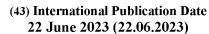
(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization

International Bureau







(10) International Publication Number WO 2023/114473 A3

- (51) International Patent Classification: *C12N 9/12* (2006.01)
- (21) International Application Number:

PCT/US2022/053174

(22) International Filing Date:

16 December 2022 (16.12.2022)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:

63/421,919

63/290,329 16 December 2021 (16.12.2021) US PCT/US2022/027024

29 April 2022 (29.04.2022) US

PCT/US2022/033199

13 June 2022 (13.06.2022) US 02 November 2022 (02.11.2022) US

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- (81) Designated States (unless otherwise indicated, for every kind of national protection available): AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BN, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CV, CZ, DE, DJ, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IQ, IR, IS, IT, JM, JO, JP, KE, KG, KH, KN, KP, KR, KW, KZ, LA, LC, LK, LR, LS, LU, LY, MA, MD, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PA, PE, PG, PH, PL, PT, QA, RO, RS,

(54) Title: RECOMBINANT REVERSE TRANSCRIPTASE VARIANTS FOR IMPROVED PERFORMANCE

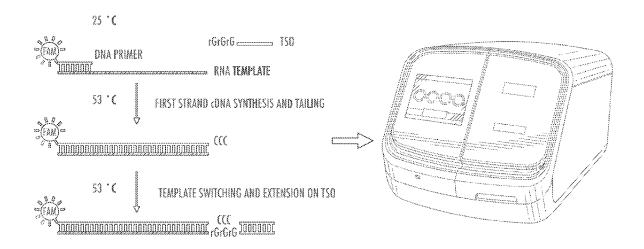


FIG. 1

(57) Abstract: The disclosure provides recombinant reverse transcriptases comprising one or more DNA binding domains conjugated to an engineered reversed transcriptase that have been modified to exhibit one or more altered reverse transcriptase related activities such as but not limited to altered template switching efficiency, altered transcription efficiency or both. The disclosure further provides compositions and kits comprising the recombinant reverse transcriptase enzymes and methods of producing, amplifying or sequencing nucleic acid molecules using these fusion reverse transcriptase enzymes.

RU, RW, SA, SC, SD, SE, SG, SK, SL, ST, SV, SY, TH, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, WS, ZA, ZM, ZW.

(84) Designated States (unless otherwise indicated, for every kind of regional protection available): ARIPO (BW, CV, GH, GM, KE, LR, LS, MW, MZ, NA, RW, SD, SL, ST, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, RU, TJ, TM), European (AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, ME, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, KM, ML, MR, NE, SN, TD, TG).

Declarations under Rule 4.17:

- as to applicant's entitlement to apply for and be granted a patent (Rule 4.17(ii))
- as to the applicant's entitlement to claim the priority of the earlier application (Rule 4.17(iii))

Published:

- with international search report (Art. 21(3))
- before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments (Rule 48.2(h))
- (88) Date of publication of the international search report: 20 July 2023 (20.07.2023)

International application No

PCT/US2022/053174

A. CLASSIFICATION OF SUBJECT MATTER

INV. C12N9/12

ADD.

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

C12N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

EPO-Internal, WPI Data, Sequence Search

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
x	WO 2018/200867 A1 (10X GENOMICS INC [US])	1,3-5,7,
	1 November 2018 (2018-11-01)	17,18,
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	22 February 2007 (2007-02-22)	47-50,
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		82-85
	sequence 35	
		
	-/	

Further documents are listed in the continuation of Box C.	X See patent family annex.
* Special categories of cited documents :	"T" later document published after the international filing date or priority
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"E" earlier application or patent but published on or after the international filing date	"X" document of particular relevance;; the claimed invention cannot be considered novel or cannot be considered to involve an inventive
"L" document which may throw doubts on priority claim(s) or which is	step when the document is taken alone
cited to establish the publication date of another citation or other special reason (as specified)	"Y" document of particular relevance;; the claimed invention cannot be considered to involve an inventive step when the document is
"O" document referring to an oral disclosure, use, exhibition or other means	combined with one or more other such documents, such combination being obvious to a person skilled in the art
"P" document published prior to the international filing date but later than the priority date claimed	"&" document member of the same patent family
Date of the actual completion of the international search	Date of mailing of the international search report
19 April 2023	23/06/2023
15 APITI 2025	237 007 2023
Name and mailing address of the ISA/	Authorized officer
European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijswijk	
Tel. (+31-70) 340-2040,	
Fax: (+31-70) 340-3016	Blanco Urgoiti, B

International application No
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Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
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	example 3	32-35,
	figure 6	37-46,
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	the whole document	
		

International application No.

INTERNATIONAL SEARCH REPORT

PCT/US2022/053174

Вох	No. I	Nucleotide and/or amino acid sequence(s) (Continuation of item 1.c of the first sheet)
1.		ard to any nucleotide and/or amino acid sequence disclosed in the international application, the international search was ut on the basis of a sequence listing:
	a	forming part of the international application as filed.
	b. X	furnished subsequent to the international filing date for the purposes of international search (Rule 13 ter.1(a)).
		x accompanied by a statement to the effect that the sequence listing does not go beyond the disclosure in the international application as filed.
2.	Ш ,	With regard to any nucleotide and/or amino acid sequence disclosed in the international application, this report has been established to the extent that a meaningful search could be carried out without a WIPO Standard ST.26 compliant sequence listing.
3.	Additiona	al comments:

International application No. PCT/US2022/053174

INTERNATIONAL SEARCH REPORT

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)
This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1. Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
Claims Nos.: because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
 As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fees, this Authority did not invite payment of additional fees.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
 No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims;; it is covered by claims Nos.: 1, 3-5, 7, 12, 17, 18, 20-85 (all partially)
The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee. The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.
No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. claims: 1, 3-5, 7, 12, 17, 18, 20-85 (all partially)

The reverse transcriptase comprising the following combination of mutations departing from SEQ ID NO:7 or 178: (a) (a) E69K, L139P, E302R, T306K, W313F, T330P, N454K; and one or more of M39V, P47L, Q91R, M66L, F155Y, D200N, D200E, H204R, G429S, L435G, L435K, P448A, D449G, H503V, D524N, T542D, E545G, D583N, H594Q, L603W, L603F, E607K, E607G, P627S, H634Y, H638G, A644V, D653H, K658R and L671P; or (b) E69K, L139P, D200N, E302R, T306K, W313F, T330P, L435G, P448A, D449G, N454K, D524N, L603W, and E607K and one or more of M39V, P47L, M66L, Q91R, F155Y, H204R, G429S, H503V, T542D, E545G, D583N, H594Q, P627S, H634Y, H638G, A644V, D653H, K658R and L671P; wherein the RT comprises the amino acid sequence of SEQ ID NO:1; fusion reverse transcriptase comprising a DNA-binding domain thereof; nucleic acid encoding the same; expression vector, host cell , methods, compositions and kits thereof.

2. claims: 13-16(completely); 1-5, 7-12, 17-85(partially)

Idem to invention 1 but related to SEQ ID NO:143

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3-30. claims: 1-12, 17-85(all partially)

invention 3 being idem to invention 1 but related to SEQ ID NO:180; invention 4 being idem to invention 1 but related to SEQ ID NO:181; [...]; invention 30 being idem to invention 1 but related to SEQ ID NO:208.

Information on patent family members

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PCT/US2022/053174

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