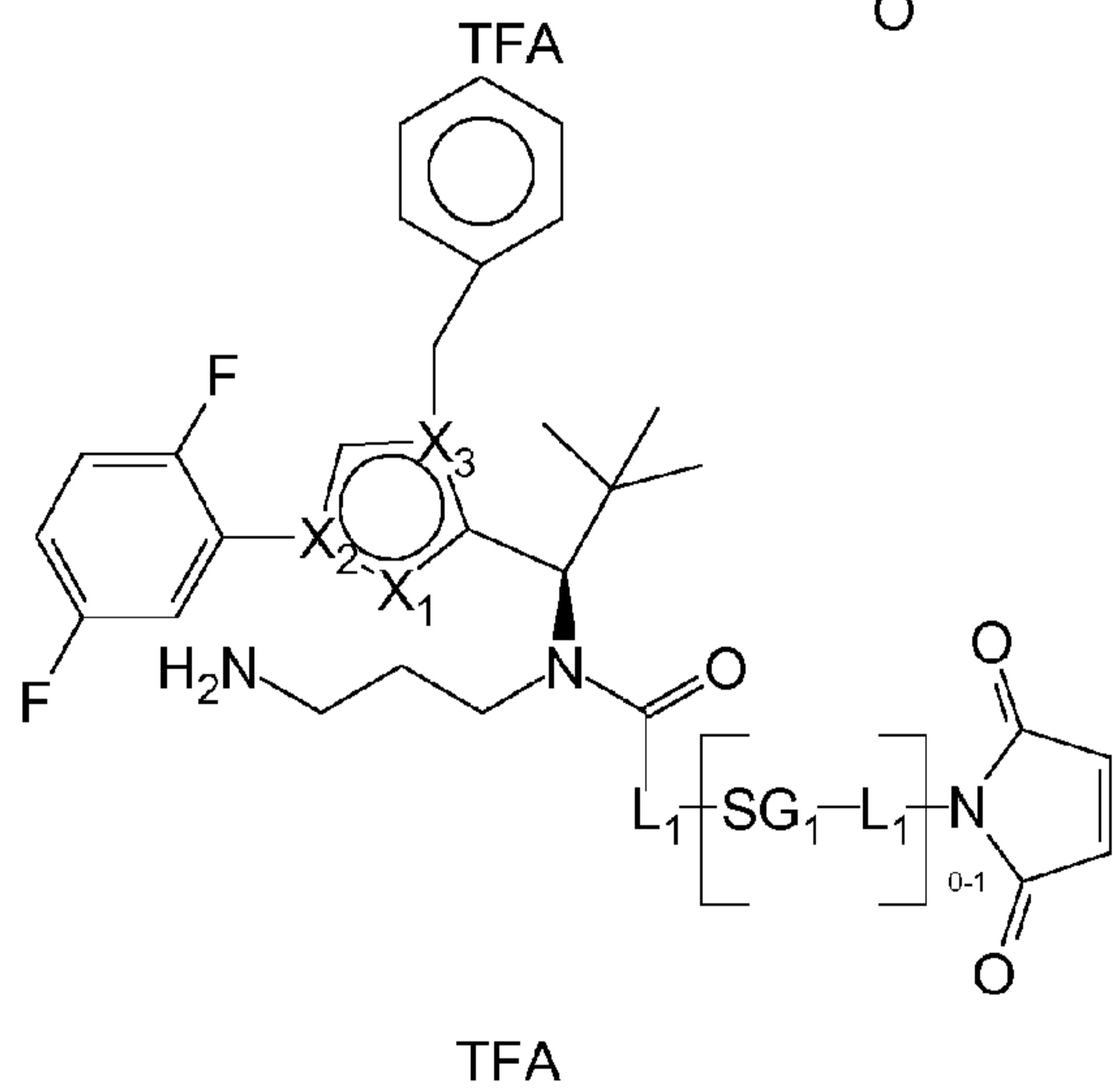
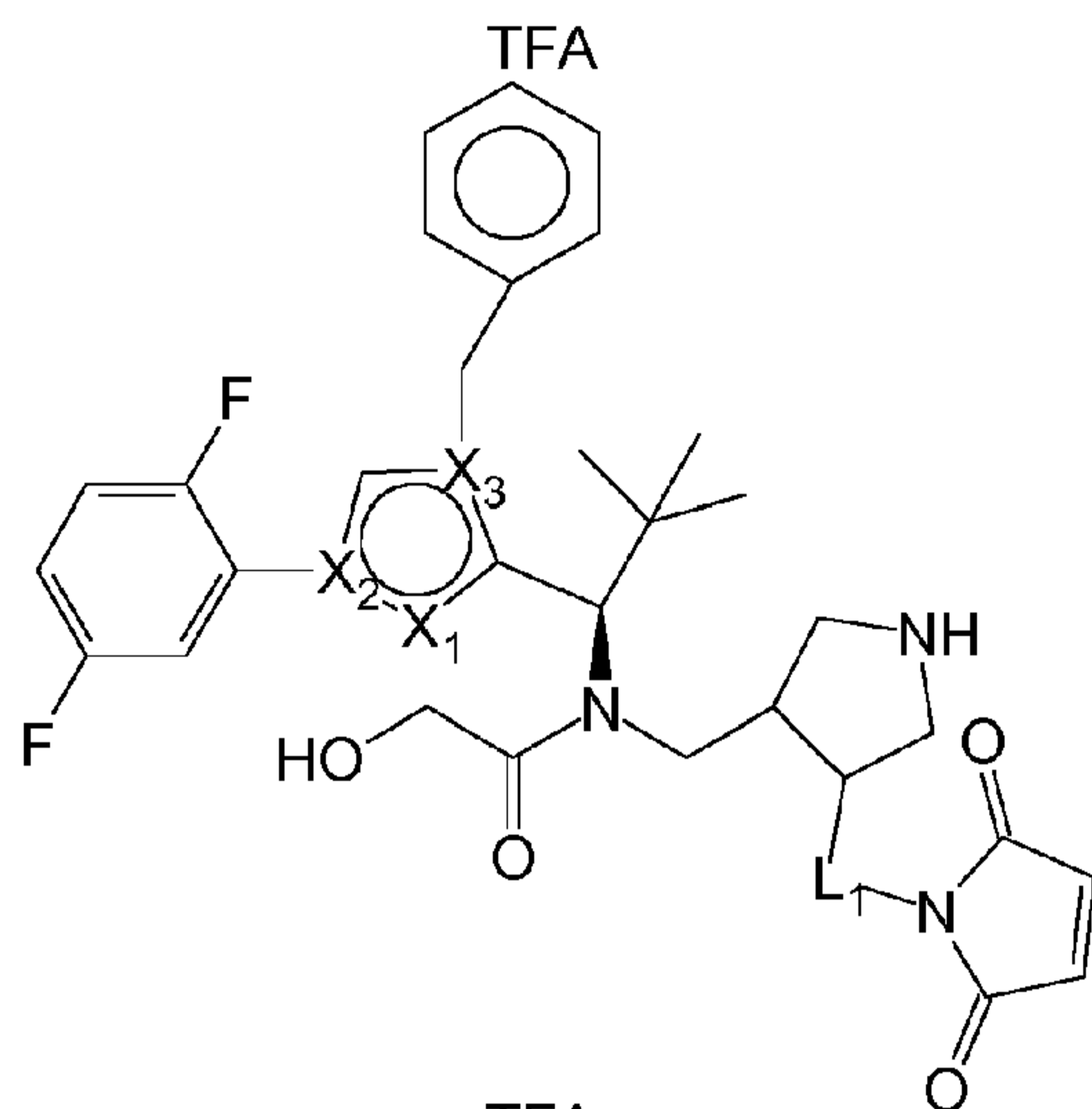
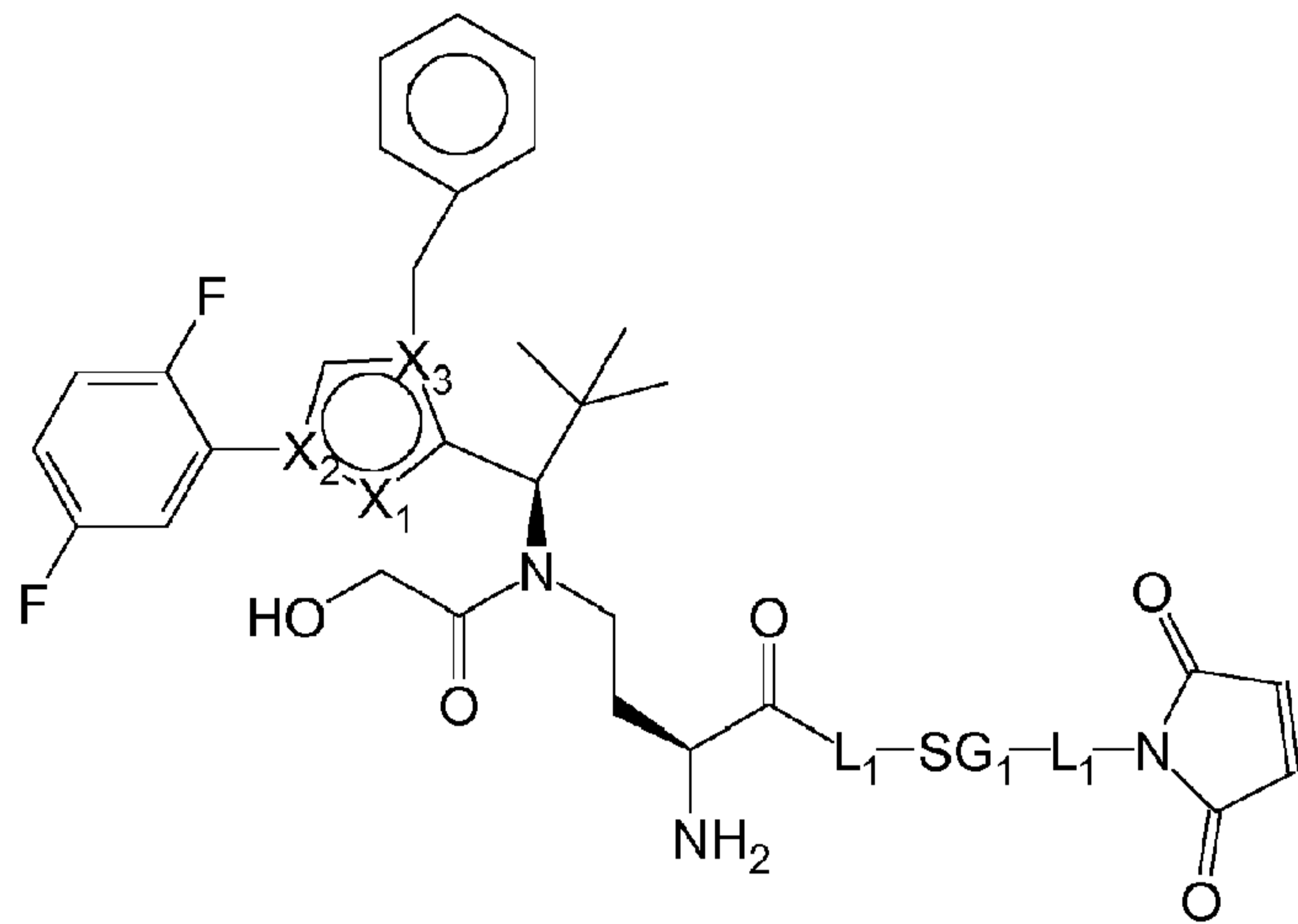
42. Fremgangsmåde til fremstilling af et konjugat ifølge krav  
 28 eller 32, idet en forbindelse med en af de følgende formler,  
 fortrinsvis i form af dens trifluoreddikesyresalt, bindes til  
 25 en cysteingrouppe i et forinden i givet fald partielt reduceret  
 binderpeptid eller -protein, idet forbindelsen fortrinsvis  
 anvendes i 2- til 12-dobbelt molært overskud i forhold til  
 binderpeptidet eller -proteinet:

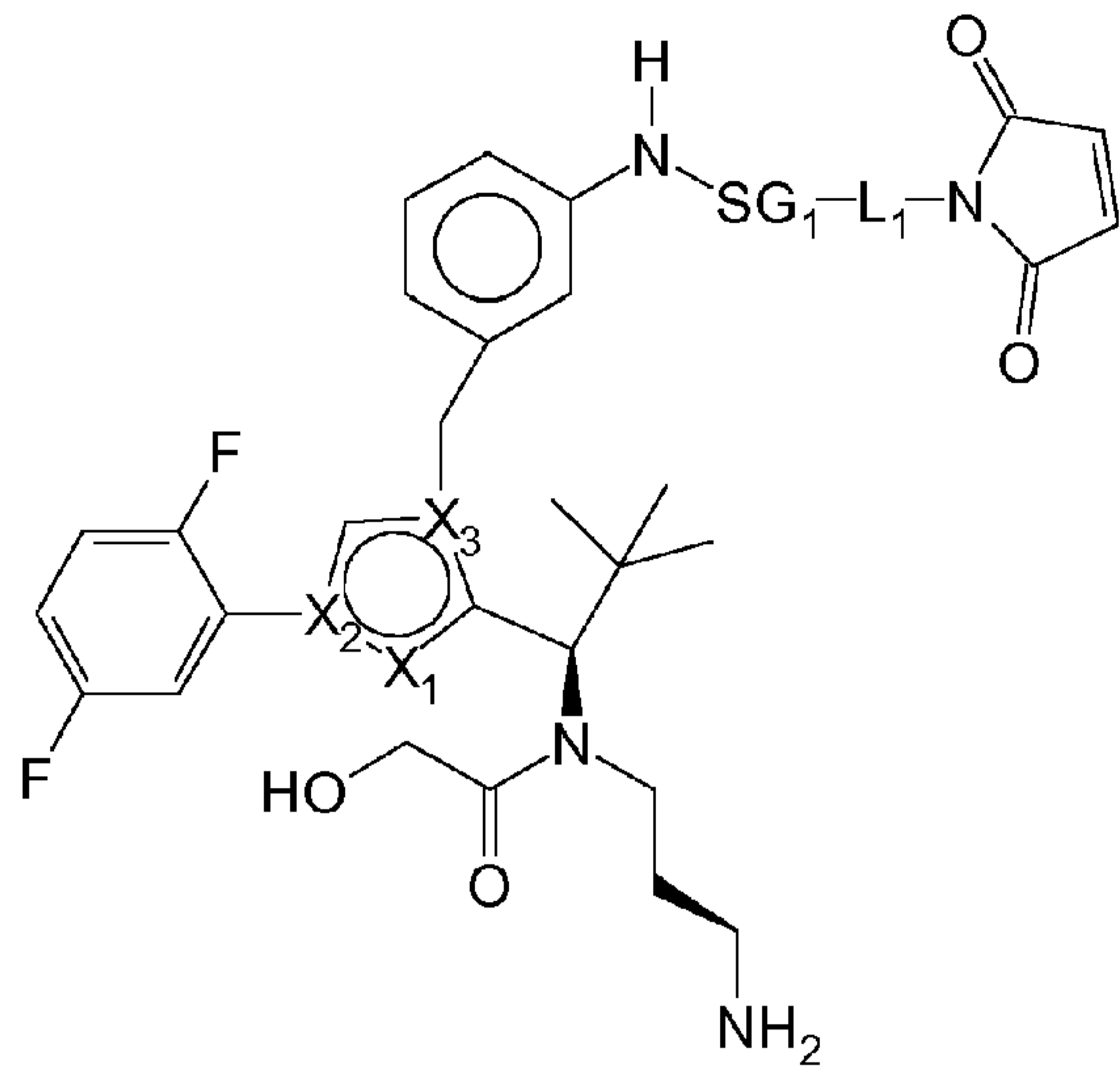












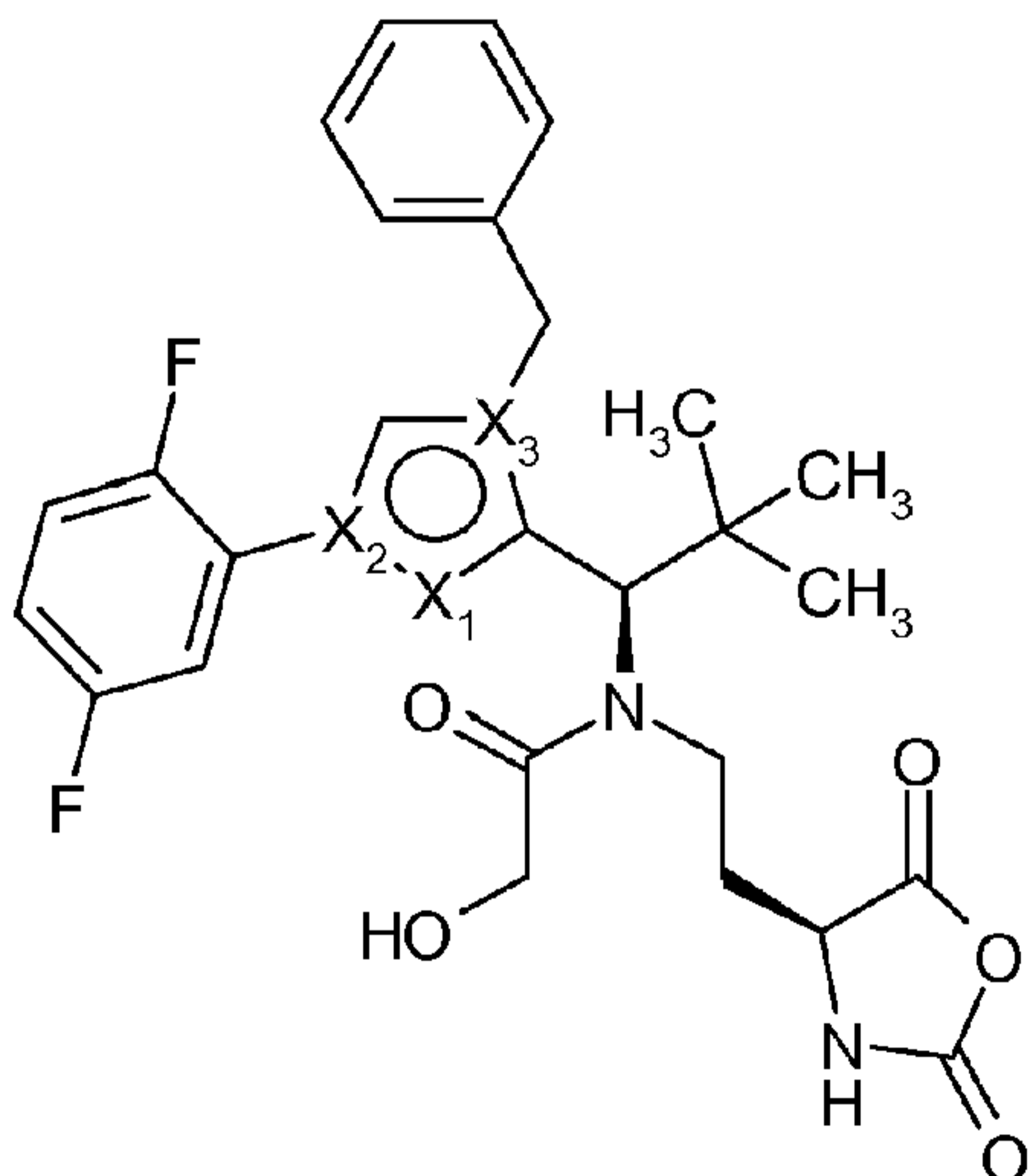
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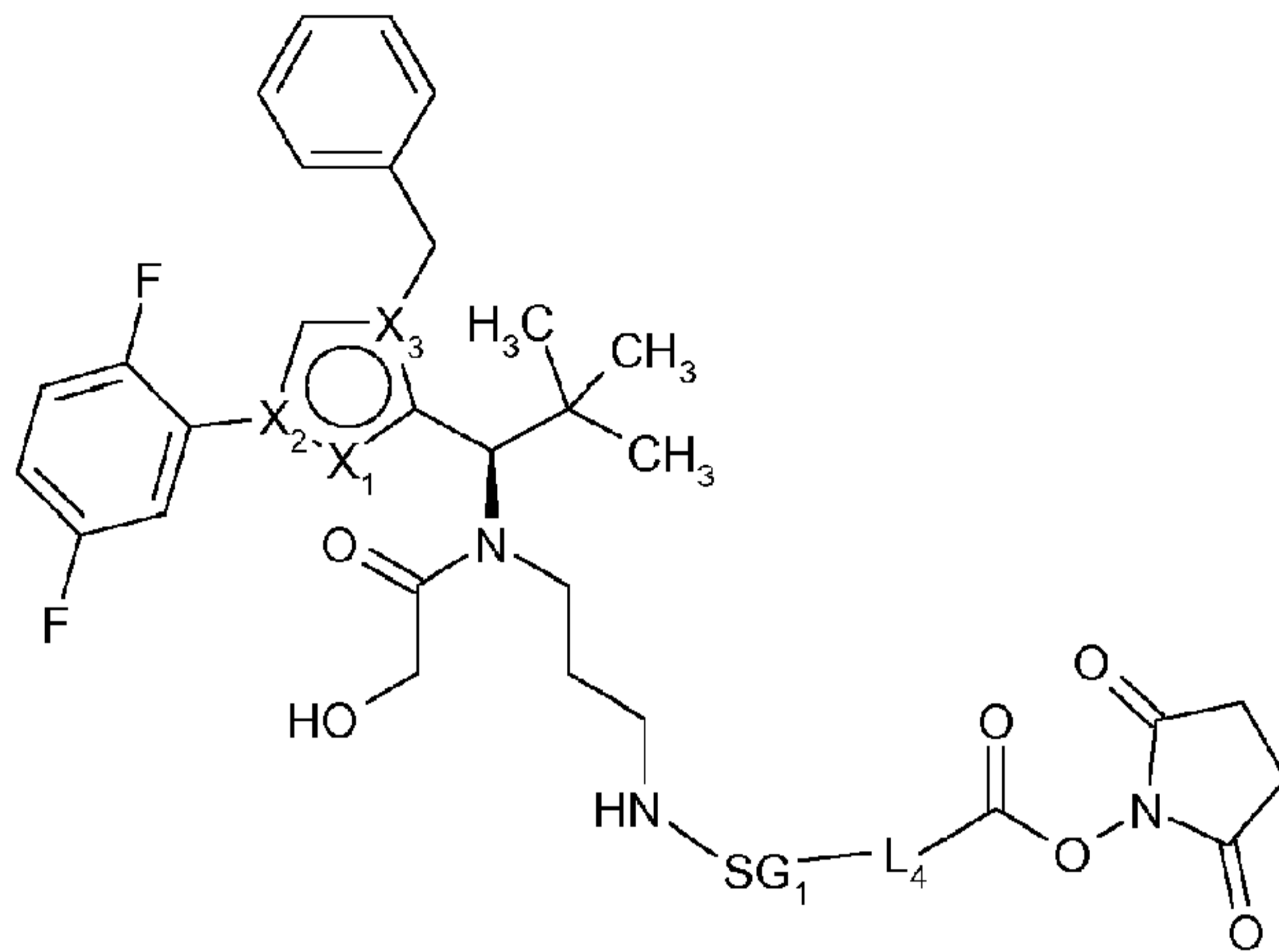
idet R er -H eller -COOH,

idet K er lineært eller forgrenet i givet fald substitueret med C<sub>1</sub>-C<sub>6</sub> alkoxy eller -OH C<sub>1</sub>-C<sub>6</sub>-alkyl, og

5 idet X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub>, SG<sub>1</sub>, L<sub>1</sub>, L<sub>2</sub>, L<sub>3</sub> og L<sub>4</sub> har samme betegnelse som i krav 28 eller 32.

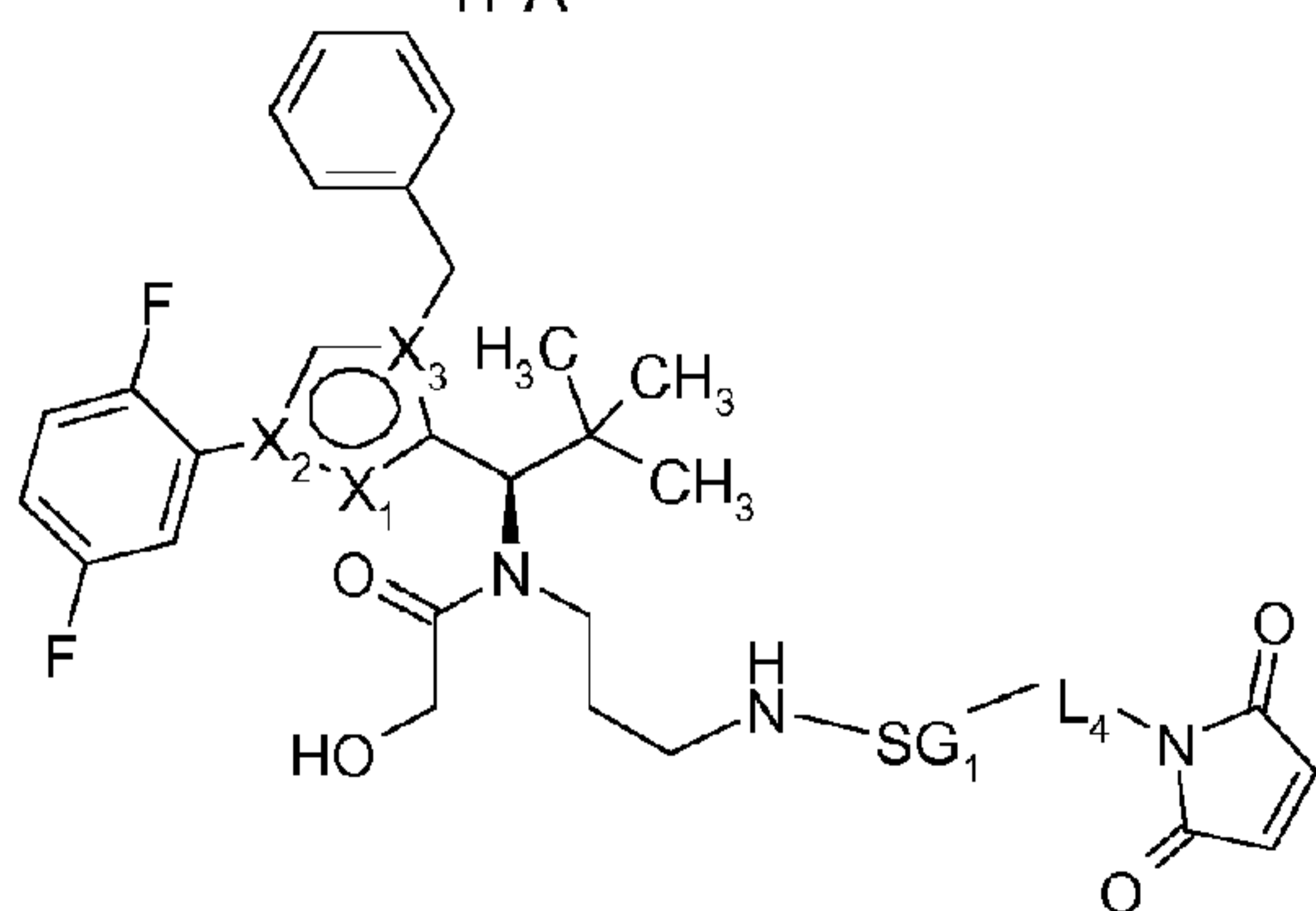
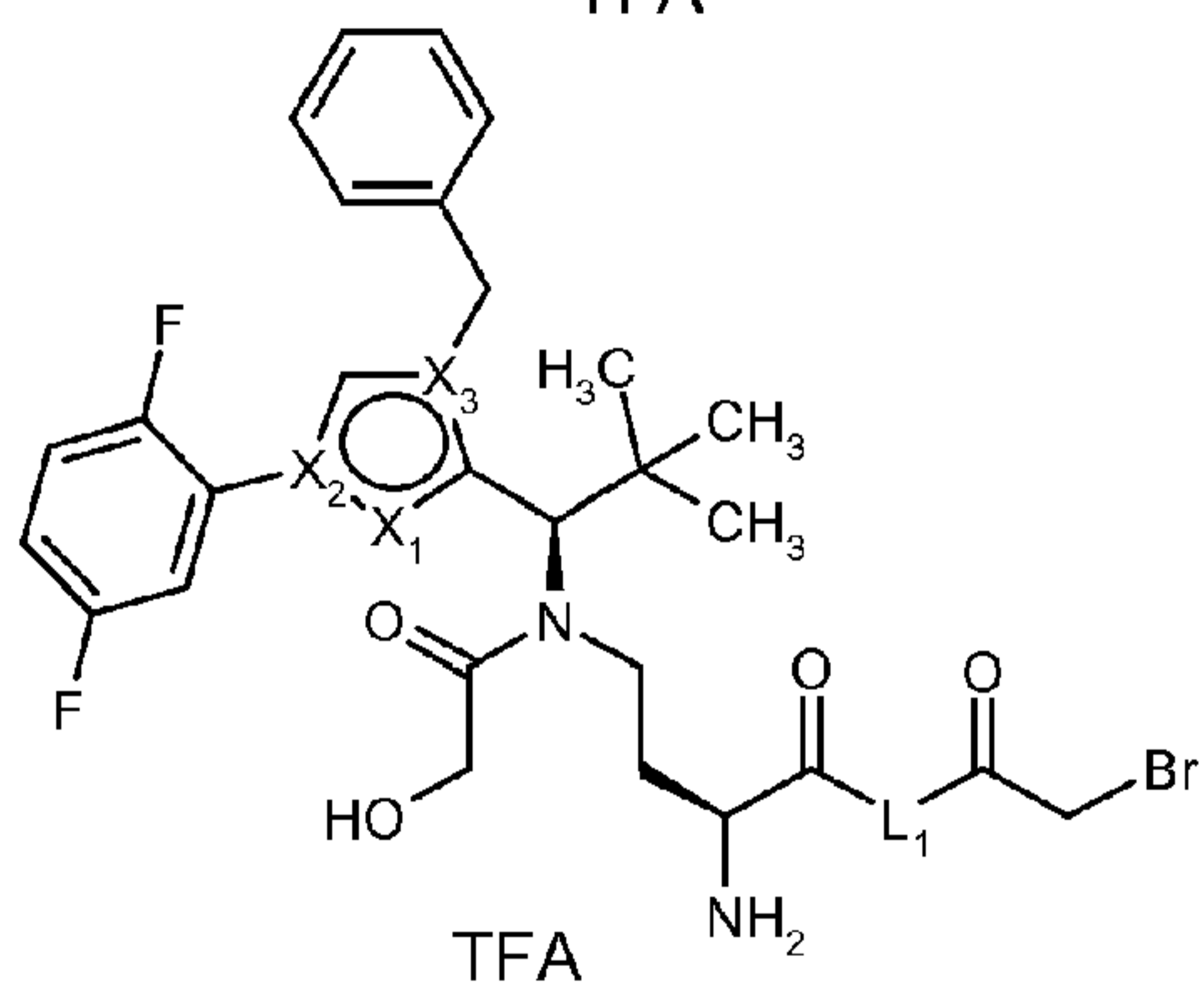
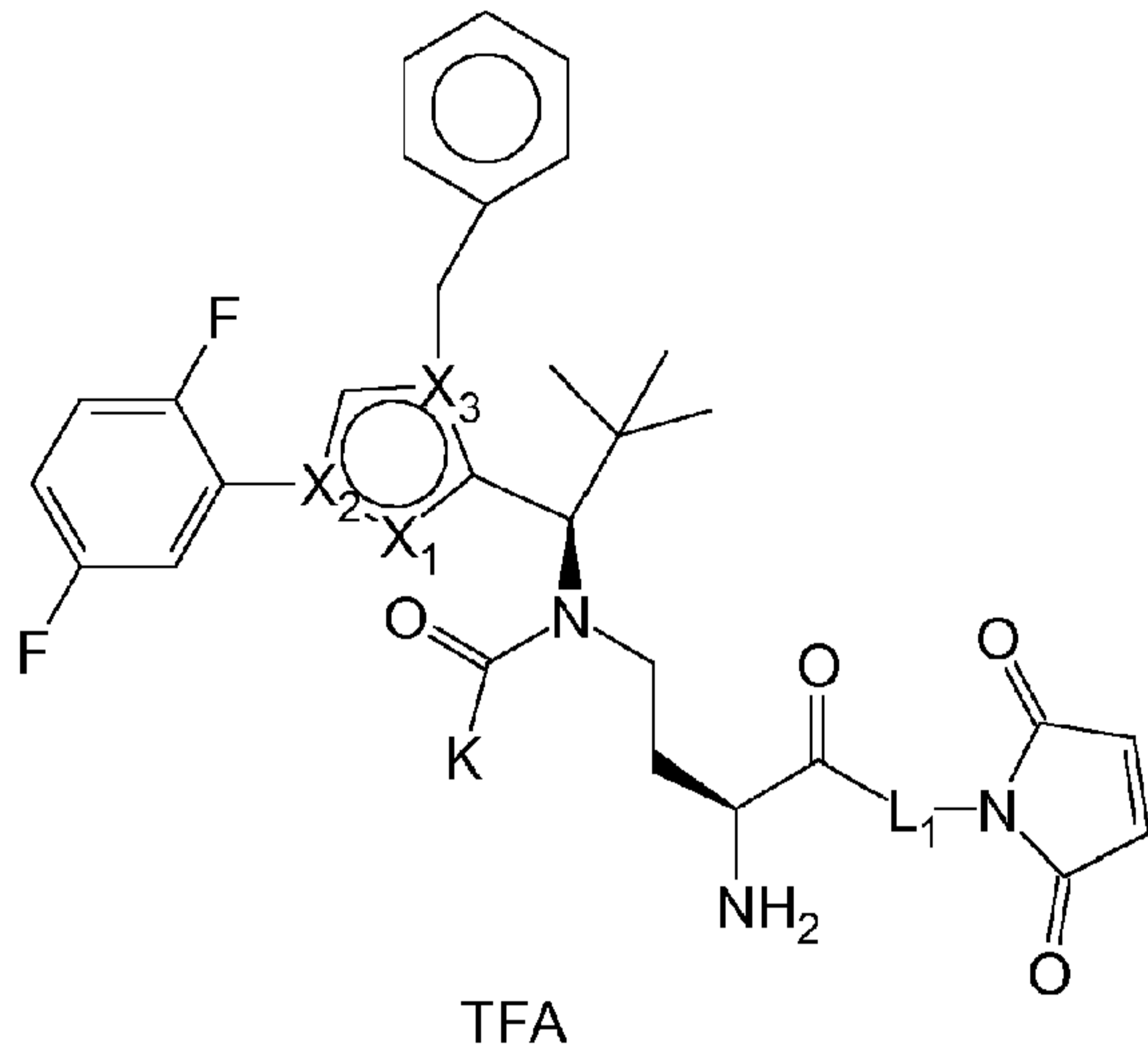
43. Fremgangsmåde til fremstilling af et konjugat ifølge krav 42, idet en forbindelse med en af de følgende formler, 10 fortrinsvis i form af dens trifluoreddikesyresalt, bindes til en lysingruppe af et binderpeptid eller -protein, idet forbindelsen fortrinsvis anvendes i 2- til 12-dobbelt molært overskud i forhold til binderpeptidet eller -proteinet:

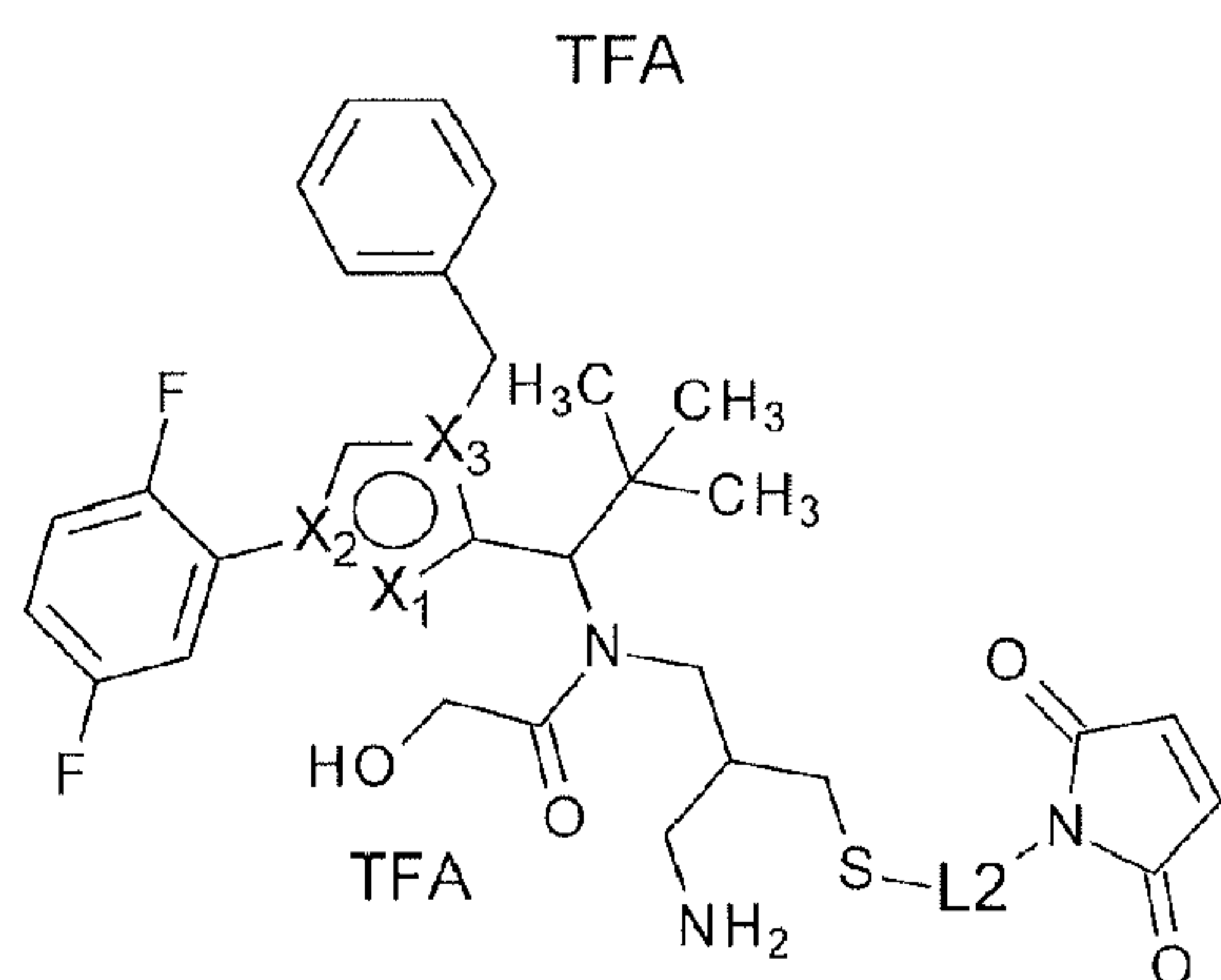
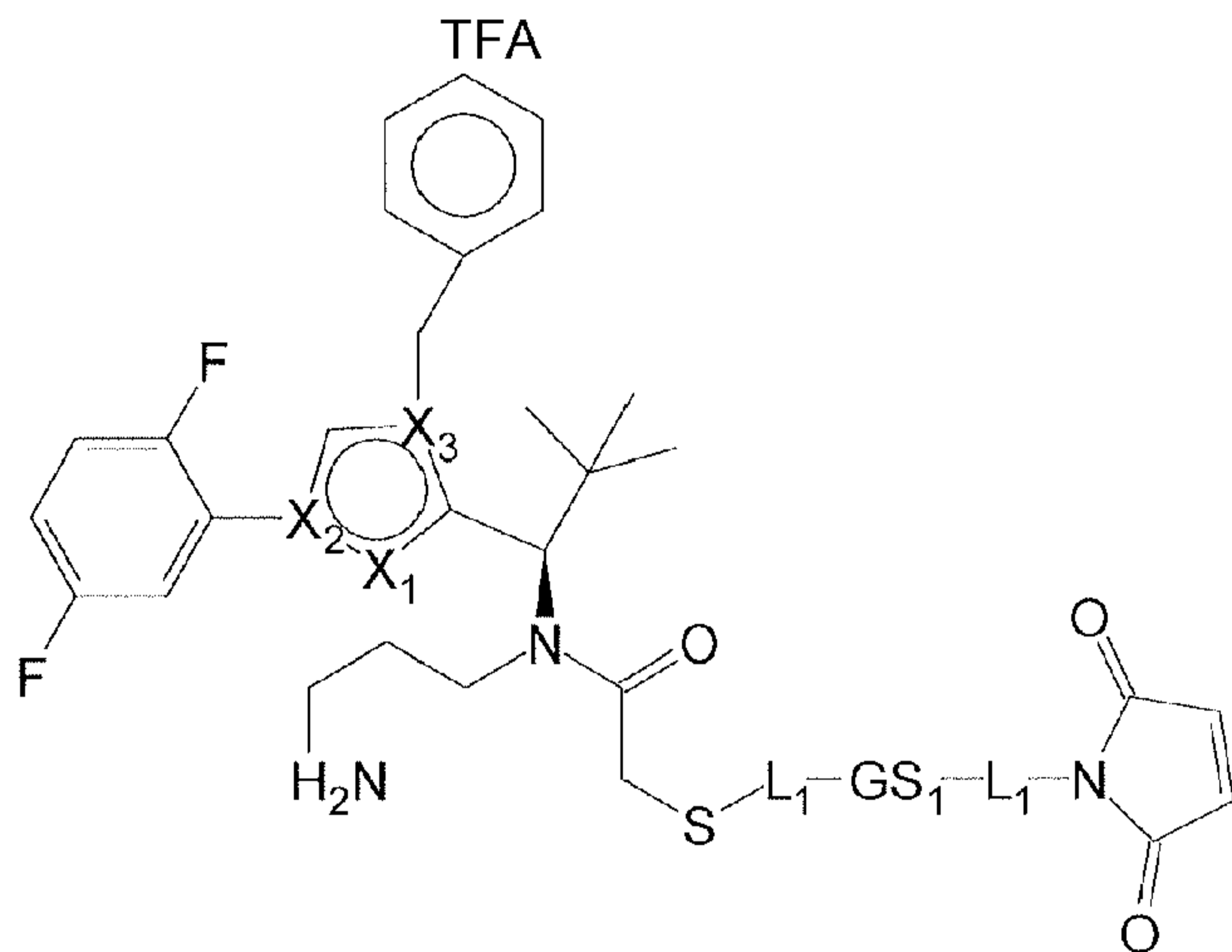
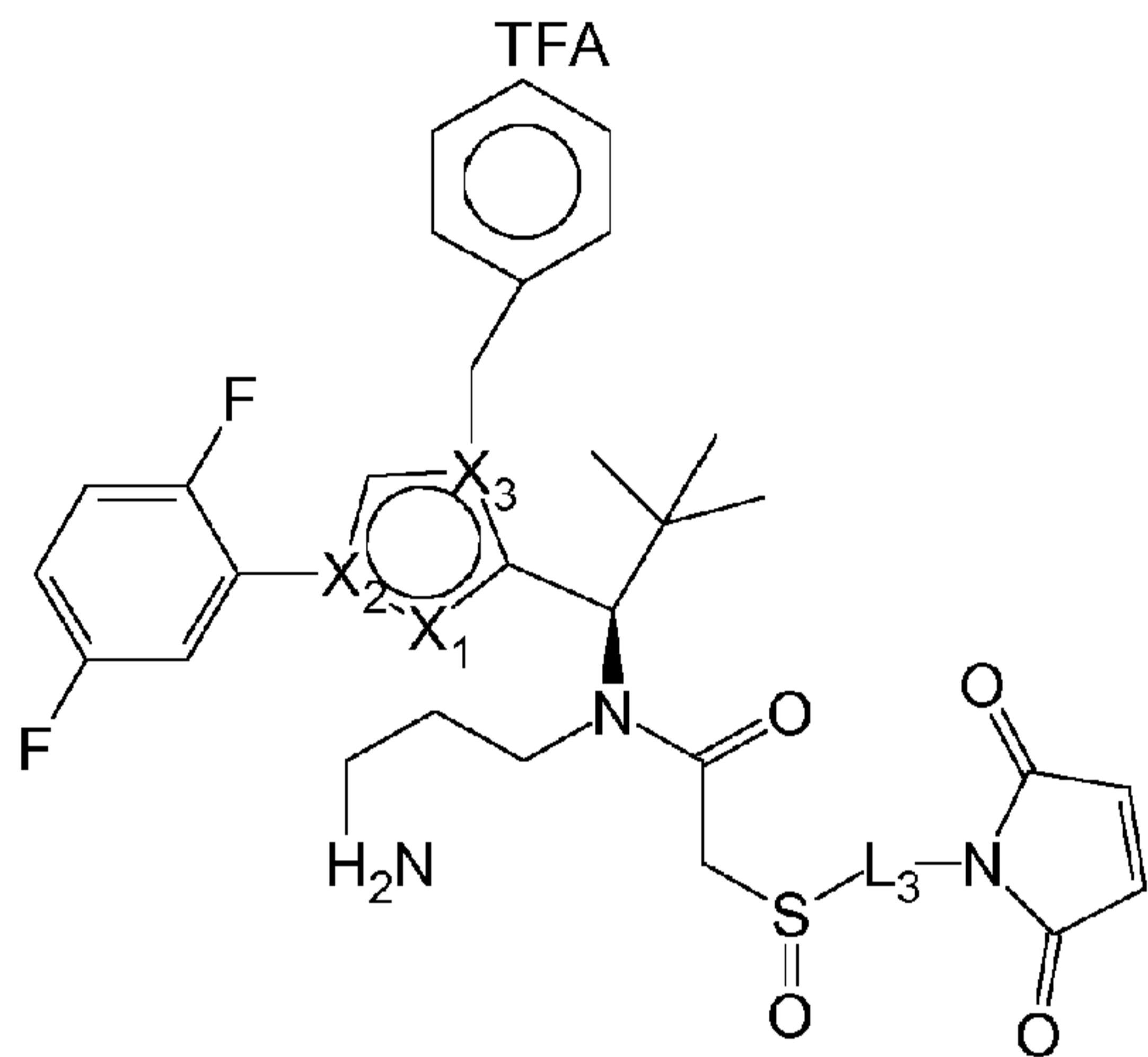
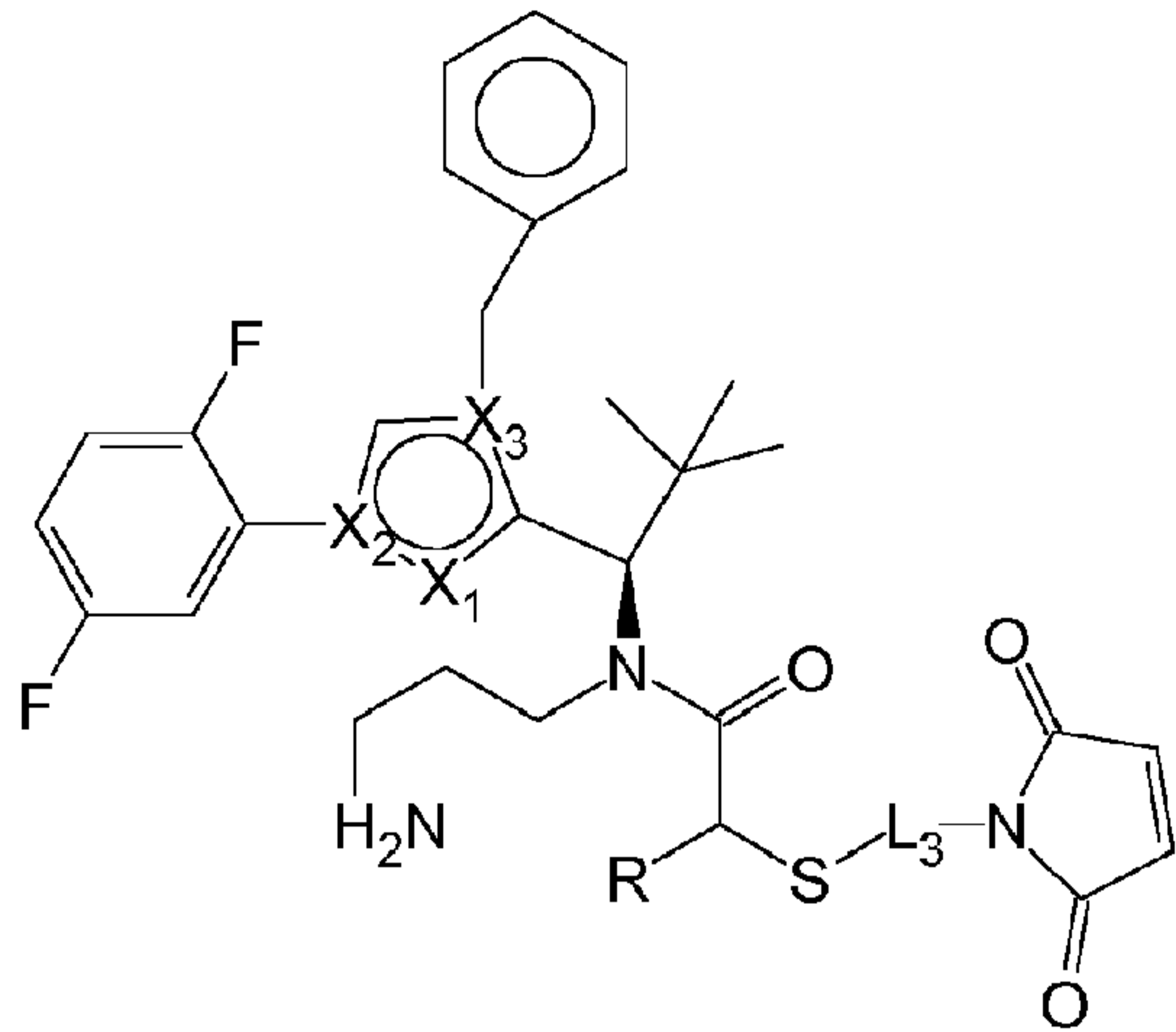


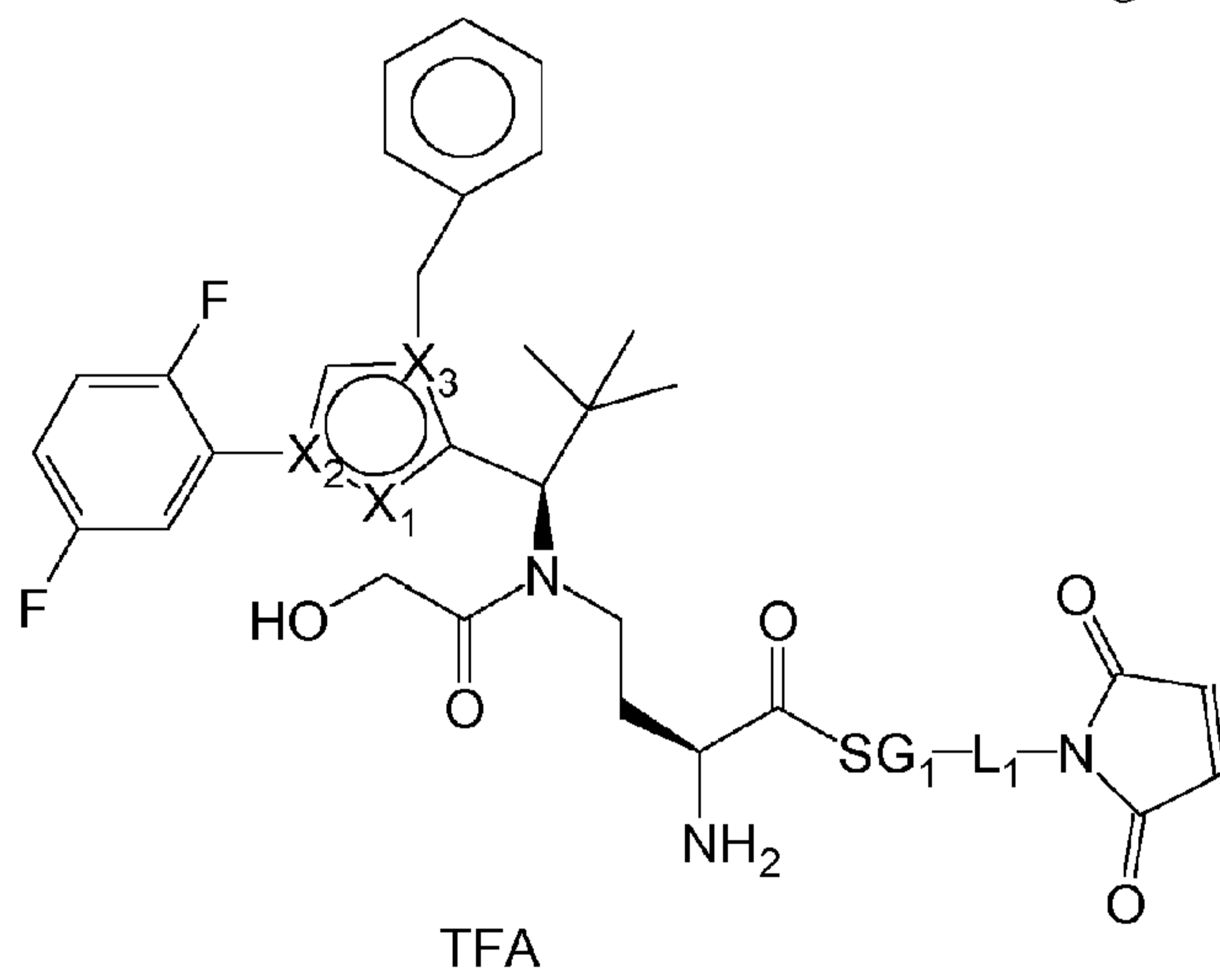
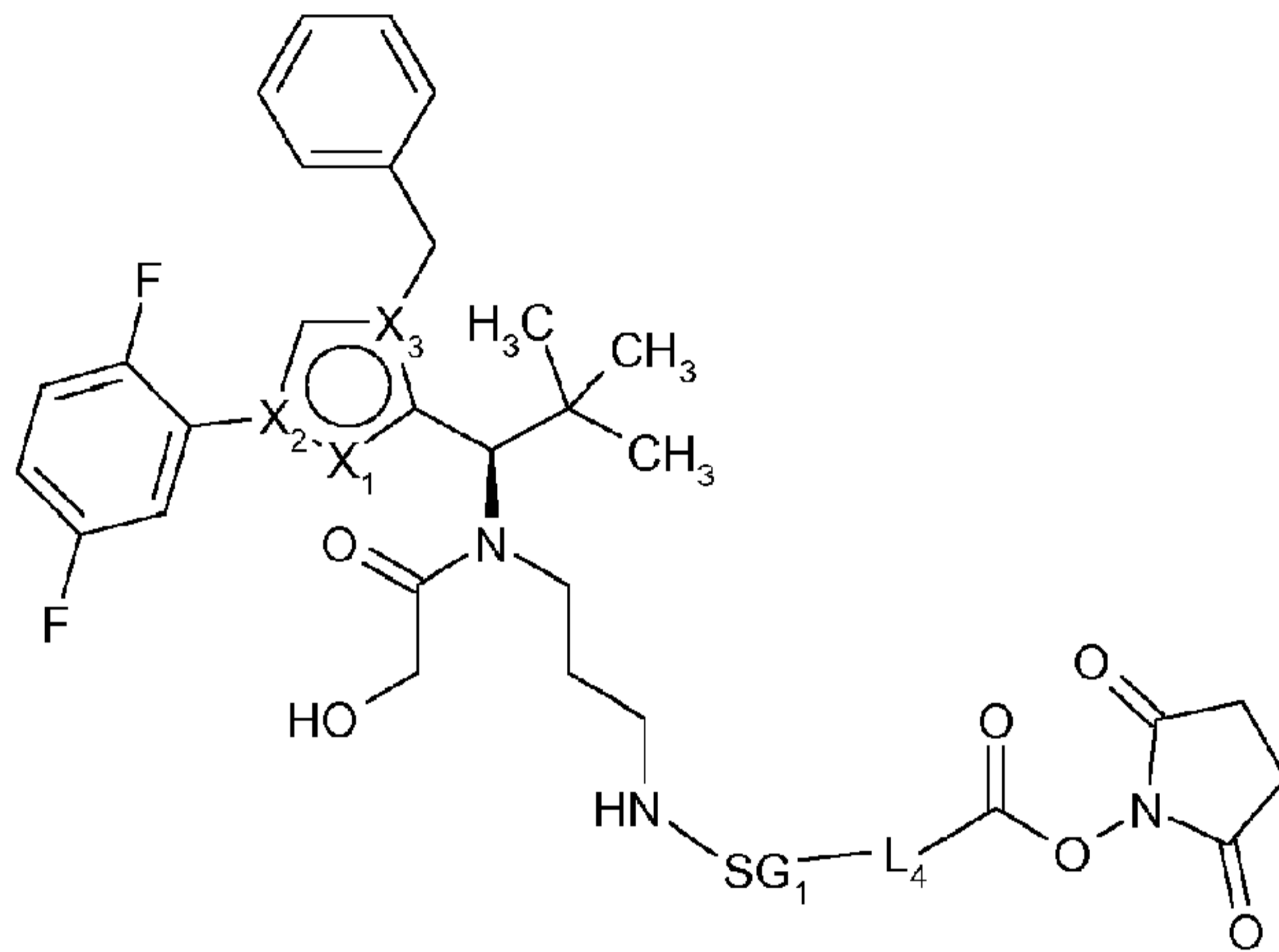
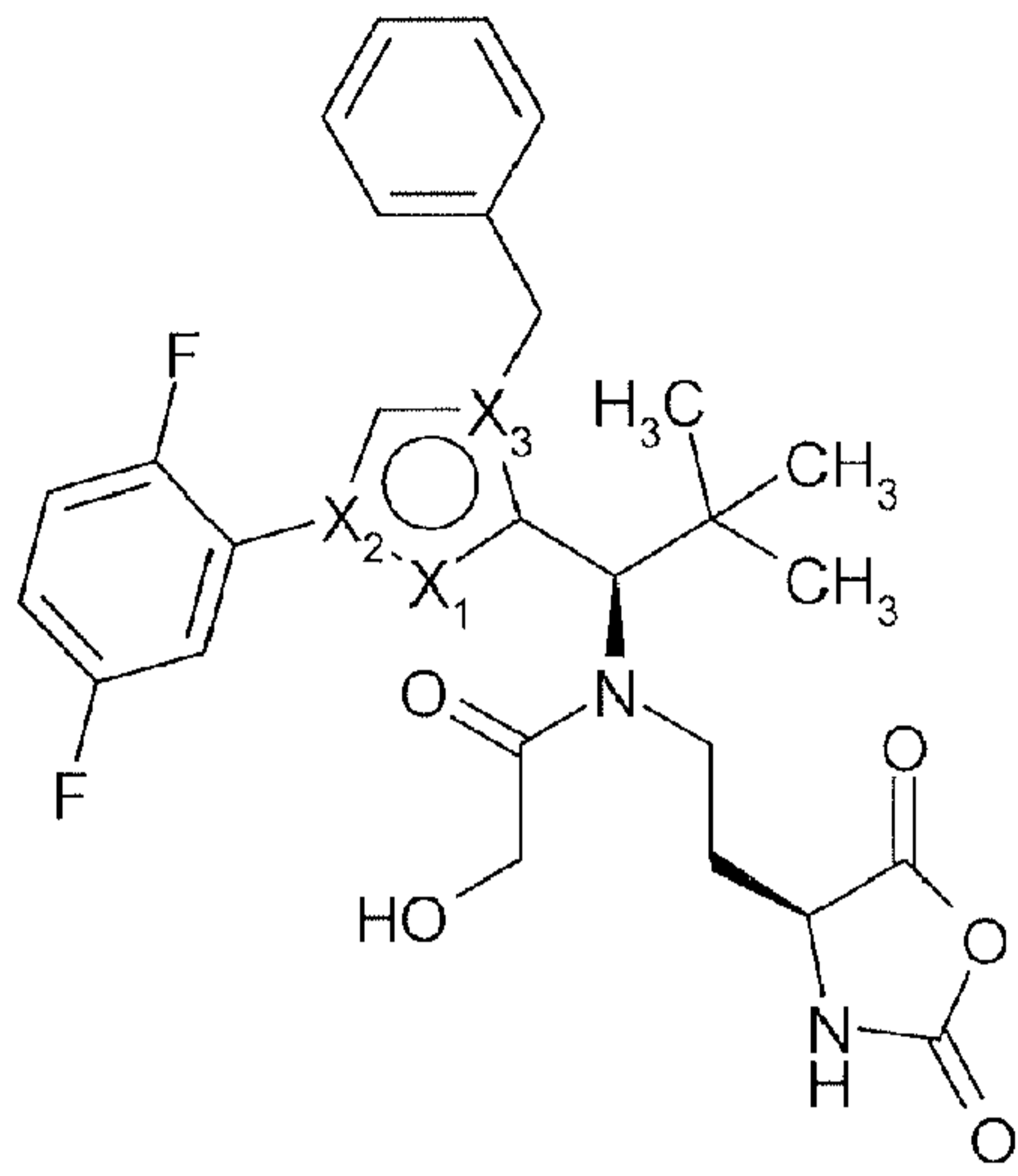


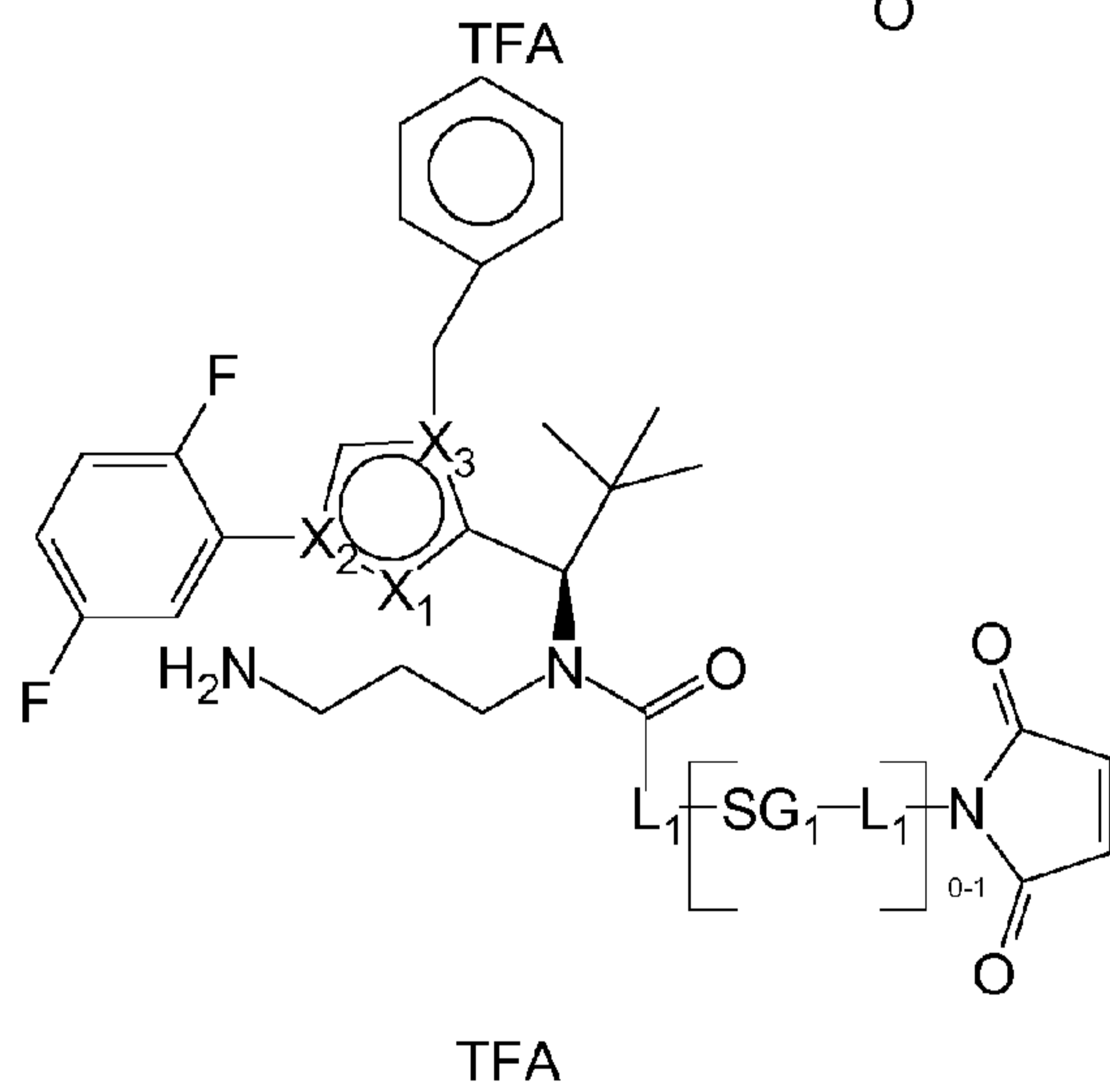
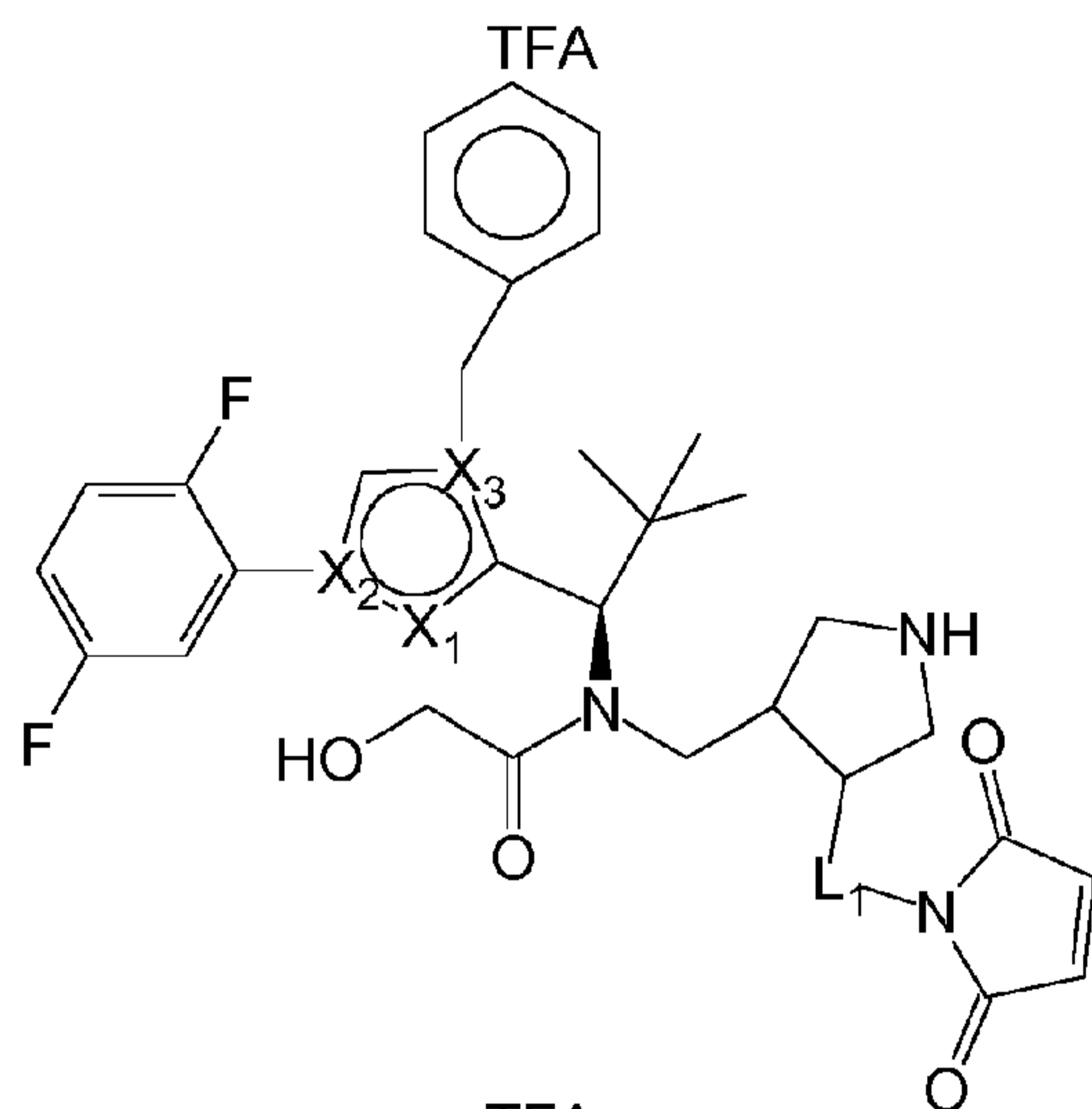
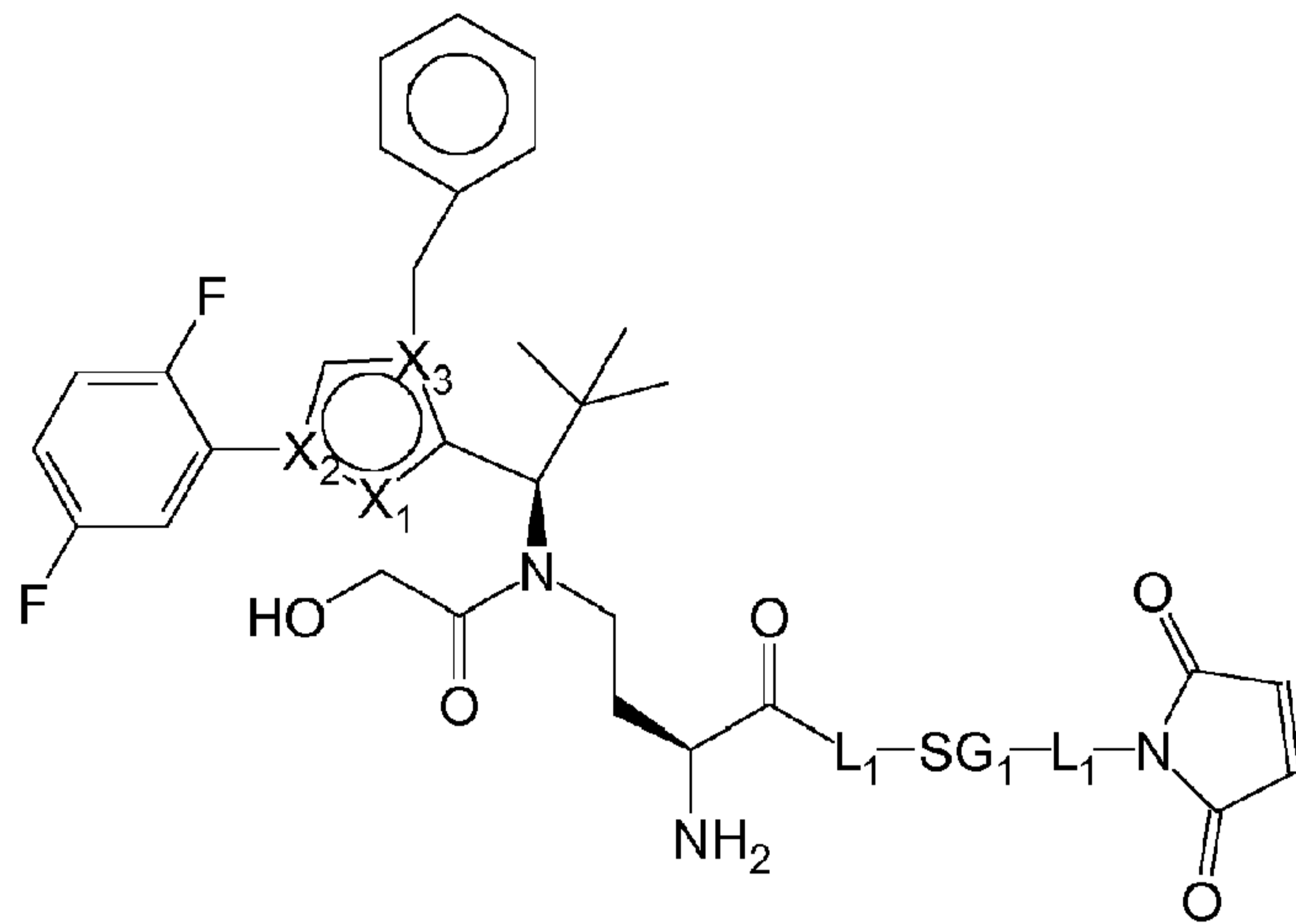
idet  $X_1$ ,  $X_2$ ,  $X_3$ ,  $SG_1$  og  $L_4$  har samme betegnelse som i krav 34.

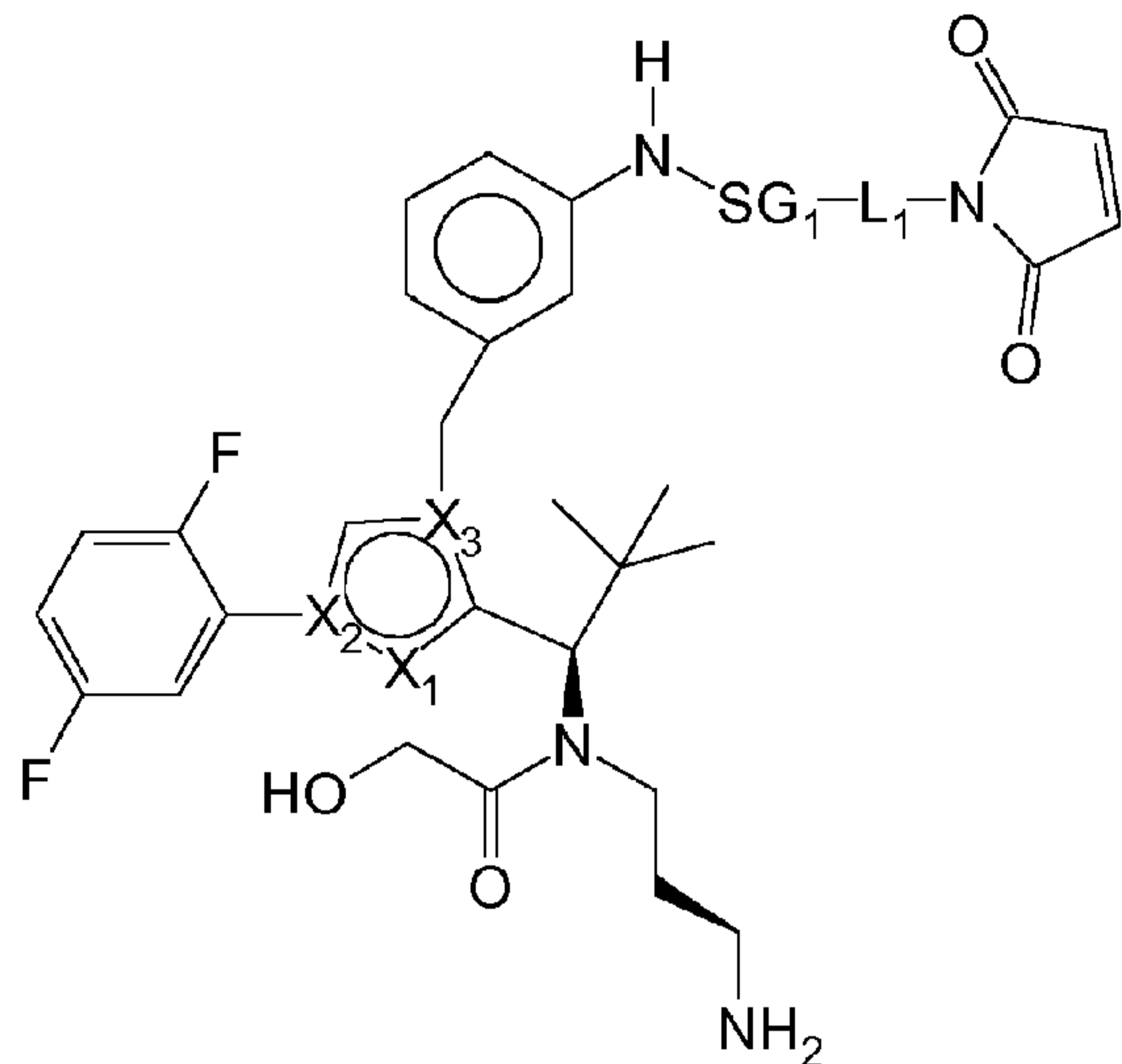
44. Forbindelse med en af de følgende formler:











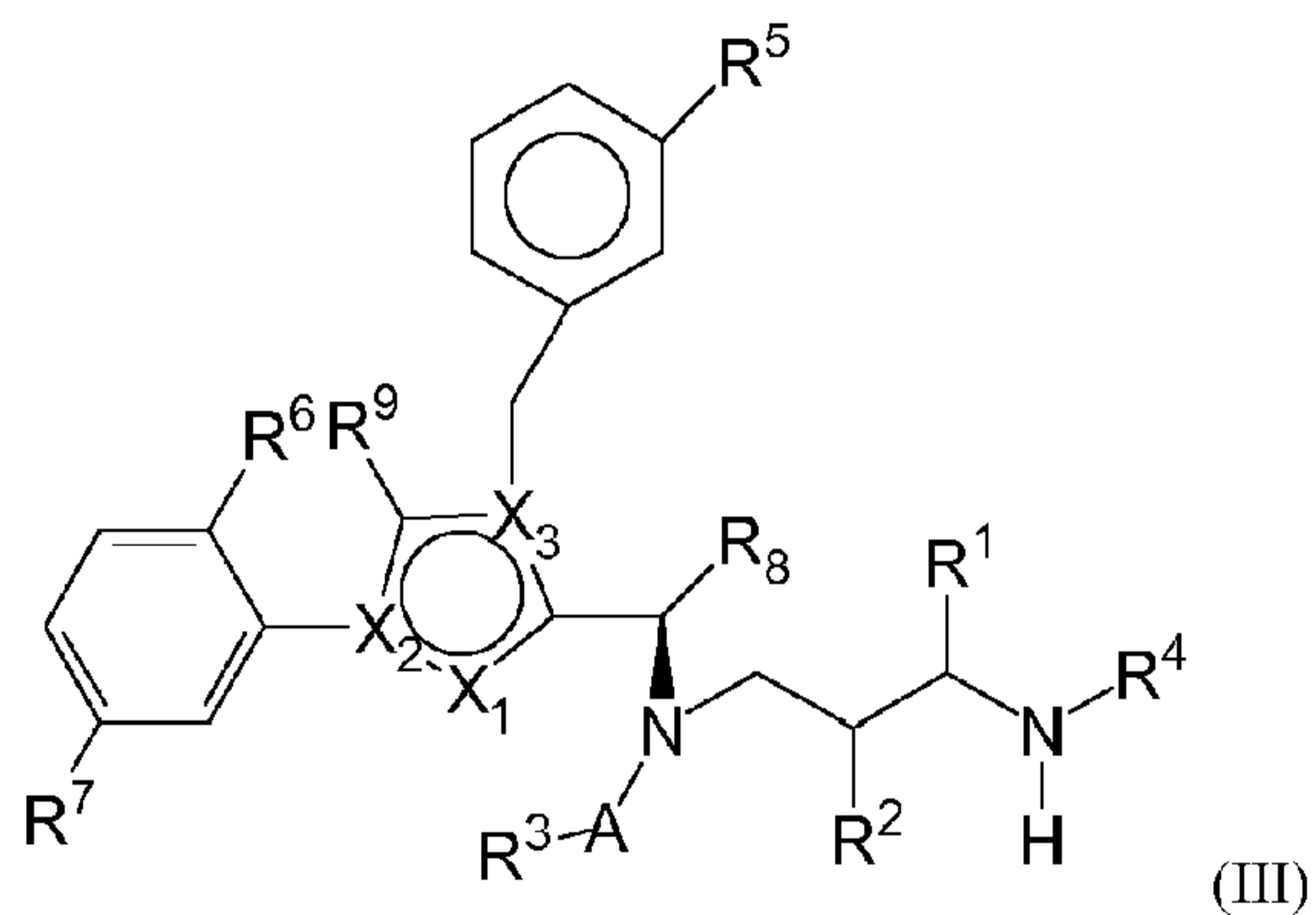
TFA

idet R er -H eller -COOH,

idet K er lineært eller forgrenet C<sub>1</sub>-C<sub>6</sub>-alkyl i givet fald substitueret med C<sub>1</sub>-C<sub>6</sub>-alkoxy eller -OH, og

5 idet X<sub>1</sub>, X<sub>2</sub>, X<sub>3</sub>, SG<sub>1</sub>, L<sub>1</sub>, L<sub>2</sub>, L<sub>3</sub> og L<sub>4</sub> har den samme betegnelse som i krav 24, 28 eller 30.

45. Forbindelser med den almene formel (III):



(III)

10 idet

X<sub>1</sub> er N, X<sub>2</sub> N og X<sub>3</sub> er C, eller X<sub>1</sub> er CH, X<sub>2</sub> C, og X<sub>3</sub> er N, eller X<sub>1</sub> er NH, X<sub>2</sub> er C, og X<sub>3</sub> er C, eller X<sub>1</sub> er CH, X<sub>2</sub> er N, og X<sub>3</sub> er C;

15 R<sup>1</sup> er -L-BINDER, H eller -(CH<sub>2</sub>)<sub>0-3</sub>Z, idet Z er -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup> eller -CO-OY<sup>3</sup>,

idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden er H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z', eller -CH(CH<sub>2</sub>W)Z', og Y<sup>3</sup> er H eller -(CH<sub>2</sub>)<sub>0-3</sub>Z', idet Z' er H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH eller -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH, idet W er H eller OH;

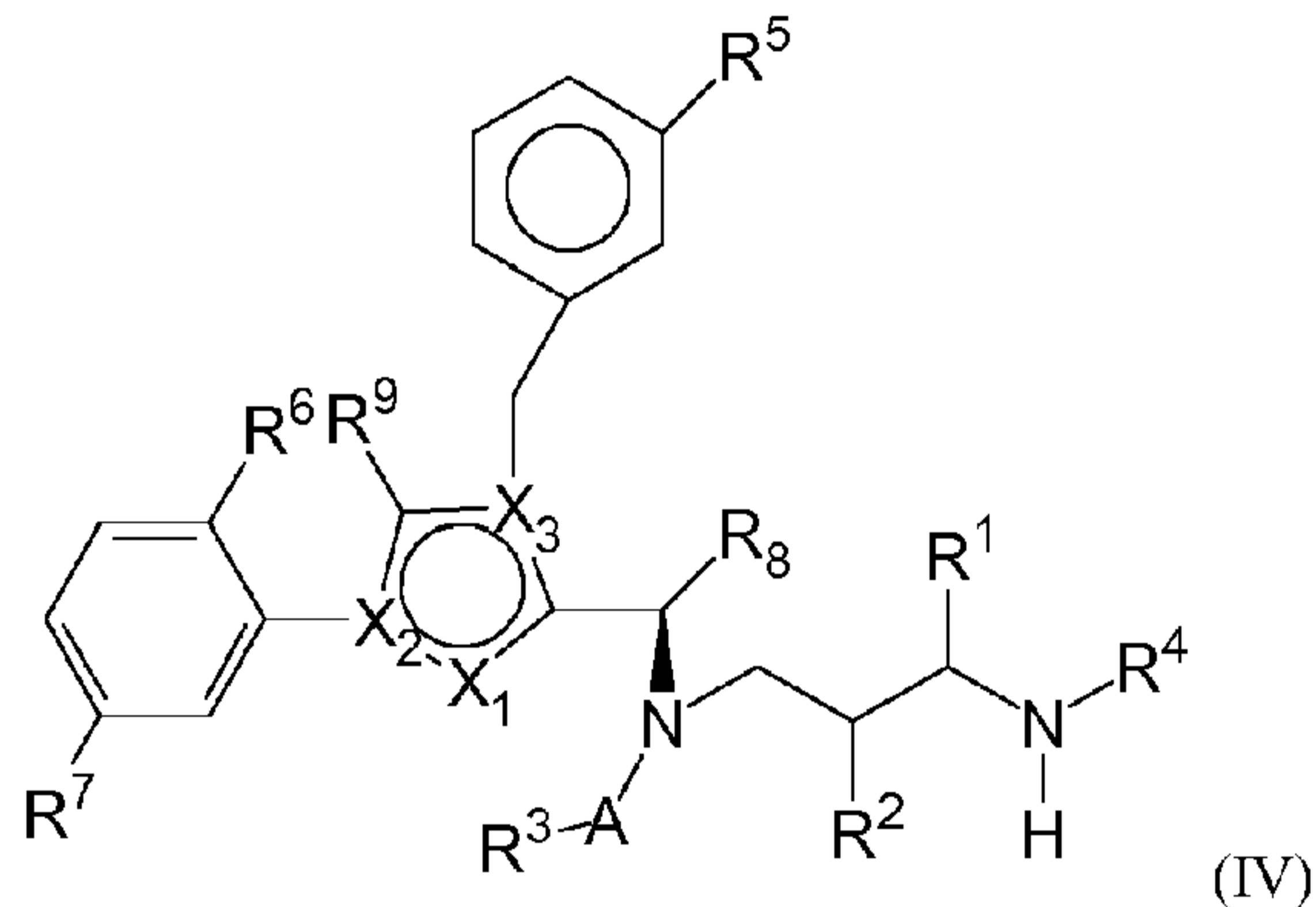
20 idet Y<sup>4</sup> uafhængigt af hinanden er lineært eller forgrenet, i givet fald med -NHCONH<sub>2</sub> substitueret C<sub>1-6</sub> alkyl eller i givet fald

med  $\text{-NH}_2$  substitueret aryl eller benzyl;  
 $R^2$  og  $R^4$  uafhængigt af hinanden er  $\text{-L-BINDER}$ , H,  $\text{-CO-CHY}^4\text{-NHY}^5$   
eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}$ , eller  $R^2$  og  $R^4$  i fællesskab (under dannelse af  
en pyrrolidinring) er  $\text{-CH}_2\text{-CHR}^{10}\text{-}$  eller  $\text{-CHR}^{10}\text{-CH}_2\text{-}$ , idet  $R^{10}$  er  
5 L-#1, H,  $\text{NH}_2$ ,  $\text{SO}_3\text{H}$ ,  $\text{COOH}$ , SH eller OH;  
idet Z er  $\text{-H}$ , halogen,  $\text{-OY}^3$ ,  $\text{-SY}^3$ ,  $\text{NHY}^3$ ,  $\text{-CO-NY}^1\text{Y}^2$  eller  $\text{-CO-OY}^3$ ,  
idet  $\text{Y}^1$  og  $\text{Y}^2$  uafhængigt af hinanden er H,  $\text{NH}_2$  eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}'$ ,  
og  $\text{Y}^3$  er H eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}'$ , idet  $\text{Z}'$  er H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  eller  $\text{COOH}$ ;  
idet  $\text{Y}^4$  uafhængigt af hinanden er lineært eller forgrenet, i  
10 givet fald med  $\text{-NHCONH}_2$  substitueret  $\text{C}_{1-6}$  alkyl eller i givet fald  
med  $\text{-NH}_2$  substitueret aryl eller benzyl, og  $\text{Y}^5$  er H eller  $\text{-CO-}$   
 $\text{CHY}^6\text{-NH}_2$ , idet  $\text{Y}^6$  er lineært eller forgrenet  $\text{C}_{1-6}$  alkyl;  
A er CO, SO,  $\text{SO}_2$ ,  $\text{SO}_2\text{NH}$  eller CNNH;  
 $R^3$  er  $\text{-L-BINDER}$  eller en i givet fald substitueret alkyl-, aryl-  
15 , heteroaryl- heteroalkyl-, heterocycloalkylgruppe, fortrinsvis  
 $\text{-L-#1}$  eller en  $\text{C}_{1-10}$ -alkyl-,  $\text{C}_{6-10}$ -aryl- eller  $\text{C}_{6-10}$ -aralkyl-,  $\text{C}_{5-}$   
 $_{10}$ -heteroalkyl-,  $\text{C}_{1-10}$ -alkyl-O- $\text{C}_{6-10}$ -aryl- eller  $\text{C}_{5-10}$ -  
heterocycloalkylgruppe, som kan være substitueret med 1-3  $\text{-OH-}$   
grupper, 1-3 halogenatomer, 1-3 halogenerede alkylgrupper (som  
20 hver især har 1-3 halogenatomer), 1-3 O-alkylgrupper, 1-3  $\text{-SH-}$   
grupper, 1-3  $\text{-S-alkylgrupper}$ , 1-3  $\text{-O-CO-alkylgrupper}$ , 1-3  $\text{-O-}$   
 $\text{CO-NH-alkylgrupper}$ , 1-3  $\text{-NH-CO-alkylgrupper}$ , 1-3  $\text{-NH-CO-NH-}$   
alkylgrupper, 1-3  $\text{-S(O)}_n\text{-alkylgrupper}$ , 1-3  $\text{-SO}_2\text{-NH-alkylgrupper}$ ,  
1-3  $\text{-NH-alkylgrupper}$ , 1-3  $\text{-N(alkyl)}_2\text{-grupper}$ , 1-3  $\text{-NH}_2\text{-grupper}$   
25 eller 1-3  $\text{-(CH}_2\text{)}_{0-3}\text{Z-}$ grupper, idet Z er  $\text{-H}$ , halogen,  $\text{-OY}^3$ ,  $\text{-SY}^3$ ,  
 $\text{-NHY}^3$ ,  $\text{-CO-NY}^1\text{Y}^2$  eller  $\text{-CO-OY}^3$ ,  $\text{Y}^1$ , og  $\text{Y}^2$  uafhængigt af hinanden  
er H,  $\text{NH}_2$ , eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}'$ , og  $\text{Y}^3$  er H,  $\text{-(CH}_2\text{)}_{0-3}\text{-CH(NHCOCH}_3\text{)Z}'$ ,  
 $\text{-(CH}_2\text{)}_{0-3}\text{-CH(NH}_2\text{)Z}'$  eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}'$ , idet  $\text{Z}'$  er H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  eller  
 $\text{COOH}$ ,  
30 (idet "alkyl" fortrinsvis betegner  $\text{C}_{1-10}$ -alkyl);  
 $R^5$  er  $\text{-L-BINDER}$ , H, F,  $\text{NH}_2$ ,  $\text{NO}_2$ , halogen, SH eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}$ ,  
idet Z er  $\text{-H}$ , halogen,  $\text{-OY}^3$ ,  $\text{-SY}^3$ ,  $\text{-NHY}^3$ ,  $\text{-CO-NY}^1\text{Y}^2$  eller  $\text{-CO-}$   
 $\text{OY}^3$ ,  
idet  $\text{Y}^1$  og  $\text{Y}^2$  uafhængigt af hinanden er H,  $\text{NH}_2$  eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}'$ ,  
35 og  $\text{Y}^3$  er H eller  $\text{-(CH}_2\text{)}_{0-3}\text{Z}'$ , idet  $\text{Z}'$  er H,  $\text{SO}_3\text{H}$ ,  $\text{NH}_2$  eller  $\text{COOH}$ ;  
idet L står for en linker, og BINDER står for binder eller  
derivat heraf, idet binderen i givet fald kan være bundet til  
flere aktivstofmolekyler,



R<sup>6</sup> og R<sup>7</sup> uafhængigt af hinanden er H, cyano, (i givet fald fluoreret) C<sub>1-10</sub>-alkyl, (i givet fald fluoreret) C<sub>2-10</sub>-alkenyl, (i givet fald fluoreret) C<sub>2-10</sub>-alkynyl, hydroxy eller halogen, R<sup>8</sup> er (i givet fald fluoreret) C<sub>1-10</sub>-alkyl, (i givet fald fluoreret) C<sub>4-10</sub>-cycloalkyl eller eventuelt substitueret oxetan; og R<sup>9</sup> er H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F eller CHF<sub>2</sub>; samt deres salte, solvater samt salte af solvaterne.

10 46. Forbindelser med den almene formel (IV):



idet

X<sub>1</sub> er CH, X<sub>2</sub> C og X<sub>3</sub> N;

R<sup>1</sup> er -L-BINDER, H eller -(CH<sub>2</sub>)<sub>0-3</sub>Z, idet Z er -H, -NHY<sup>3</sup>, -OY<sup>3</sup>, -SY<sup>3</sup>, halogen, -CO-NY<sup>1</sup>Y<sup>2</sup> eller -CO-OY<sup>3</sup>,

idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden er H, NH<sub>2</sub>, -(CH<sub>2</sub>CH<sub>2</sub>O)<sub>0-3</sub>-(CH<sub>2</sub>)<sub>0-3</sub>Z' eller -CH(CH<sub>2</sub>W)Z', og Y<sup>3</sup> er H eller -(CH<sub>2</sub>)<sub>0-3</sub>Z', idet Z' er H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, -NH-CO-CH<sub>2</sub>-CH<sub>2</sub>-CH(NH<sub>2</sub>)COOH eller -(CO-NH-CHY<sup>4</sup>)<sub>1-3</sub>COOH; idet W er H eller OH;

20 idet Y<sup>4</sup> uafhængigt af hinanden er lineært eller forgrenet, i givet fald med -NHCONH<sub>2</sub> substitueret C<sub>1-6</sub> alkyl eller i givet fald med -NH<sub>2</sub> substitueret aryl eller benzyl;

R<sup>2</sup> og R<sup>4</sup> uafhængigt af hinanden er -L-BINDER, H, -CO-CHY<sup>4</sup>-NHY<sup>5</sup> eller -(CH<sub>2</sub>)<sub>0-3</sub>Z, eller R<sup>2</sup> og R<sup>4</sup> i fællesskab (under dannelse af en pyrrolidinring) er -CH<sub>2</sub>-CHR<sup>10</sup>- eller -CHR<sup>10</sup>-CH<sub>2</sub>-, idet R<sup>10</sup> er L-#1, H, NH<sub>2</sub>, SO<sub>3</sub>H, COOH, SH eller OH; idet Z er -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> eller -CO-OY<sup>3</sup>,

idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden er H, NH<sub>2</sub> eller -(CH<sub>2</sub>)<sub>0-3</sub>Z', og Y<sup>3</sup> er H eller -(CH<sub>2</sub>)<sub>0-3</sub>Z', idet Z' er H, SO<sub>3</sub>H, NH<sub>2</sub> eller COOH;

30 idet Y<sup>4</sup> uafhængigt af hinanden er lineært eller forgrenet, i givet fald med -NHCONH<sub>2</sub> substitueret C<sub>1-6</sub> alkyl eller i givet

fald med  $-NH_2$  substitueret aryl eller benzyl, og  $Y^5$  er H eller  $-CO-CHY^6-NH_2$ , idet  $Y^6$  er lineært eller forgrenet  $C_{1-6}$  alkyl;  
A er CO, SO,  $SO_2$ ,  $SO_2NH$  eller CNNH;  
 $R^3$  er -L-BINDER eller en i givet fald substitueret alkyl-, aryl-,  
5 , heteroaryl- heteroalkyl-, heterocycloalkylgruppe,  
fortrinsvis -L-#1 eller en  $C_{1-10}$ -alkyl-,  $C_{6-10}$ -aryl- eller  $C_{6-10}$ -  
aralkyl-,  $C_{5-10}$ -heteroalkyl-,  $C_{1-10}$ -alkyl-O- $C_{6-10}$ -aryl- eller  $C_{5-10}$ -  
heterocycloalkylgruppe,  
som kan være substitueret med 1-3 -OH-grupper, 1-3  
10 halogenatomer, 1-3 halogenerede alkylgrupper (som hver især har  
1-3 halogenatomer), 1-3 O-alkylgrupper, 1-3 -SH-grupper, 1-3 -  
S-alkylgrupper, 1-3 -O-CO-alkylgrupper, 1-3 -O-CO-NH-  
alkylgrupper, 1-3 -NH-CO-alkylgrupper, 1-3-NH-CO-NH-  
alkylgrupper, 1-3 -S(O)<sub>n</sub>-alkylgrupper, 1-3 -SO<sub>2</sub>-NH-alkylgrupper,  
15 1-3-NH-alkylgrupper, 1-3 -N(alkyl)<sub>2</sub>-grupper, 1-3 -NH<sub>2</sub>-grupper  
eller 1-3 -(CH<sub>2</sub>)<sub>0-3</sub>Z-grupper, idet Z er -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>,  
-NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> eller -CO-OY<sup>3</sup>, idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af  
hinanden er H, NH<sub>2</sub> eller -(CH<sub>2</sub>)<sub>0-3</sub>Z', og Y<sup>3</sup> er H, -(CH<sub>2</sub>)<sub>0-3</sub>-  
CH(NHCOCH<sub>3</sub>)Z', -(CH<sub>2</sub>)<sub>0-3</sub>-CH(NH<sub>2</sub>)Z' eller -(CH<sub>2</sub>)<sub>0-3</sub>Z', idet Z' er H,  
20 SO<sub>3</sub>H, NH<sub>2</sub> eller COOH,  
(idet "alkyl" fortrinsvis betegner  $C_{1-10}$ -alkyl);  
 $R^5$  er -L-BINDER, H, F, NH<sub>2</sub>, NO<sub>2</sub>, halogen, SH eller -(CH<sub>2</sub>)<sub>0-3</sub>Z,  
idet Z er -H, halogen, -OY<sup>3</sup>, -SY<sup>3</sup>, -NHY<sup>3</sup>, -CO-NY<sup>1</sup>Y<sup>2</sup> eller -CO-  
OY<sup>3</sup>,  
25 idet Y<sup>1</sup> og Y<sup>2</sup> uafhængigt af hinanden er H, NH<sub>2</sub> eller -(CH<sub>2</sub>)<sub>0-3</sub>Z',  
og Y<sup>3</sup> er H eller -(CH<sub>2</sub>)<sub>0-3</sub>Z', idet Z' er H, SO<sub>3</sub>H, NH<sub>2</sub> eller COOH;  
idet L står for en linker, og BINDER står for binder eller  
derivat heraf, idet binderen kan være bundet til flere  
aktivstofmolekyler,  
30  $R^6$  og  $R^7$  uafhængigt af hinanden er H, cyano, (i givet fald  
fluoreret)  $C_{1-10}$ -alkyl, (i givet fald fluoreret)  $C_{2-10}$ -alkenyl, (i  
givet fald fluoreret)  $C_{2-10}$ -alkynyl, hydroxy eller halogen,  
 $R^8$  er (i givet fald fluoreret)  $C_{1-10}$ -alkyl, (i givet fald  
fluoreret)  $C_{4-10}$ -cycloalkyl eller eventuelt substitueret oxetan;  
35 og  
 $R^9$  er H, F, CH<sub>3</sub>, CF<sub>3</sub>, CH<sub>2</sub>F eller CHF<sub>2</sub>;  
samt deres salte, solvater samt salte af solvaterne.

47. Forbindelse ifølge krav 46, idet Z er Cl eller Br.
48. Forbindelse ifølge krav 46, idet  $R^1$  er  $-(CH_2)_{0-3}Z$ , idet Z er  $-CO-NY^1Y^2$ , idet  $Y^2$  er  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$ , og  $Y^1$  er H,  $NH_2$  eller  
5  $-(CH_2CH_2O)_{0-3}-(CH_2)_{0-3}Z'$ .
49. Forbindelse ifølge et eller flere af kravene 46-48, idet  $Y^1$  er H,  $Y^2$  er  $-(CH_2CH_2O)_3-CH_2CH_2Z'$ , og  $Z'$  er  $-COOH$ .
- 10 50. Forbindelse ifølge et eller flere af kravene 46-48, idet  $Y^1$  er H,  $Y^2$  er  $-CH_2CH_2Z'$ , og  $Z'$  er  $-(CONHCHY^4)_2COOH$ .
51. Forbindelse ifølge krav 46, idet  $Y^4$  er lineært eller forgrenet, i givet fald med  $-NHCONH_2$  substitueret  $C_{1-6}$  alkyl.  
15
52. Forbindelse ifølge krav 46, idet der vælges i det mindste en repræsentant for  $Y^4$  fra i-propyl eller  $-CH_3$ .
53. Forbindelse ifølge et eller flere af kravene 46 og 48, idet  
20  $Y^1$  er H,  $Y^2$  er  $-CH_2CH_2Z'$ ,  $Z'$  er  $-CONHCHY^4COOH$ , og  $Y^4$  er i givet fald med  $-NH_2$  substitueret aryl eller benzyl.
54. Forbindelse ifølge krav 53, idet  $Y^4$  er aminobenzyl.
- 25 55. Forbindelse ifølge krav 46, idet  $R^2$  er  $-(CH_2)_{0-3}Z$ , og Z er  $-SY^3$ .
56. Forbindelse ifølge krav 46, idet  $R^4$  er  $-CO-CHY^4-NHY^5$ , og  $Y^5$  er H.  
30
57. Forbindelse ifølge krav 46 idet  $R^4$  er  $-CO-CHY^4-NHY^5$ , og  $Y^5$  er  $CO-CHY^6-NH_2$ .
58. Forbindelse ifølge et eller flere af kravene 56 og 57, idet  
35  $Y^4$  er lineært eller forgrenet, i givet fald med  $-NHCONH_2$  substitueret  $C_{1-6}$  alkyl.
59. Forbindelse ifølge et eller flere af kravene 46 til 58,

idet forbindelsen har en af de følgende formler:

N-(3-aminopropyl)-N-[(1R)-1-[4-benzyl-1-(2,5-difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl]-2-hydroxyacetamid;

N-(3-aminopropyl)-N-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]-2-hydroxyacetamid;

(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]-N-methylbutanamid(1:1);

N-(3-aminopropyl)-N-[(1S)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]acetamid.

N-(3-aminopropyl)-N-[(1S)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl]-2-hydroxyacetamid;

S-[1-(2-[[2-[(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]amino)ethyl]amino)-2-oxoethyl]-2,5-dioxopyrrolidin-3-yl]-L-cystein:

N-[(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]-beta-alanyl-L-alanyl-N-[4-(3-[(2R)-2-amino-2-carboxyethyl]sulfanyl)-2,5-dioxopyrrolidin-1-yl]phenyl]-N<sup>5</sup>-carbamoyl-L-ornithinamid;

S-(1-[2-[(N-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]-beta-alanyl)amino]ethyl]-2,5-dioxopyrrolidin-3-yl)-L-cystein;

S-[1-(2-[[2-[(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]amino)ethyl]amino)-2-oxoethyl]-2,5-dioxopyrrolidin-3-yl]-L-cystein;

N-[(2S)-2-amino-4-[(1R)-1-[4-benzyl-1-(2,5-difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]-beta-alanyl-L-alanyl-N-[4-(3-[(2R)-2-amino-2-carboxyethyl]sulfanyl)-2,5-dioxopyrrolidin-1-yl]phenyl]-N<sup>5</sup>-carbamoyl-L-ornithinamid;

S-(1-[2-[(N-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl](glycoloyl)amino]butanoyl]-beta-alanyl)amino]ethyl]-2,5-dioxopyrrolidin-3-yl)-L-cystein;

N-[6-(3-{{(2R)-2-amino-2-carboxyethyl}sulfanyl}-2,5-dioxopyrrolidin-1-yl)hexanoyl]-L-valyl-N<sup>5</sup>-carbamoyl-L-ornithyl-N<sup>6</sup>-{(2S)-2-amino-4-{{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-L-  
 5 lysin;  
 S-[1-(2-{{2-{{(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl}amino)-2-oxoethyl)-2,5-dioxopyrrolidin-3-yl]-L-cystein;  
 10 S-(2-{{2-{{(2S)-2-amino-4-{{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl}amino)-2-oxoethyl)-L-cystein;  
 S-{{1-[6-(2-{{(2S)-2-amino-4-{{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}hydrazino)-6-oxohexyl]-2,5-dioxopyrrolidin-3-yl}-L-cystein;  
 15 N-[19-(3(R/S)-{{(2R)-2-amino-2-carboxyethyl}sulfanyl}-2,5-dioxopyrrolidin-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-R/S-{{2-[[3-aminopropyl]{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}homocystein;  
 20 S-{{(3R1S)-1-[2-{{(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl]-2,5-dioxopyrrolidin-3-yl}-L-cystein;  
 25 N-[19-(3(R/S)-{{(2R)-2-amino-2-carboxyethyl}sulfanyl}-2,5-dioxopyrrolidin-1-yl)-17-oxo-4,7,10,13-tetraoxa-16-azanonadecan-1-oyl]-R/S-{{2-[[3-aminopropyl]{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-imidazol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}homocystein;  
 30 S-[[3(R/S)-1-(2-{{6-{{2-[[3-aminopropyl]-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-oxoethyl}sulfanyl)hexanoyl}amino)ethyl)-2,5-dioxopyrrolidin-3-yl]-L-cystein;  
 35 S-{{1-[2-{{[[1(R,3S)-3-{{(2S)-2-amino-4-{{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)cyclopentyl}car

bonyl}amino)ethyl]-2,5-dioxopyrrolidin-3-yl} -L-cystein;  
 S-(2-{[2-({(2S)-2-amino-4-[-{(1R)-1-[1-benzyl-4-(2,5-  
 difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl}amino)-2-  
 5 oxoethyl)-L-cystein;  
 N<sup>6</sup>-(N-{(2S)-2-amino-4-[-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-  
 beta-alanyl)-N<sup>2</sup>-{N-[6-(3-{{(2R)-2-amino-2-  
 carboxyethyl}sulfanyl)-2,5-dioxopyrrolidin-1-yl)hexanoyl]-L-  
 10 valyl-L-alanyl}-L-lysin;  
 N-[2-({(2S)-2-amino-4-[-{(1R)-1-[1-benzyl-4-(2,5-  
 difluorophenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl}(glycoloyl)amino]butanoyl}amino)ethyl)-L-  
 glutamin;  
 15 N<sup>6</sup>-(N-{(2S)-2-amino-4-[-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}-  
 beta-alanyl)-L-lysin;  
 N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
 1H-pyrazol-3-yl]-2,2-dimethylpropyl} acetamid;  
 20 N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
 1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-methoxyacetamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
 1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2,4-difluorobenzamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
 25 1H-pyrazol-3-yl]-2,2-dimethylpropyl}-4-methylbenzamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
 1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}-3,3,3-trifluoropropanamid;  
 30 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-fluorbenzamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}acetamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorophenyl)-  
 35 1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-  
 (trifluoromethyl)benzamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorophenyl)-  
 1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamid;

N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl}-2-ethoxyacetamid  
 (2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butansyre;  
 5 (2S)-2-amino-N-(2-aminoethyl)-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanamid;  
 4-[(2-[[2-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl} amino) ethyl] amino}-2-oxoethyl) amino]-3-[(2R)-2-amino-2-carboxyethyl] sulfanyl}-4-oxobutansyre;  
 10 4-[(2-[[2-[(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl} amino) ethyl] amino}-2-oxoethyl) amino]-2-[(2R)-2-amino-2-carboxyethyl] sulfanyl}-4-oxobutansyre;  
 15 N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl}-beta-alanin;  
 20 N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl}-L-serin;  
 N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl}-L-alanin;  
 25 N-{(2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} (glycoloyl) amino]butanoyl}glycin;  
 30 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-methylbenzamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-4-(methylsulfanyl)benzamid;  
 35 (2S)-N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-hydroxypropanamid;  
 N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-

1H-pyrrol-2-yl]-2,2-dimethylpropyl}-2-(  
 (methylsulfanyl)acetamid;  
 (2S)-N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-  
 difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl} -2-  
 5 hydroxypropanamid;  
 Methyl-4-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-  
 difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl} amino] -4-  
 oxobutanoat;  
 4-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-  
 10 pyrrol-2-yl]-2,2-dimethylpropyl} amino] -4-oxobutansyre;  
 (2R)-22-[(3R/S)-3-{{(2R)-2-amino-2-carboxyethyl}sulfanyl}-2,5-  
 dioxopyrrolidin-1-yl]-2-[(2-[(3-aminopropyl){(1R)-1-[1-  
 benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl}amino]-2-oxoethyl)sulfanyl)methyl]-4,20-dioxo-  
 15 7,10,13,16-tetraoxa-3,19-diazadocosan-1-syre;  
 4-amino-N-(3-aminopropyl)-N-{(1R)-1-[4-benzyl-1-(2,5-  
 difluorphenyl)-1H-pyrazol-3-yl]-2,2-dimethylpropyl} benzamid;  
 N-Acetyl-S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-  
 difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-  
 20 oxoethyl}-L-cystein;  
 N-Acetyl-S-[2-([3-(L-alanyl-amino)propyl]{(1R)-1-[1-benzyl-4-  
 (2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}  
 amino)-2-oxoethyl]-L-cystein;  
 (2S)-N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-  
 25 difluorphenyl)-1H-pyrrol-2-yl]-2,2-  
 dimethylpropyl}tetrahydrofuran-2-carboxamid;  
 3-({2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl} amino] -2-  
 oxoethyl}sulfanyl)propansyre;  
 30 S-{2-[(3-aminopropyl){(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-  
 1H-pyrrol-2-yl]-2,2-dimethylpropyl}amino]-2-  
 oxoethyl}homocystein;  
 4-amino-N-(3-aminopropyl)-N-{(1R)-1-[1-benzyl-4-(2,5-  
 difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}benzamid;  
 35 4-[(2-{{(2R)-2-((2S)-2-amino-4-[(1R)-1-[1-benzyl-4-(2,5-  
 difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}  
 (glycoloyl)amino]butanoyl}amino)-2-carboxyethyl}amino] -2-  
 oxoethyl}amino]-3-{{(2R)-2-amino-2-carboxyethyl}sulfanyl}-4-



oxobutansyre;

4-[(2-[[[(2R)-2-({(2S)-2-amino-4-[[[(1R)-1-[1-benzyl-4-(2,5-difluorphenyl)-1H-pyrrol-2-yl]-2,2-dimethylpropyl}(glycoloyl)amino]butanoyl}amino)-2-carboxyethyl]amino]-2-oxoethyl)amino]-2-[[[(2R)-2-amino-2-carboxyethyl]sulfanyl]-4-oxobutansyre.

60. Farmaceutisk præparat omfattende et konjugat ifølge et eller flere af kravene 1 til 41 eller en forbindelse ifølge krav 46-59 i kombination med et inert, ikke-toksisk, farmaceutisk egnet hjælpestof.

61. Konjugat ifølge et eller flere af kravene 1 til 41 eller forbindelse ifølge krav 46-59 til anvendelse i en fremgangsmåde til behandling og/eller profylakse af sygdomme.

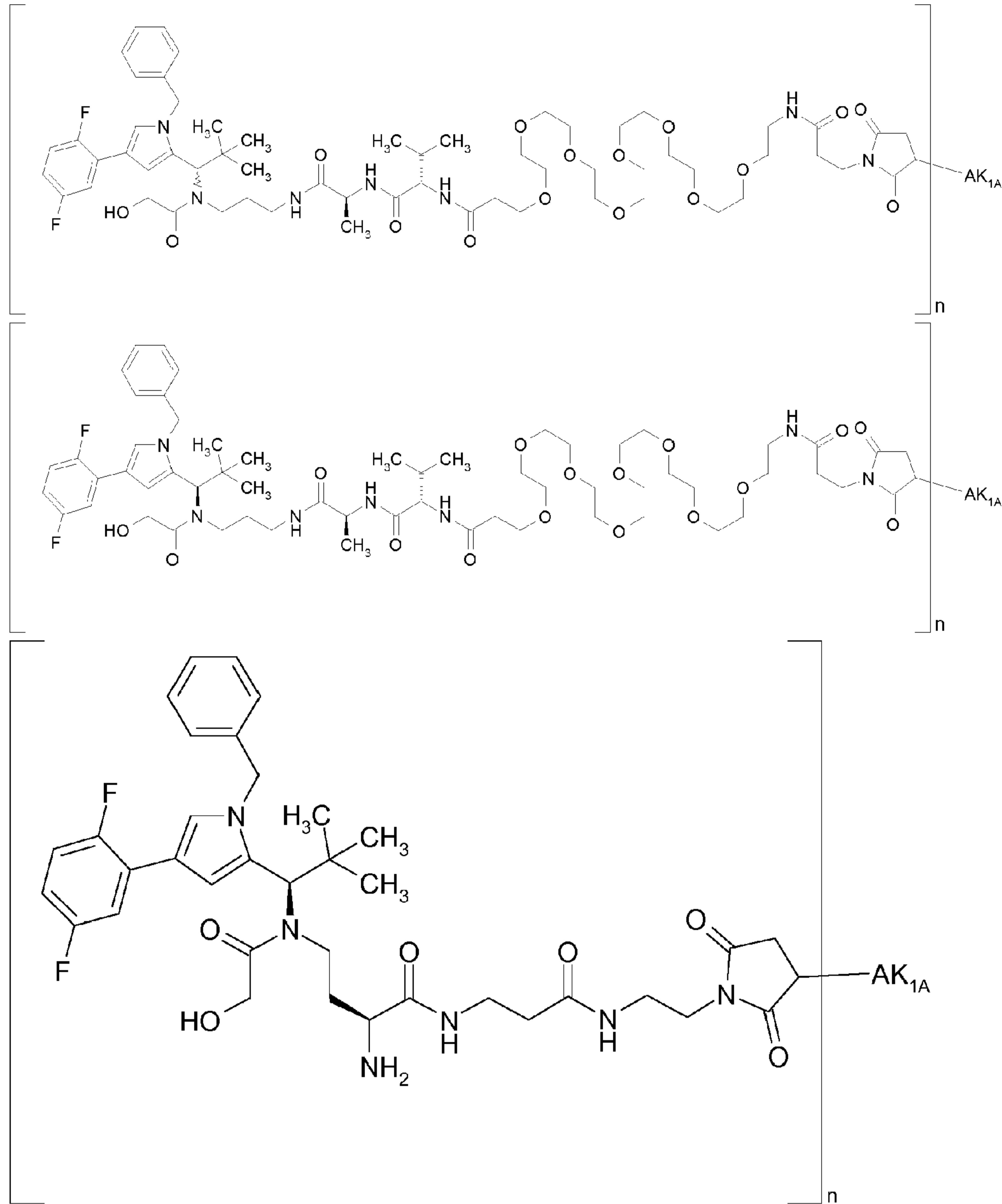
62. Konjugat ifølge et eller flere af kravene 1 til 41 eller forbindelse ifølge krav 46-59 til anvendelse i en fremgangsmåde til behandling af hyperproliferative og/eller angiogene sygdomme.

63. Konjugat ifølge et eller flere af kravene 1 til 41 eller forbindelse ifølge krav 46-59, idet  $R^{3a}$  eller  $R^3$  er -L-BINDER eller en substitueret alkyl-, aryl- eller heteroarylgruppe, fortrinsvis -L-#1 eller en  $C_{1-10}$ -alkyl-,  $C_{6-10}$ -aryl- eller  $C_{6-10}$ -aralkylgruppe eller  $C_{5-10}$ -heteroalkyl-, som kan være substitueret med -OH, O-alkyl, SH, S-alkyl, O-CO-alkyl, O-CO-NH-alkyl, NH-CO-alkyl, NH-CO-NH-alkyl,  $S(O)_n$ -alkyl,  $SO_2$ -NH-alkyl, NH-alkyl,  $N(alkyl)_2$ ,  $NH_2$  eller  $-(CH_2)_{0-3}Z$ , idet Z er -H, halogen,  $-OY^3$ ,  $-SY^3$ ,  $-NHY^3$ ,  $-CO-NY^1Y^2$  eller  $-CO-OY^3$ , idet  $Y^1$  og  $Y^2$  uafhængigt af hinanden er H,  $NH_2$  eller  $-(CH_2)_{0-3}Z'$ , og  $Y^3$  er H eller  $-(CH_2)_{0-3}Z'$ , idet  $Z'$  er H,  $SO_3H$ ,  $NH_2$  eller  $COOH$ , (idet "alkyl" fortrinsvis betegner  $C_{1-10}$ -alkyl).

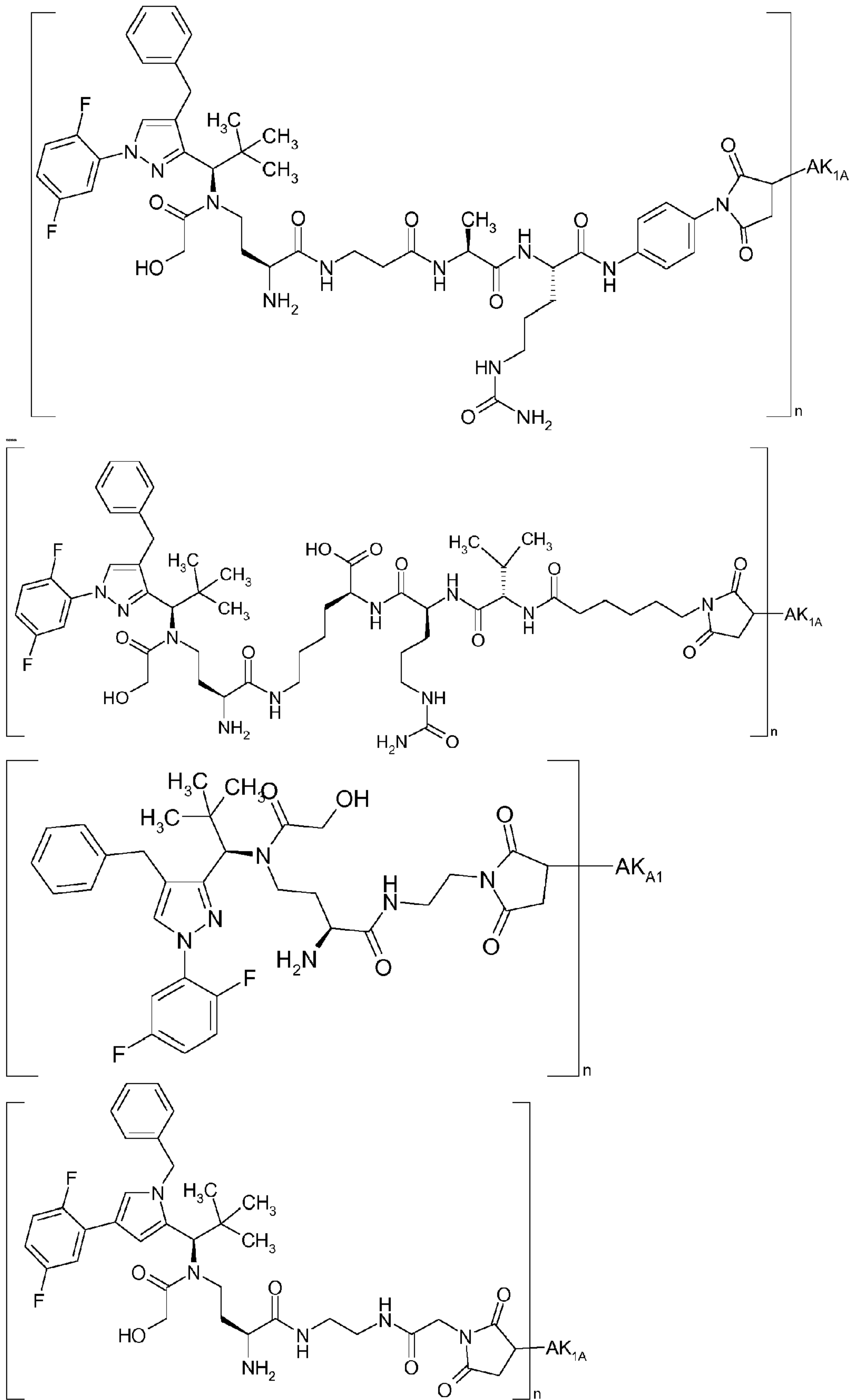
64. Konjugat ifølge et eller flere af kravene 1 til 41 eller forbindelse ifølge et af kravene 46 til 59 til anvendelse i en fremgangsmåde til behandling og/eller profylakse af hyperproliferative og/eller angiogene sygdomme hos mennesker og

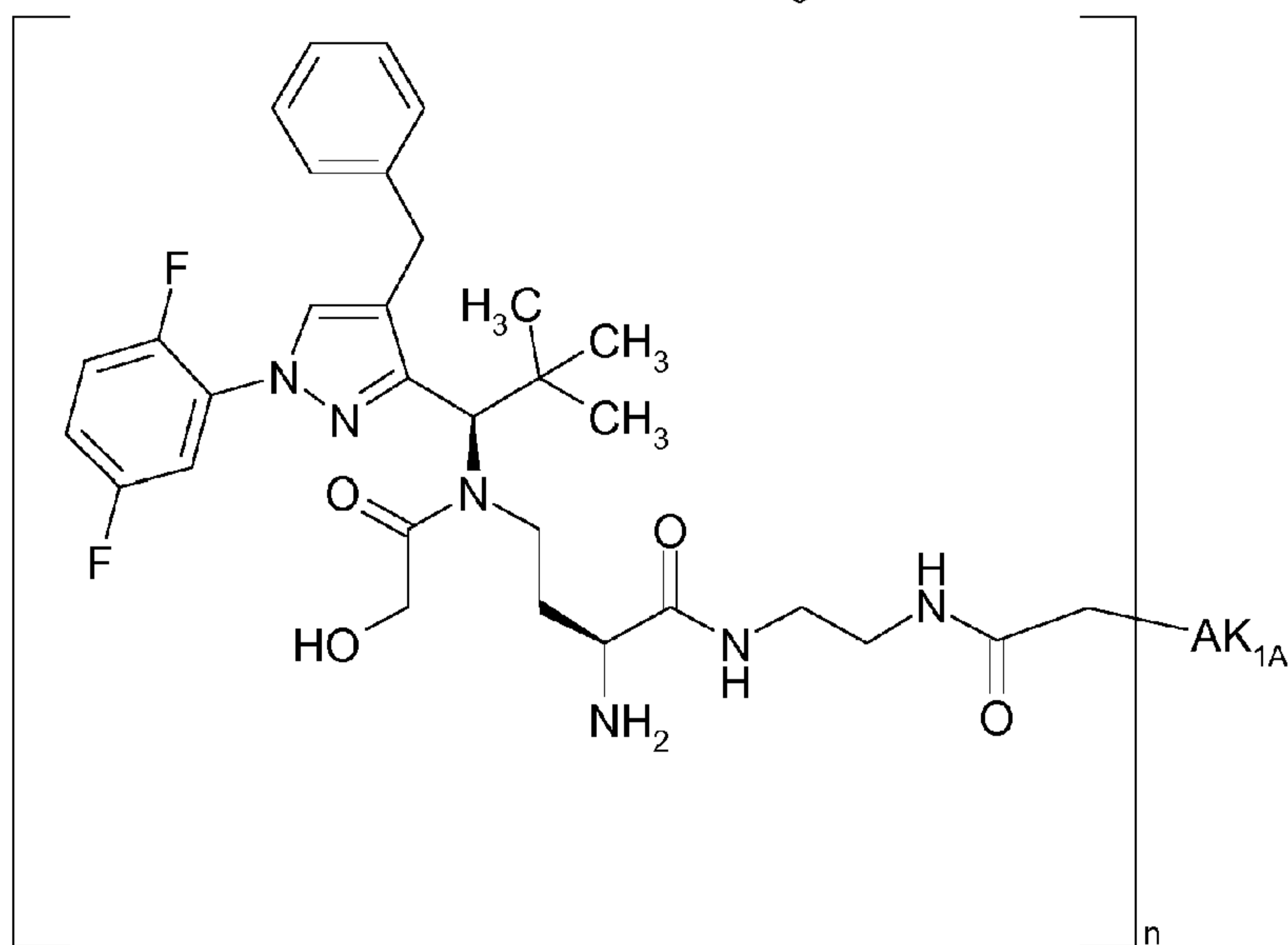
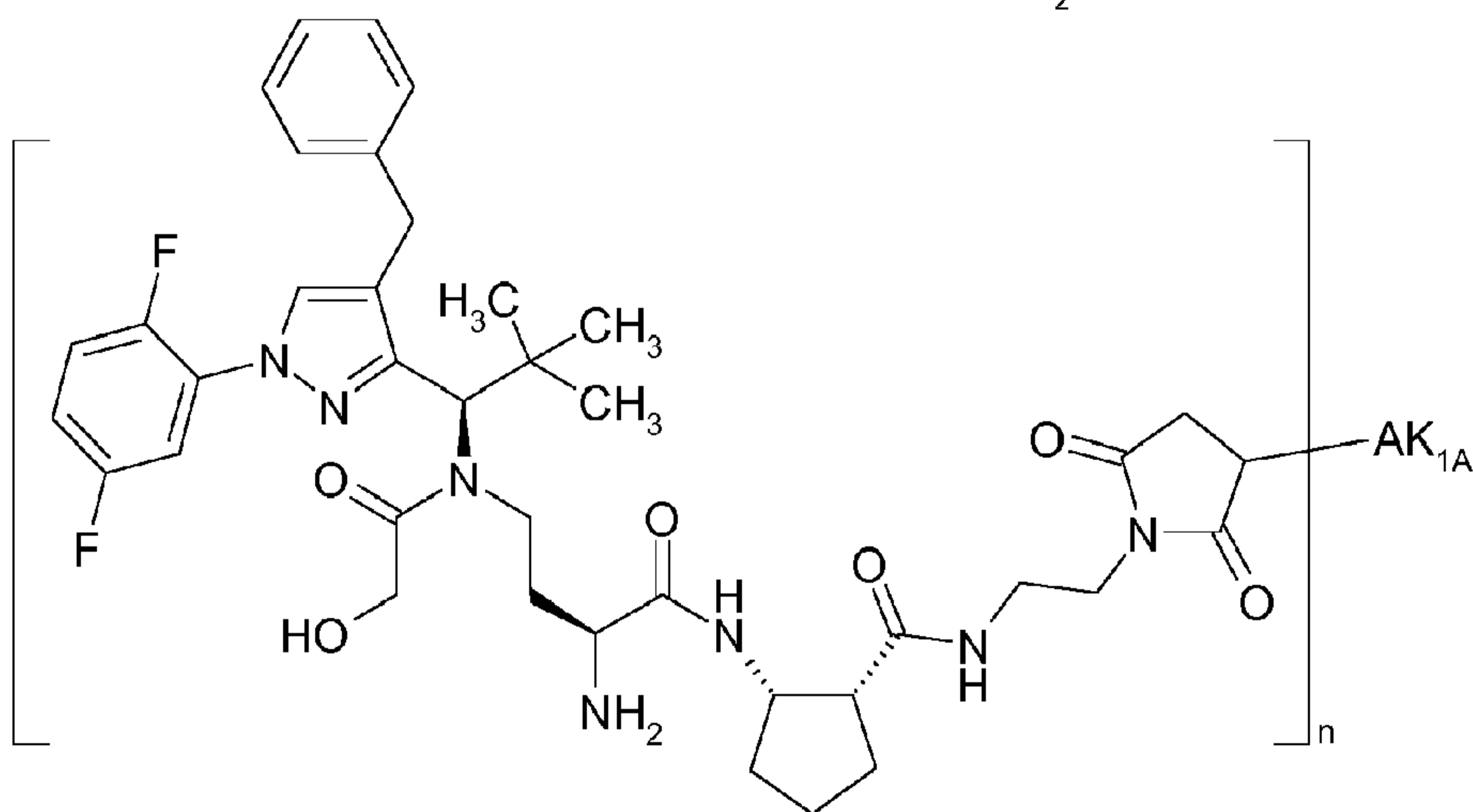
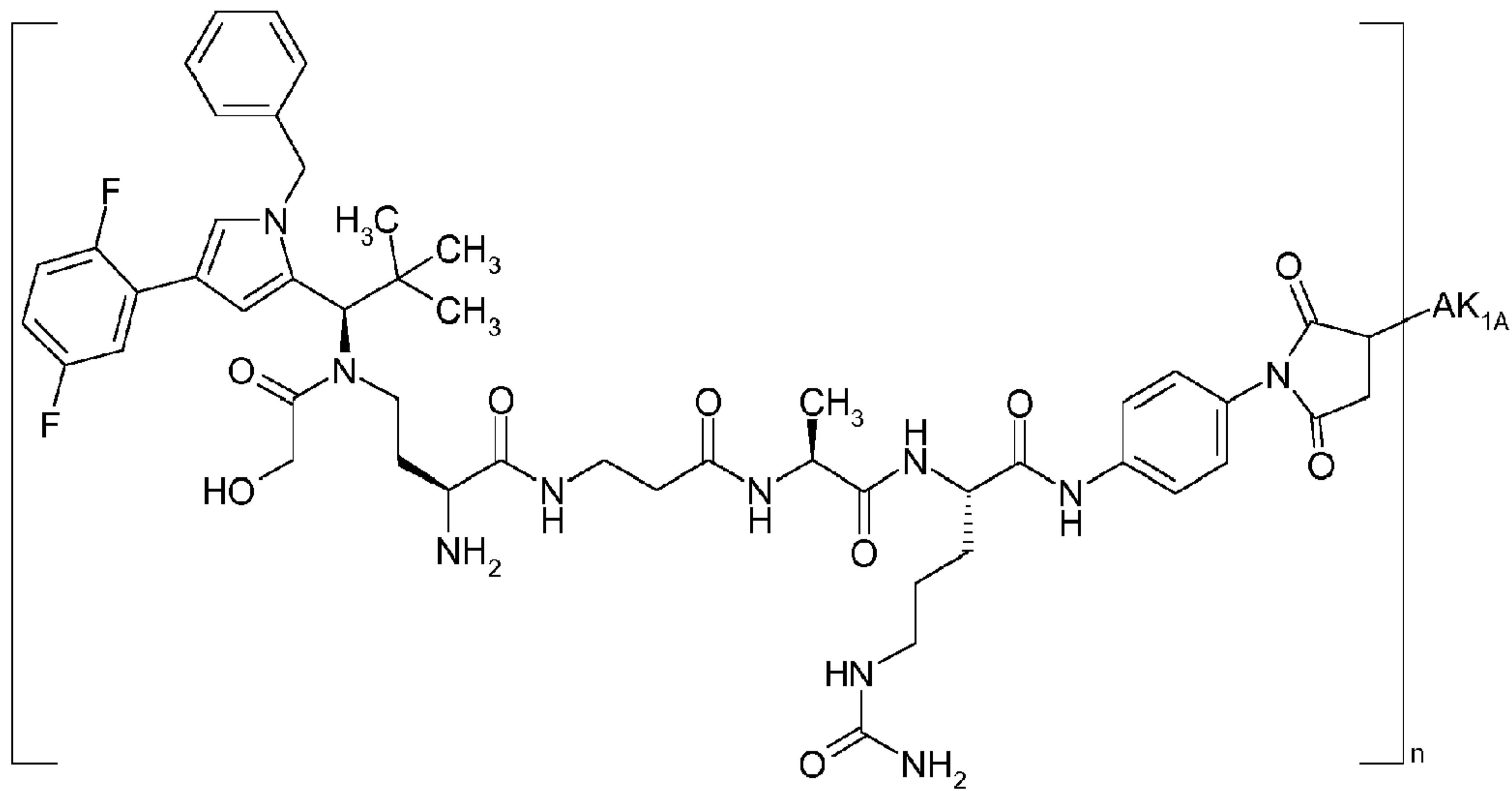
dyr under anvendelse af en virksom mængde.

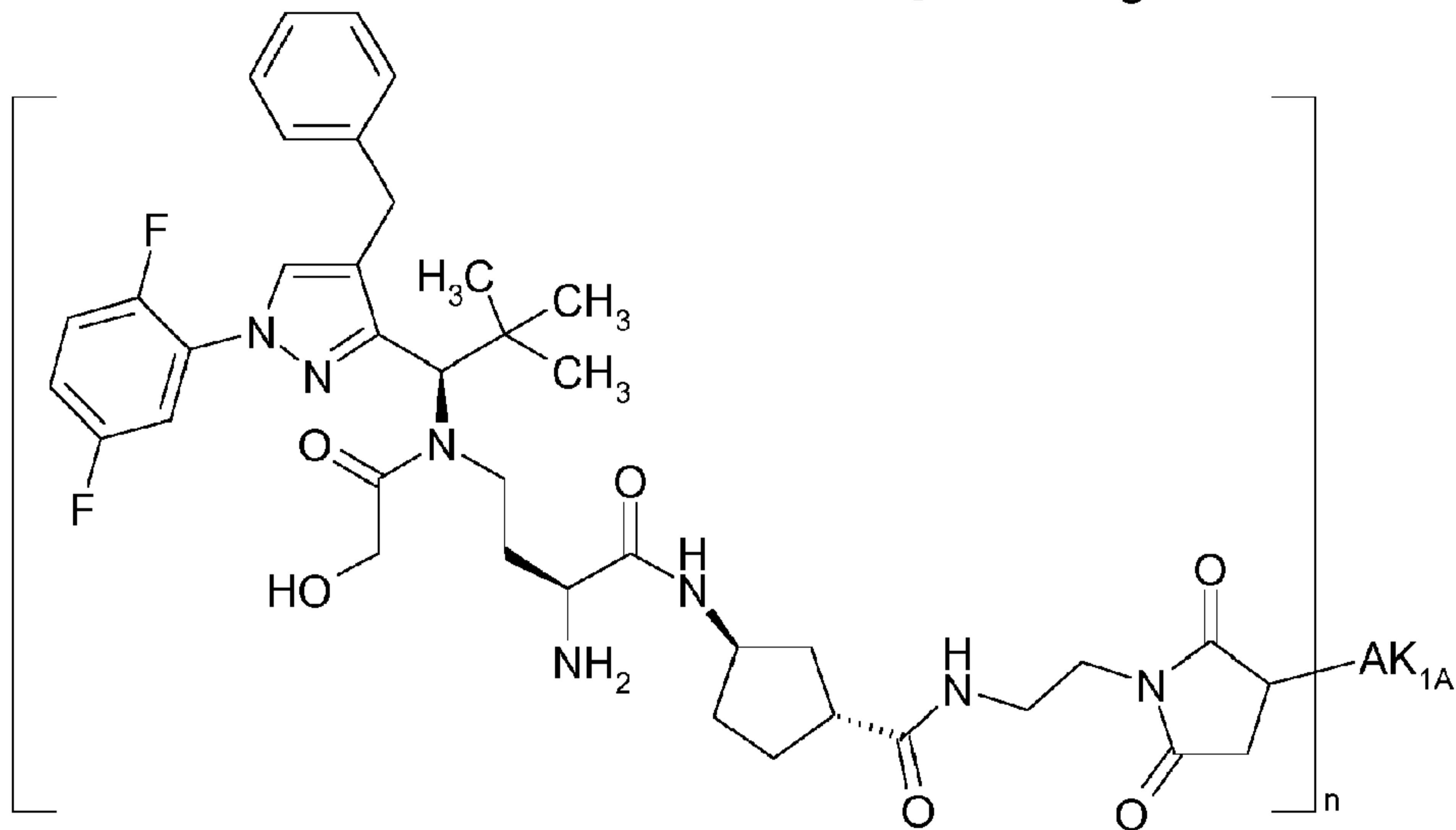
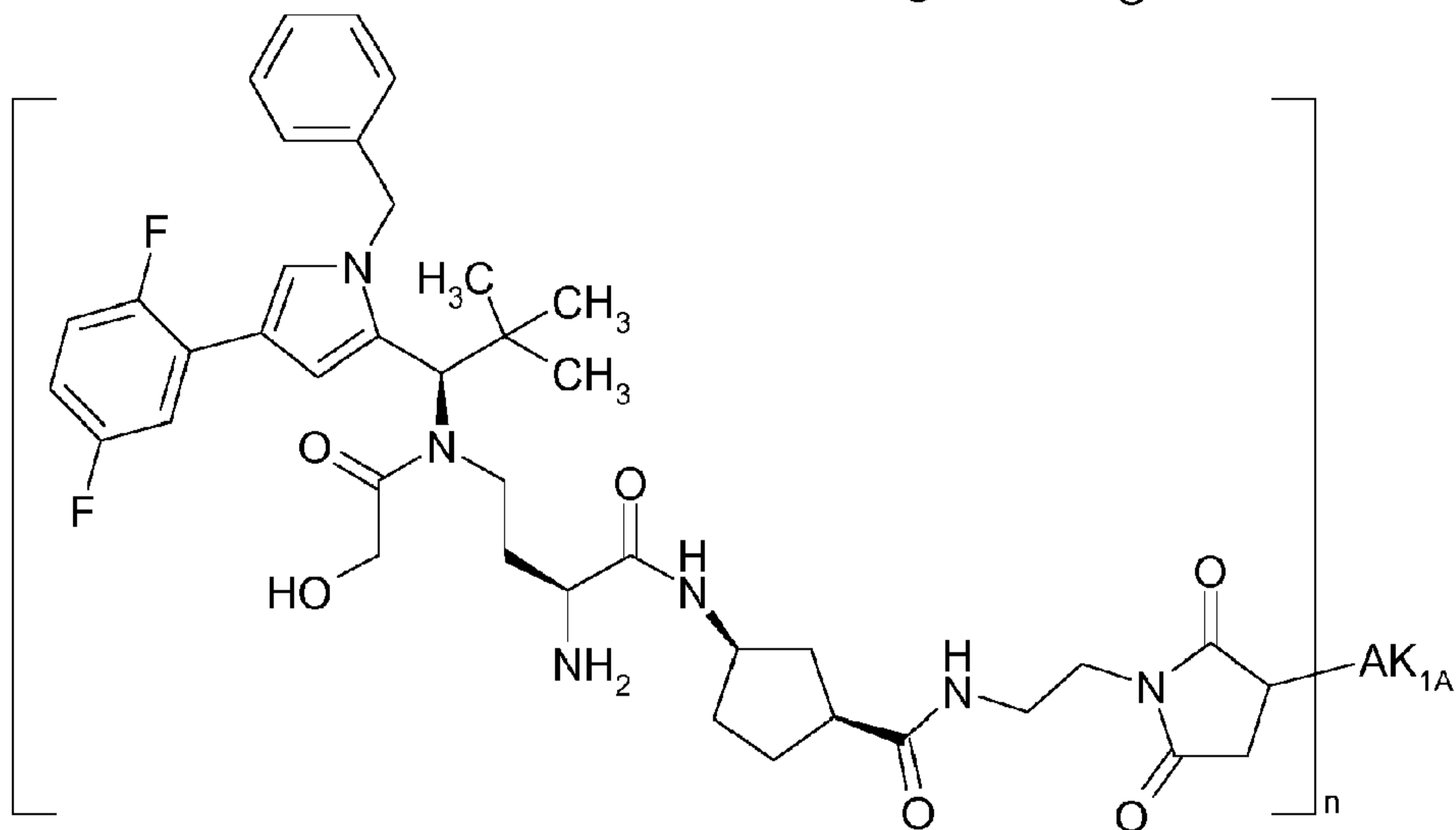
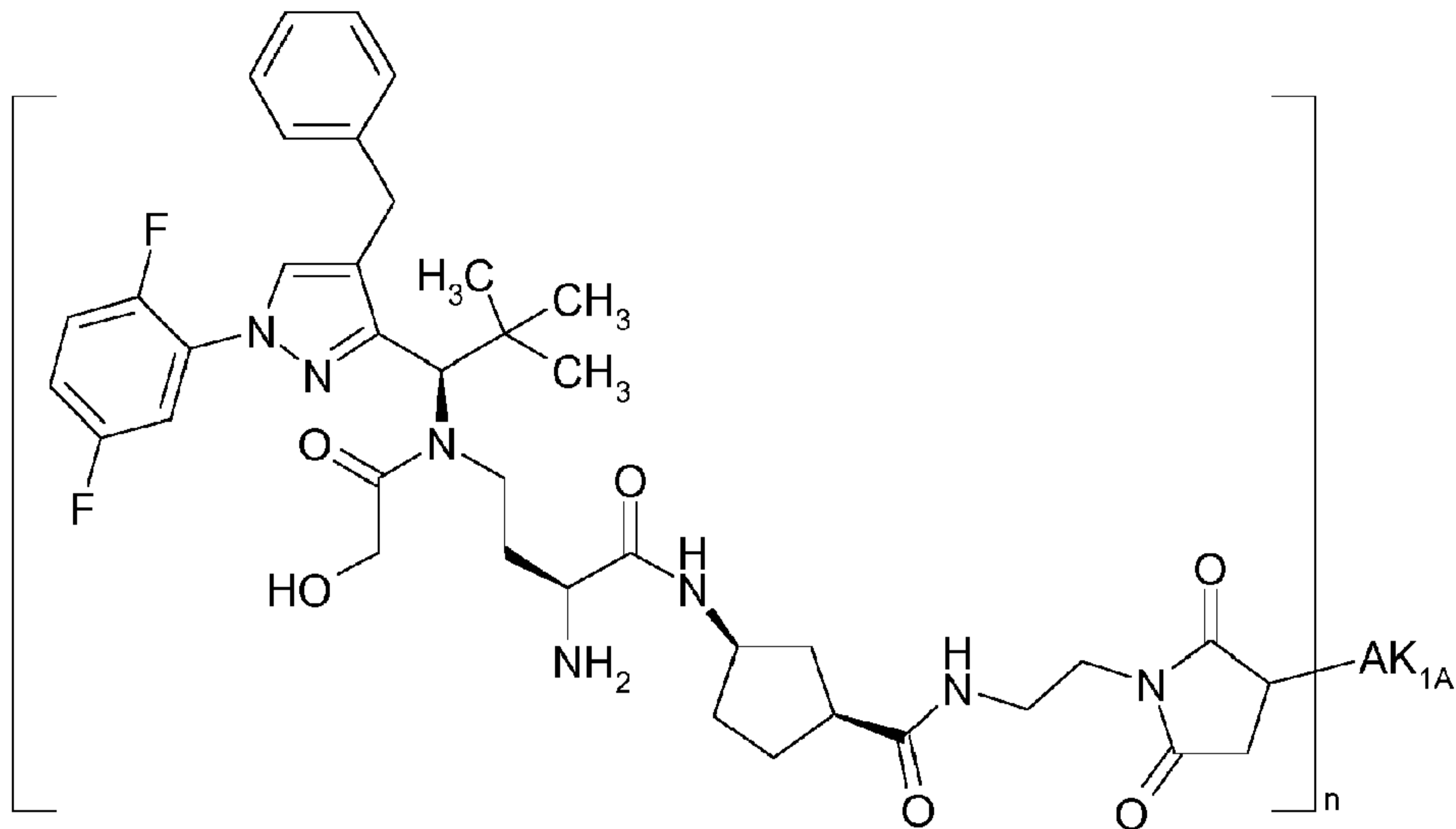
65. Antistof-konjugat ifølge en af de følgende formler, idet  $n$  er et tal fra 1 til 20, og AK<sub>1A</sub> eller AK<sub>2A</sub> er et antistof:

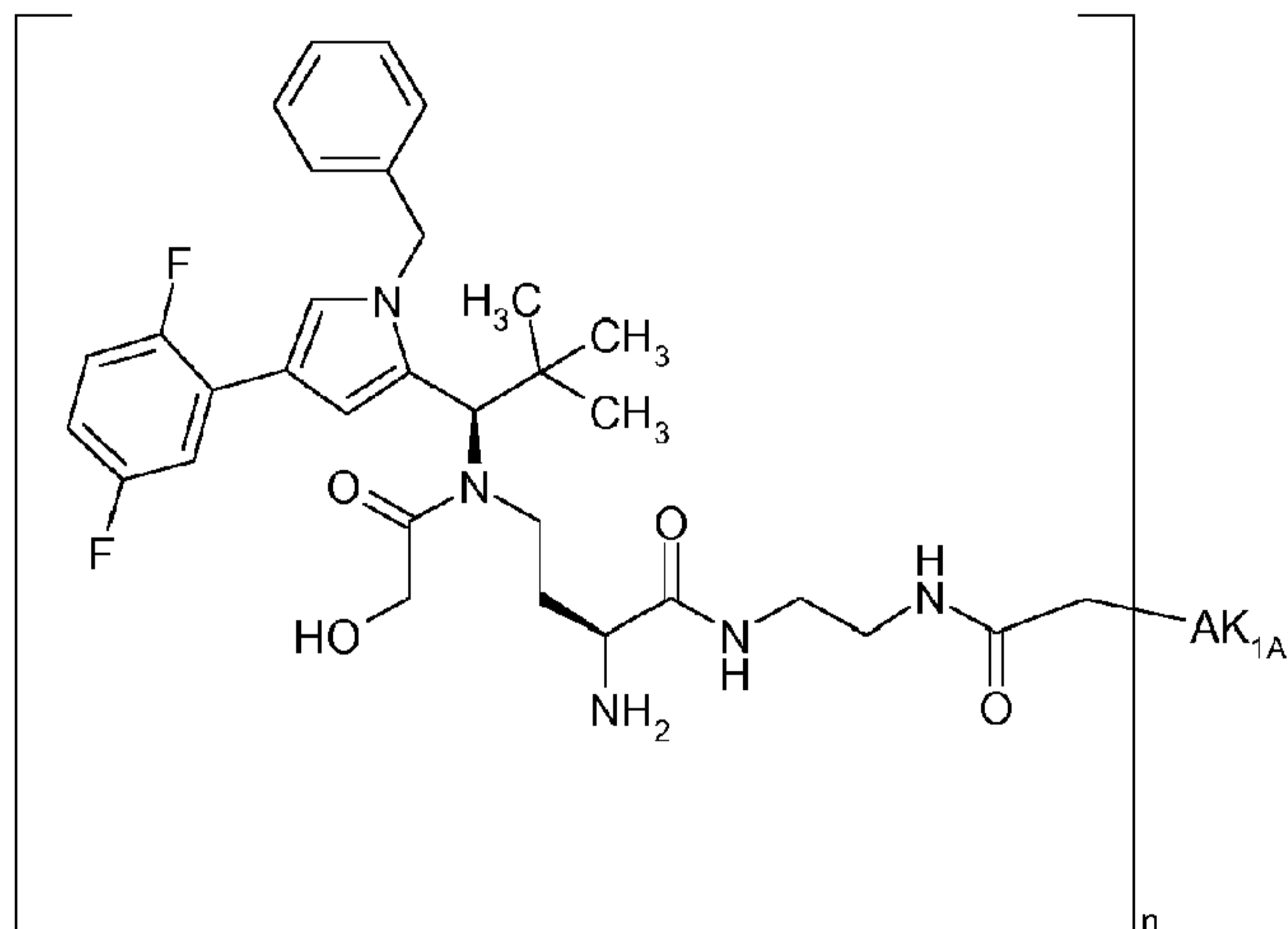
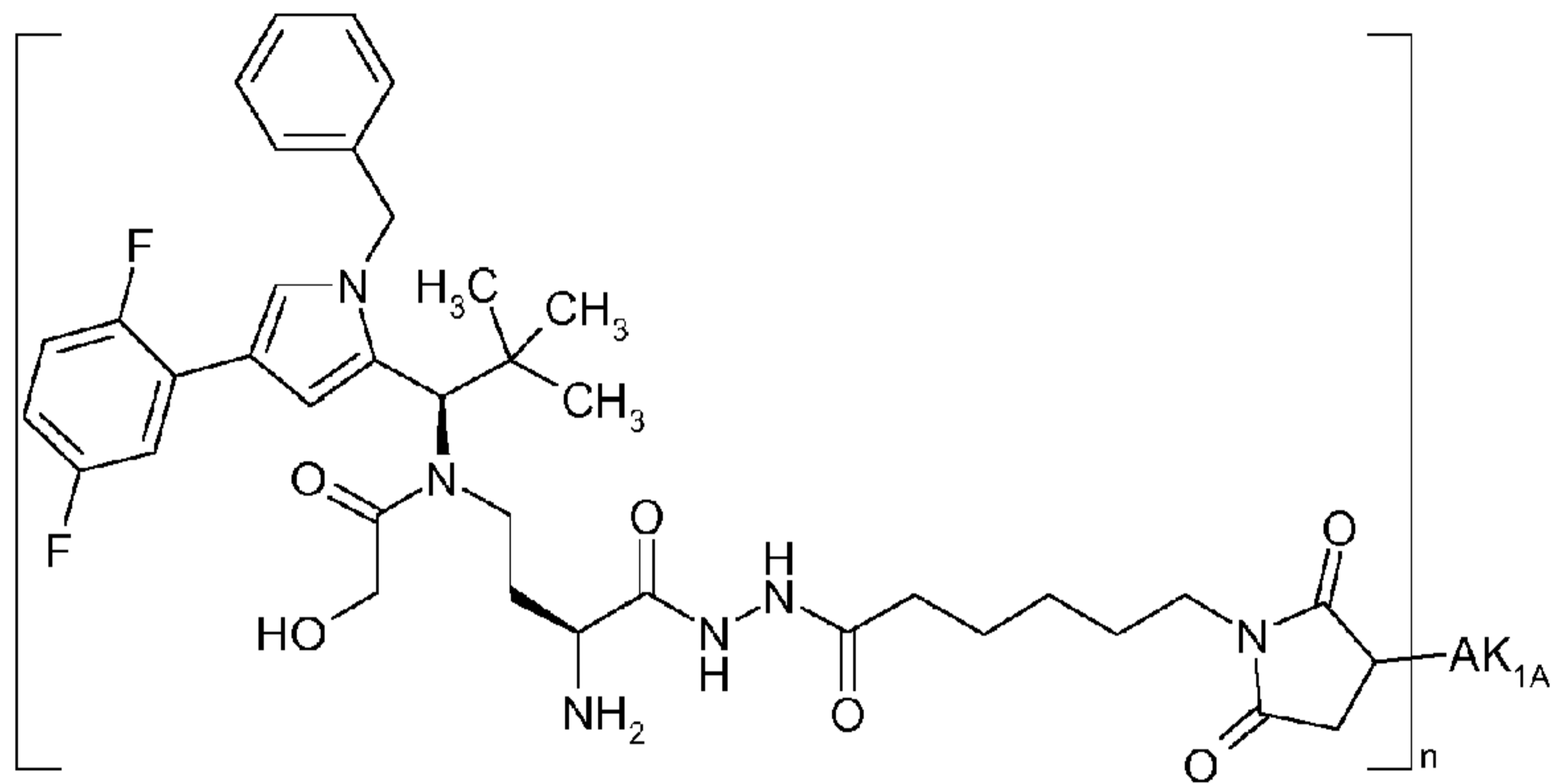
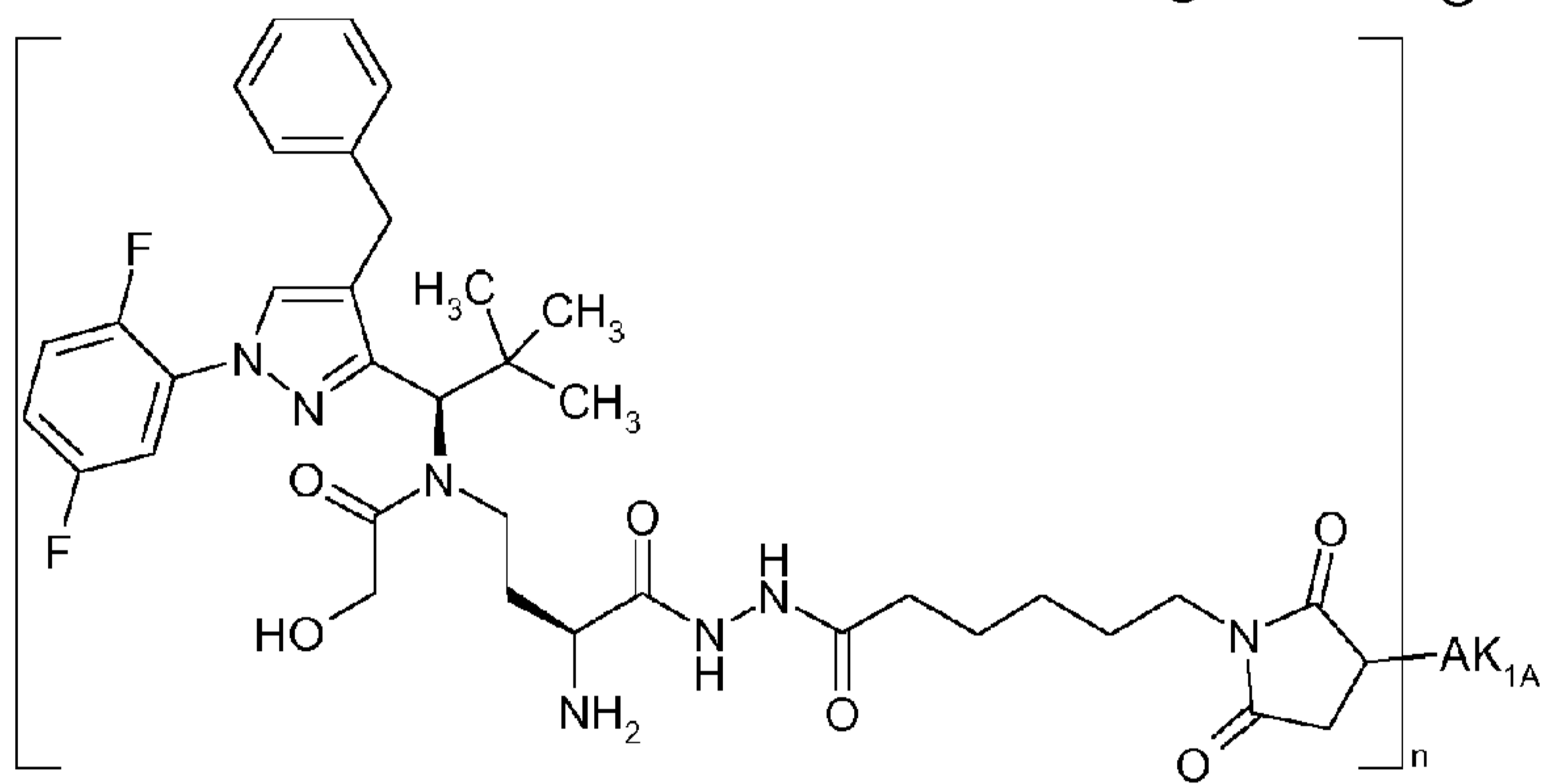
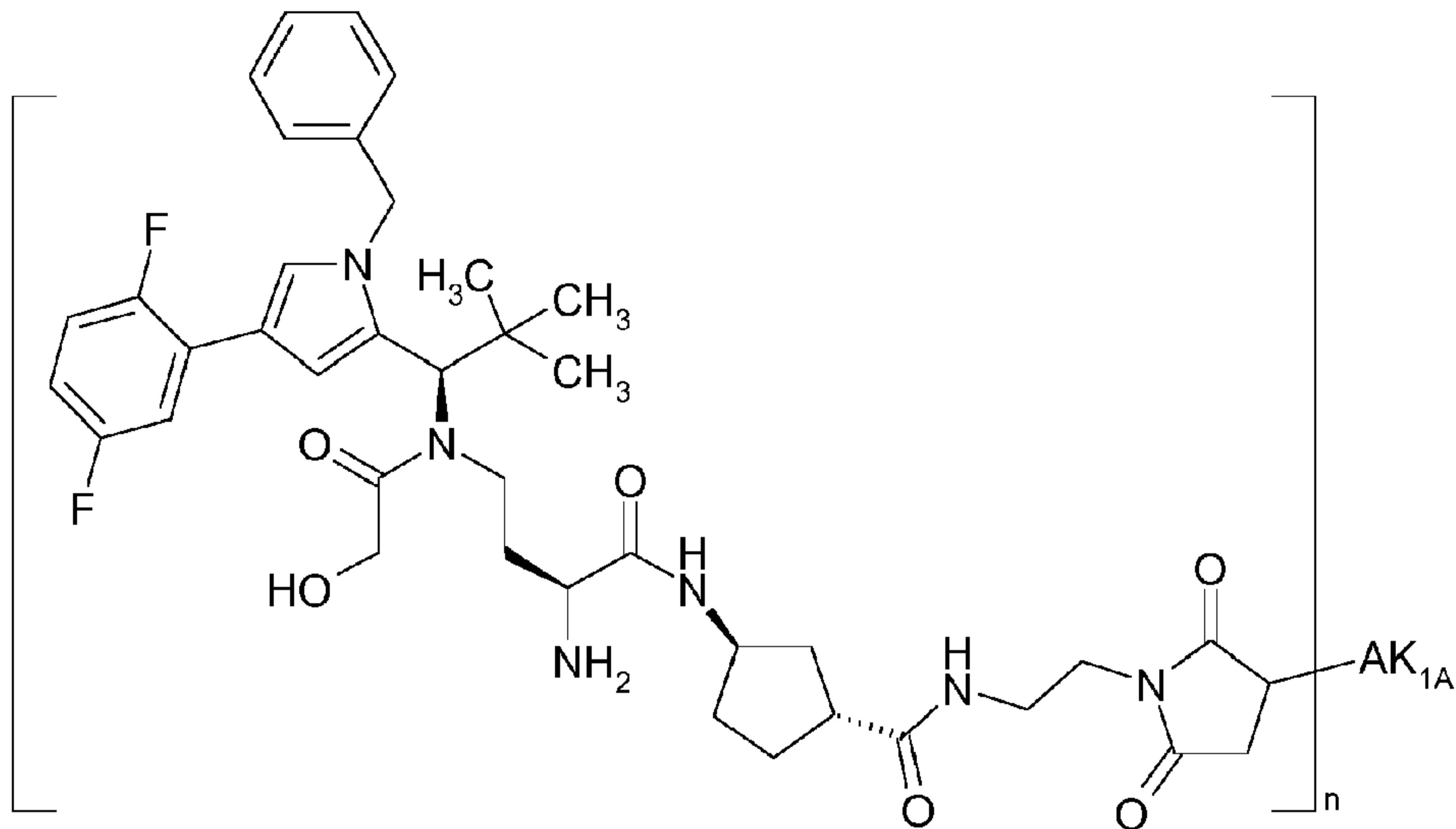


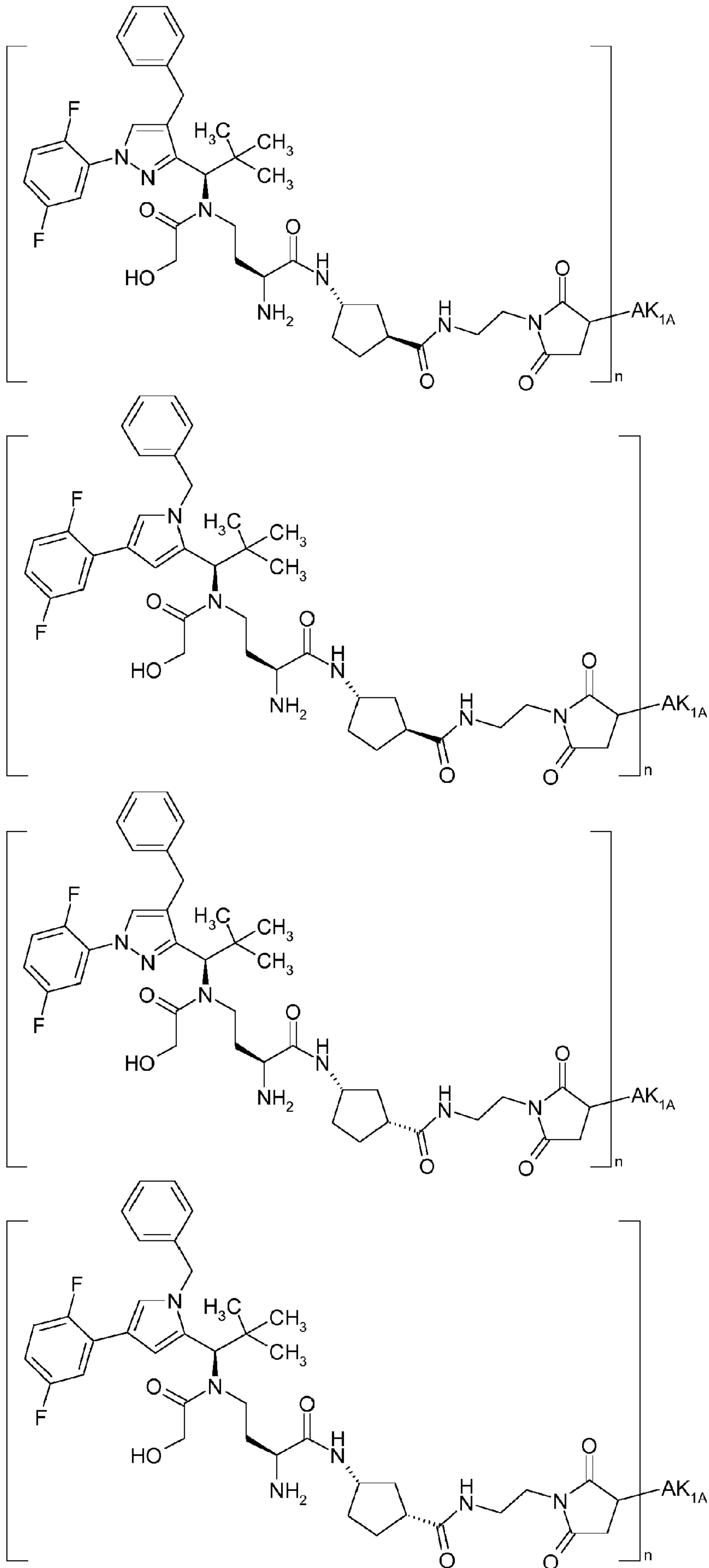




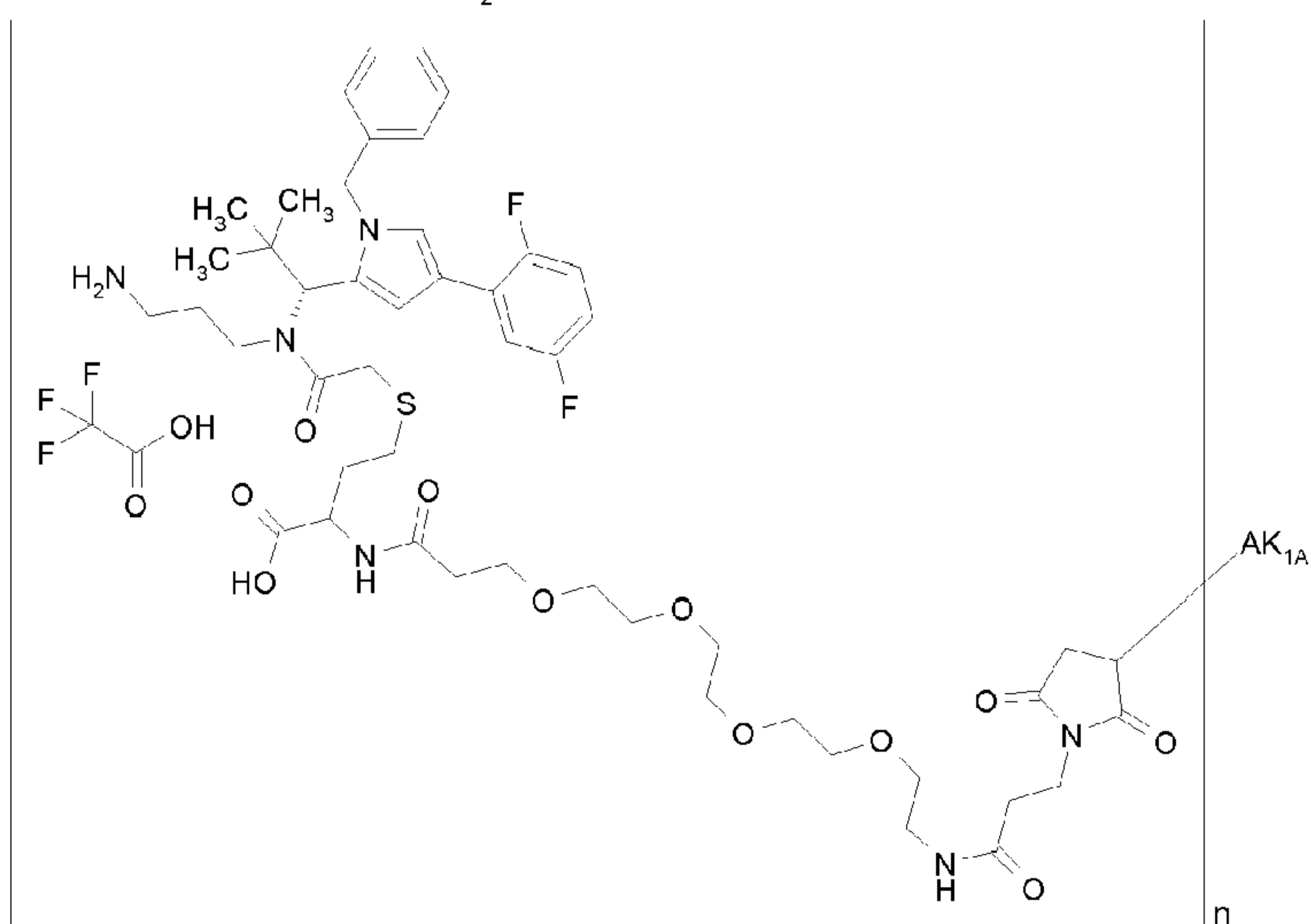
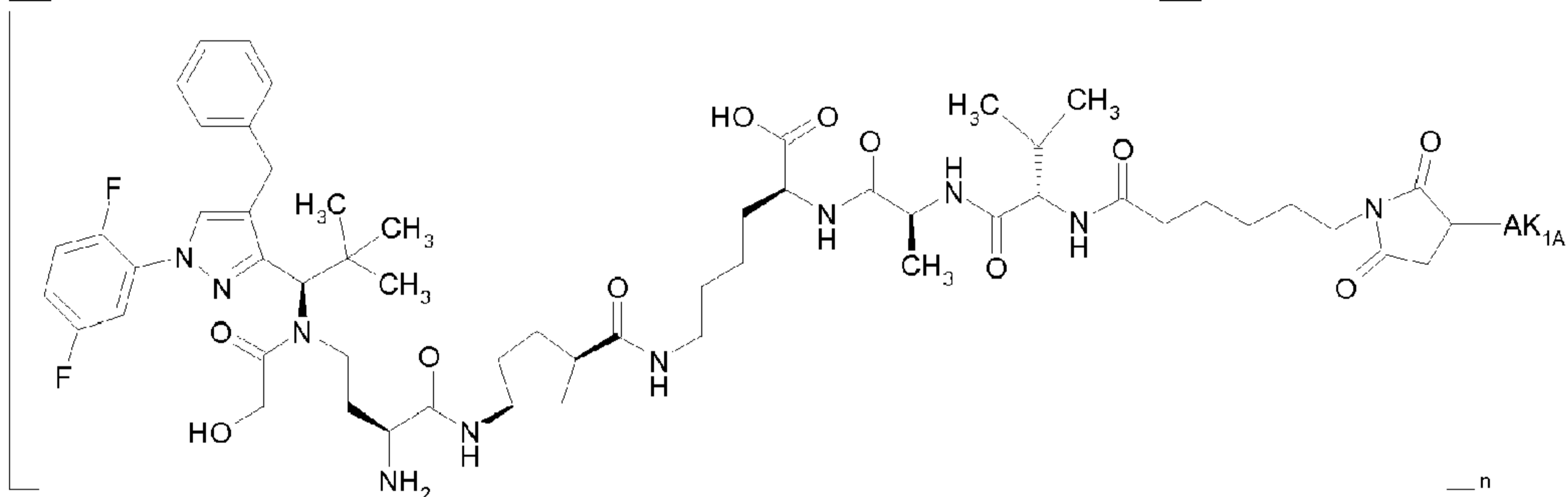
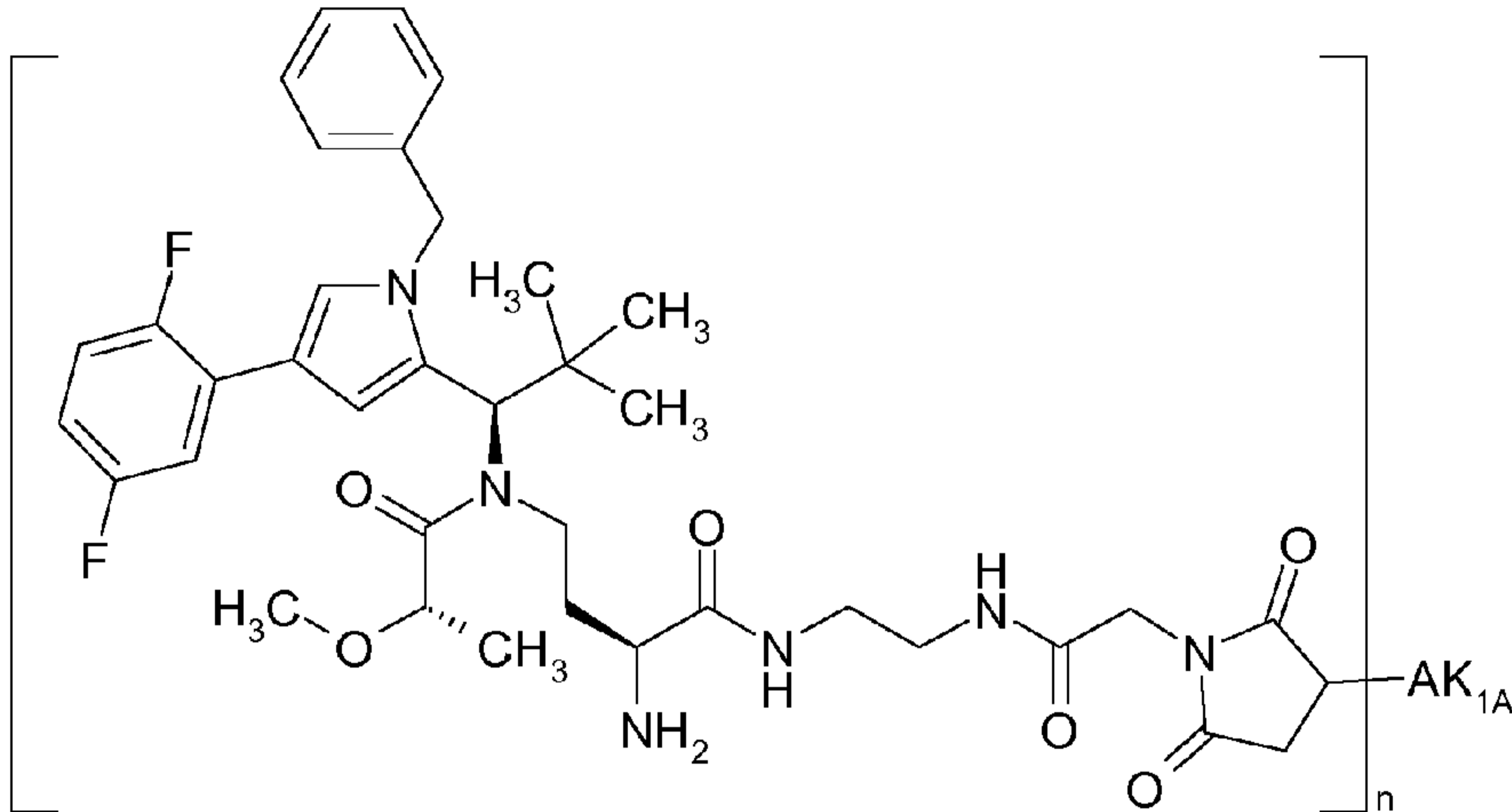
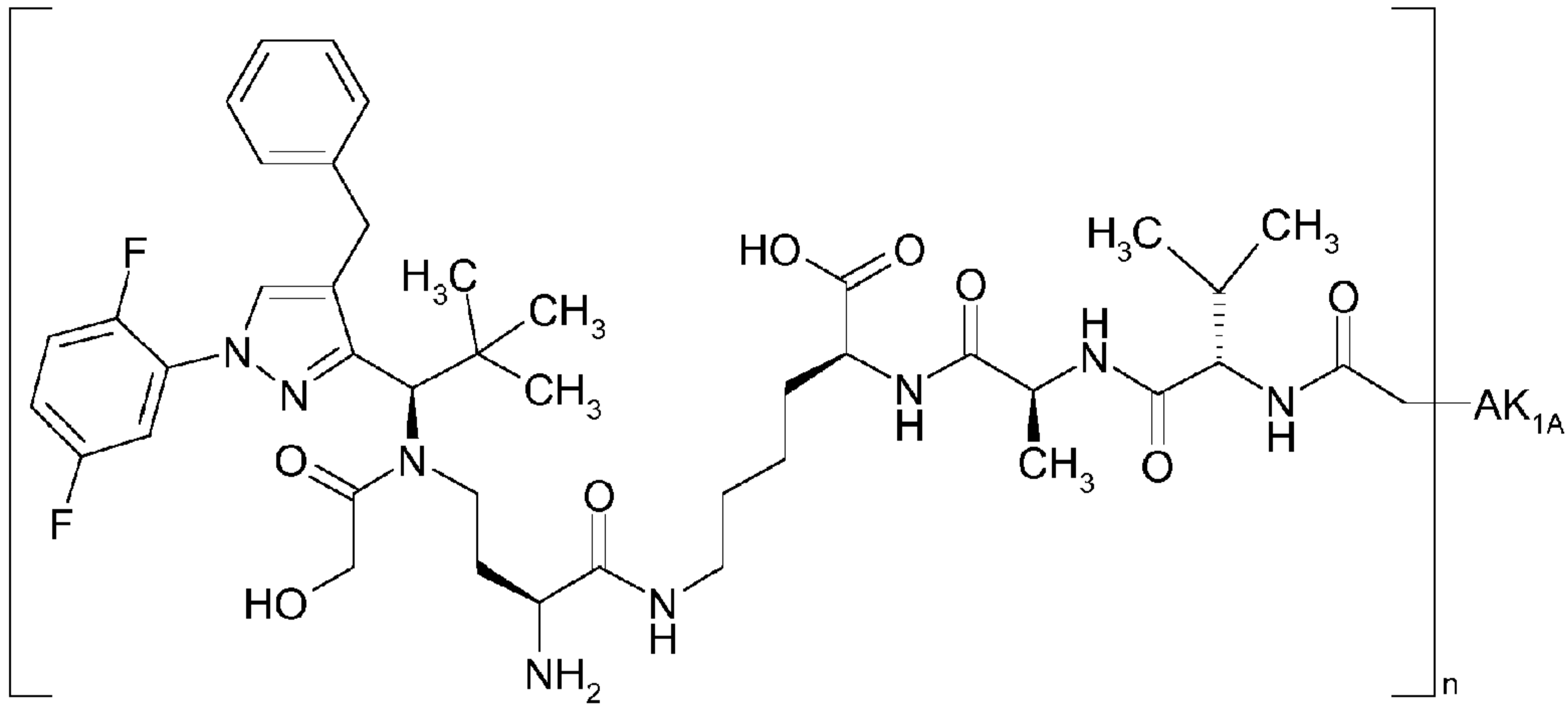


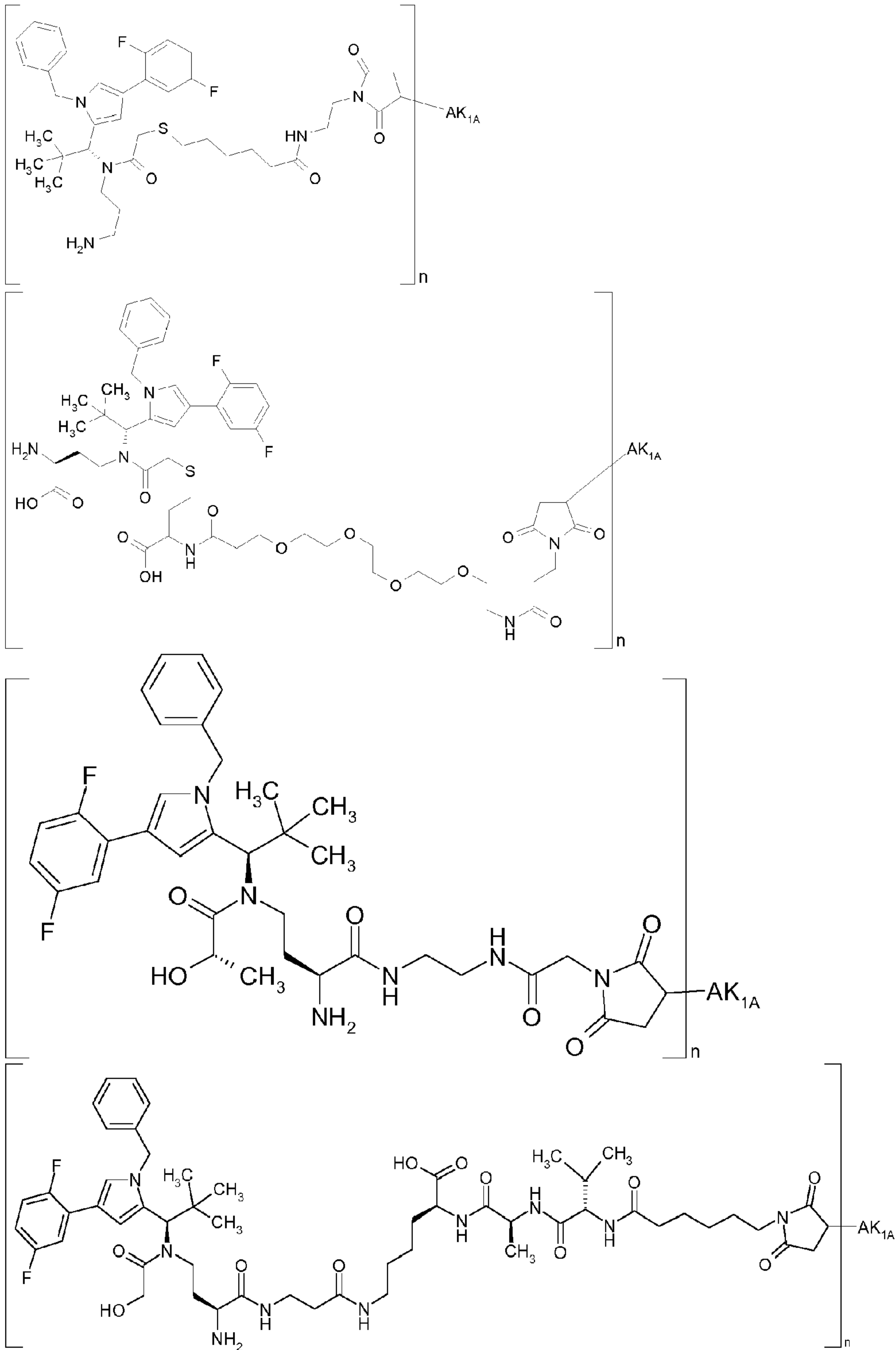


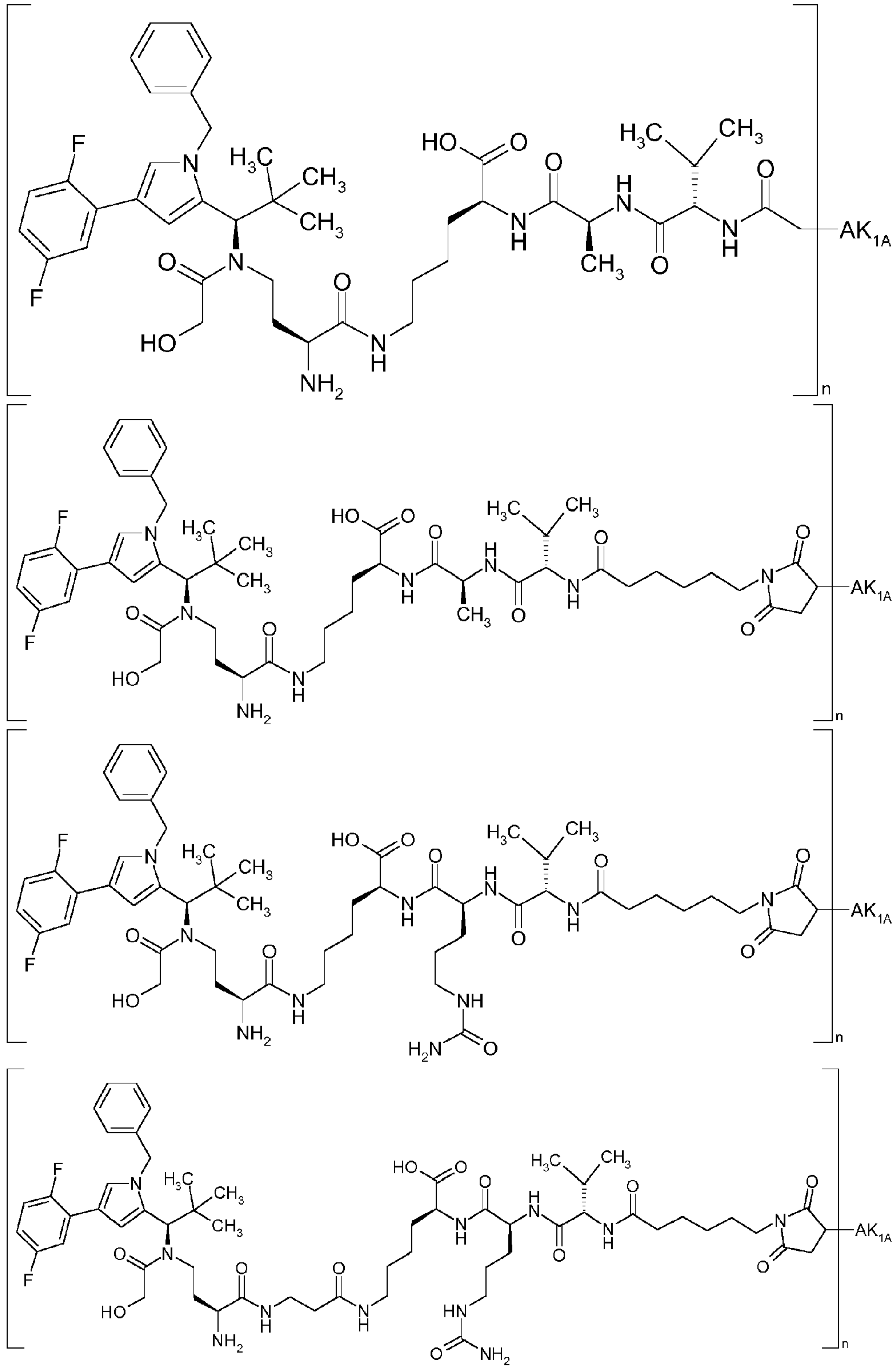




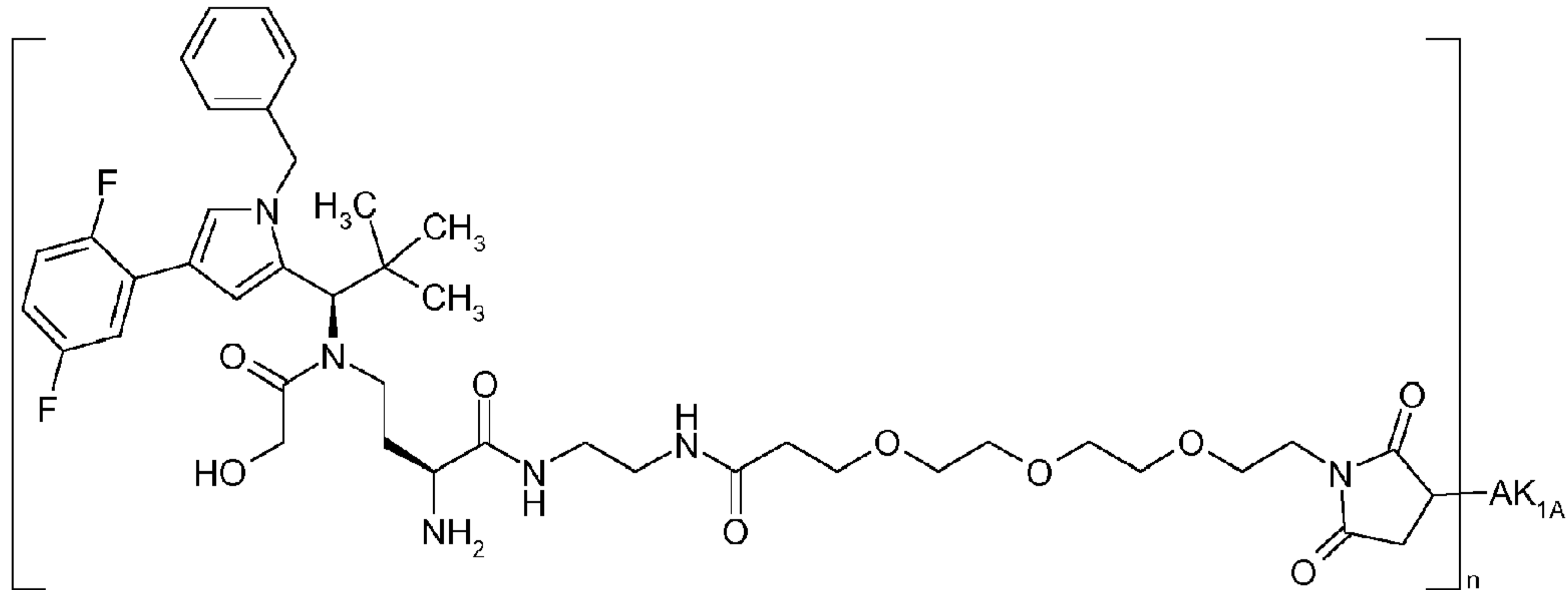
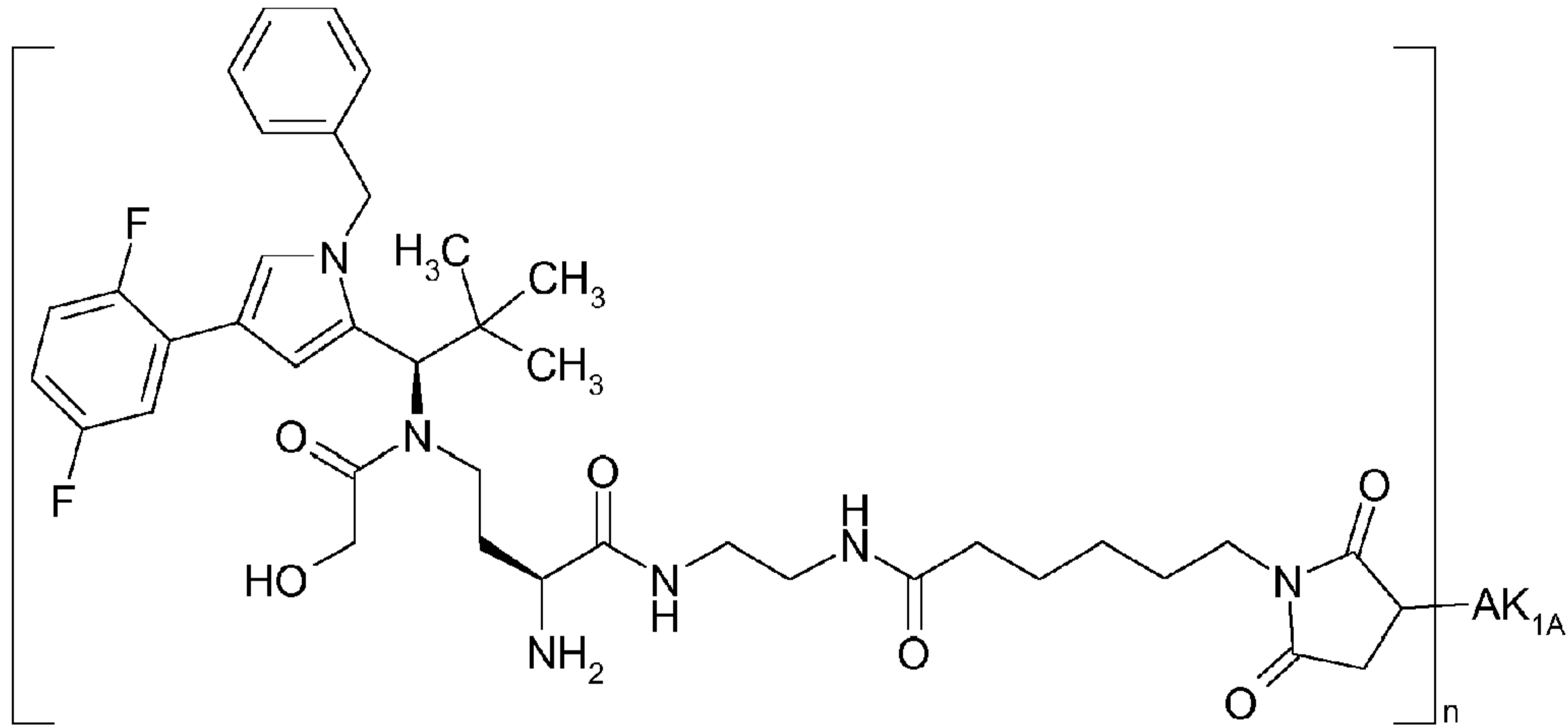
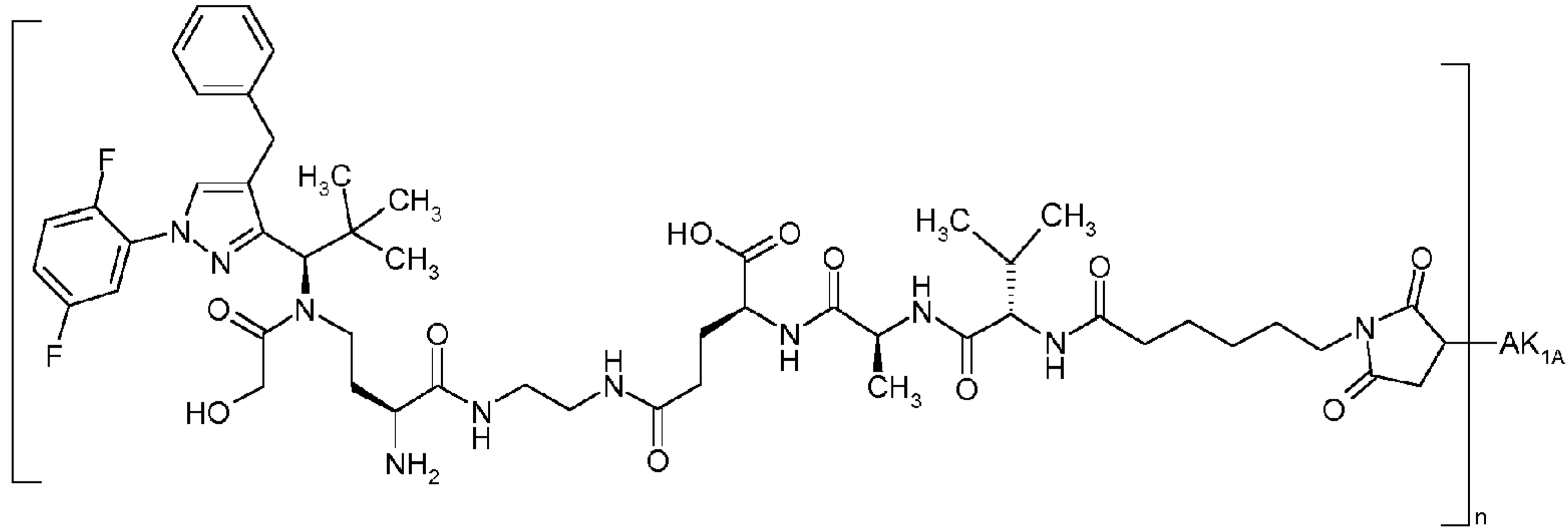
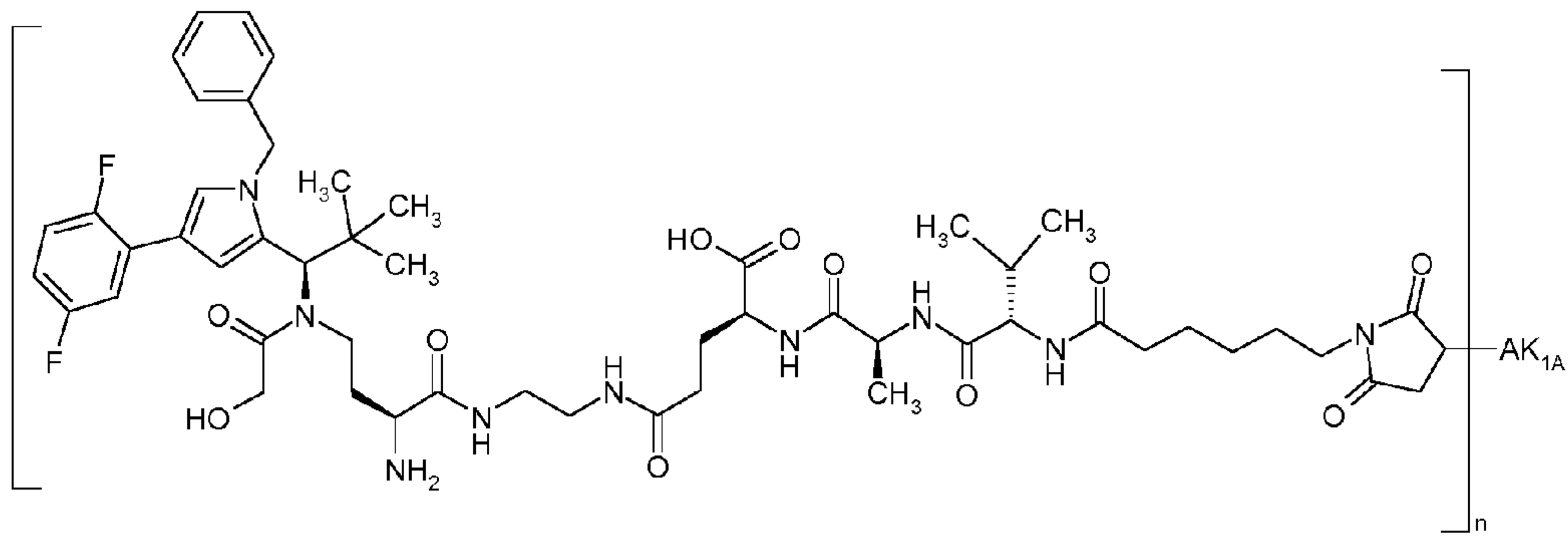


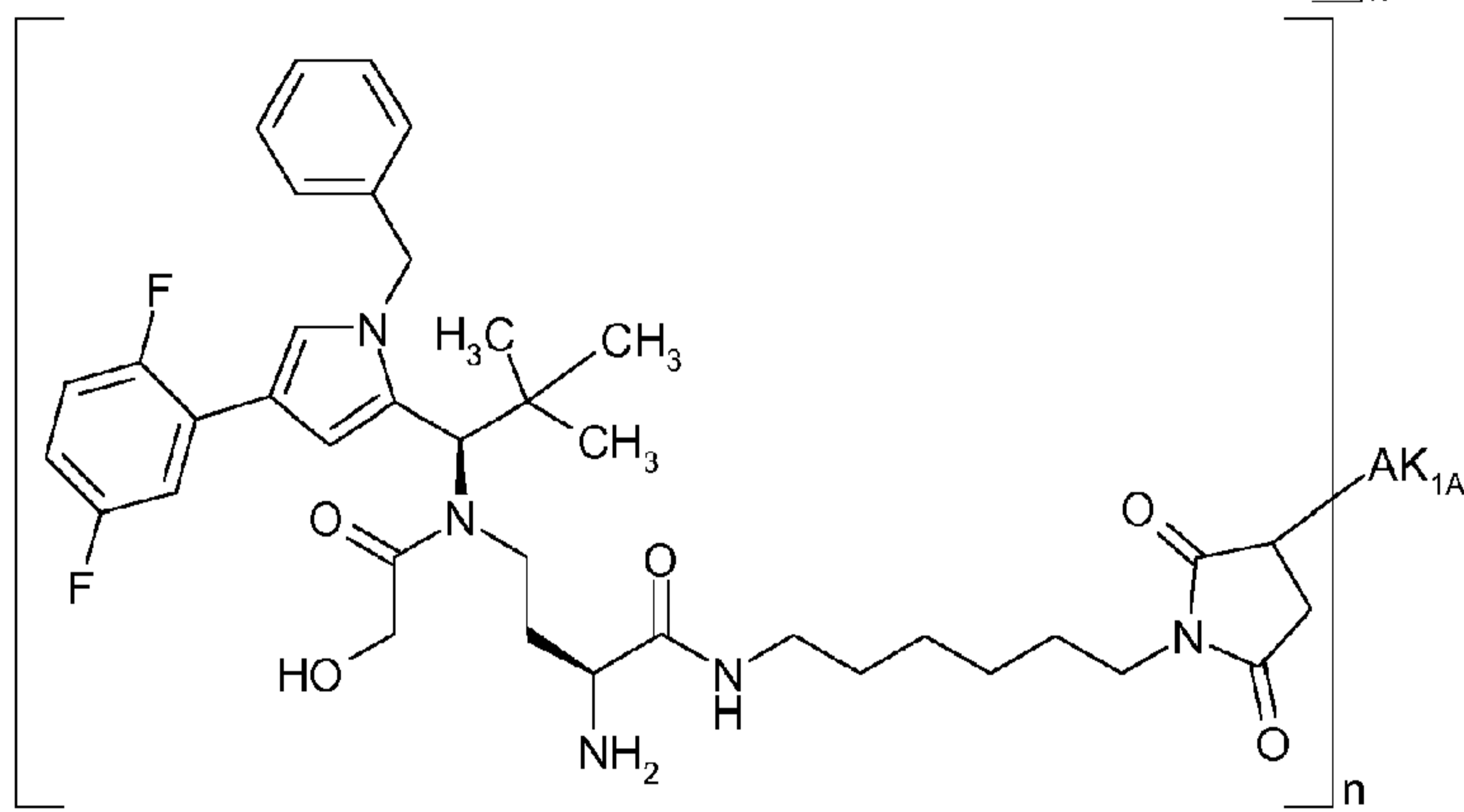
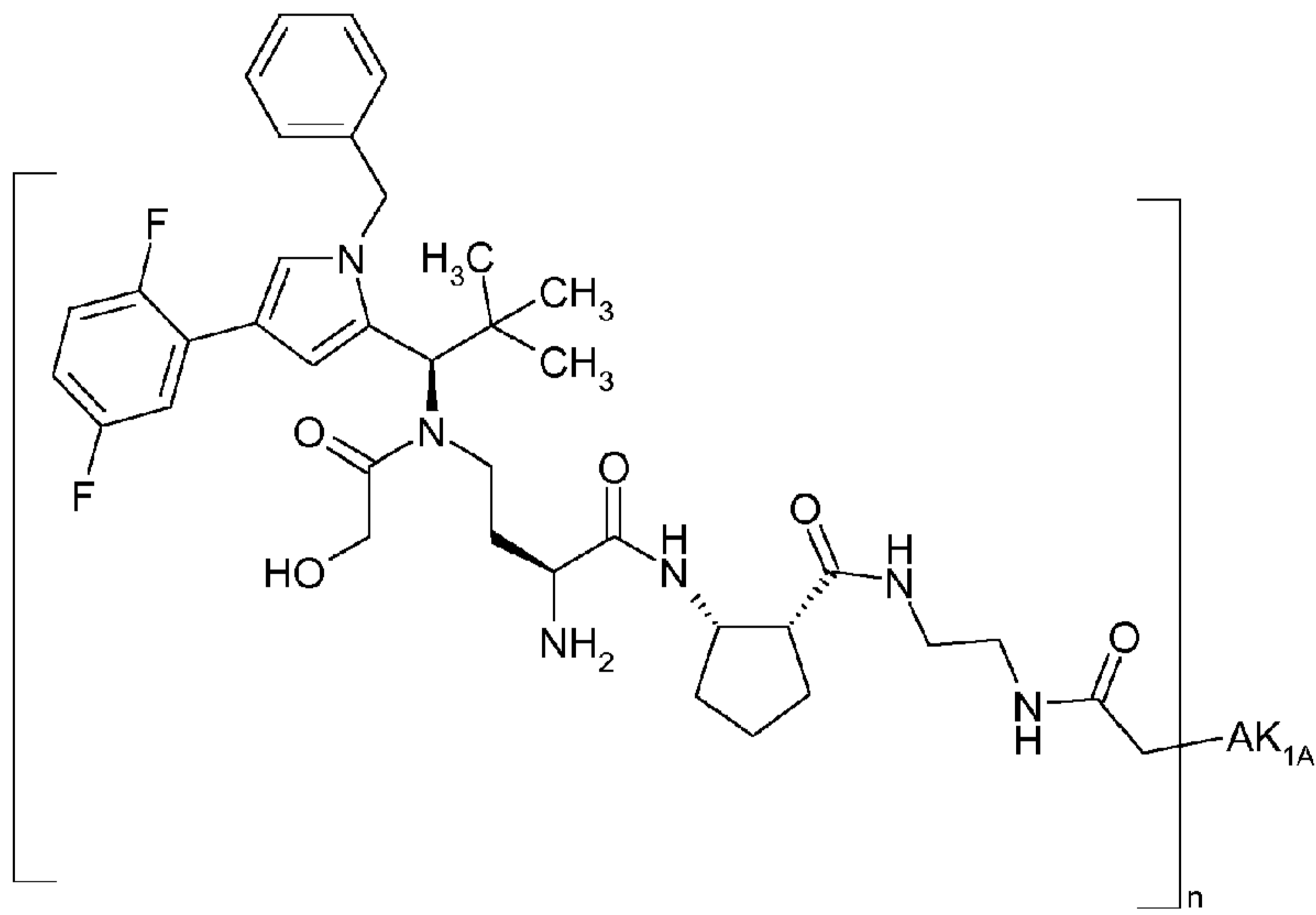
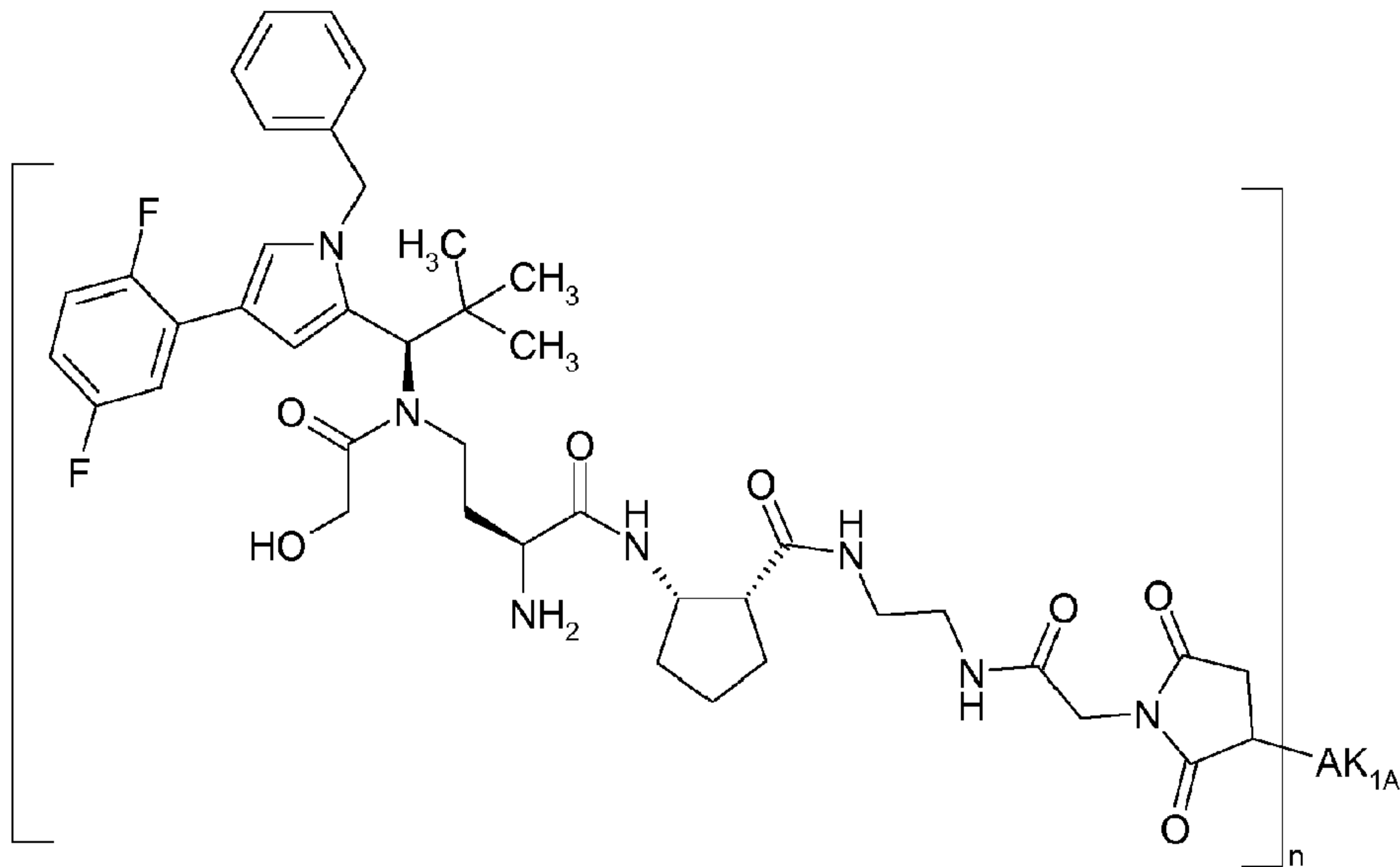


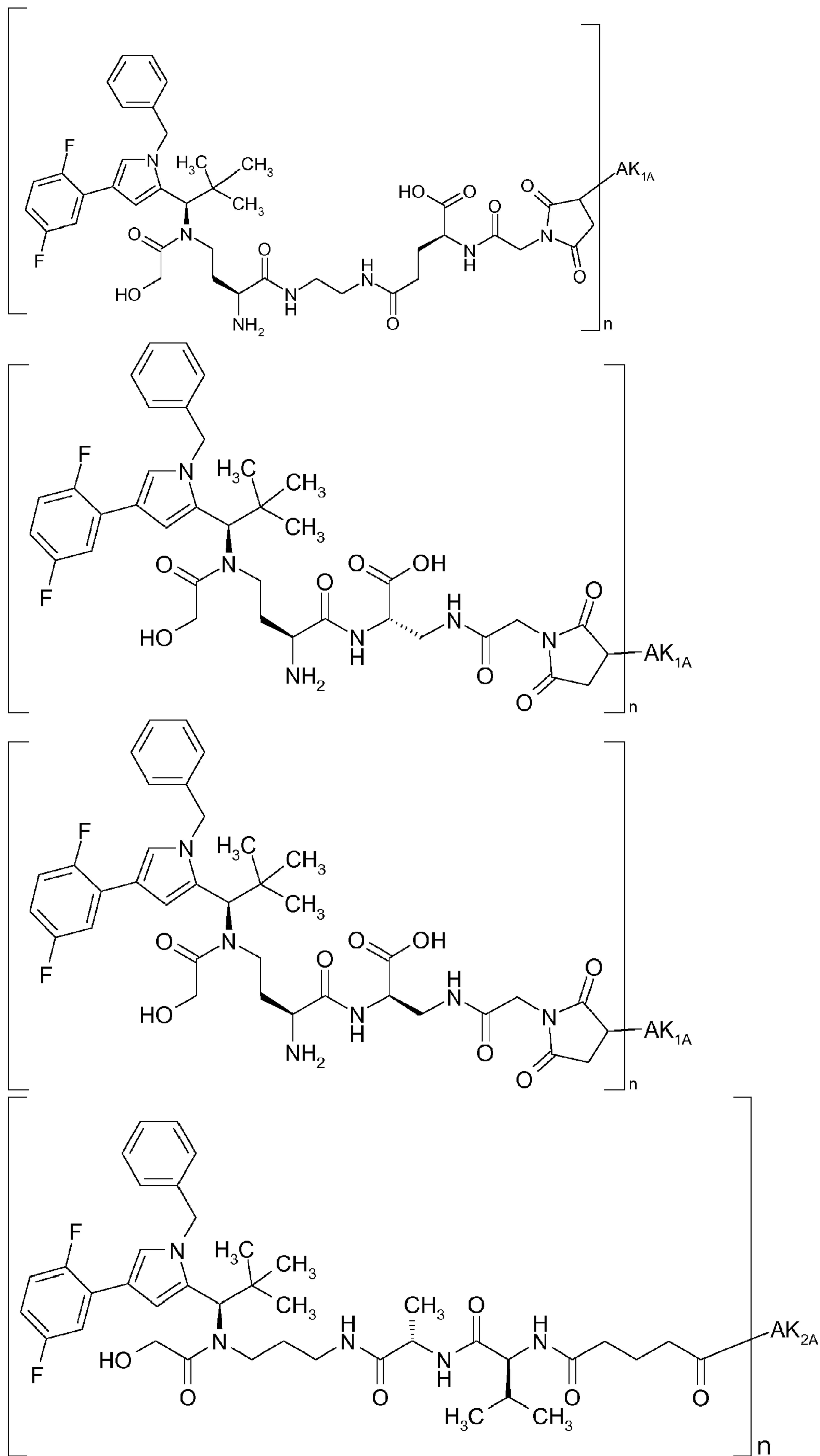


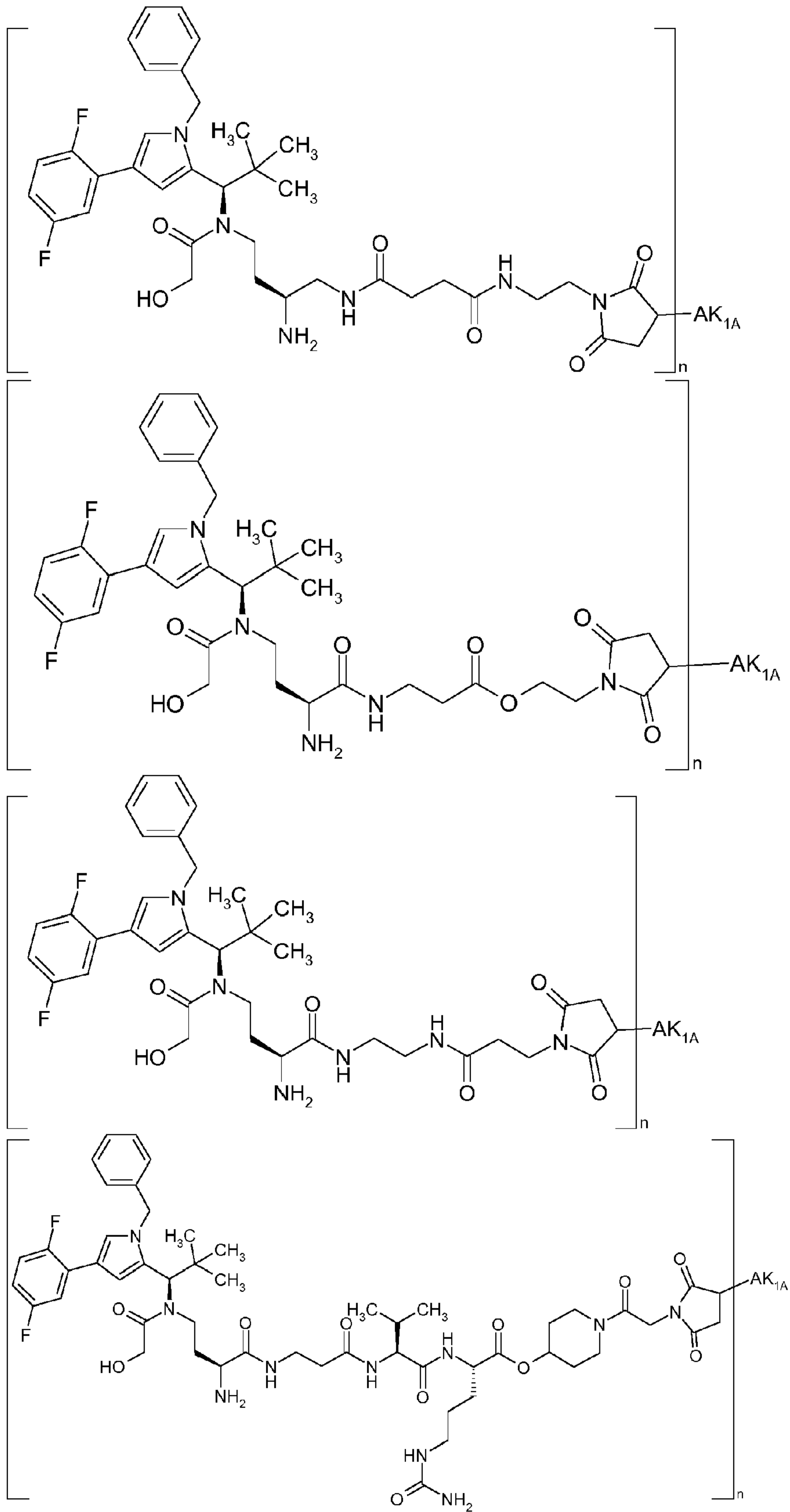






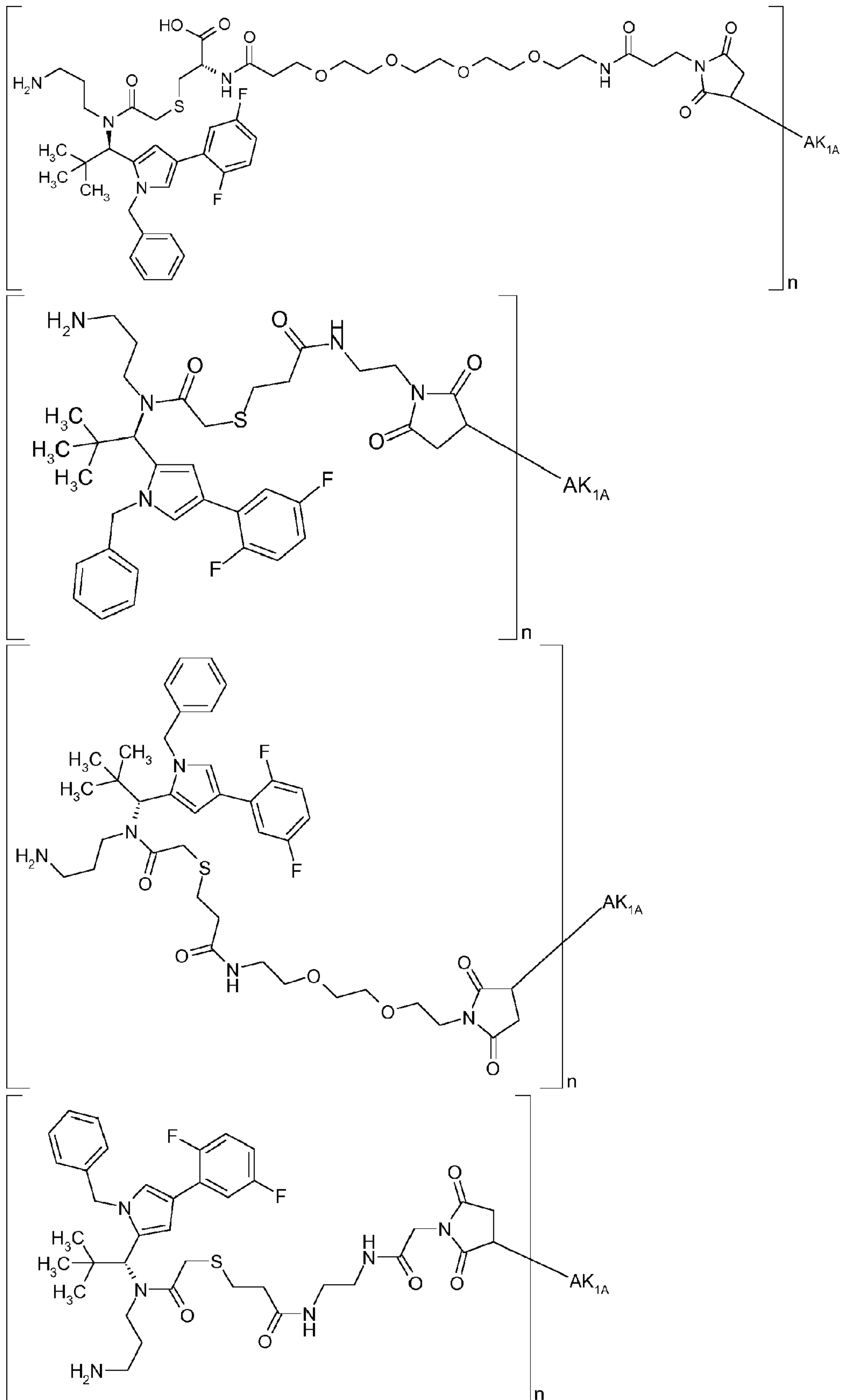


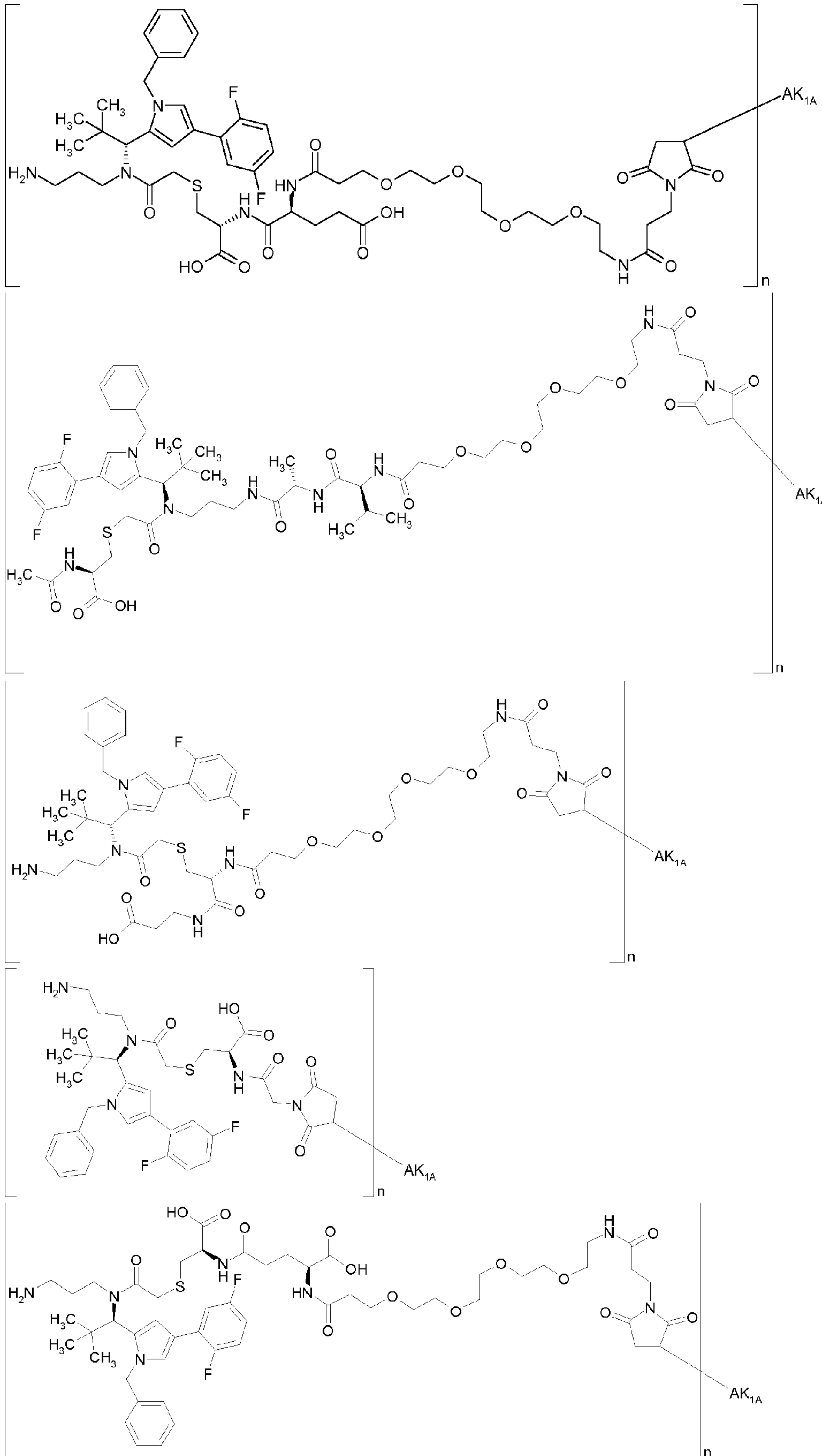


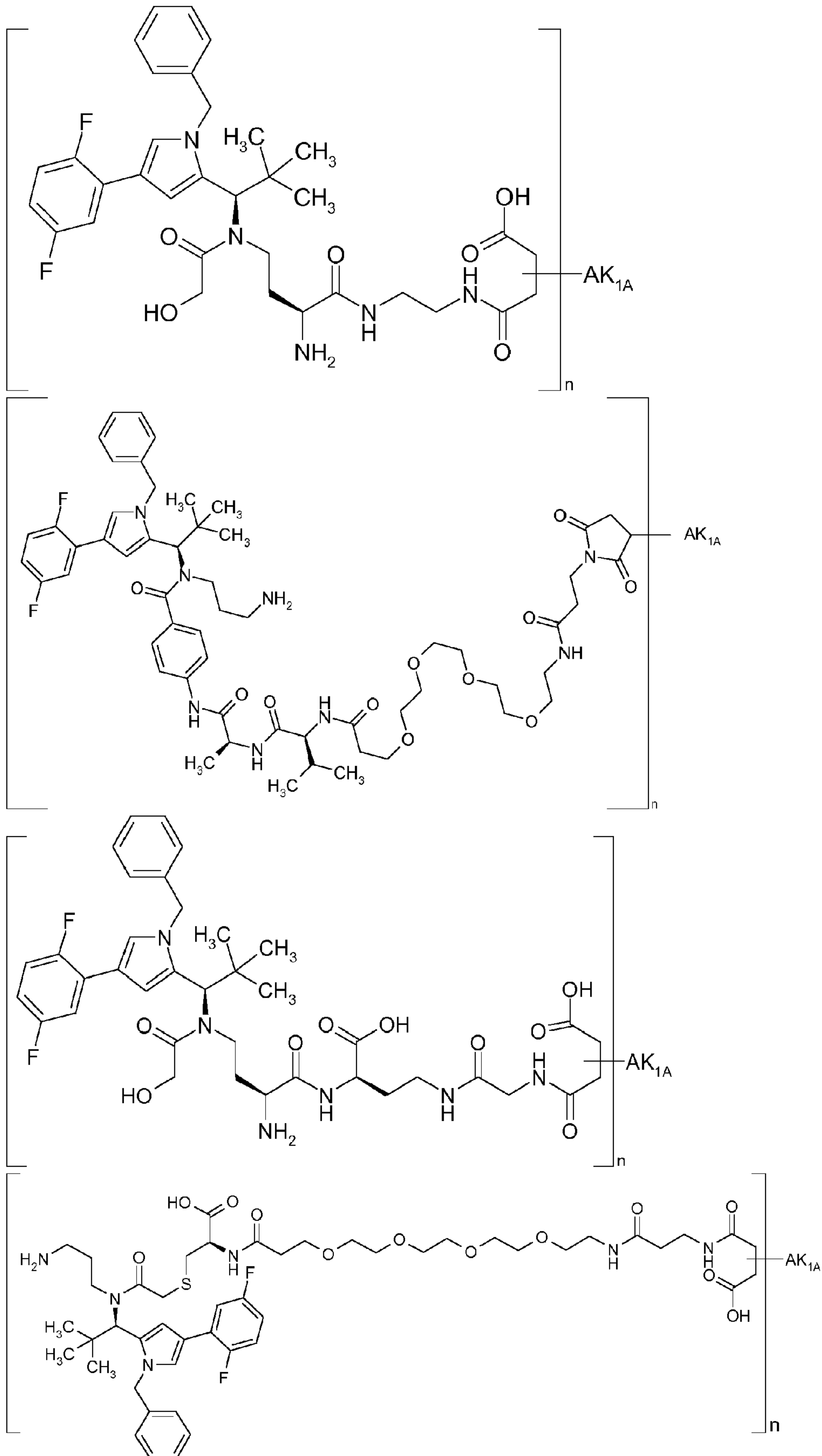


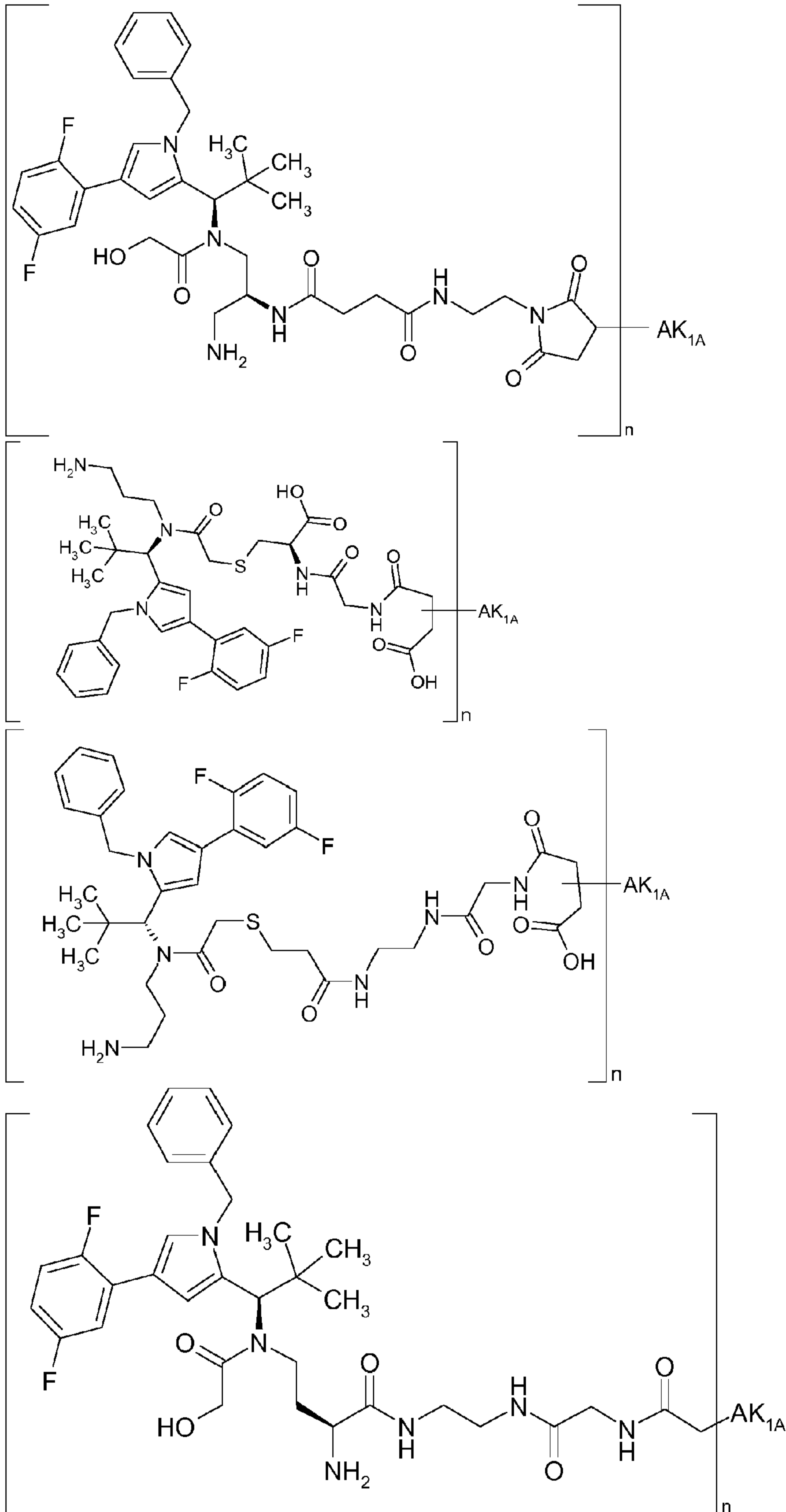


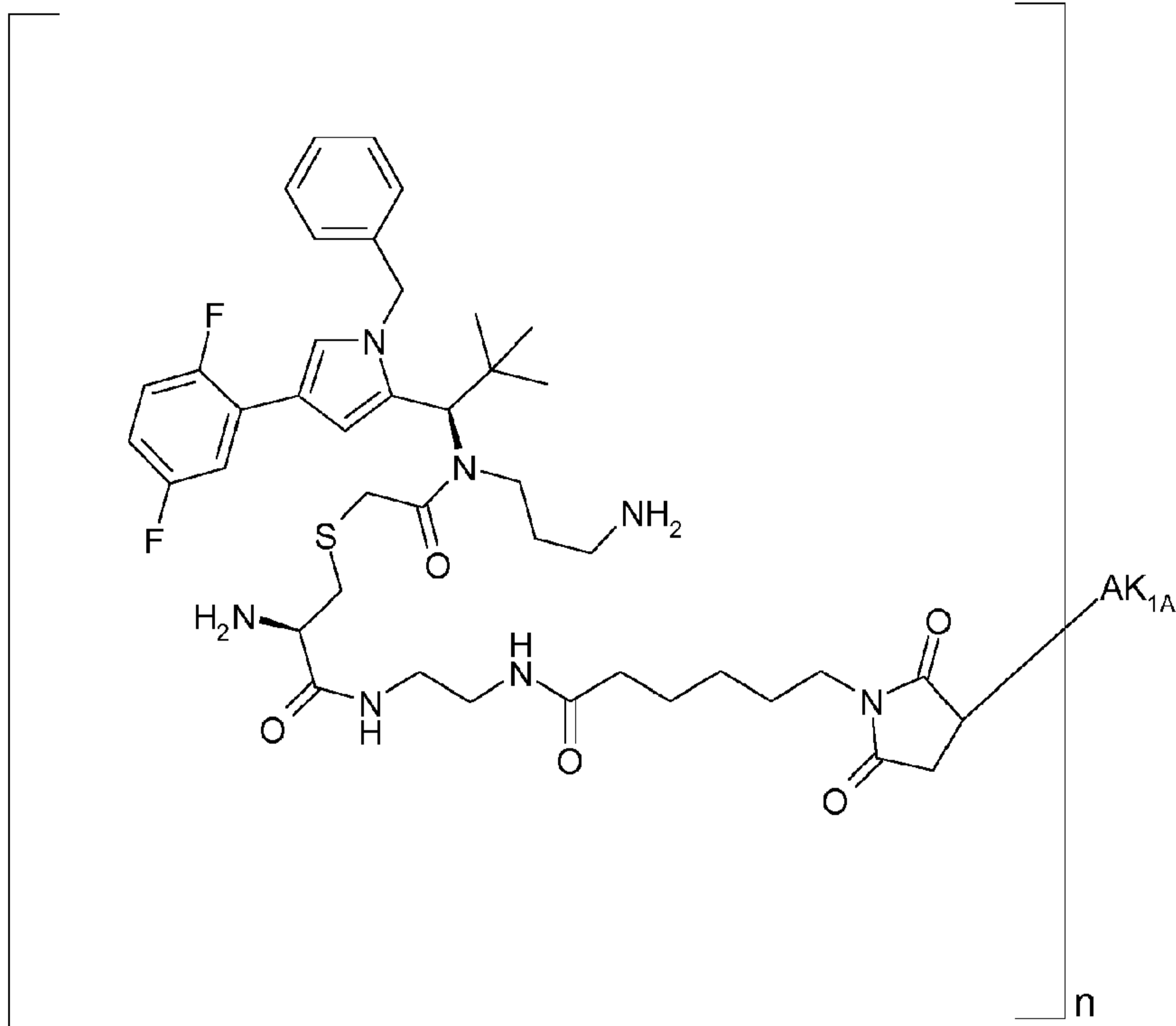
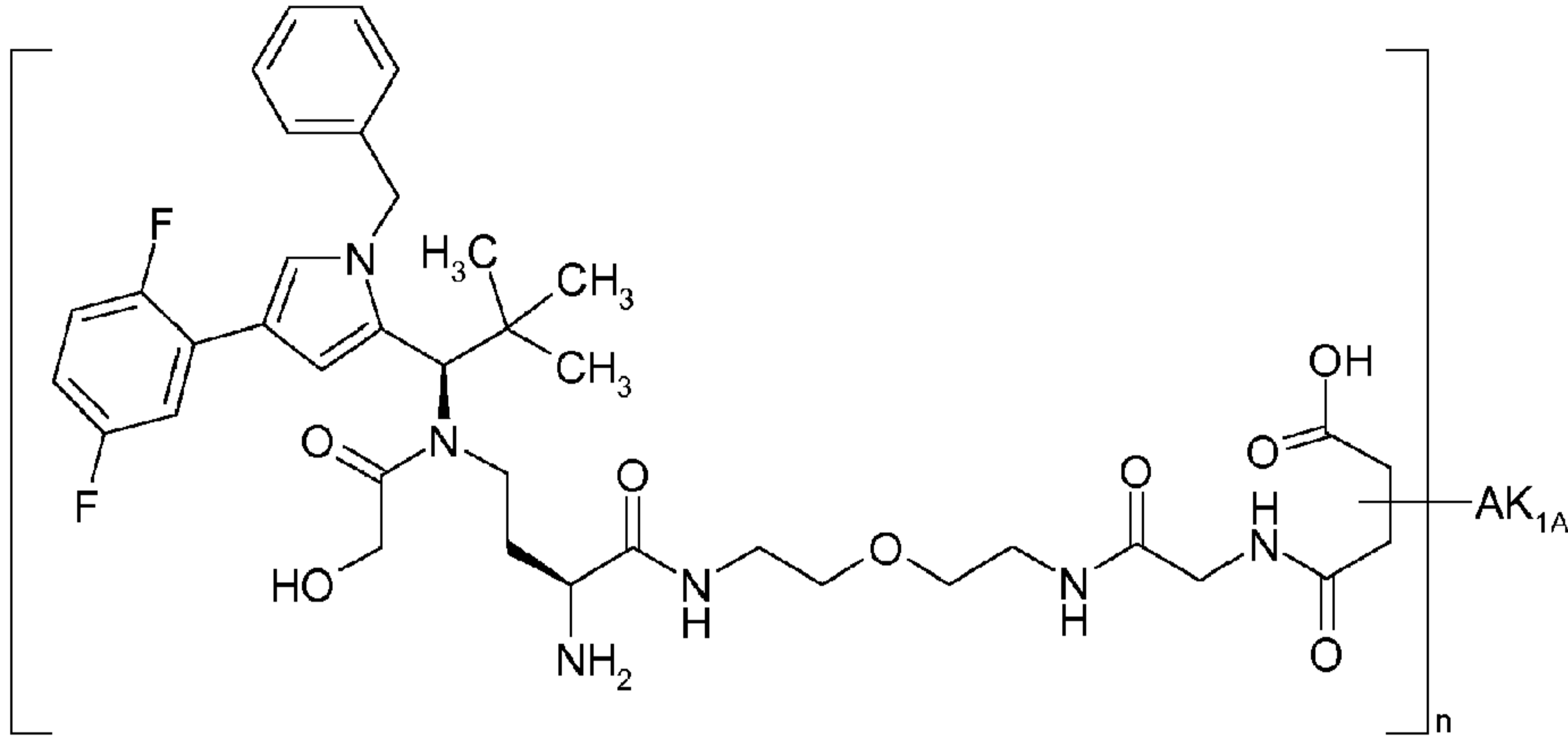
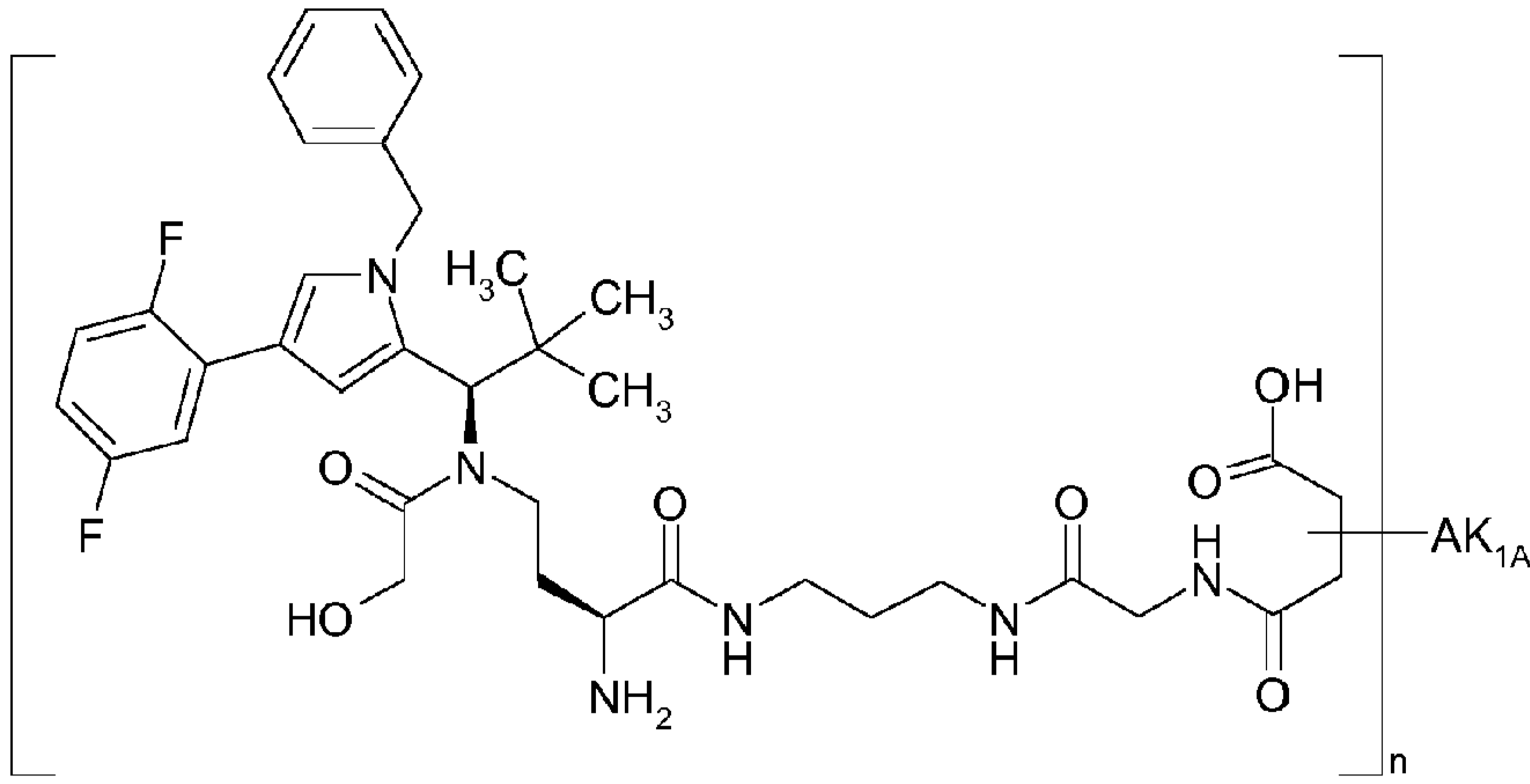


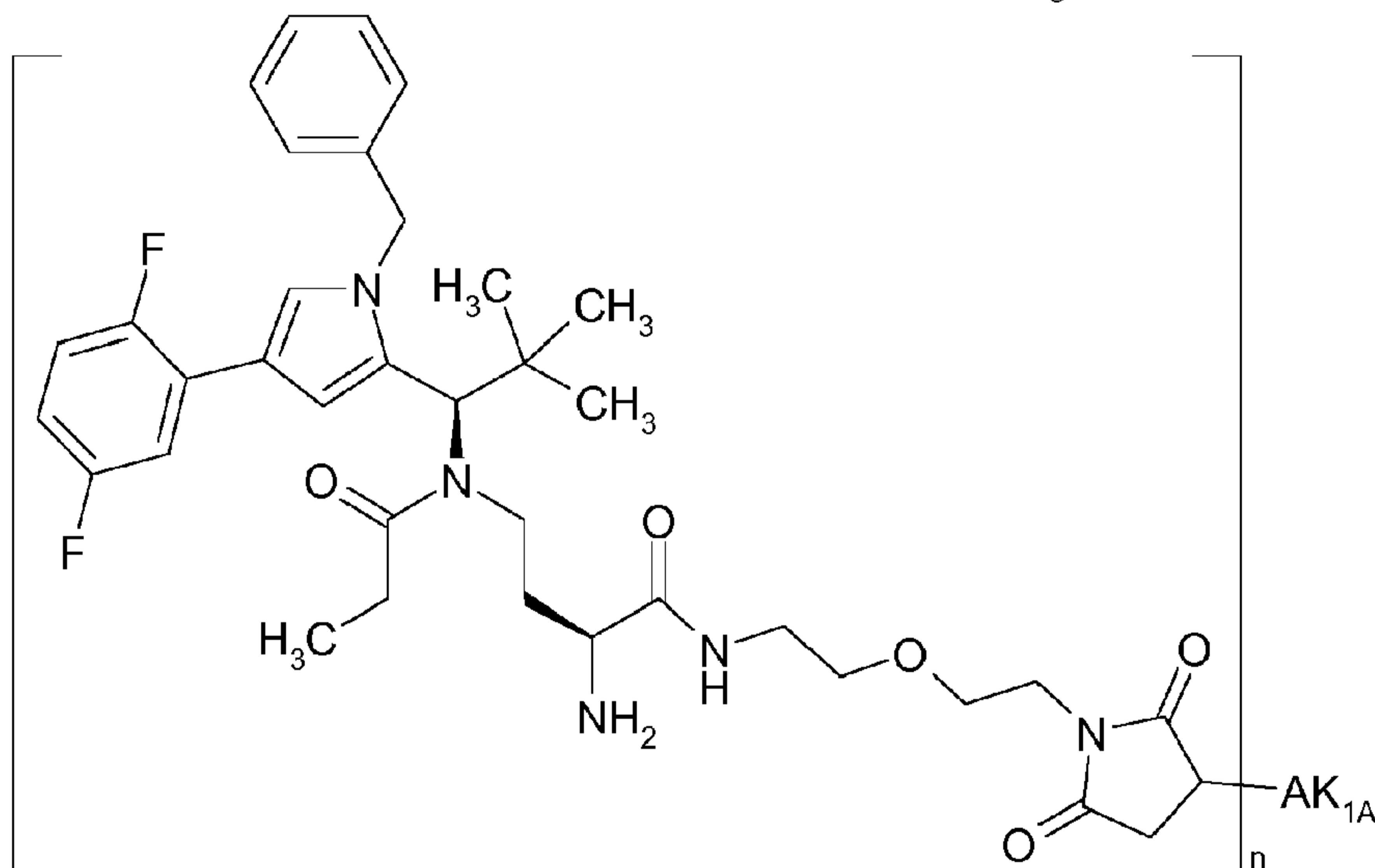
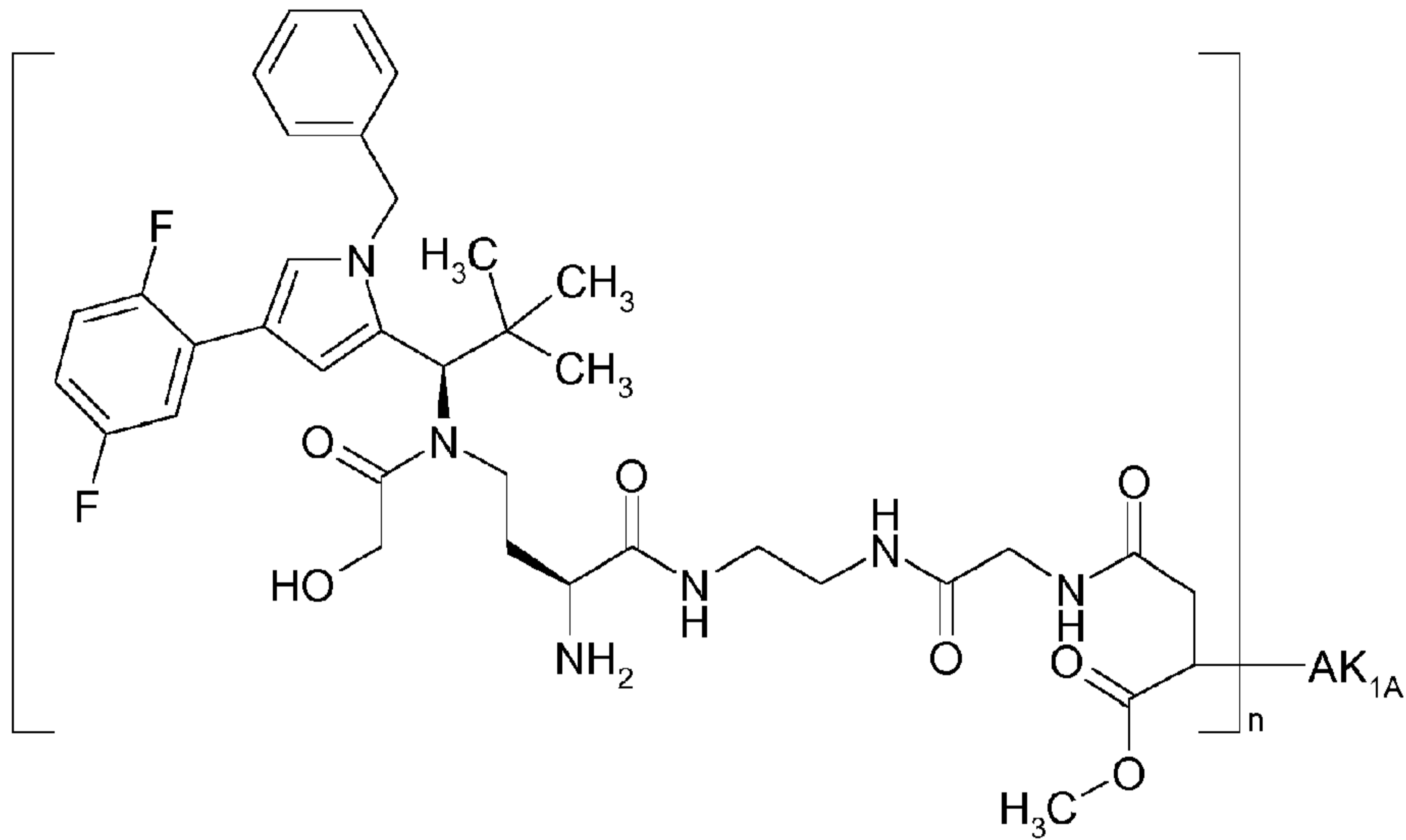
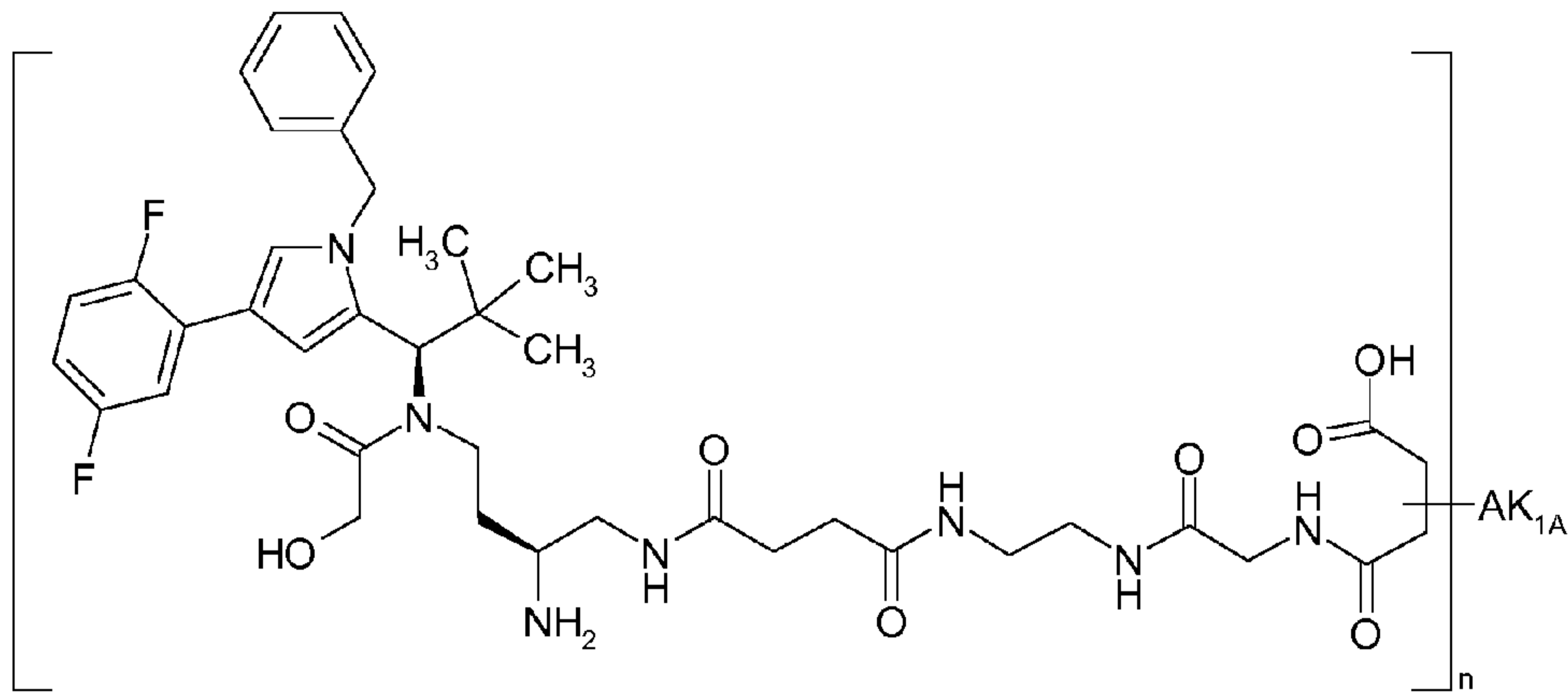


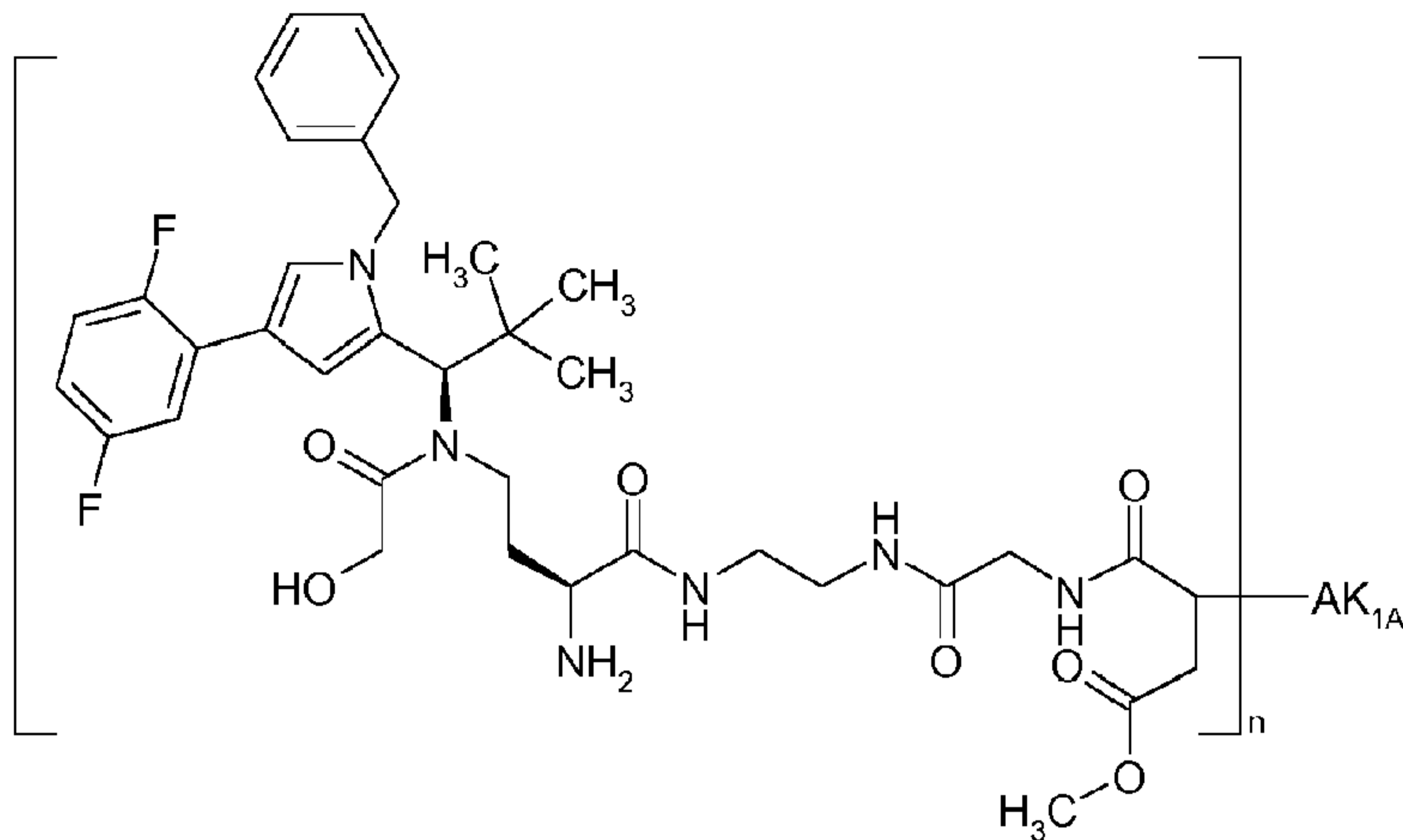




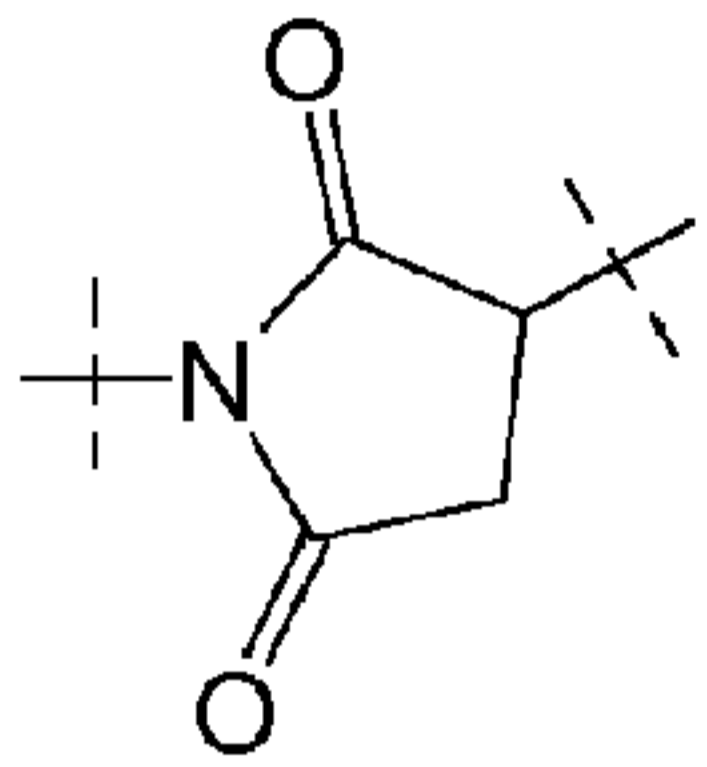








idet ADC, såfremt det via en gruppe



er forbundet med antistoffet, i stedet for denne gruppe også kan  
5 foreligge i det mindste delvist i form af de hydrolyserede  
åbenkædede ravsyreamider sammenknyttet med antistoffet.

66. Konjugat ifølge krav 65, idet antistoffet er et humant,  
10 humaniseret eller kimært monoklonalt antistof eller et antigen-  
bindende fragment heraf.

67. Konjugat ifølge krav 66, idet antistoffet er et anti-HER2-  
antistof, et anti-EGFR-antistof, et anti-TWEAKR-antistof eller  
et antigen-bindende fragment heraf.

15

68. Konjugat ifølge krav 67, idet anti-TWEAKR-antistoffet  
specifikt binder til aminosyren D i position 47 (D47) af TWEAKR  
(SEQ ID NO:169), fortrinsvis anti-TWEAKR-antistoffet TPP-2090.





Fig. 2:

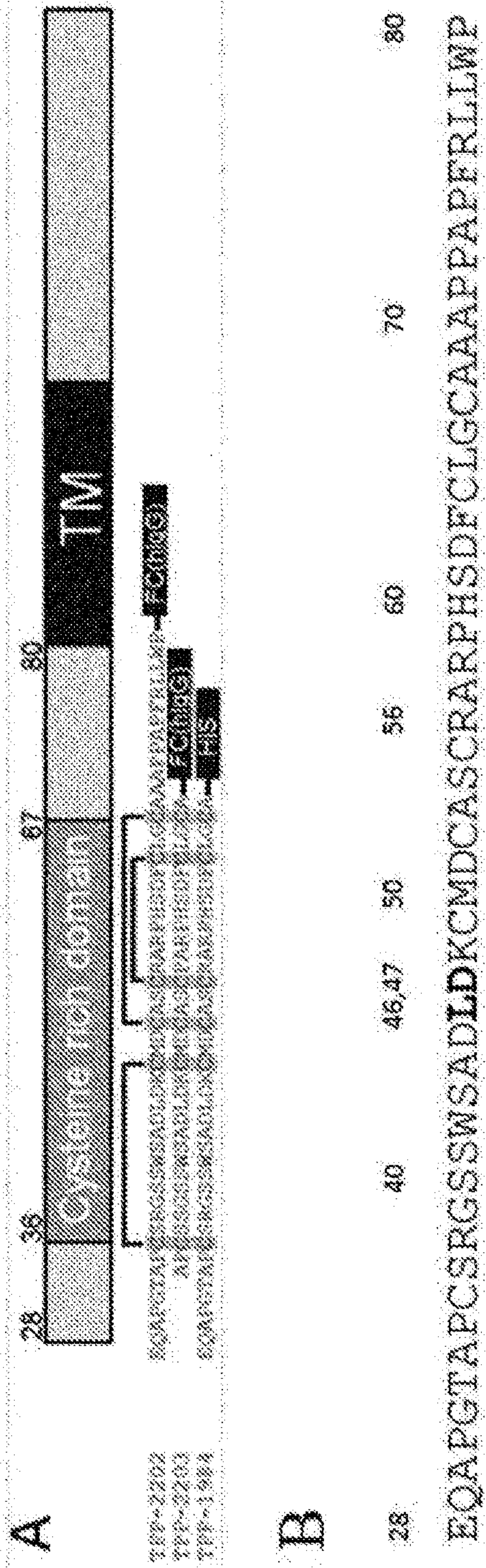


Fig. 3:

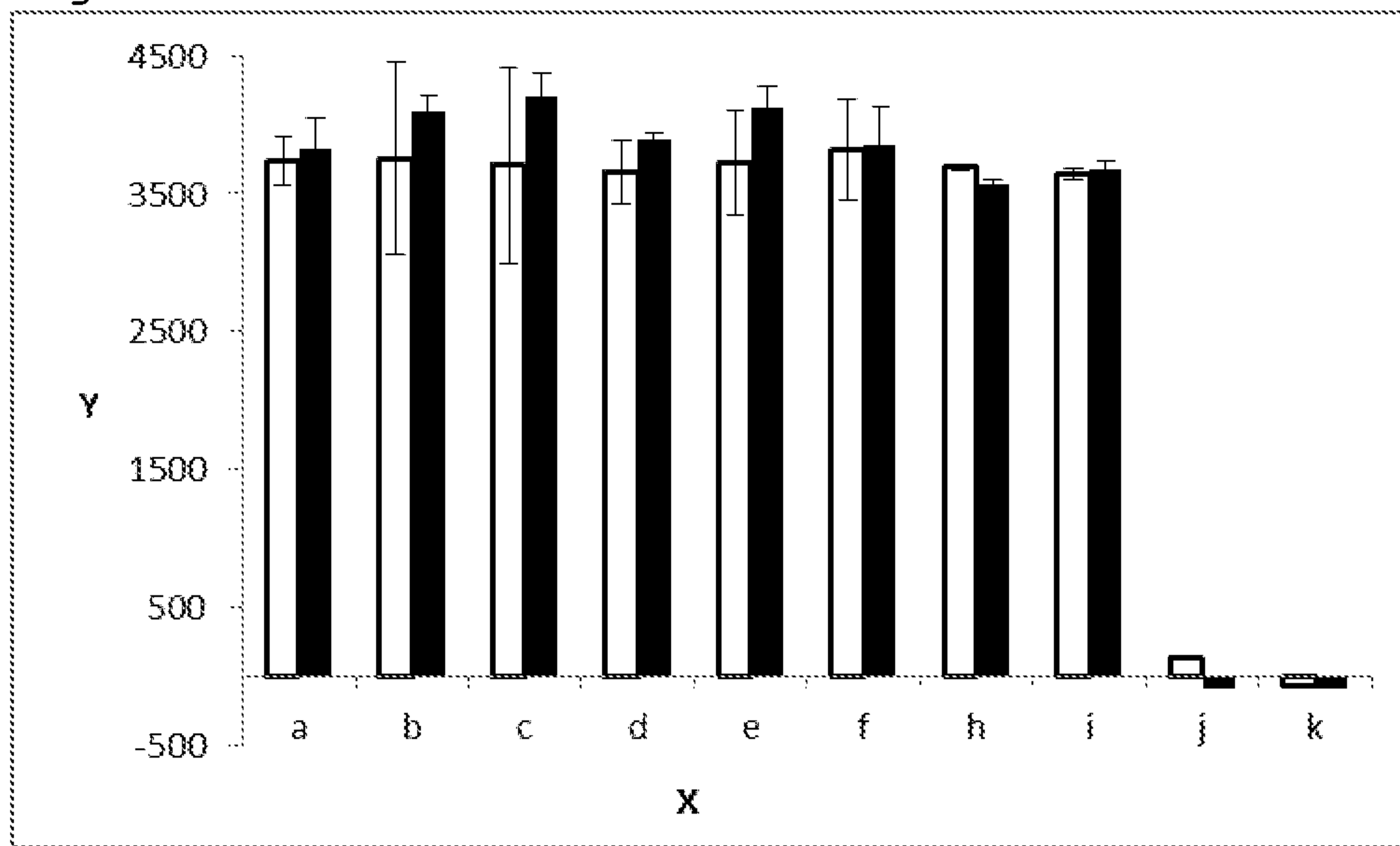


Fig. 4:

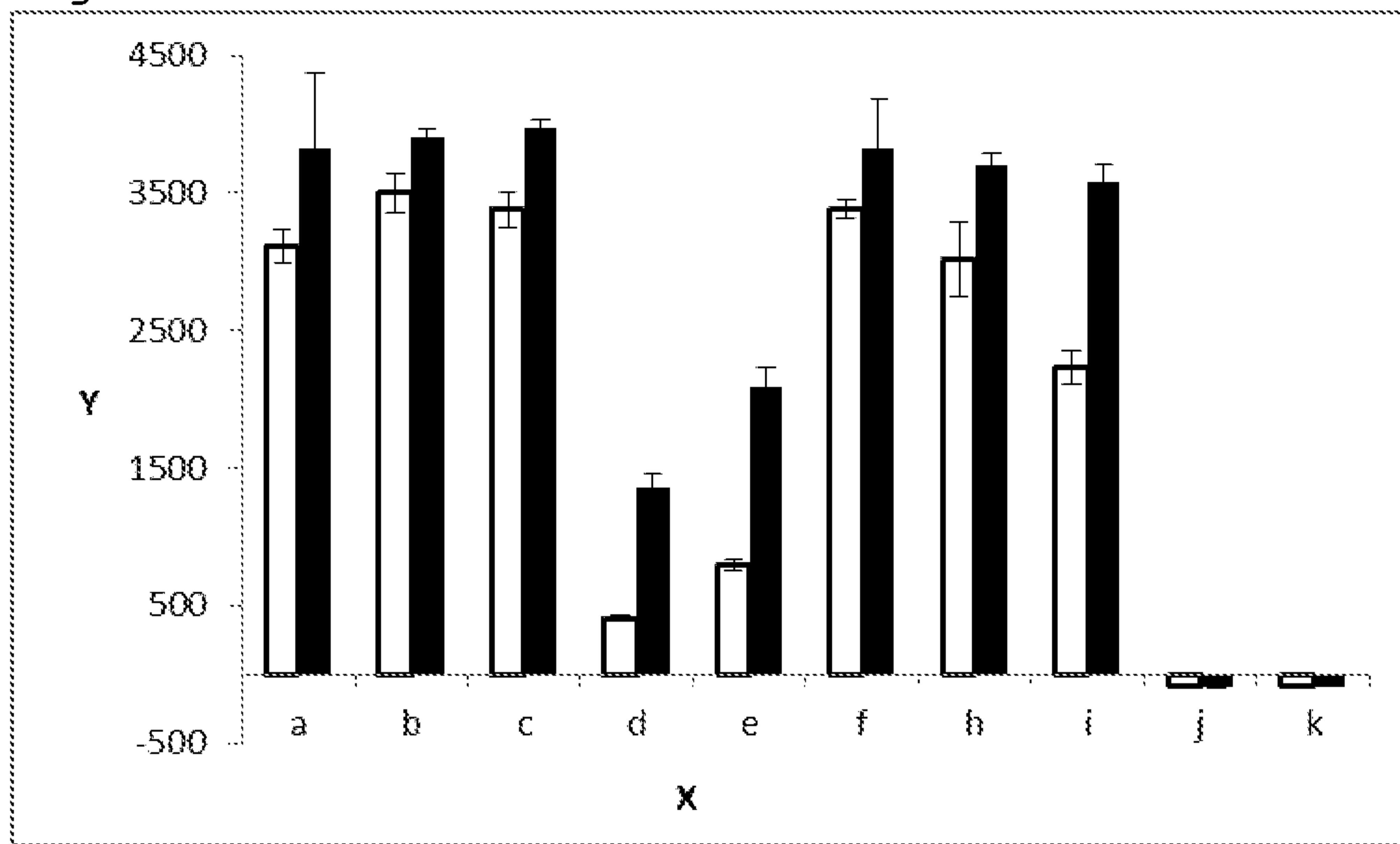


Fig. 5:

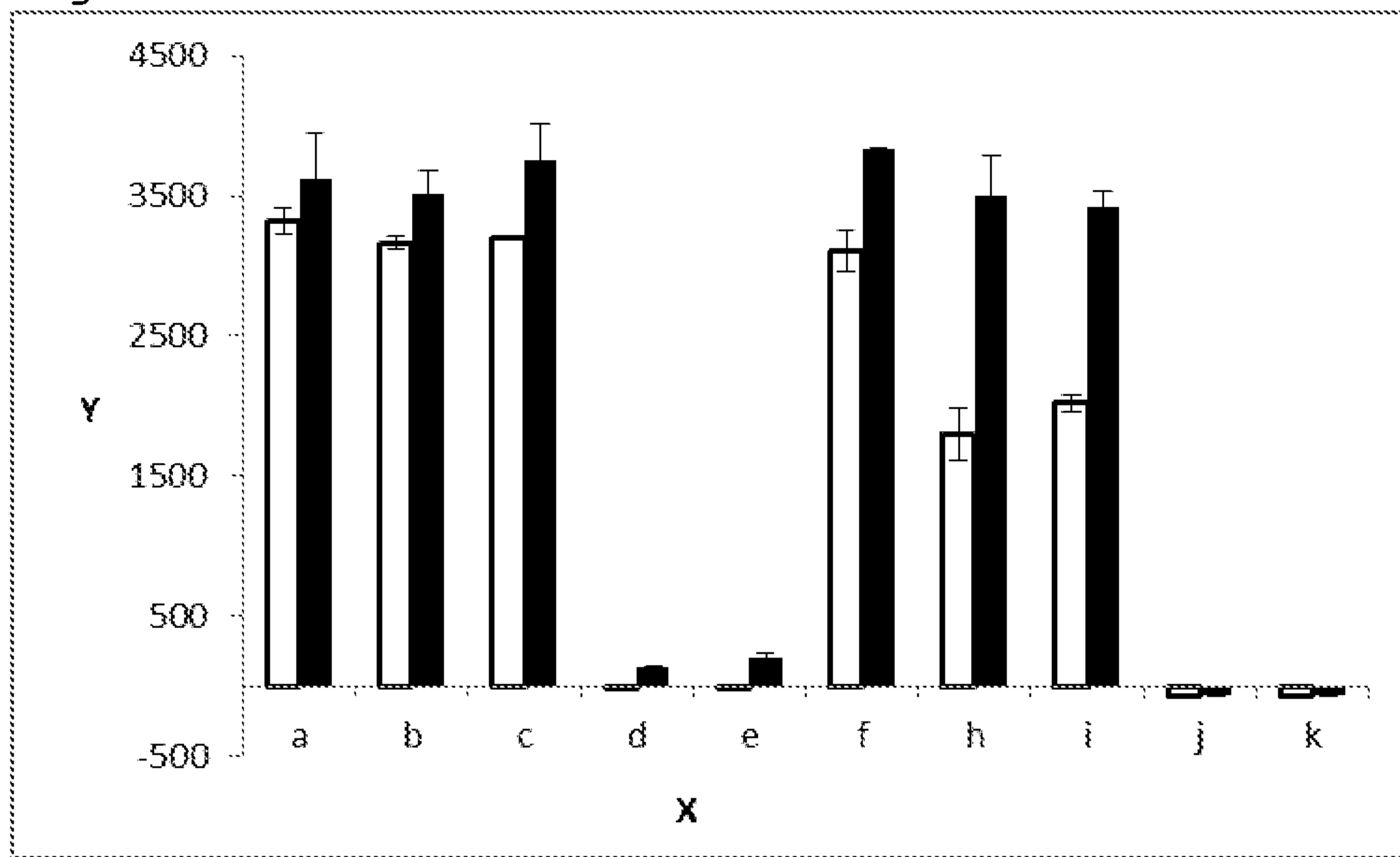
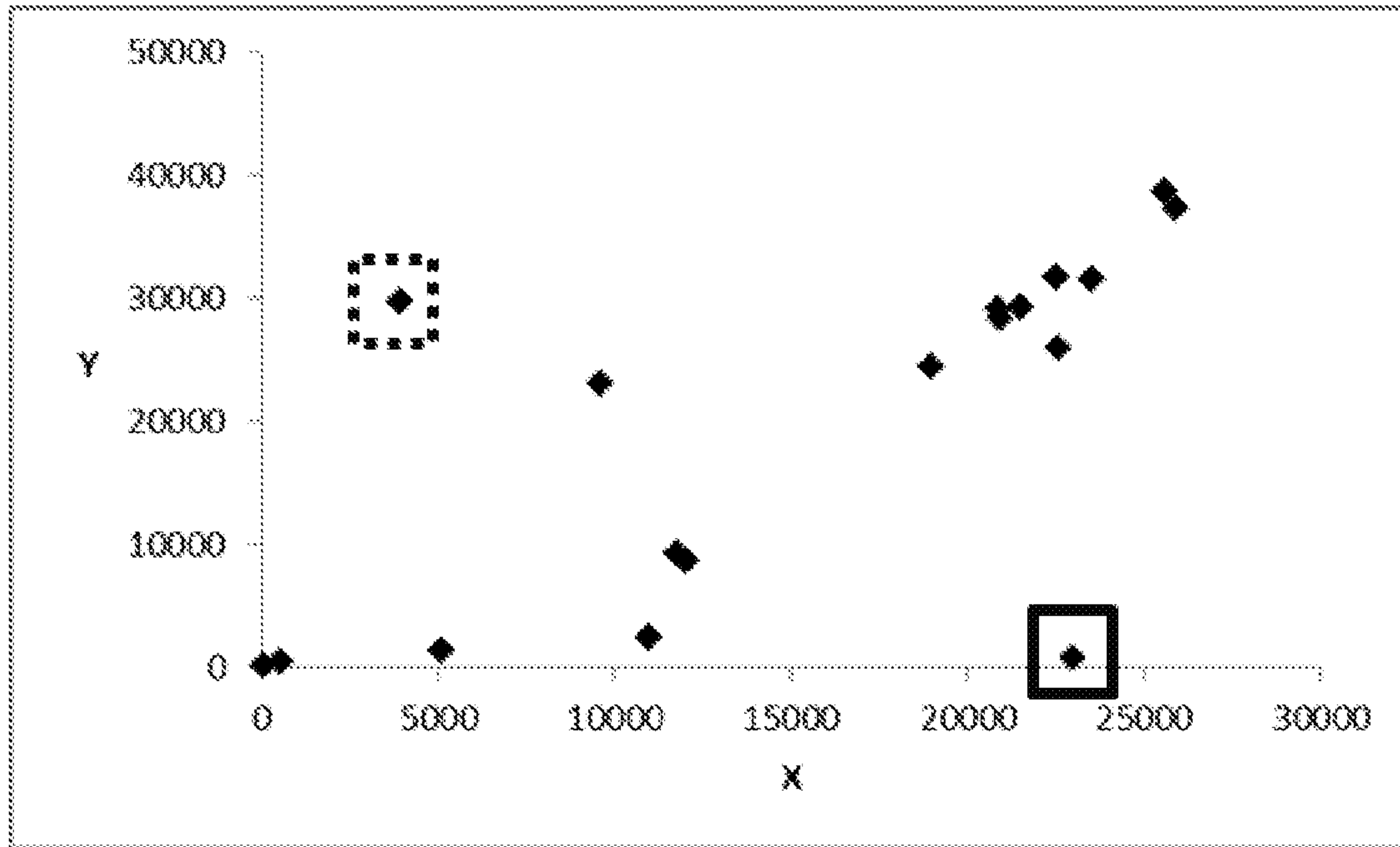


Fig. 6:

A



B

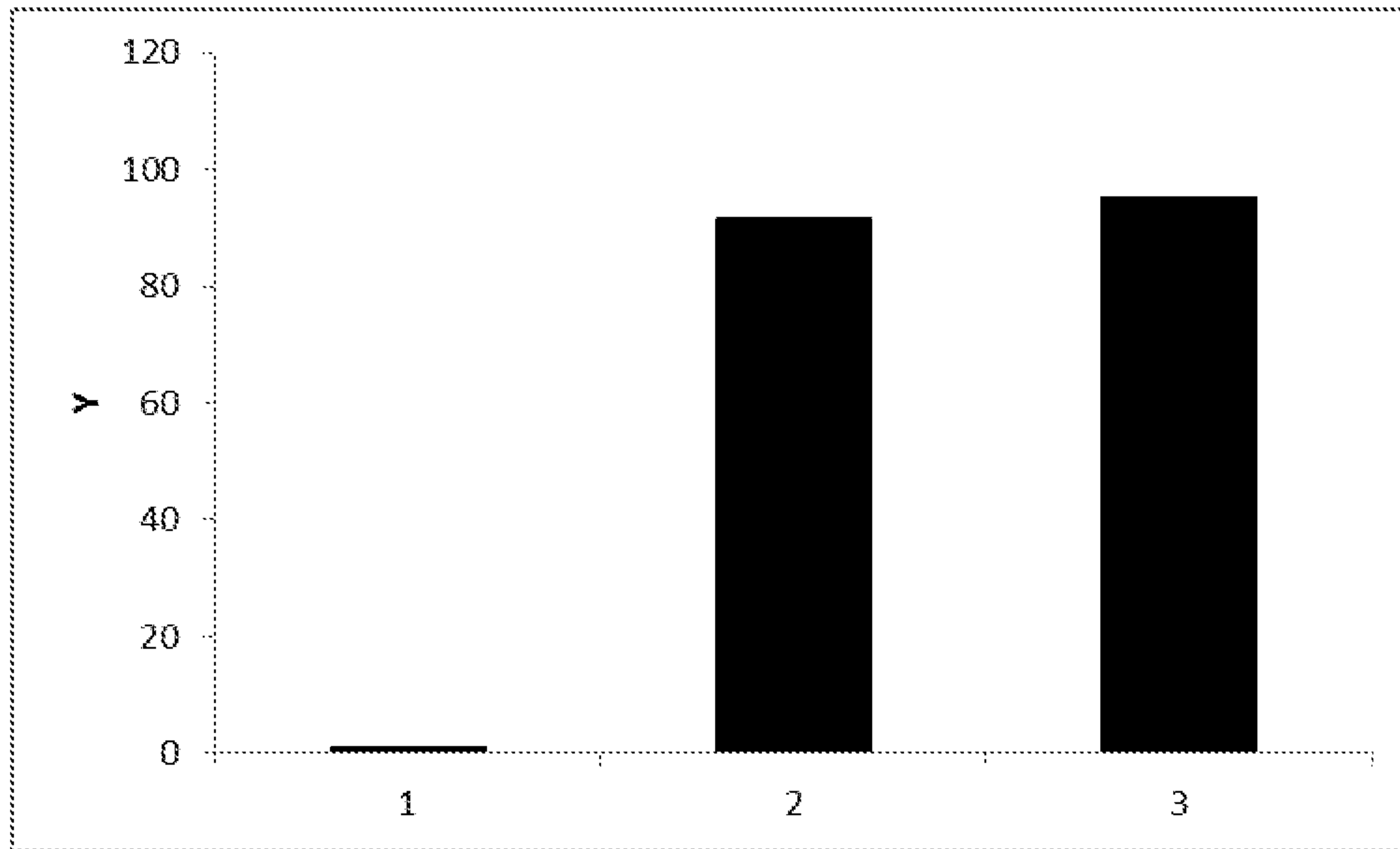


Fig. 7:

