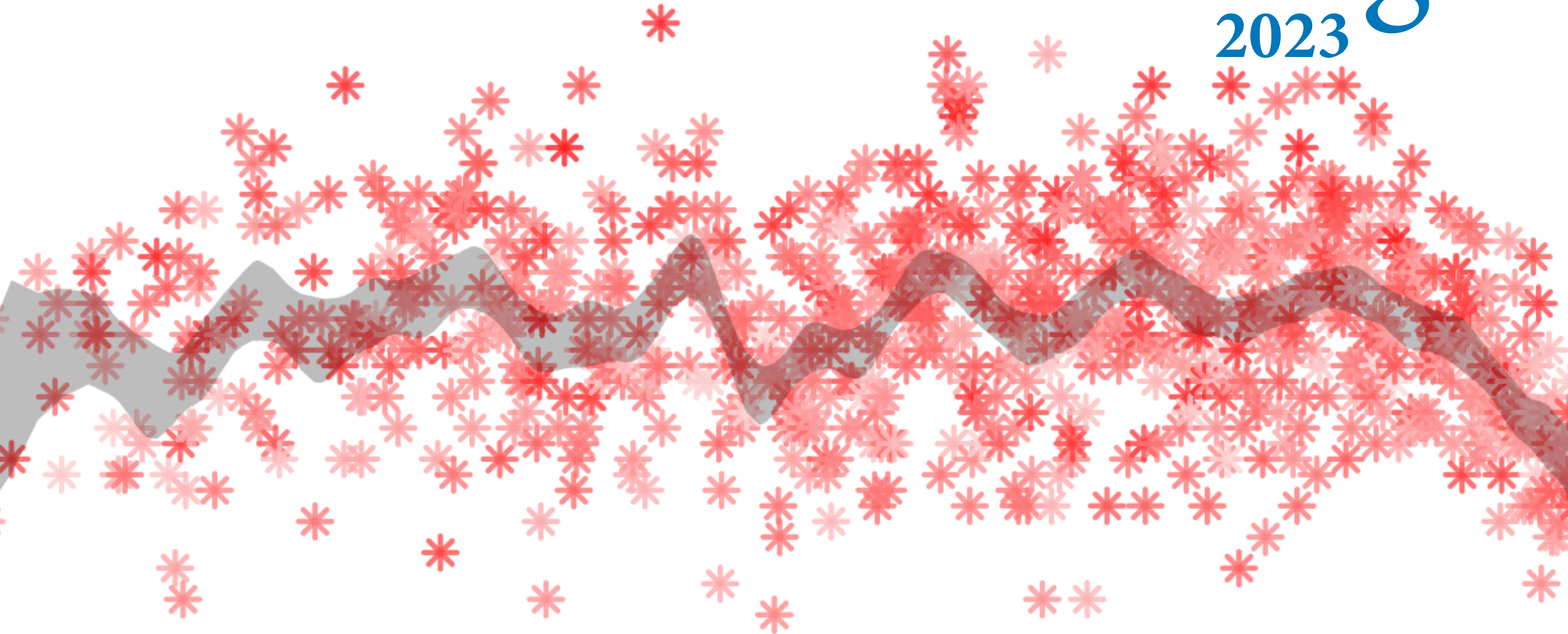


Statistical Rethinking

2023



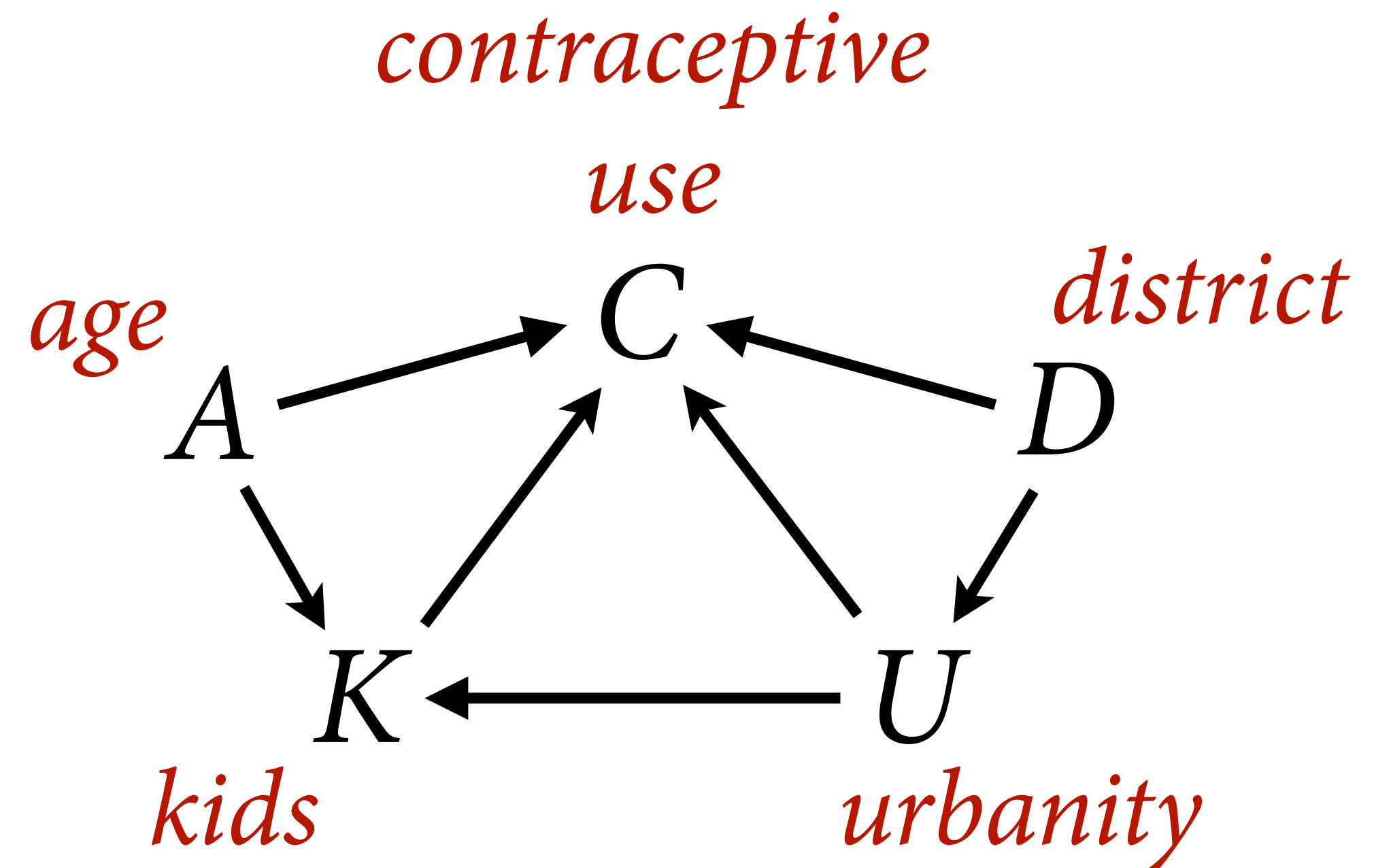
14. Correlated Features

Bangladesh 1989, Party Like It's

Estimand 1: C in each district

Estimand 2: Effect of U

Estimand 3: Effects of K and A

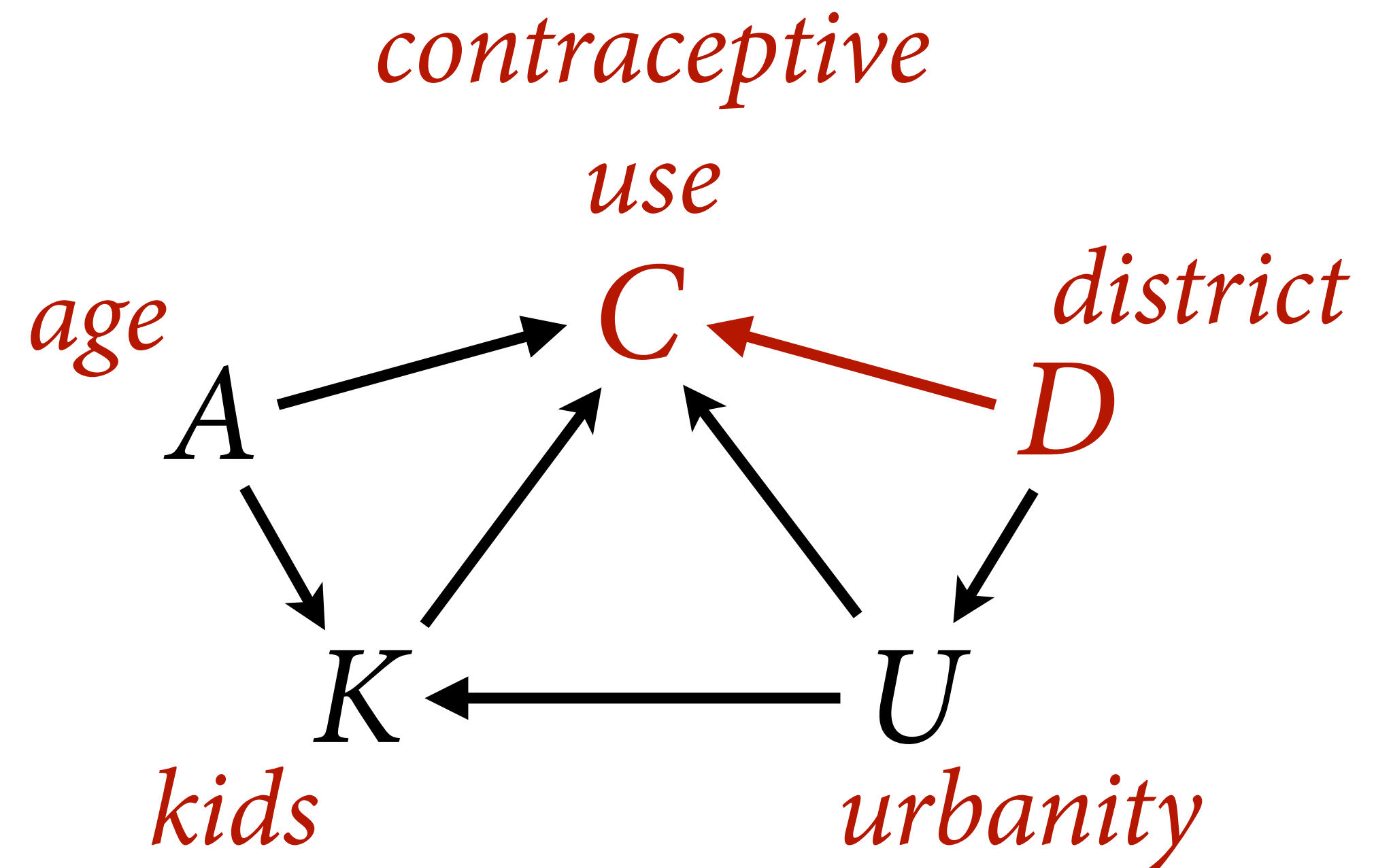


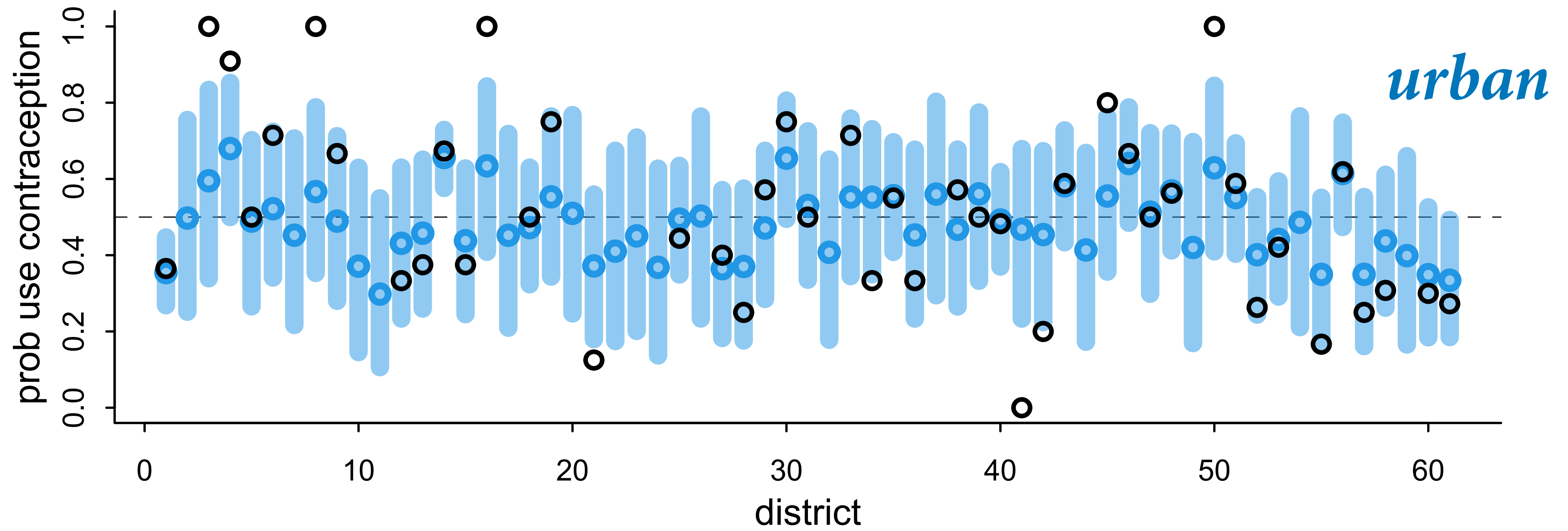
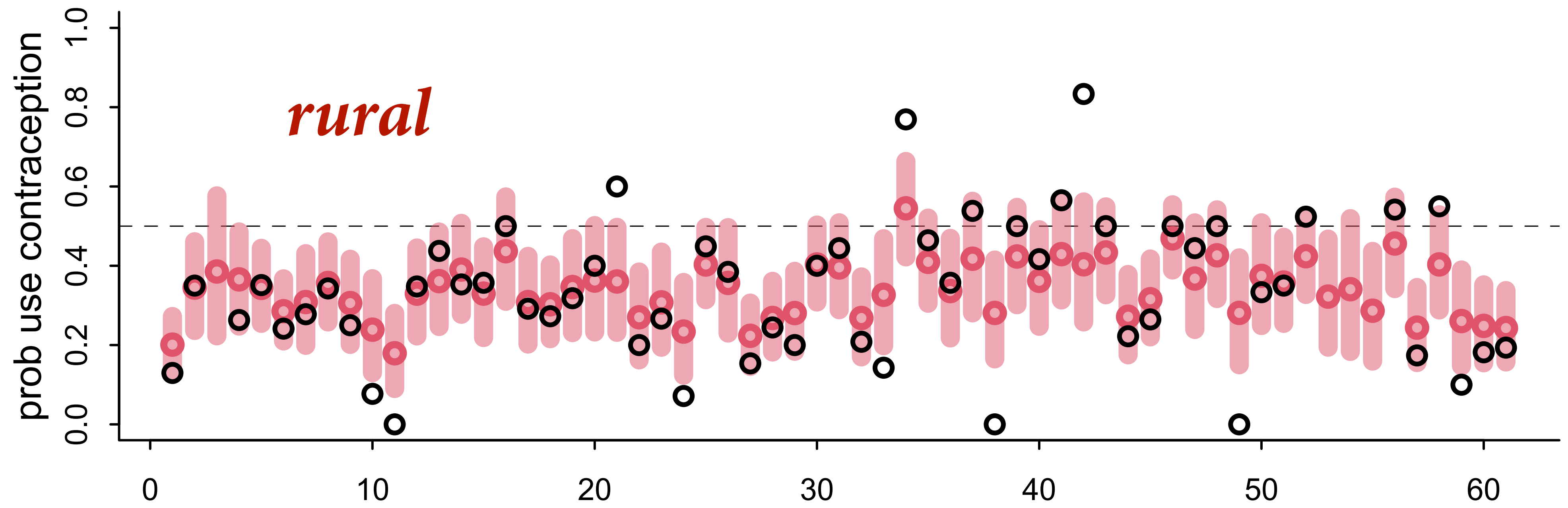
Bangladesh 1989, Party Like It's

Estimand 1: C in each district

Estimand 2: Effect of U

Estimand 3: Effects of K and A



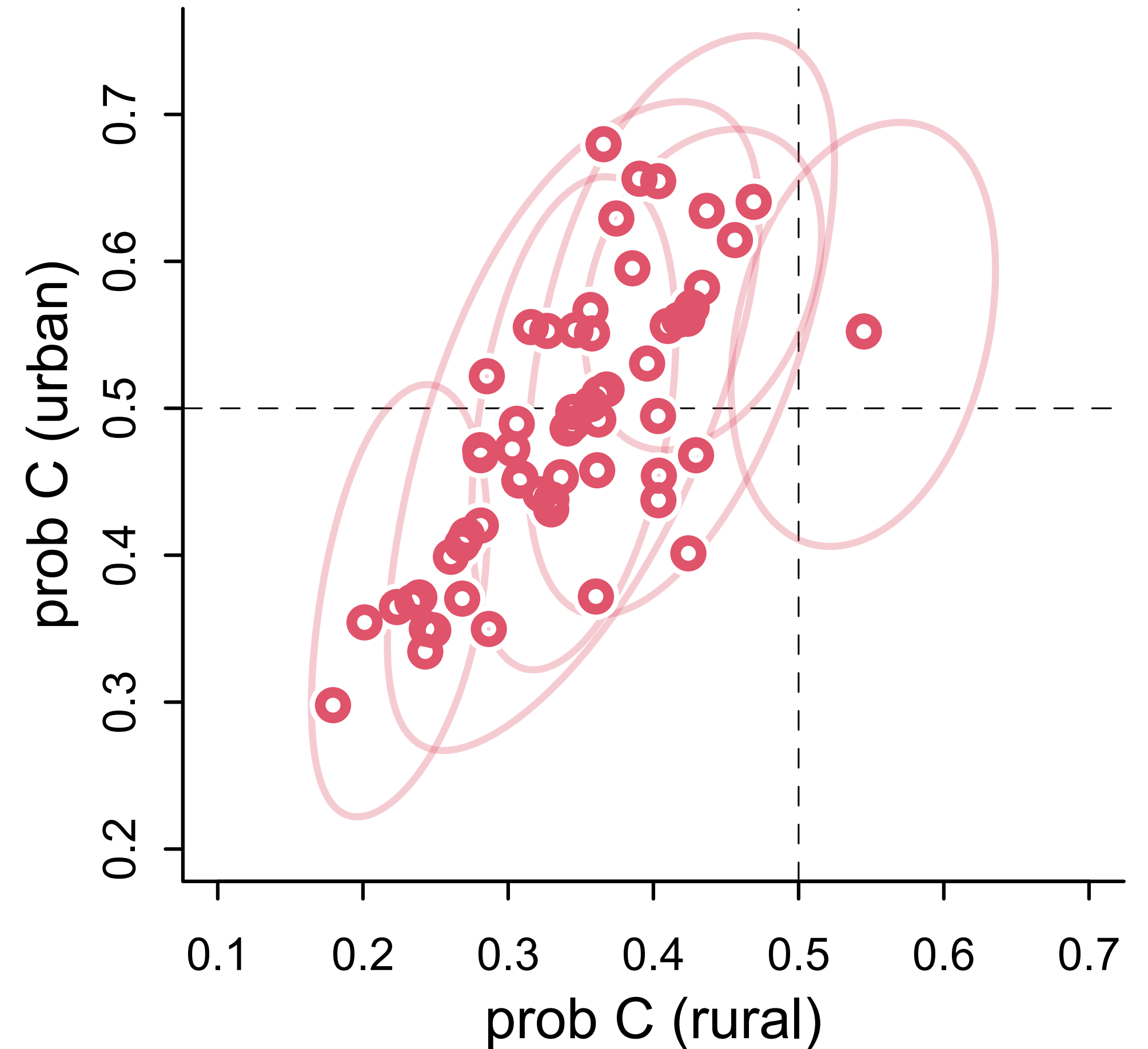


Multilevel Adventures

Clusters: Kinds of groups in the data (districts)

Features: Aspects of the model (parameters) that vary by cluster (rural, urban)

There is useful information to transfer across features



Adding Correlated Features

One prior distribution for each cluster

One feature: One-dimensional distribution

$$\alpha_j \sim \text{Normal}(\bar{\alpha}, \sigma)$$

Two features: Two-D distribution

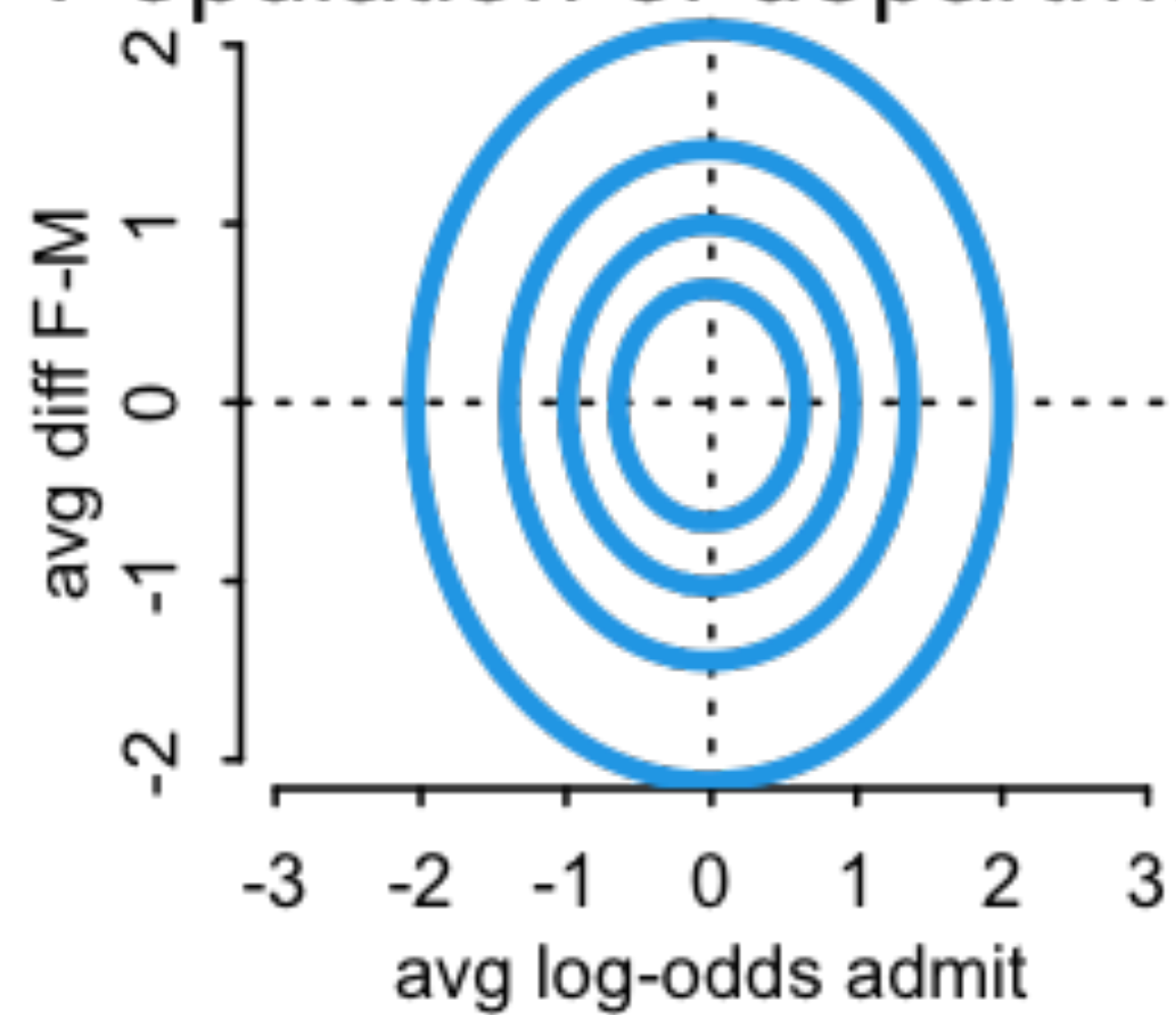
$$[\alpha_j, \beta_j] \sim \text{MVNormal}([\bar{\alpha}, \bar{\beta}], \Sigma)$$

N features: N -dimensional distribution

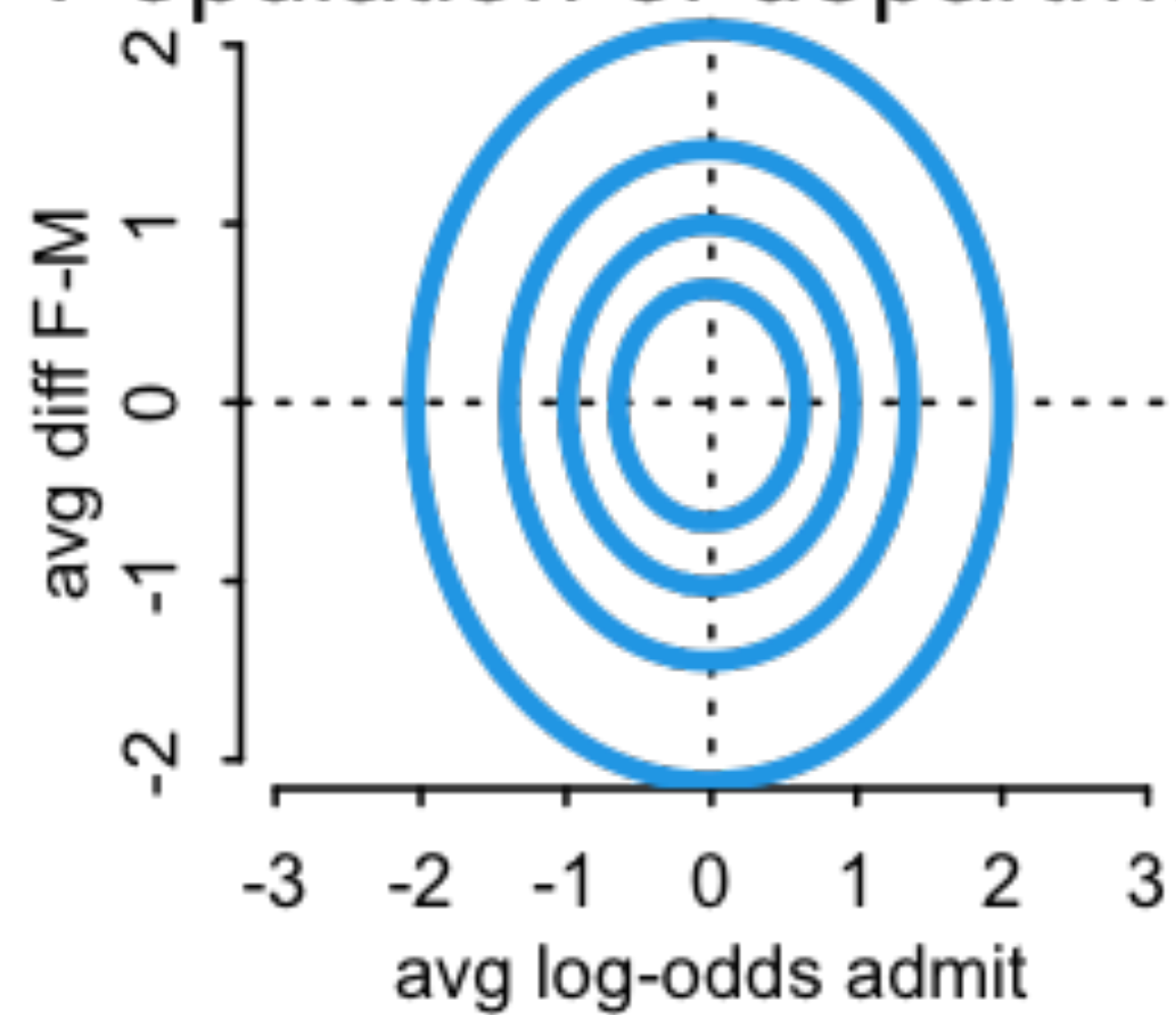
$$\alpha_{j,1..N} \sim \text{MVNormal}(A, \Sigma)$$

Hard part: Learning associations

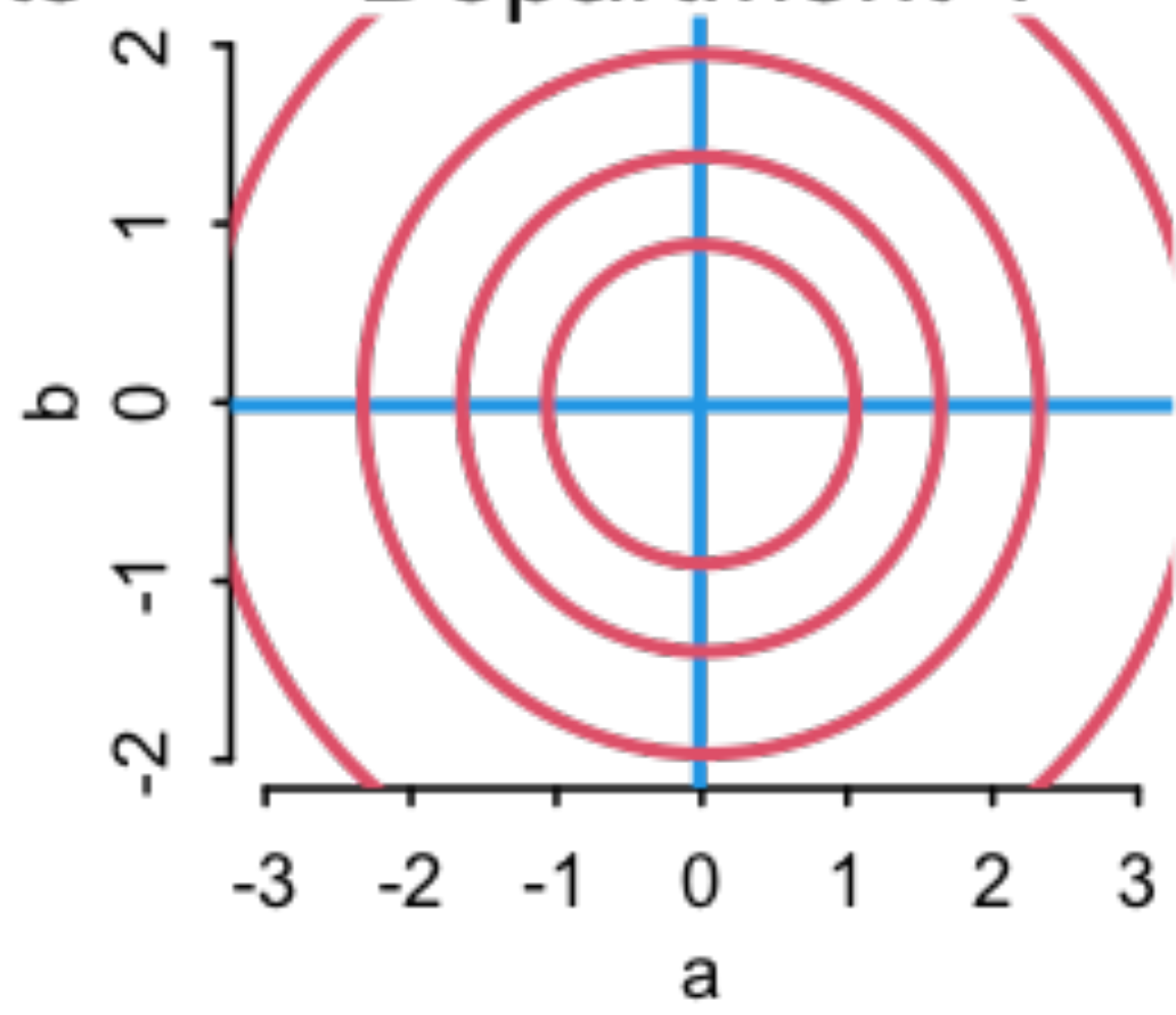
Population of departments



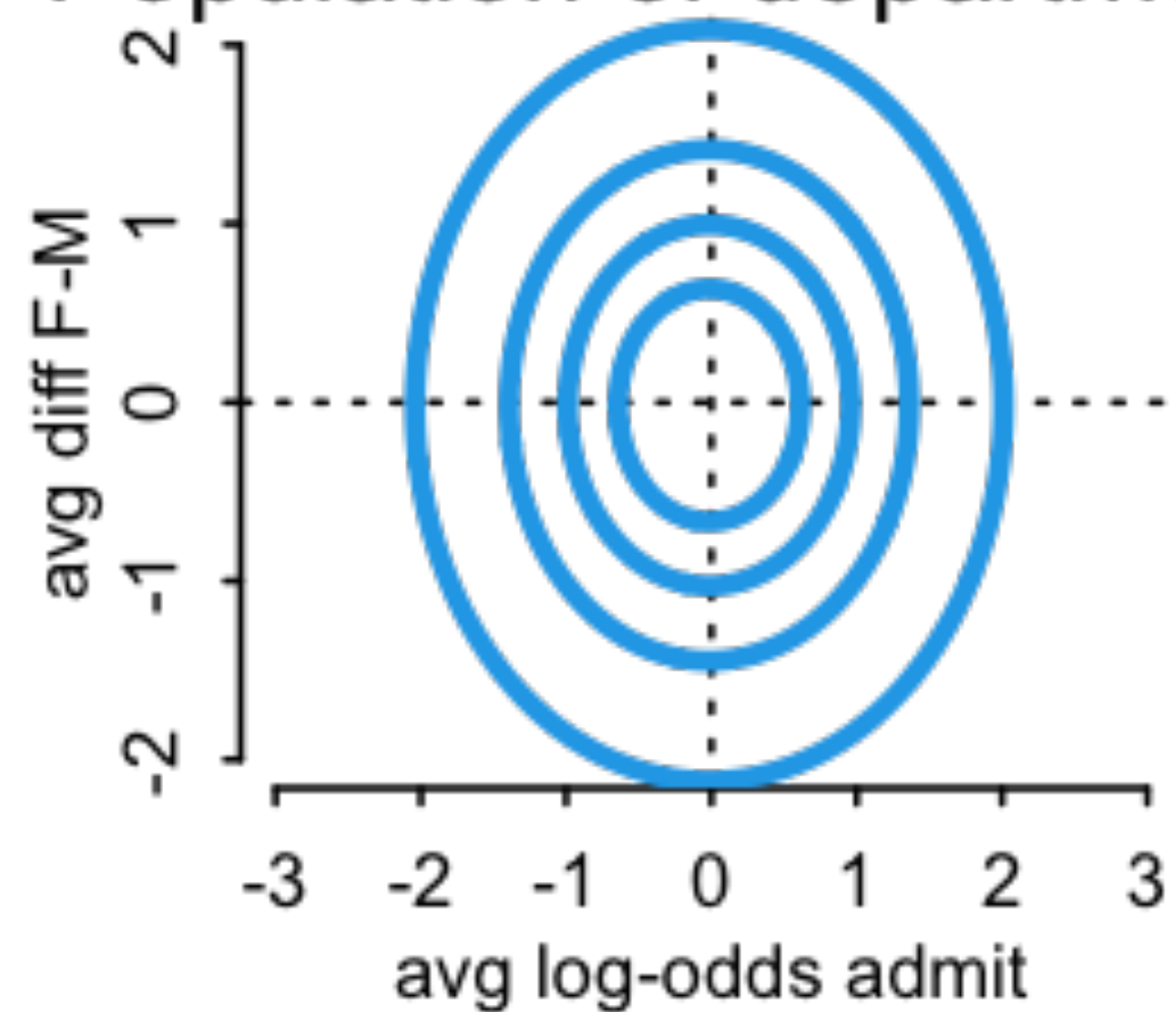
Population of departments



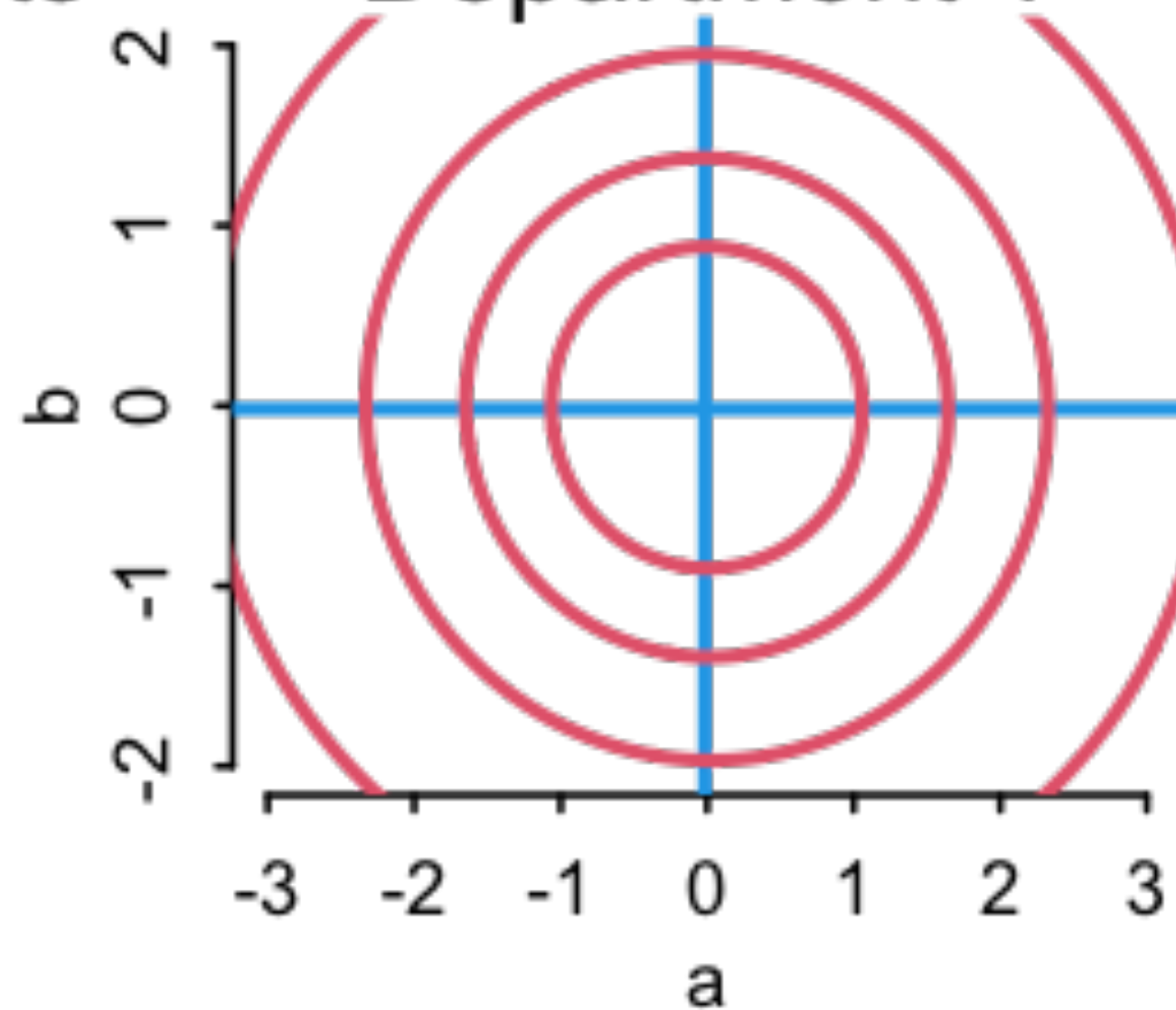
Department 1



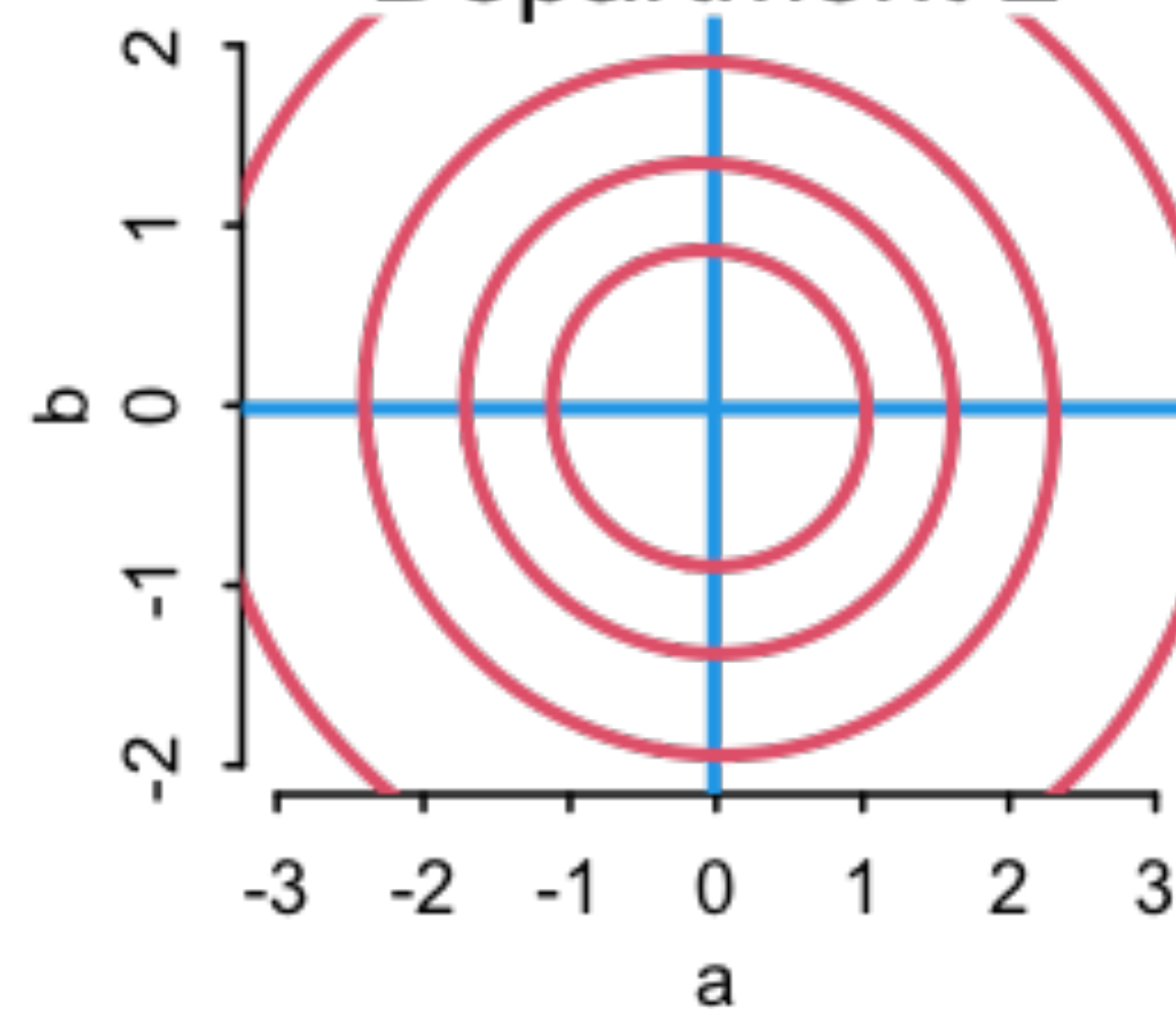
Population of departments



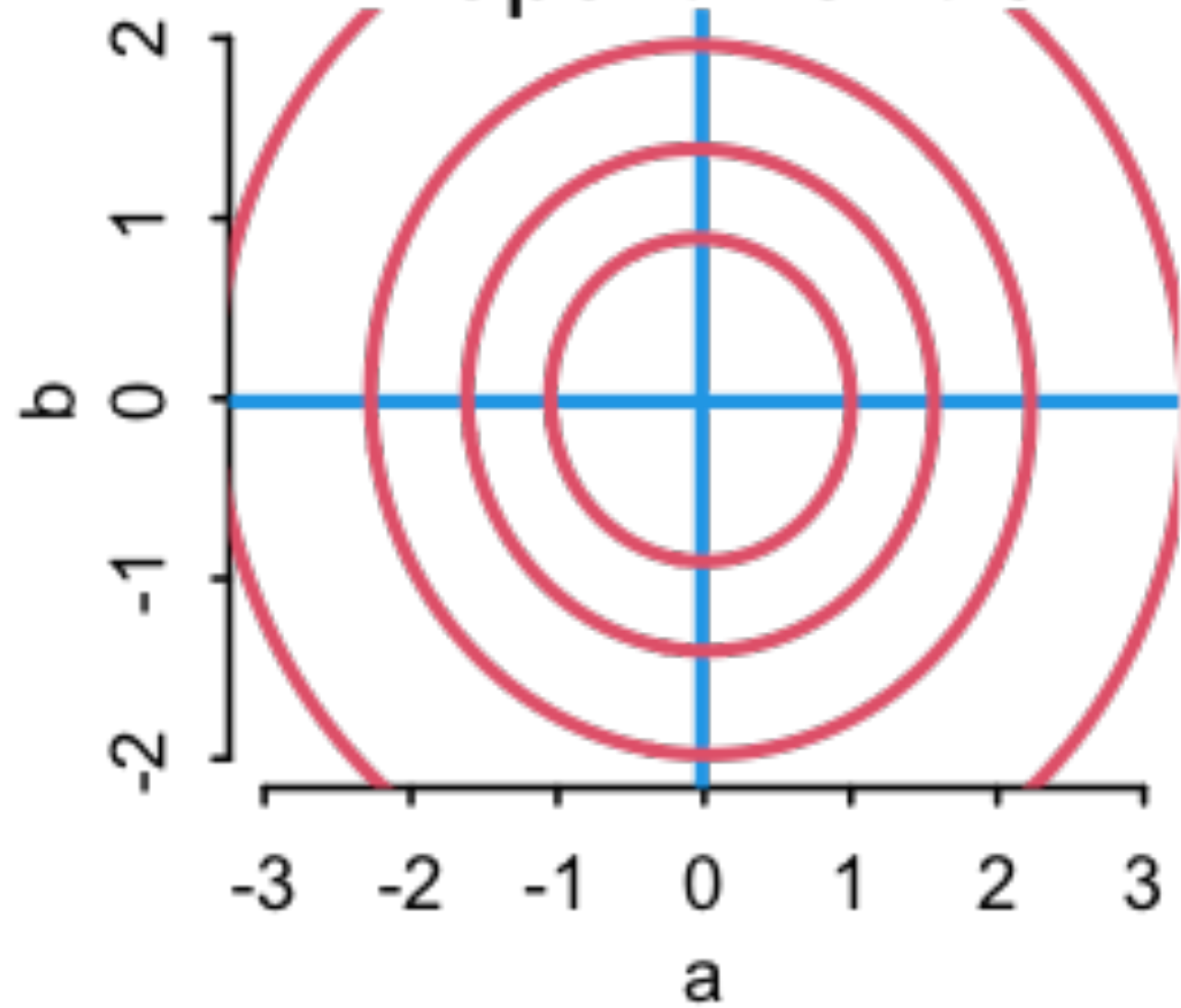
Department 1



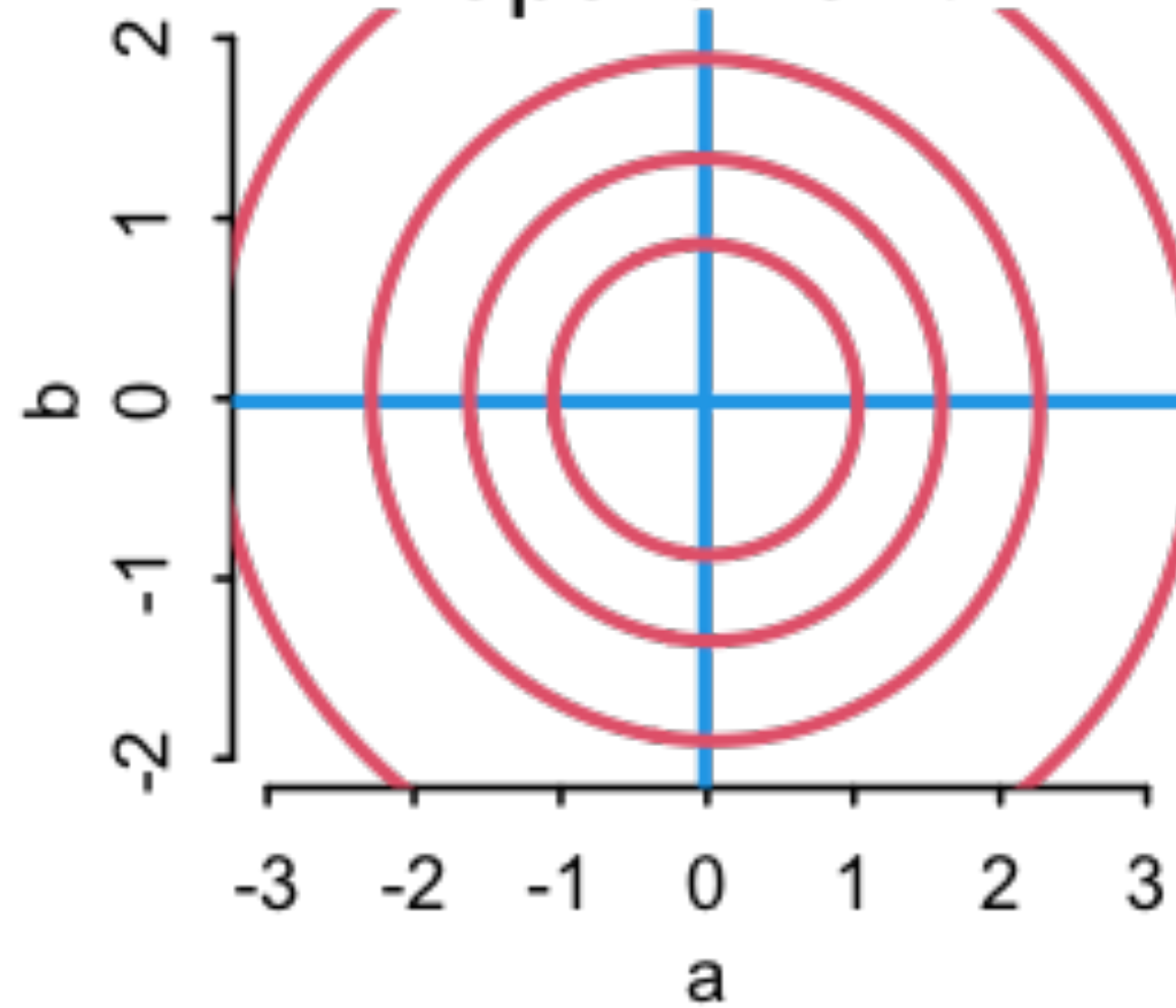
Department 2



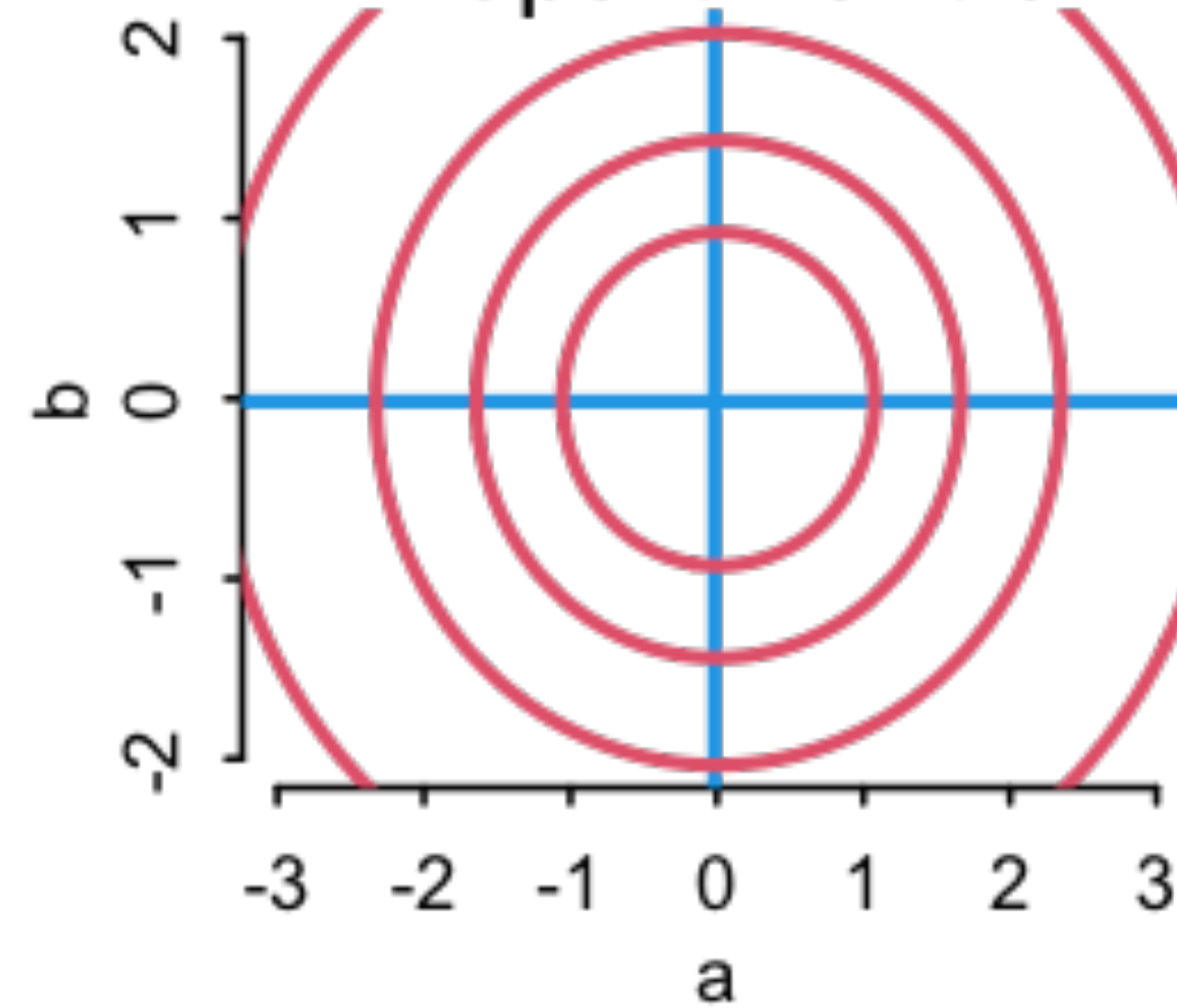
Department 3



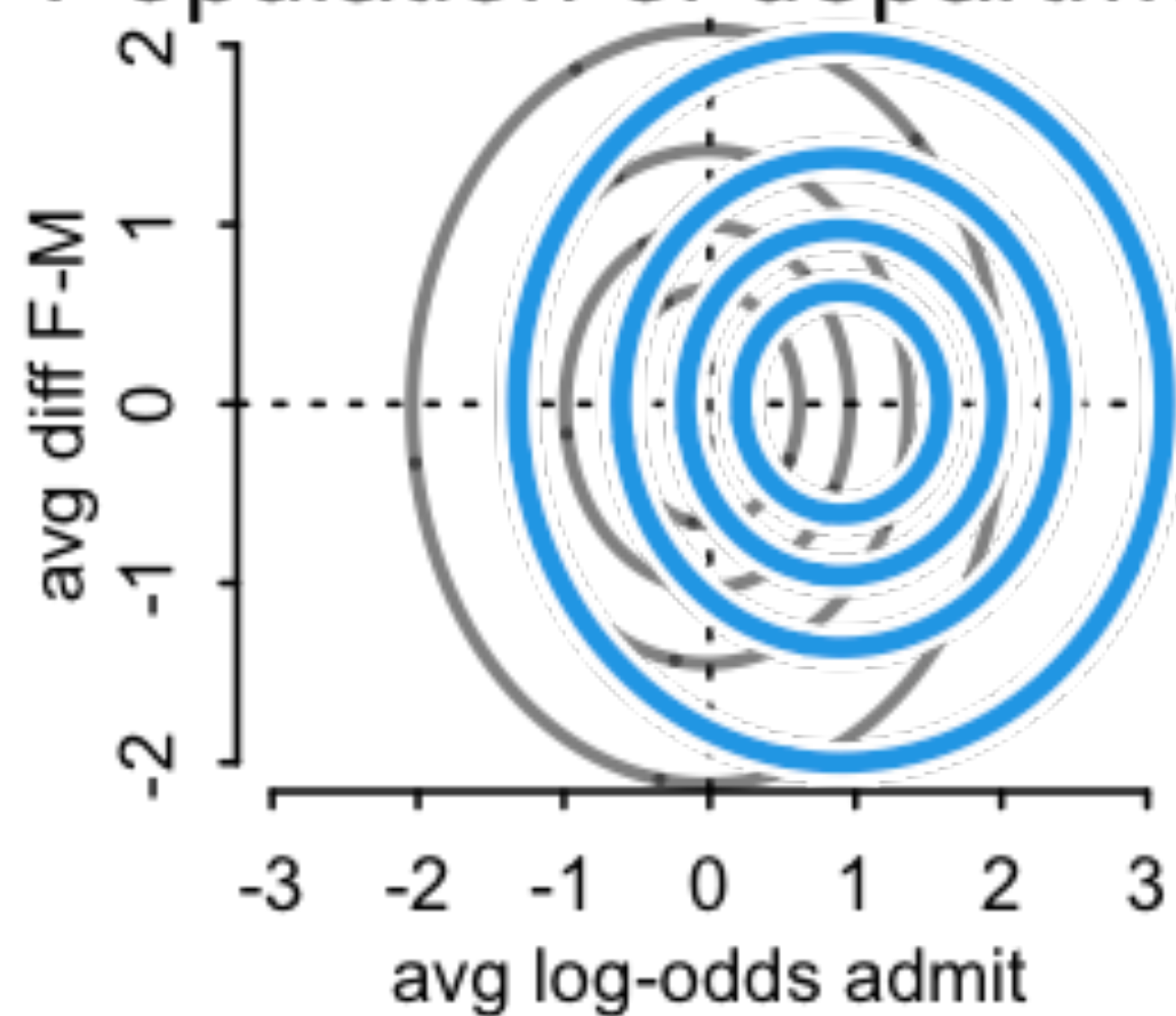
Department 4



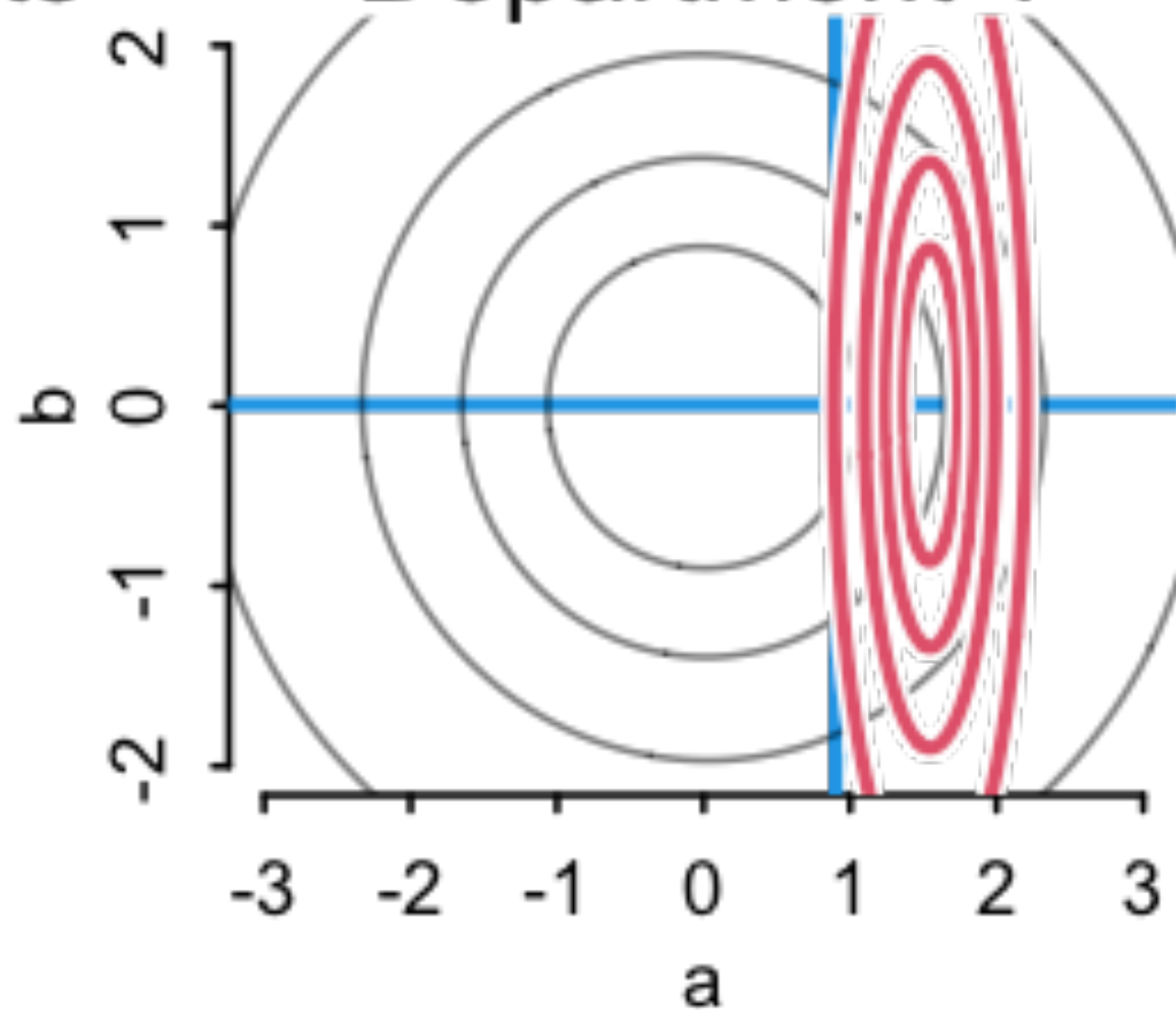
Department 5



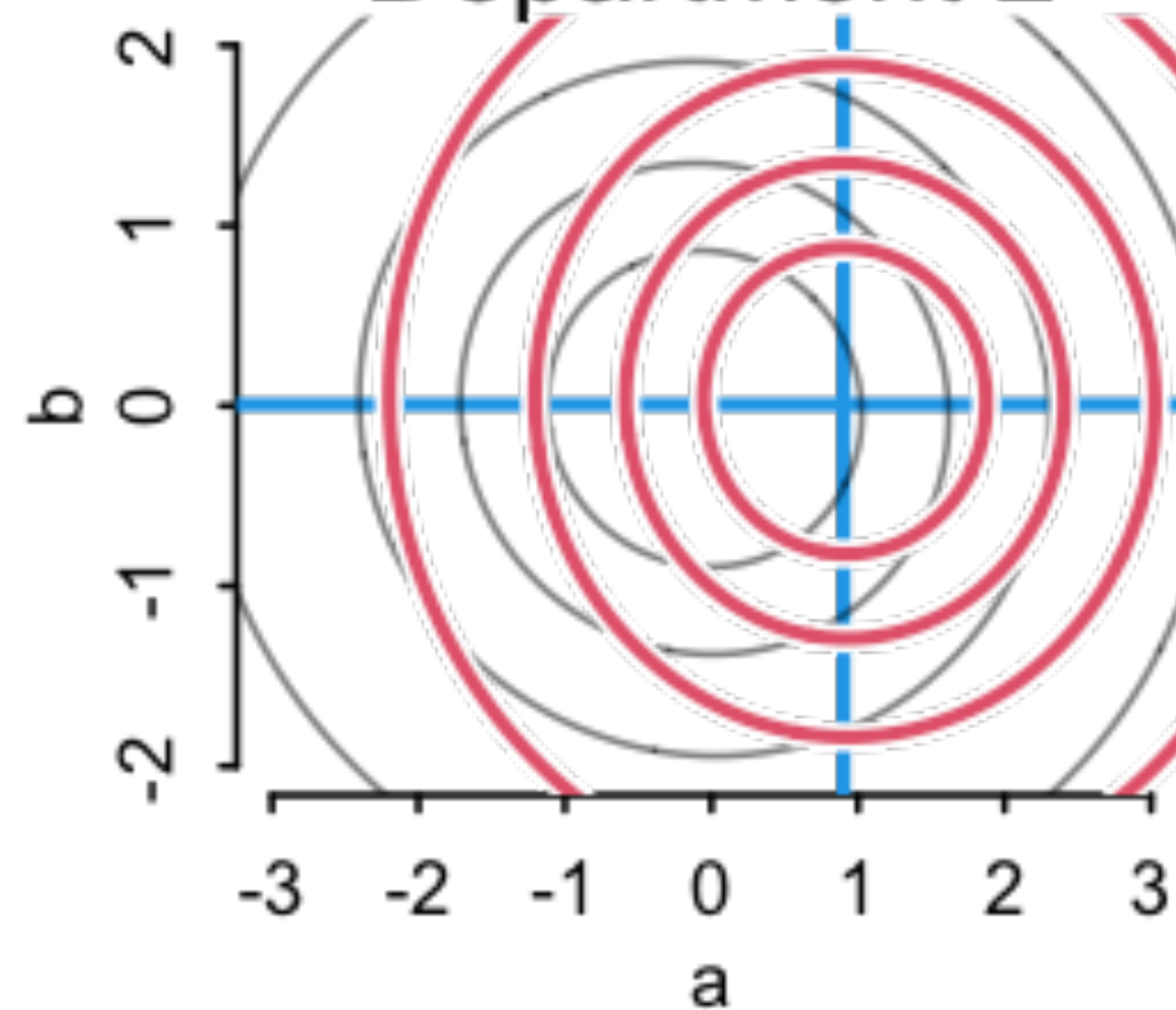
Population of departments



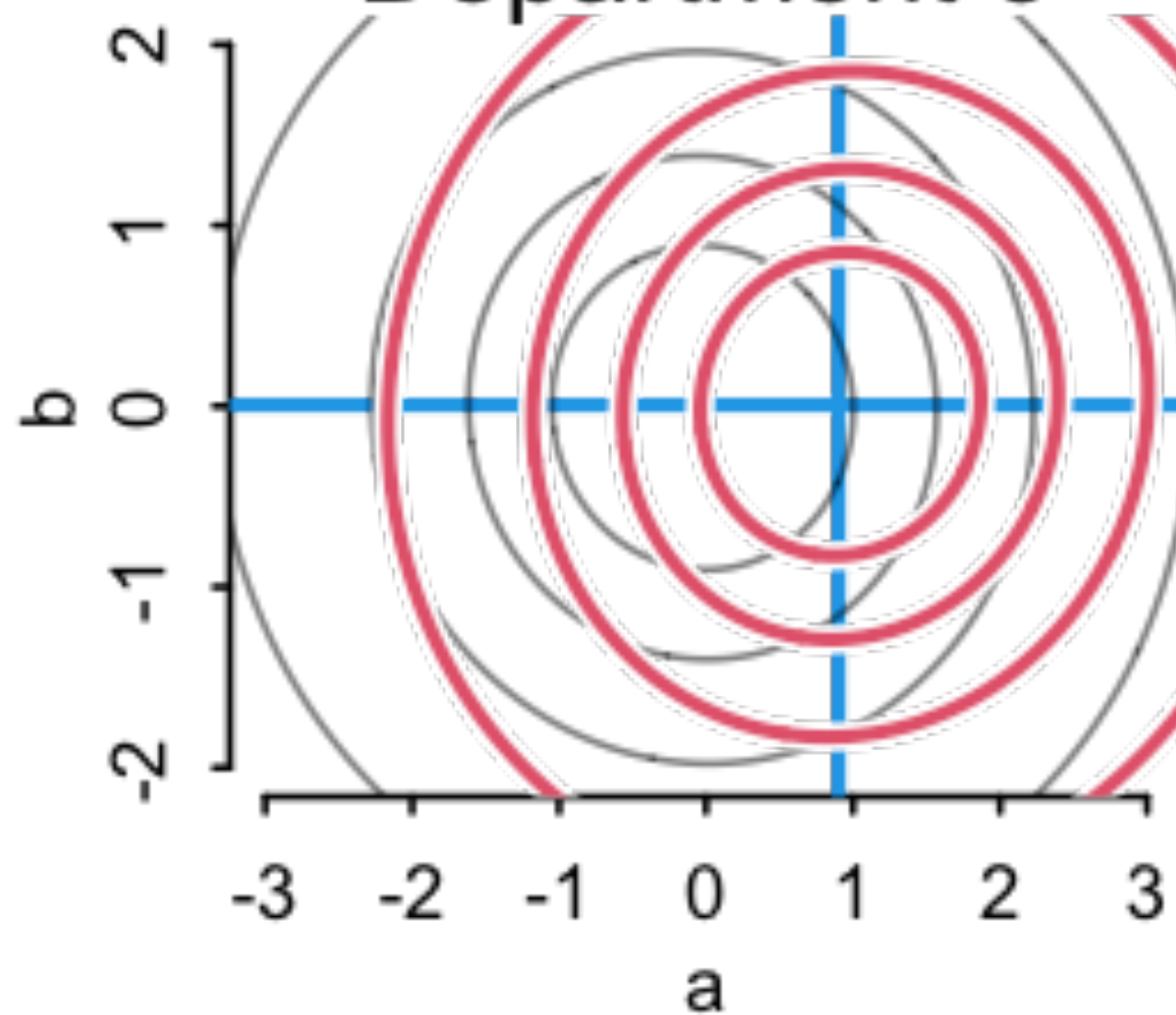
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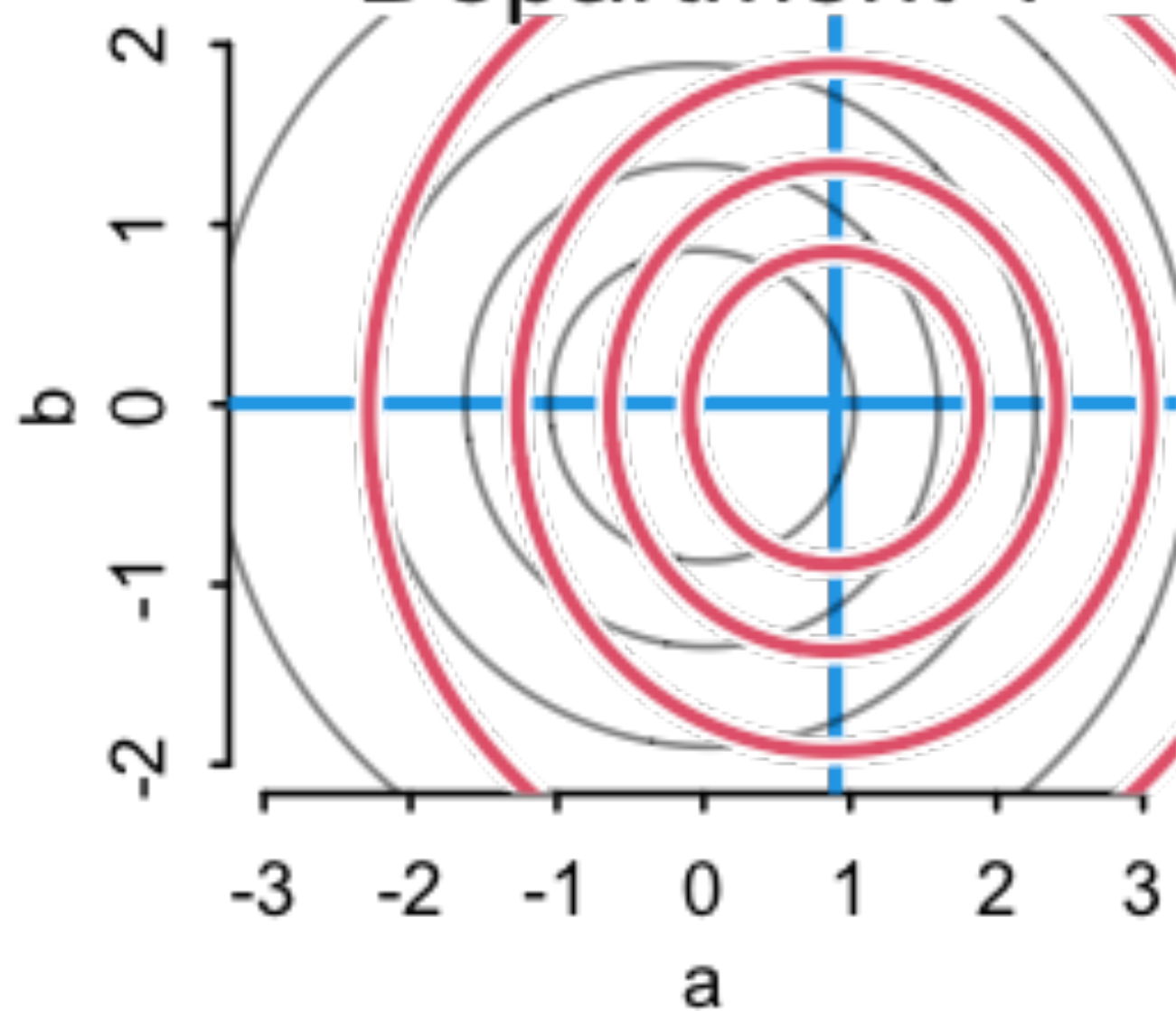
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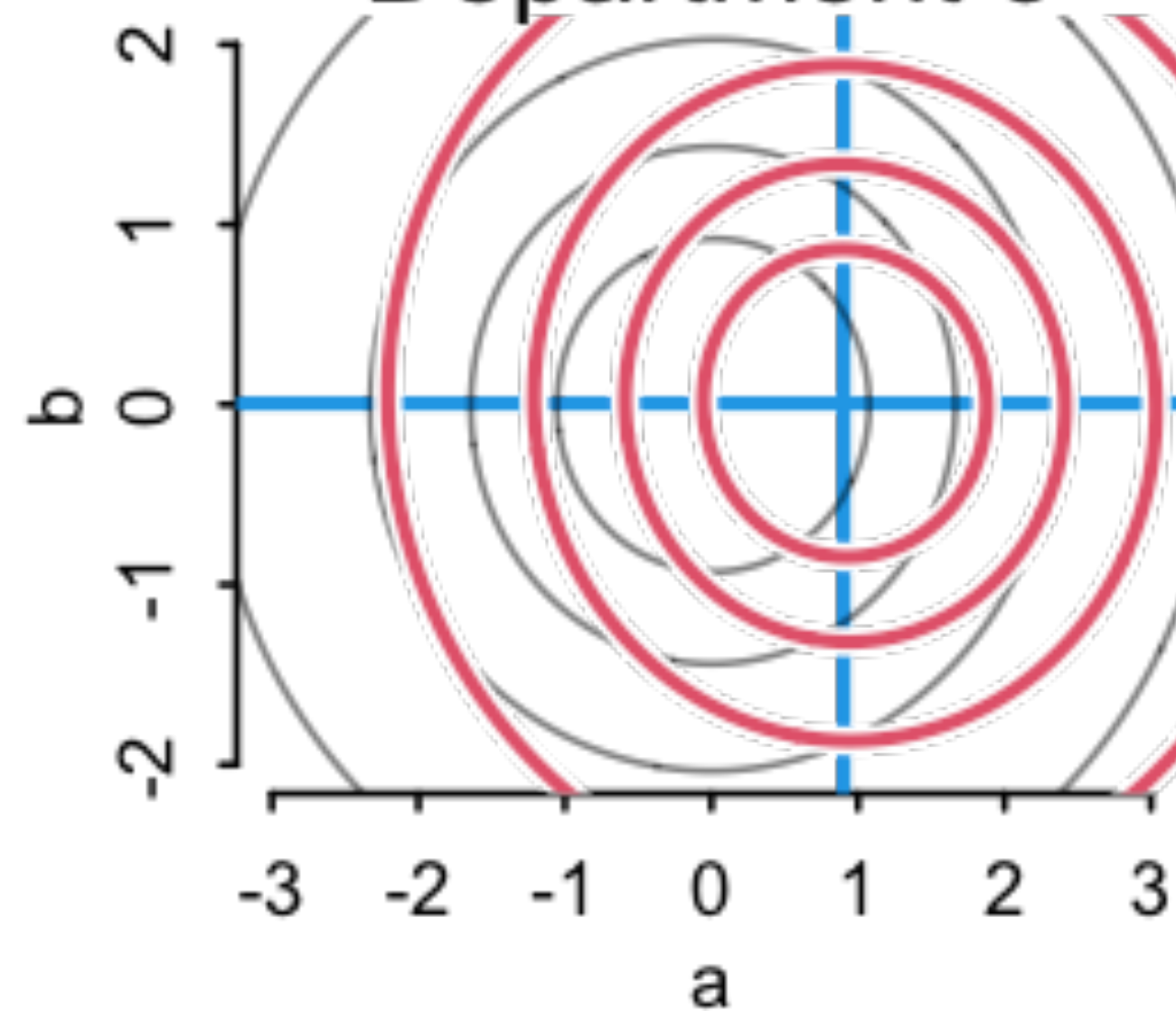
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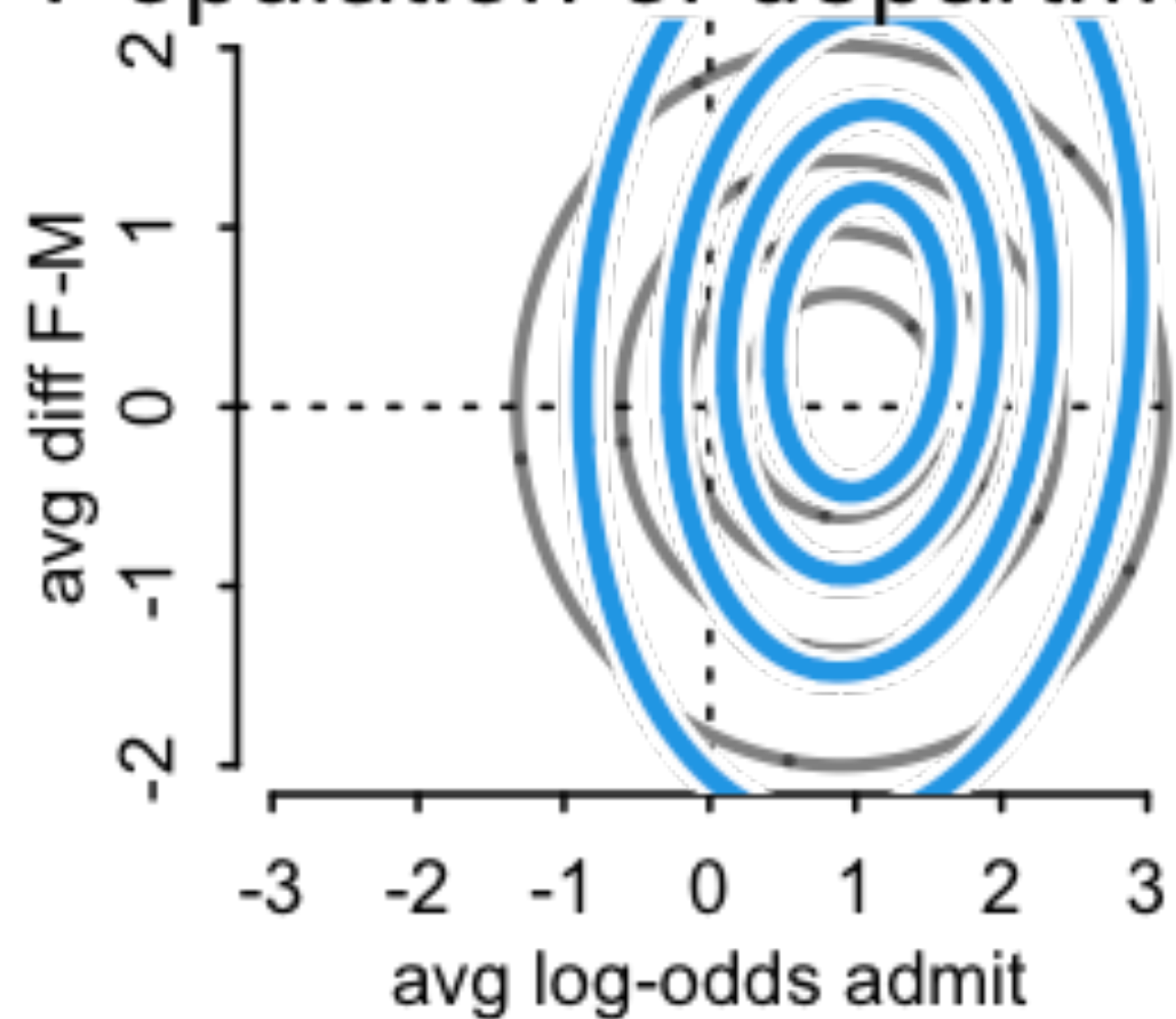
Department 4



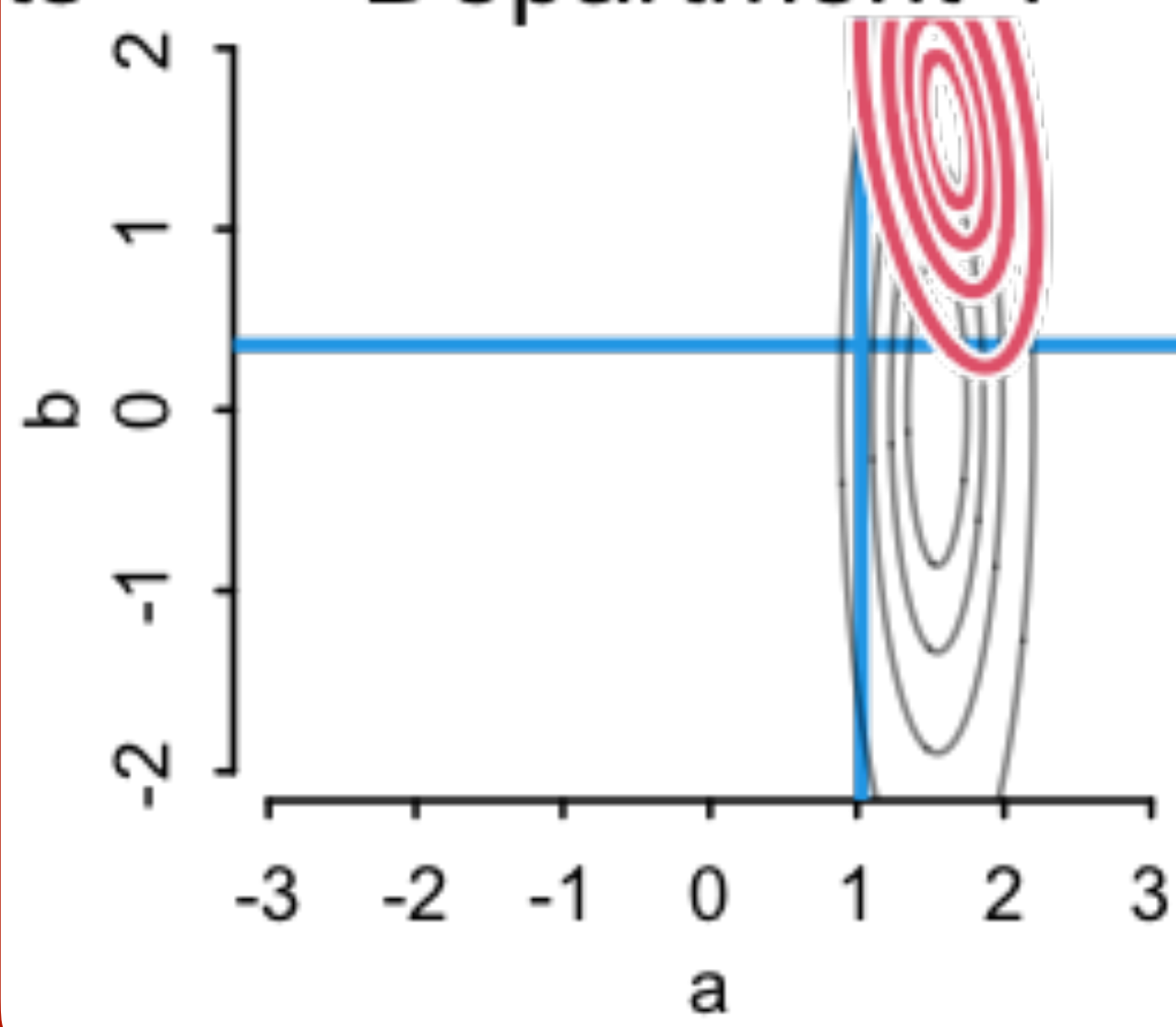
Department 5



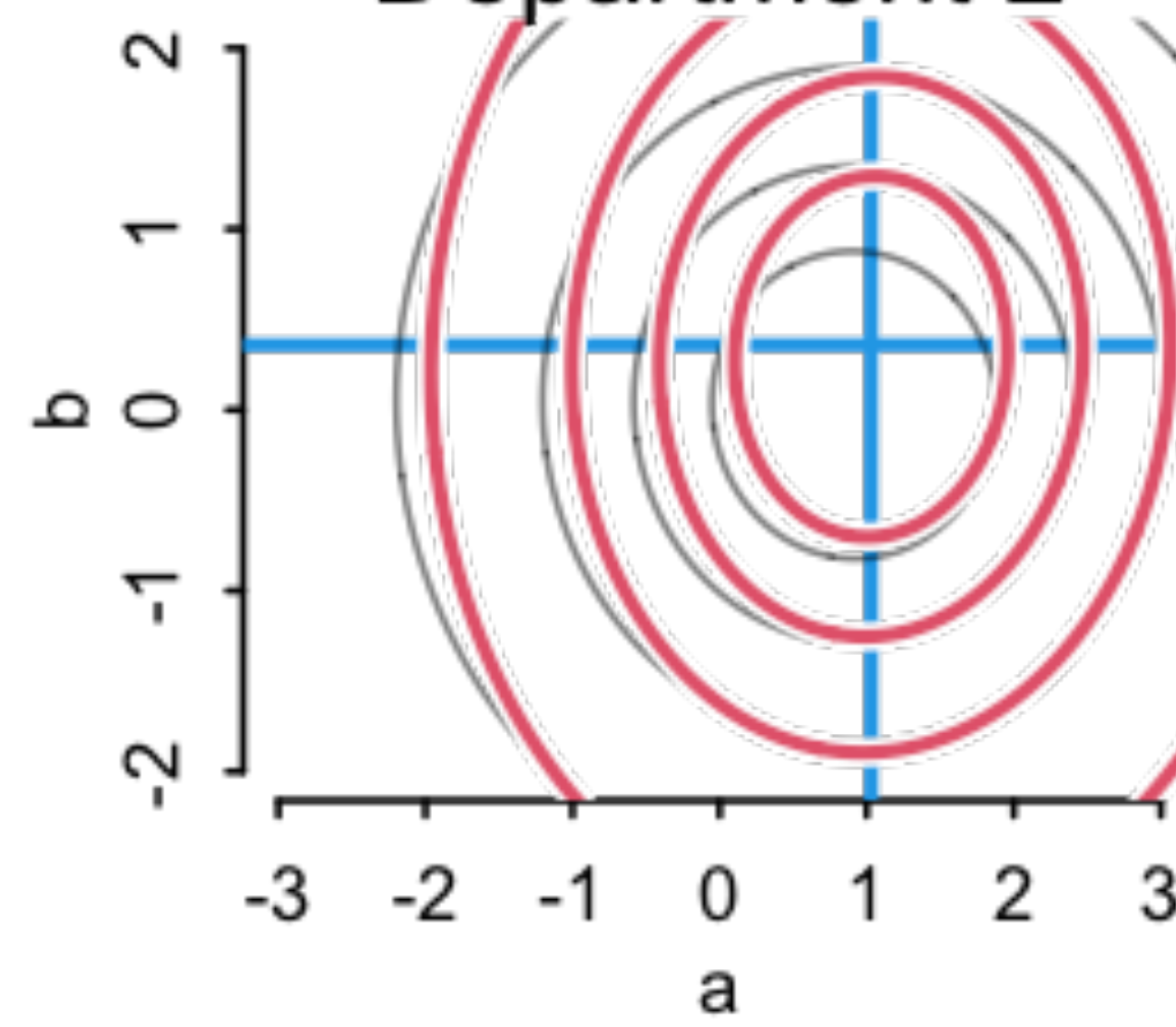
Population of departments



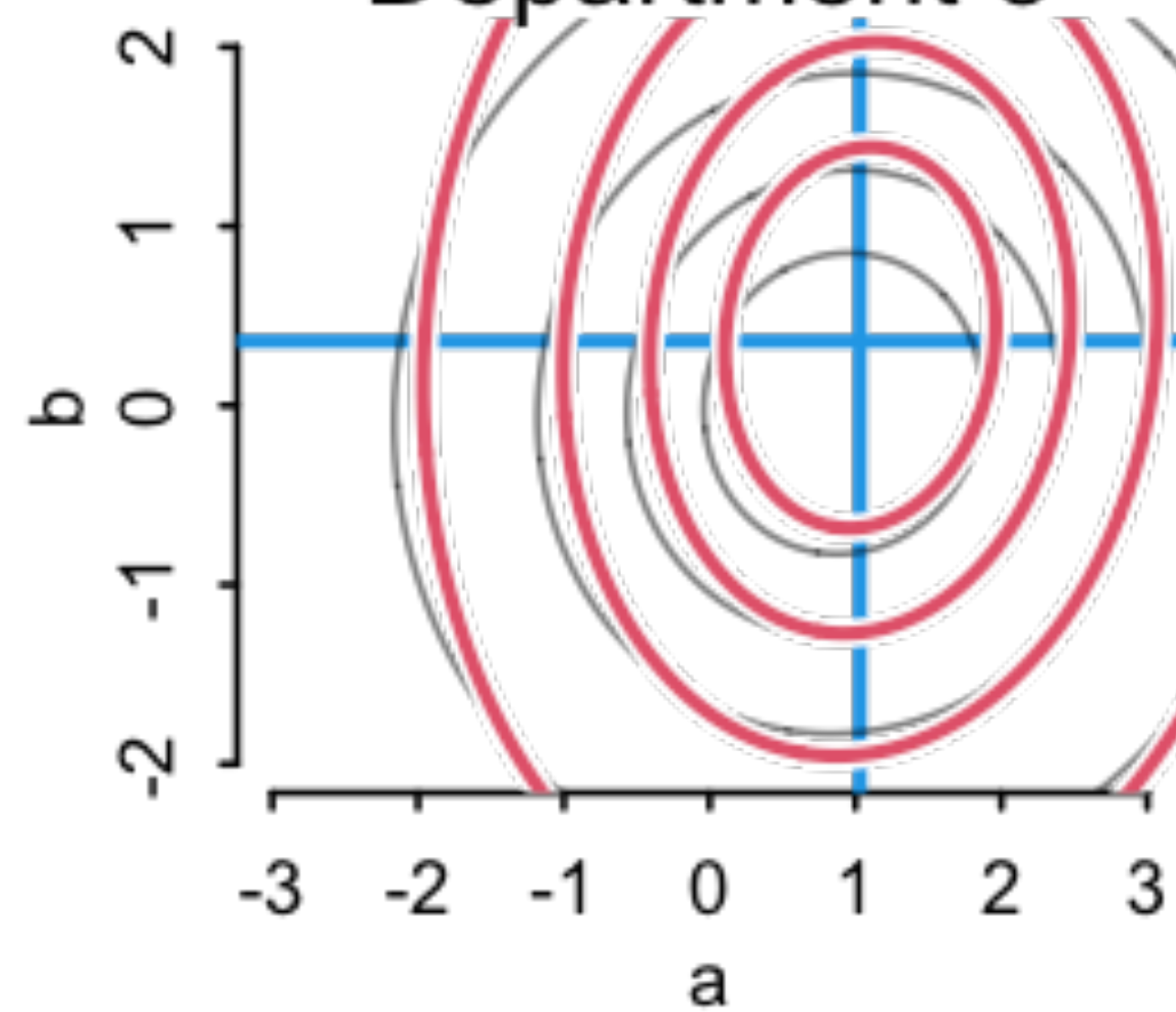
Department 1



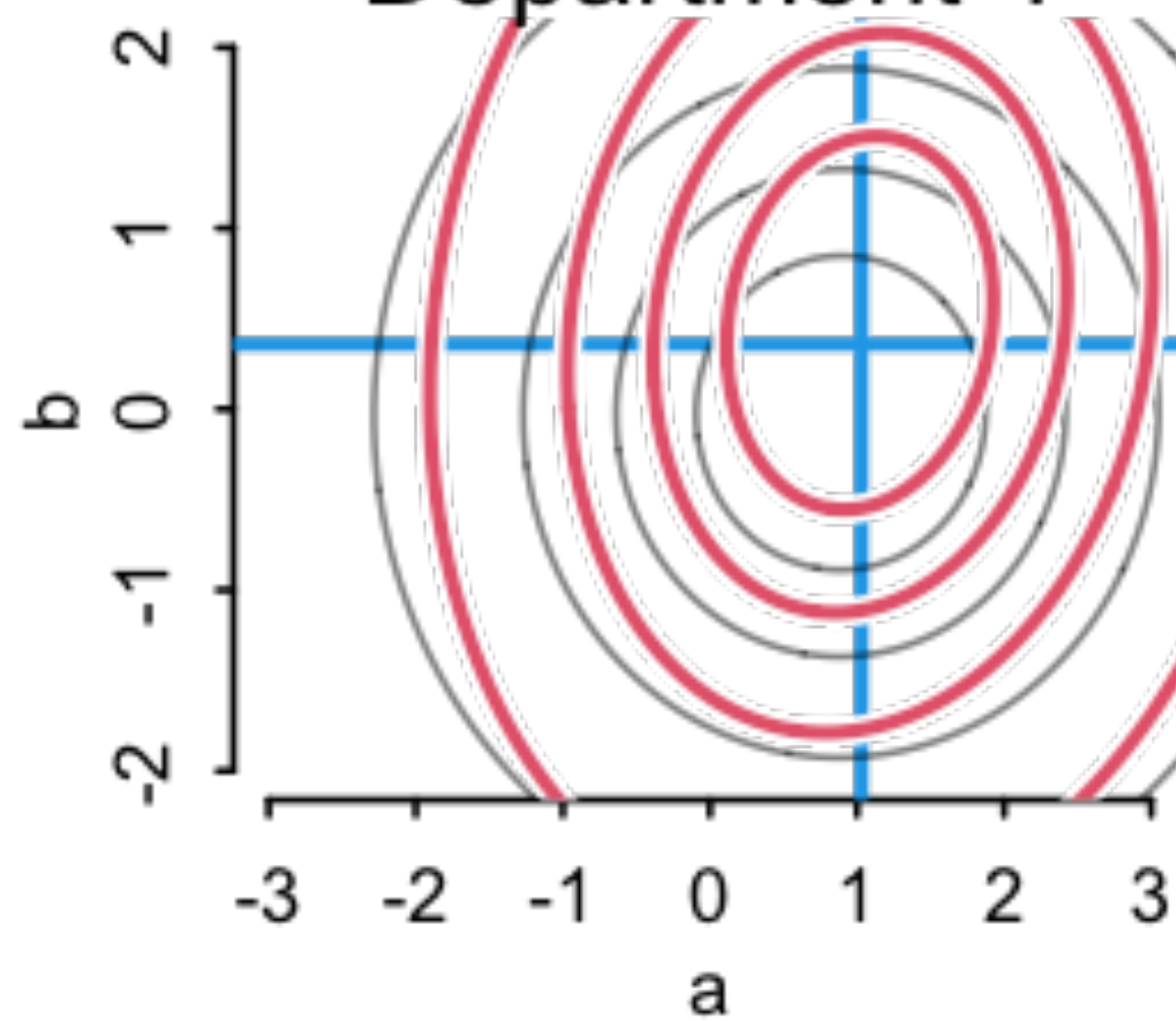
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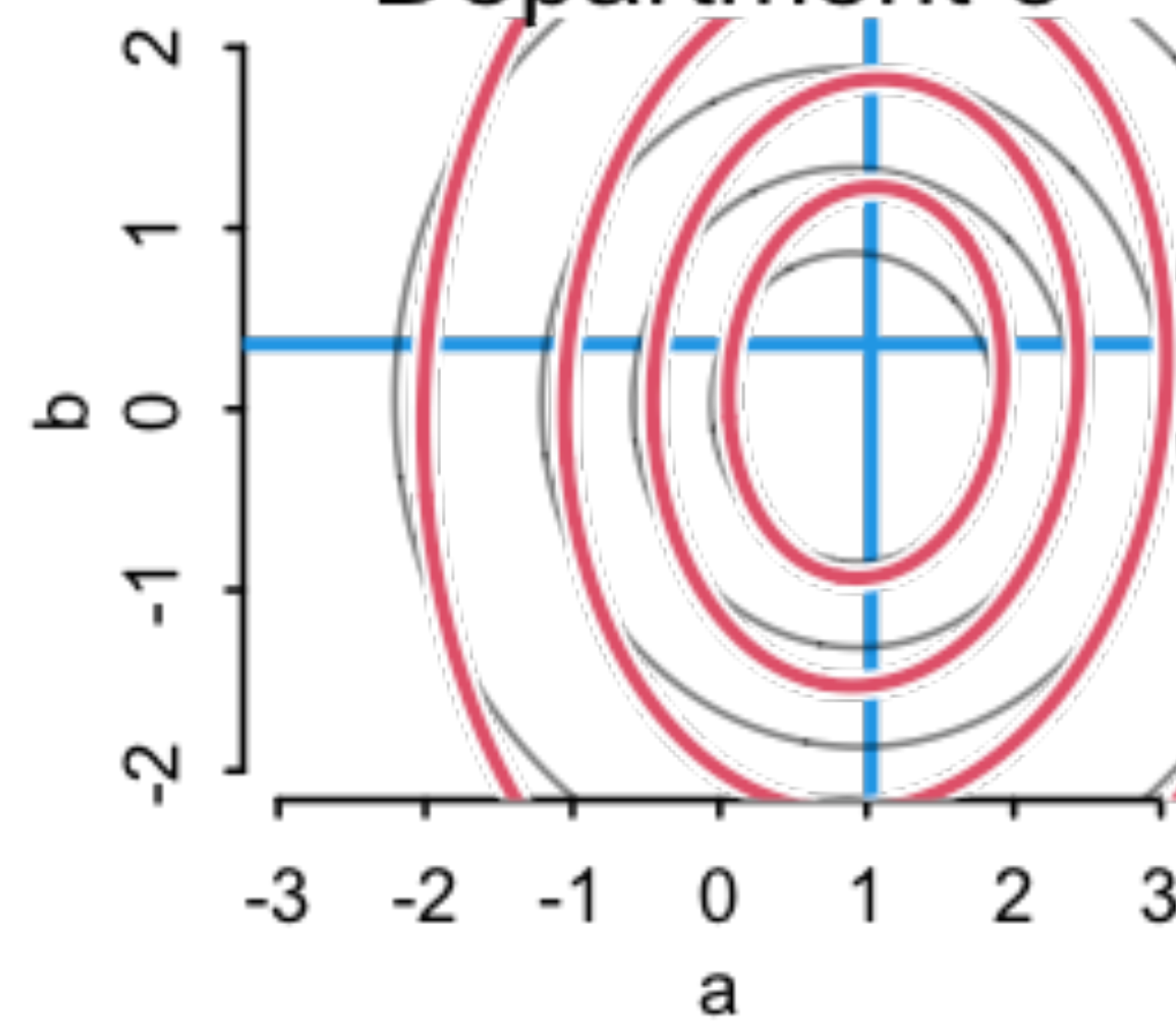
Department 3



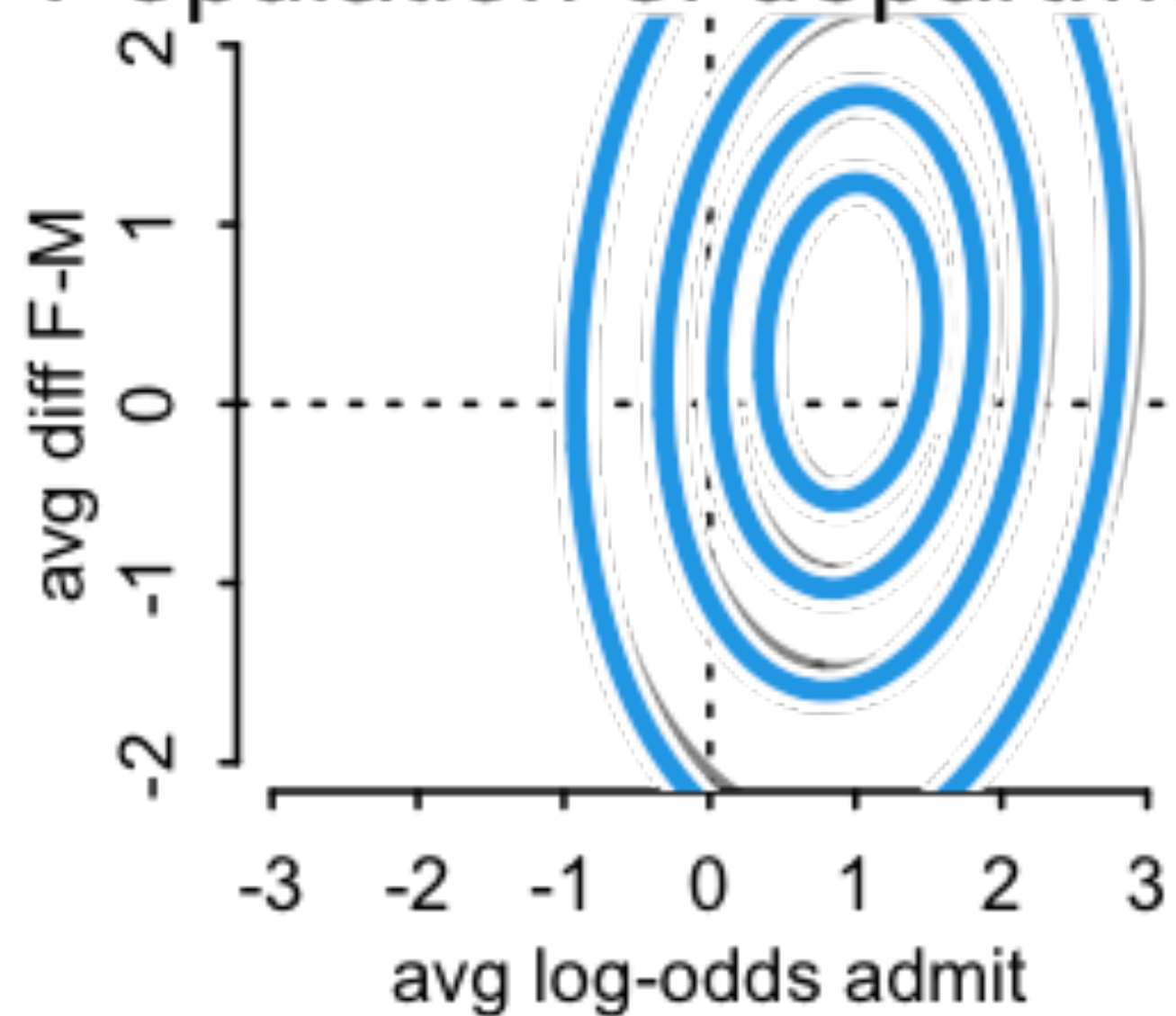
Department 4



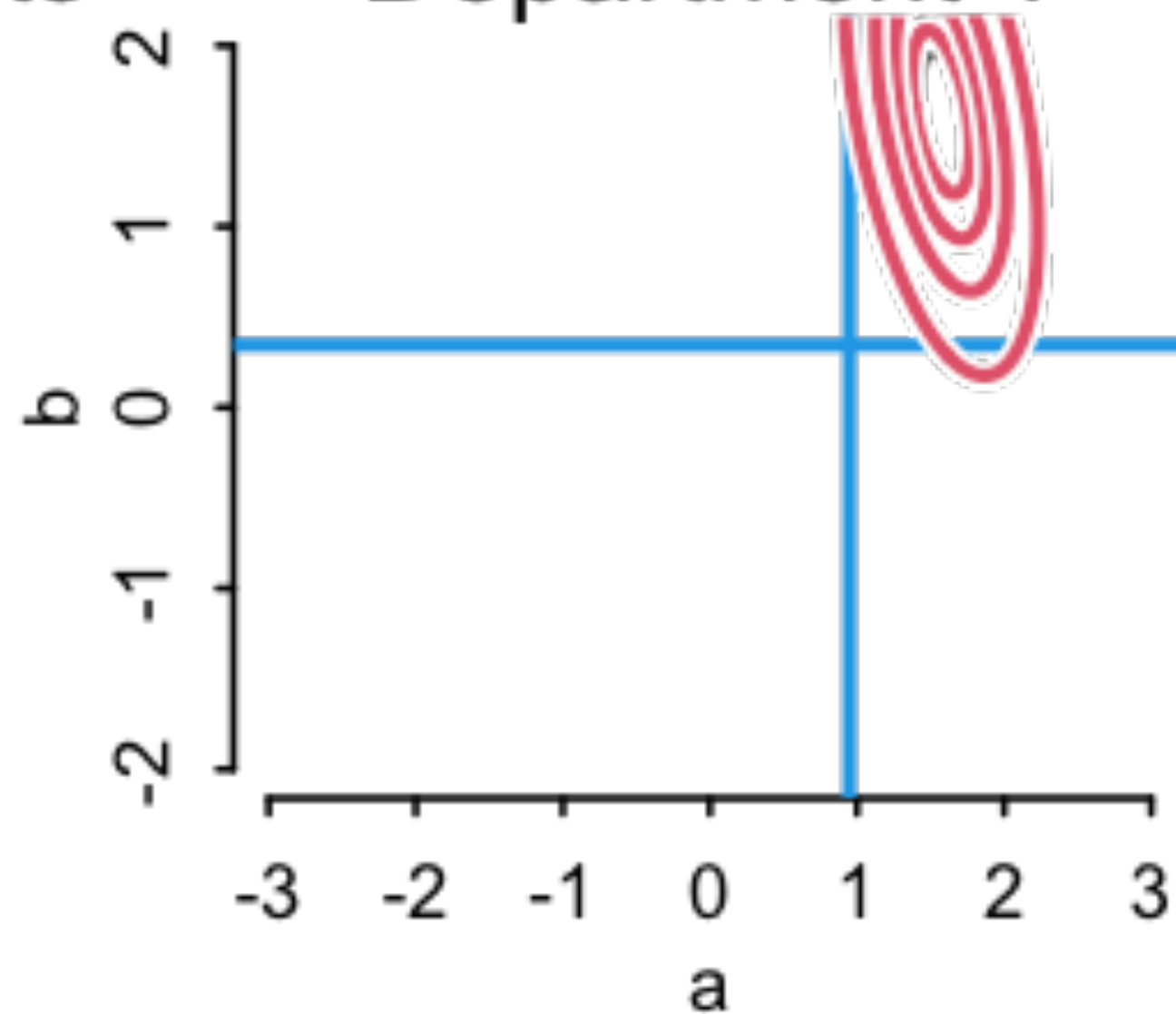
Department 5



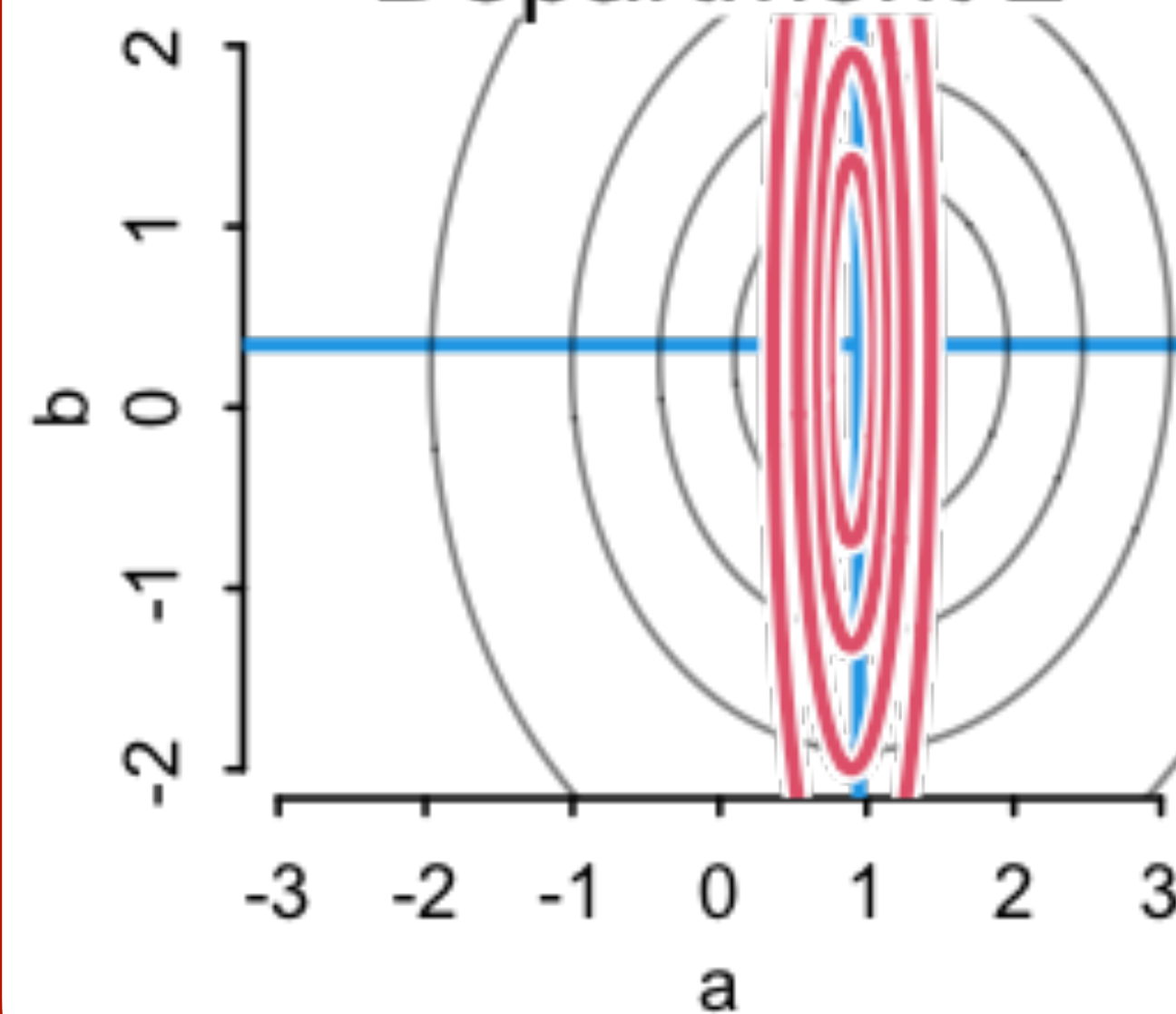
Population of departments



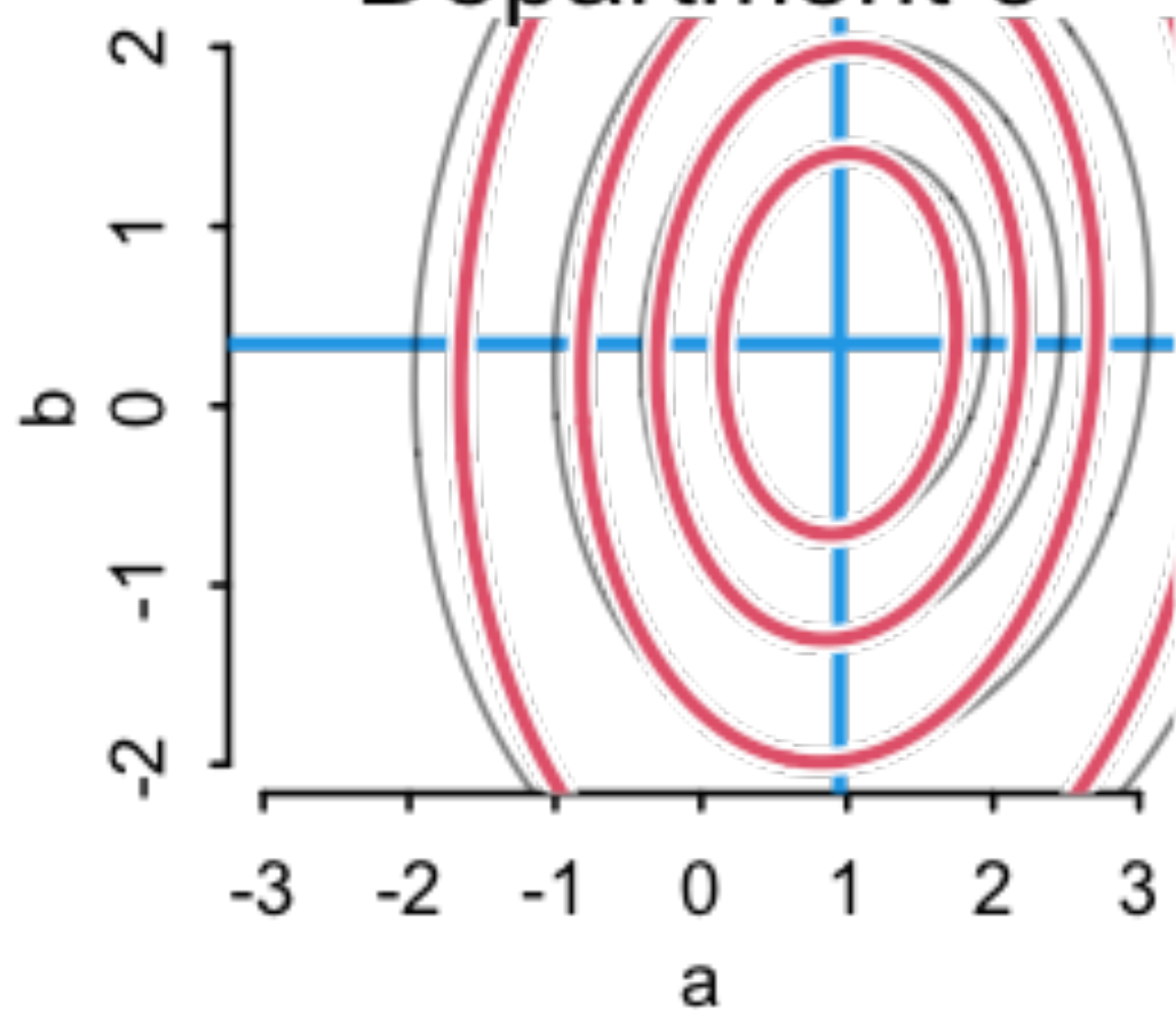
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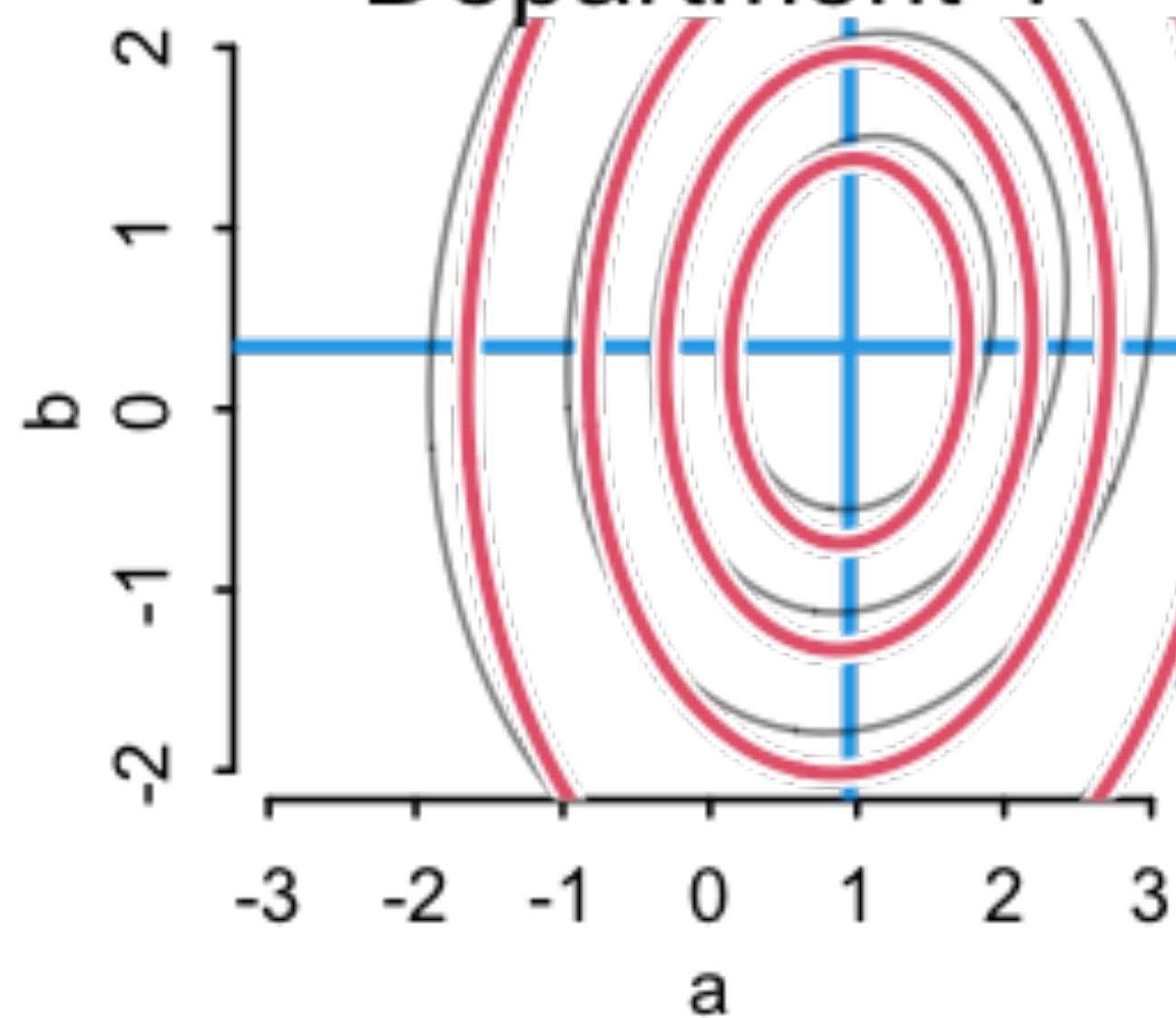
Department 2



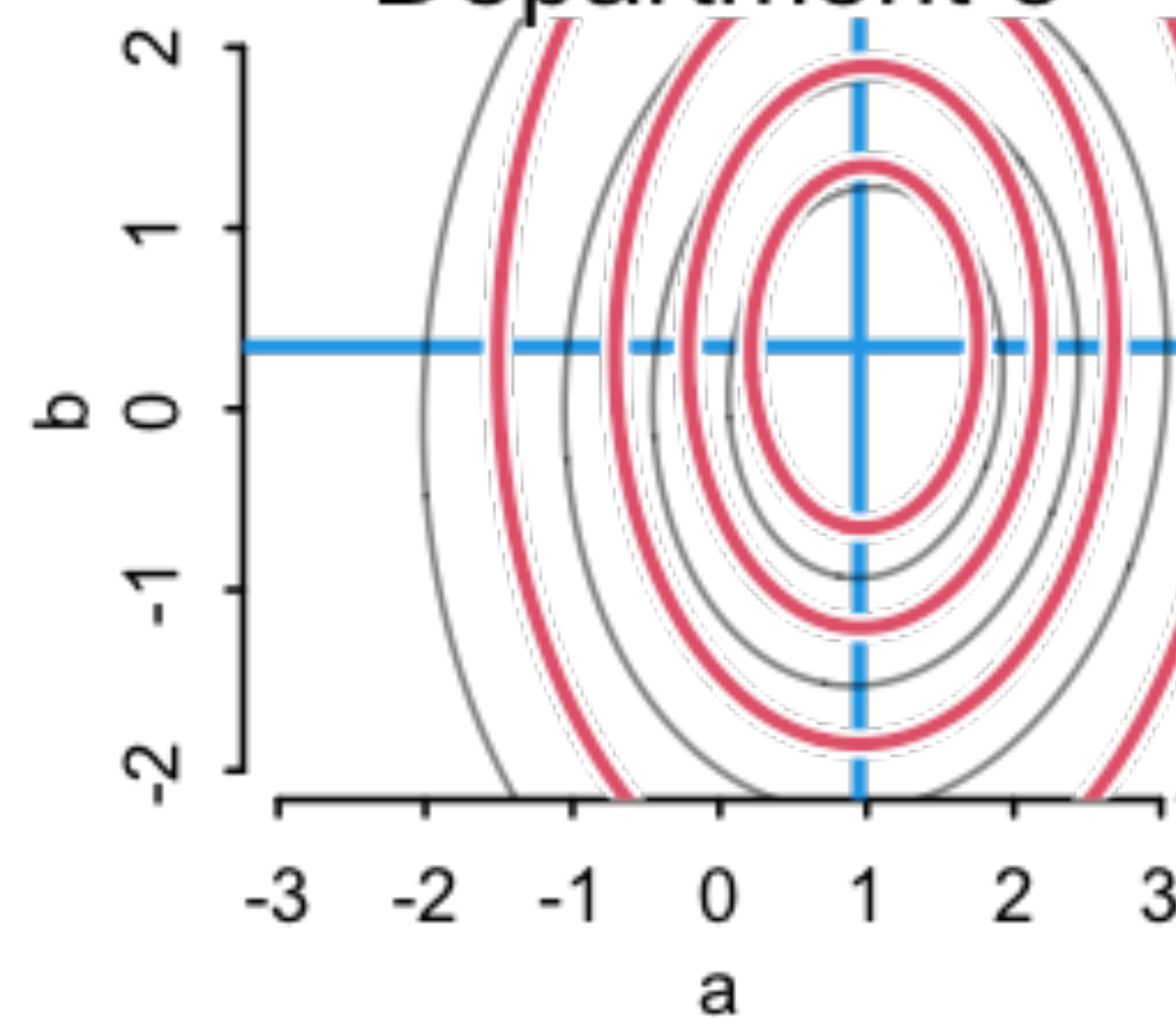
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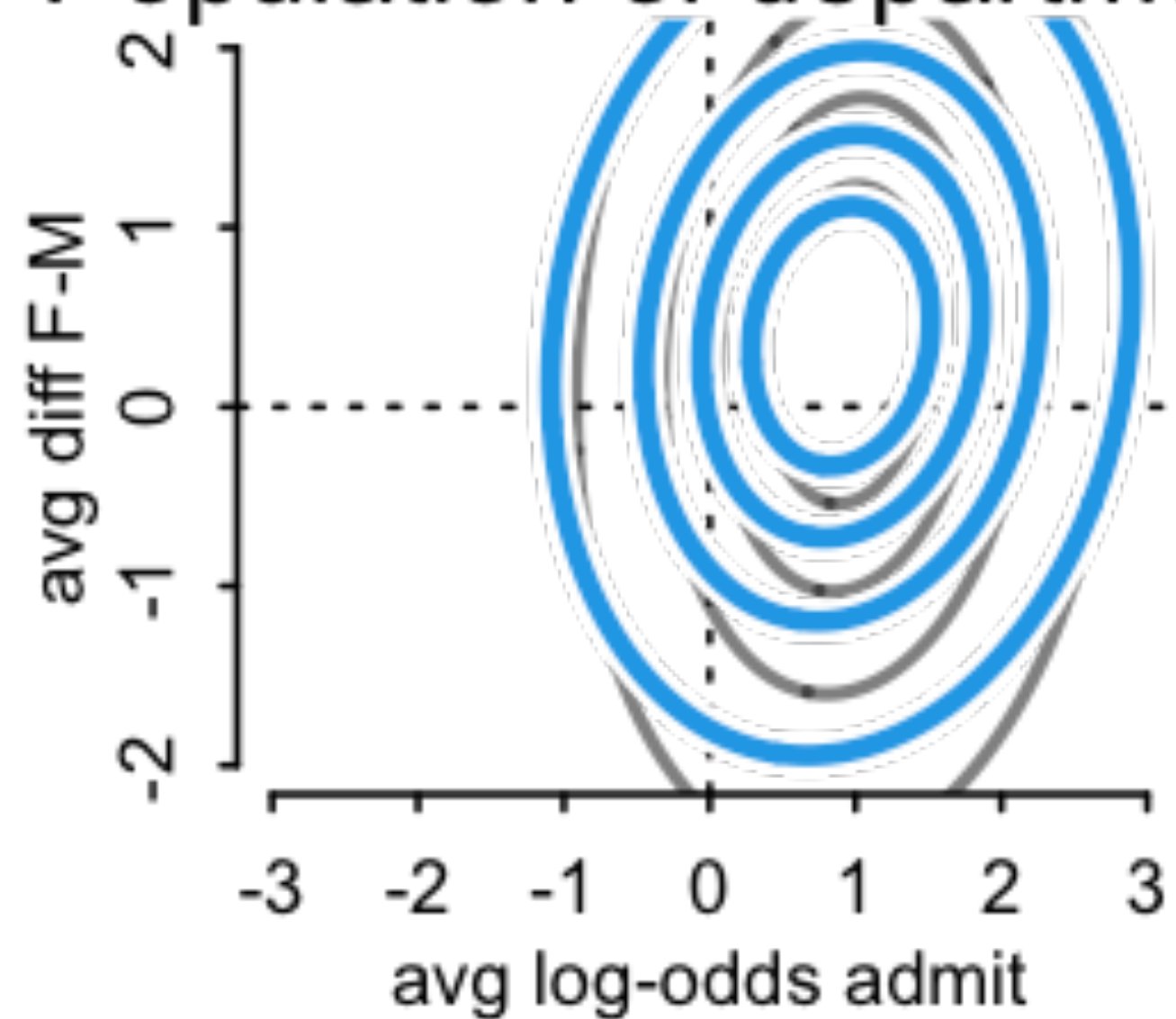
Department 4



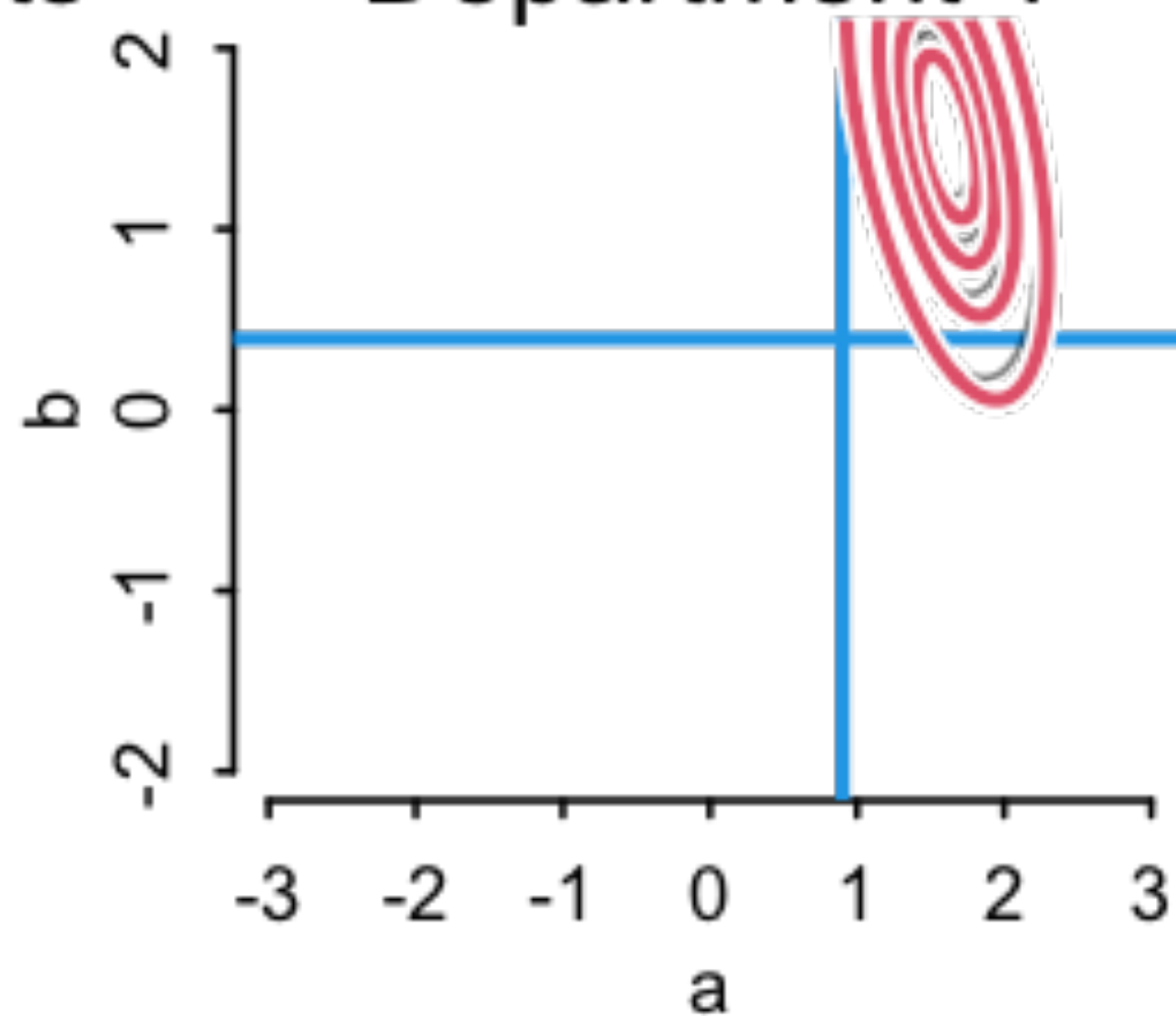
Department 5



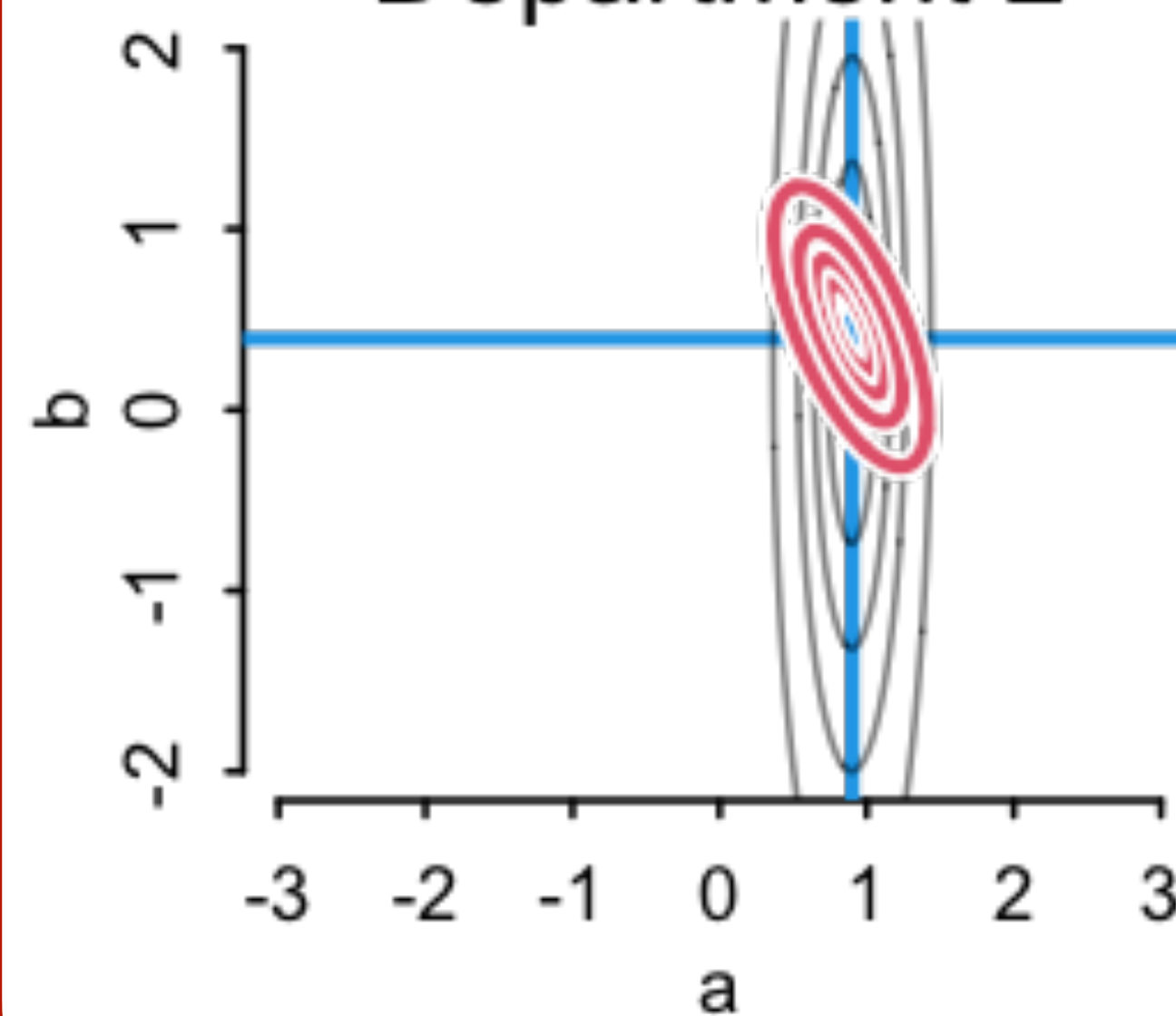
Population of departments



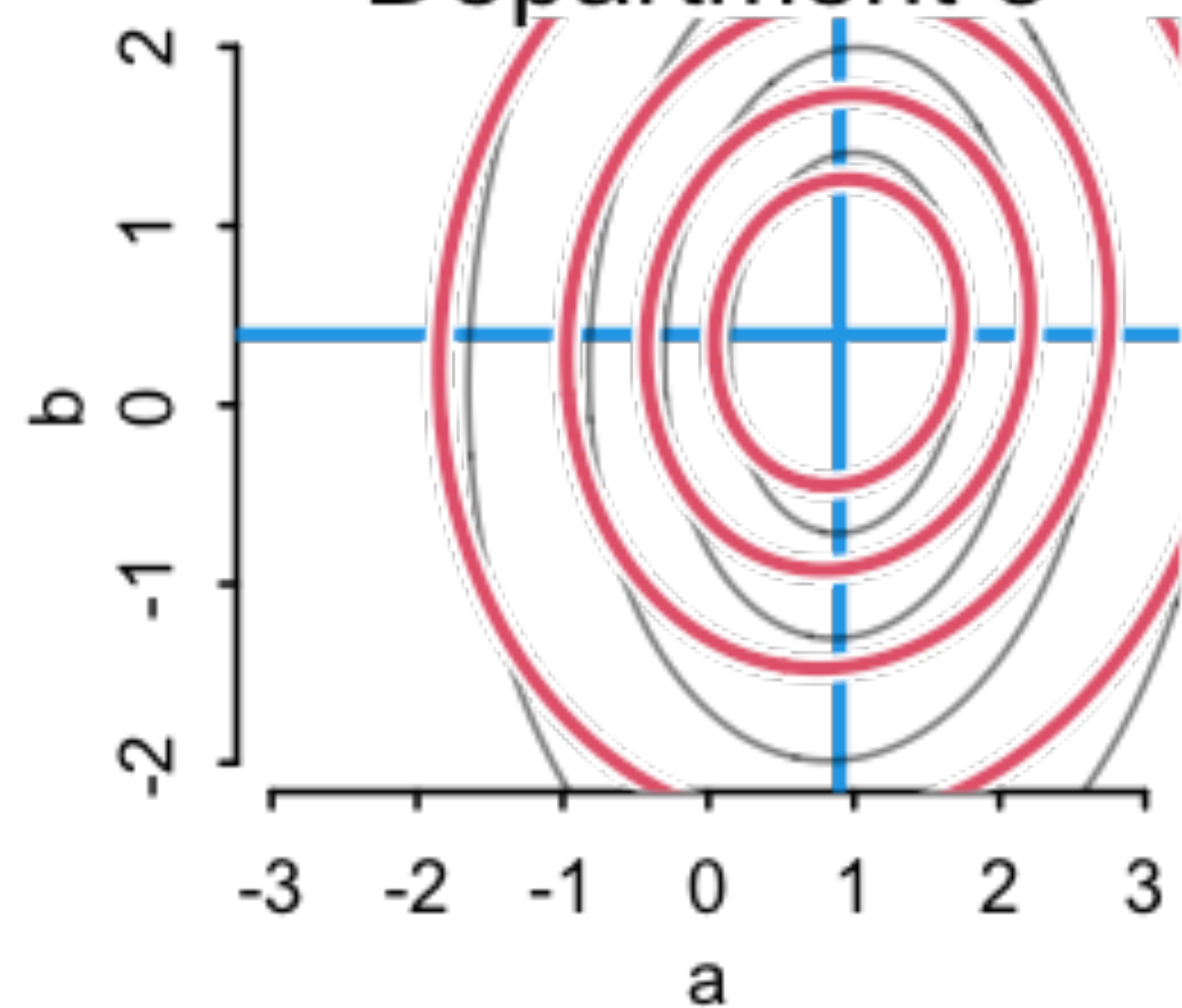
Department 1



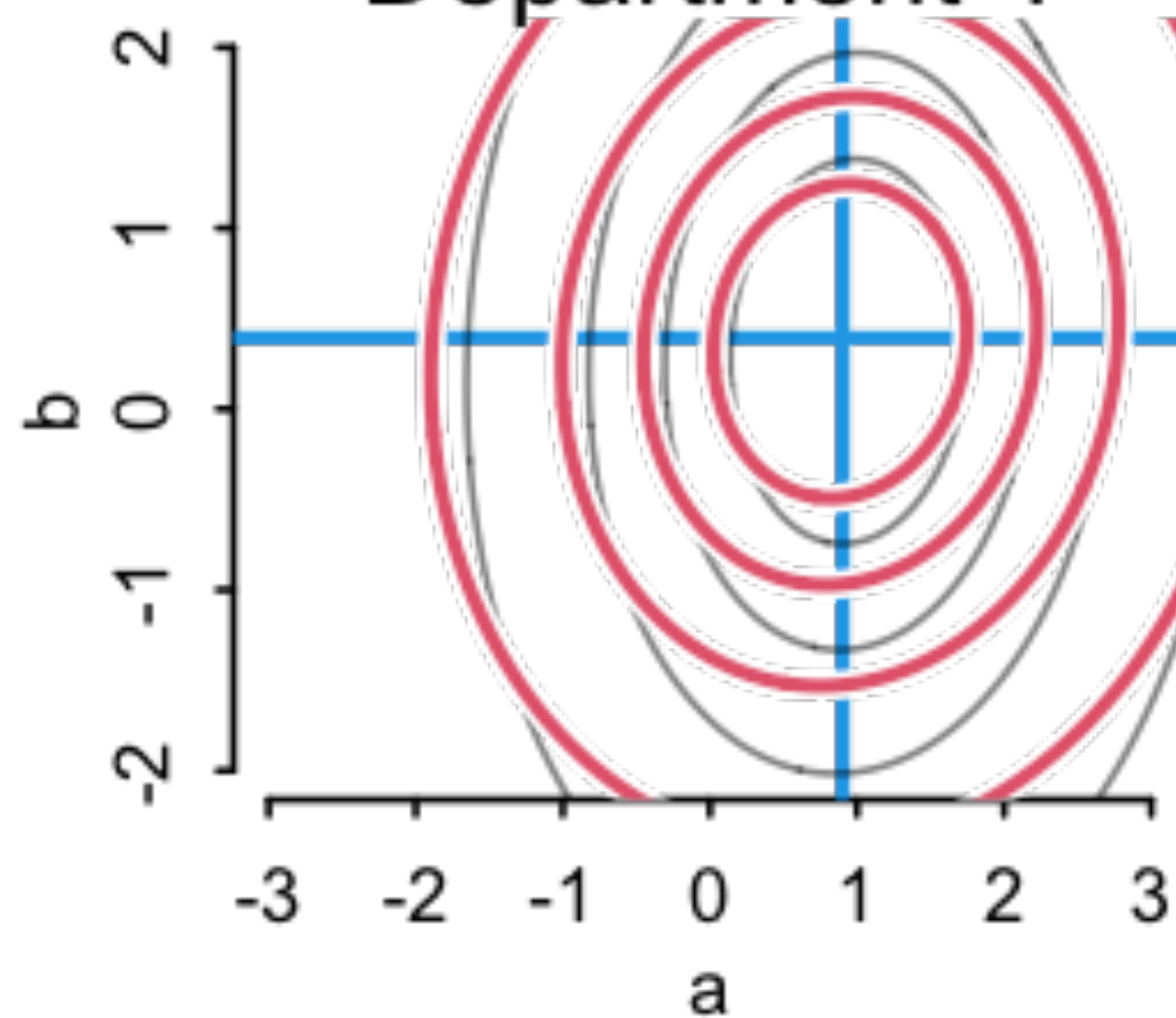
Department 2



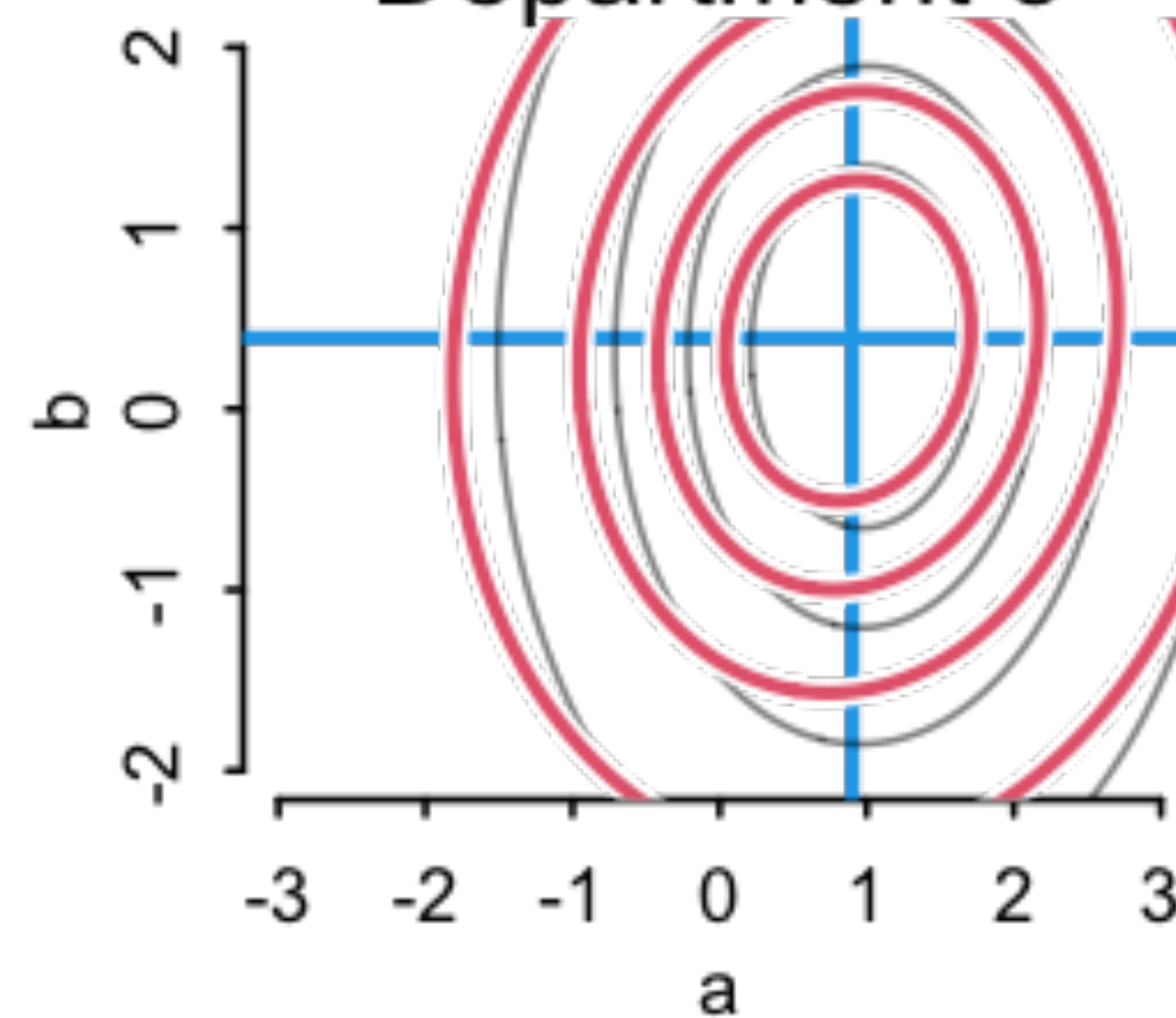
Department 3



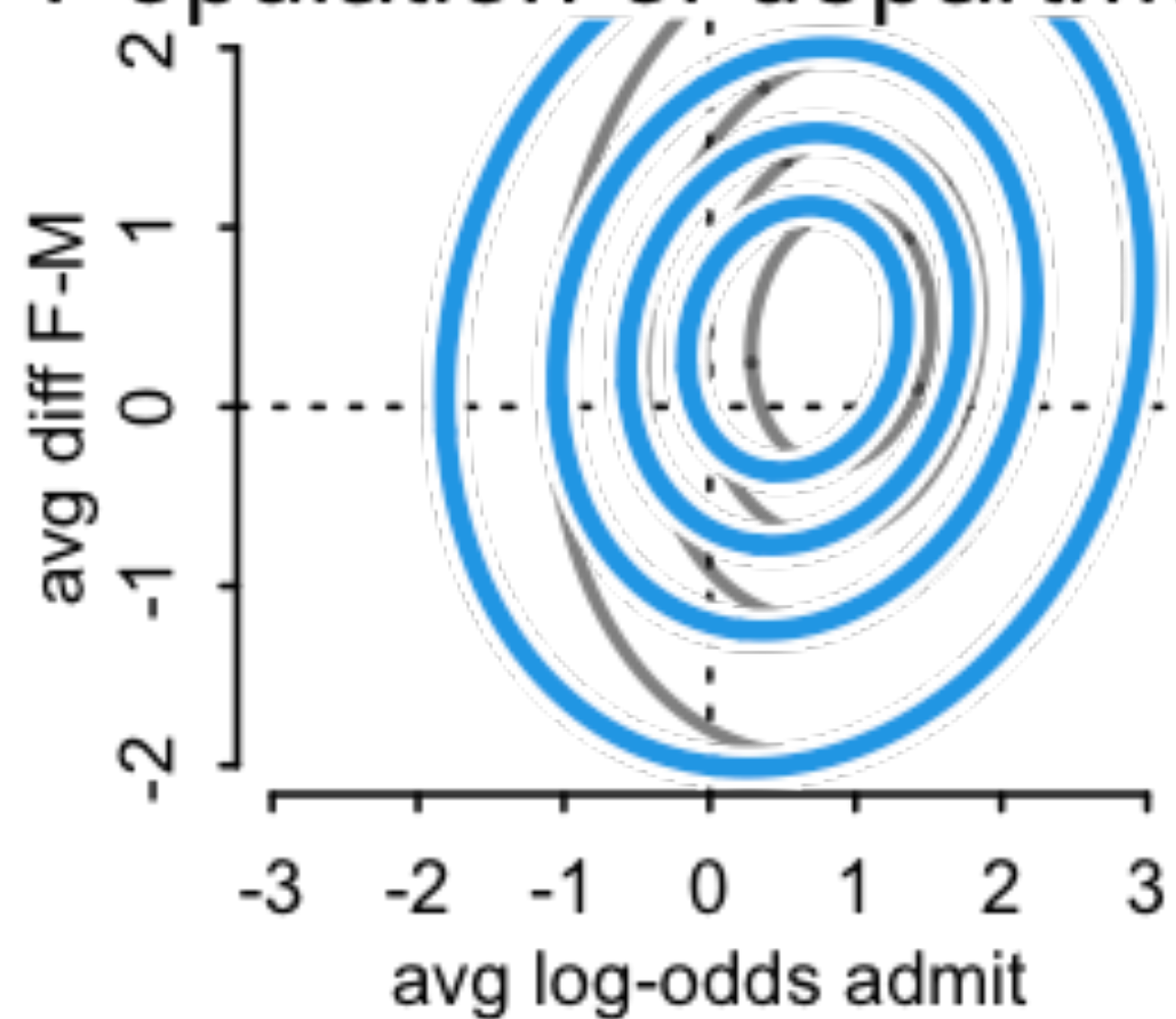
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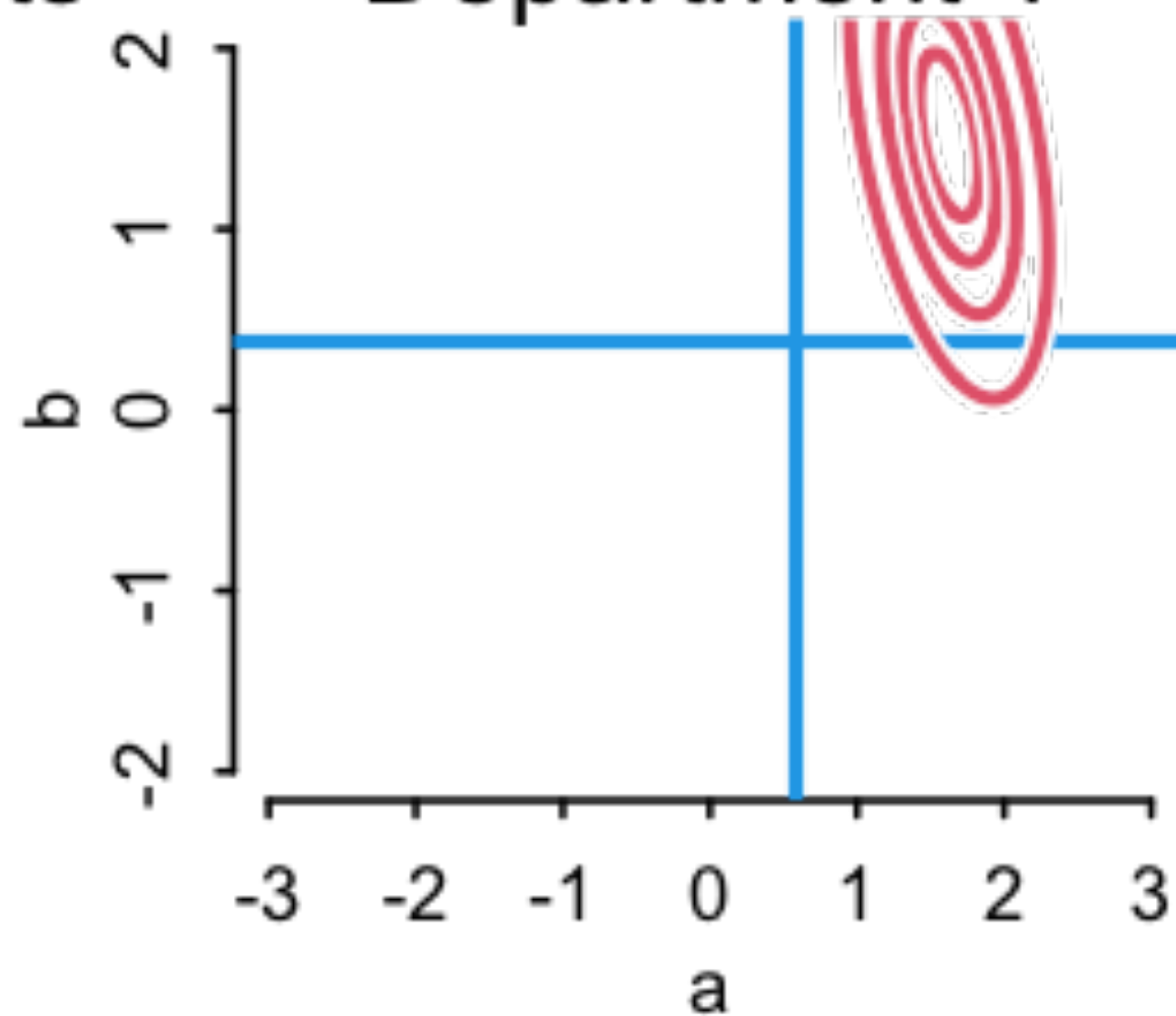
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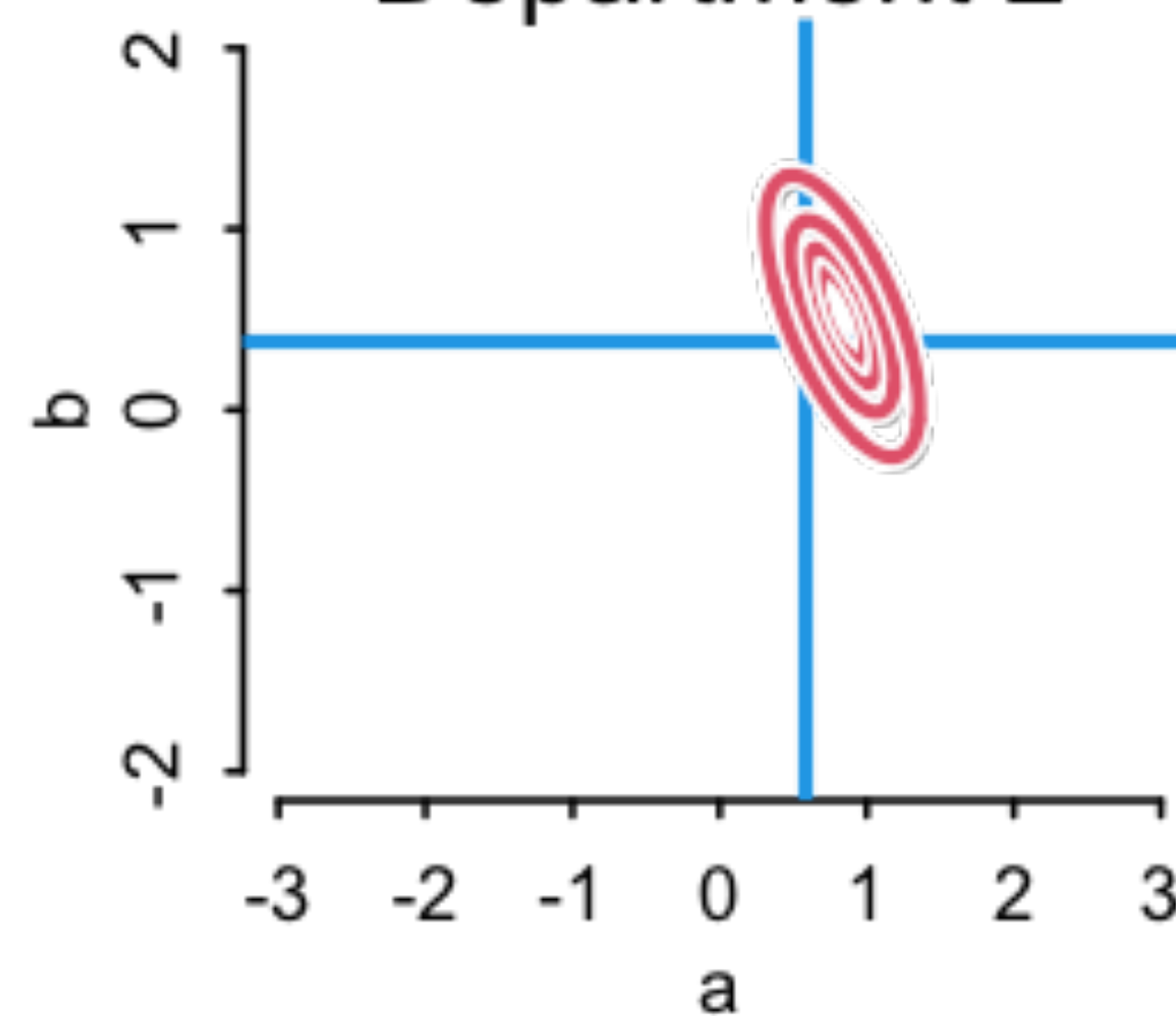
Population of departments



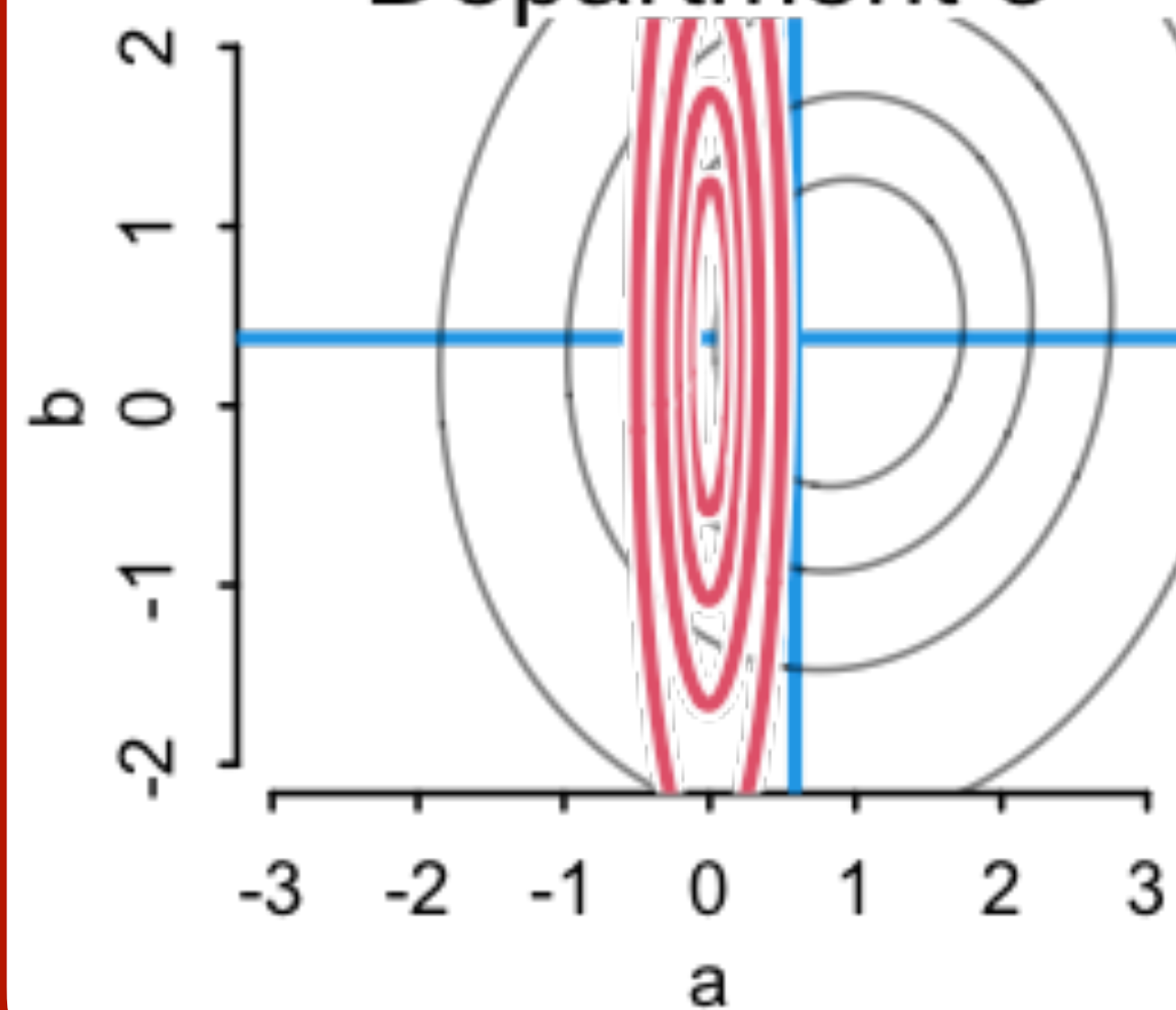
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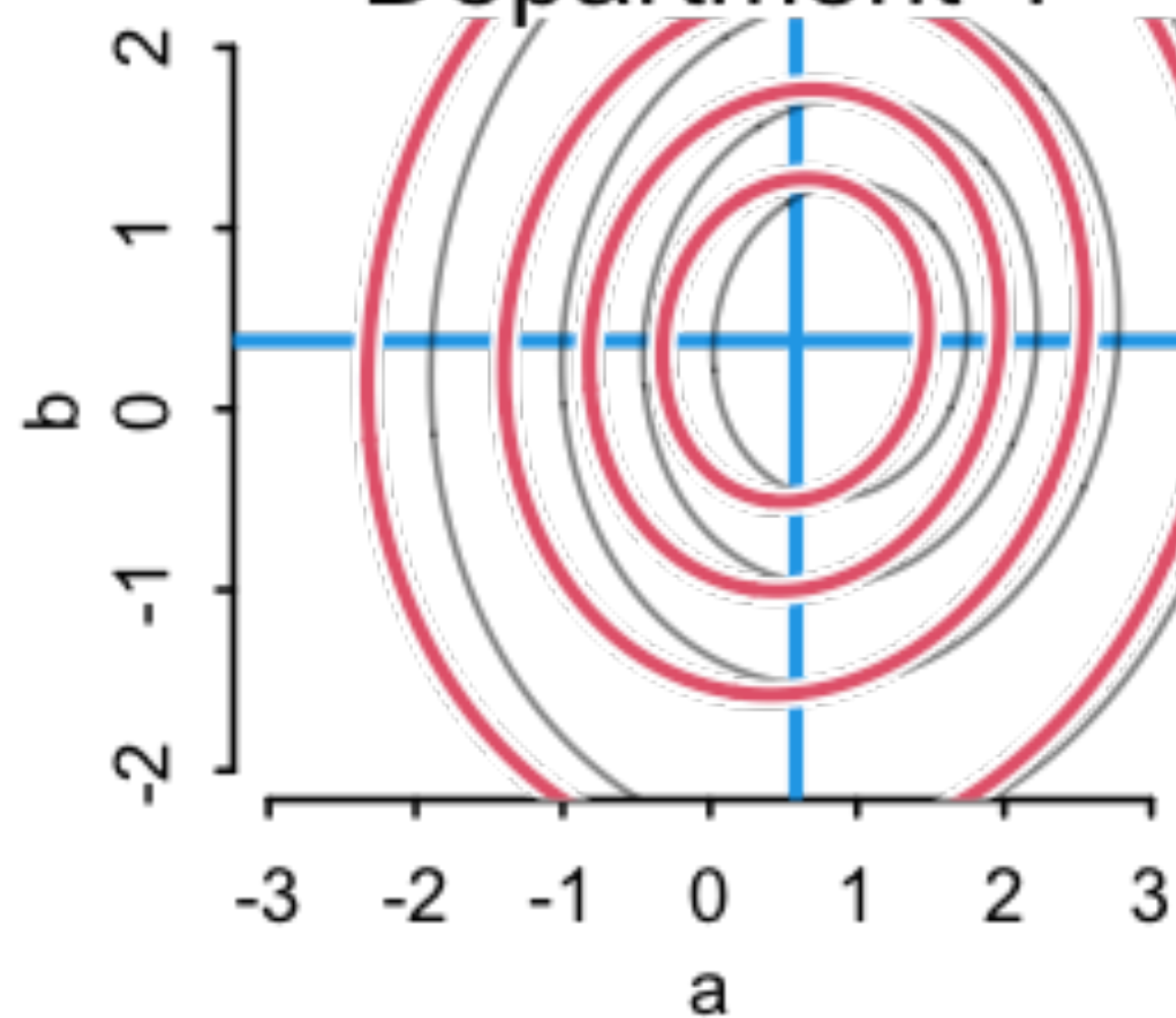
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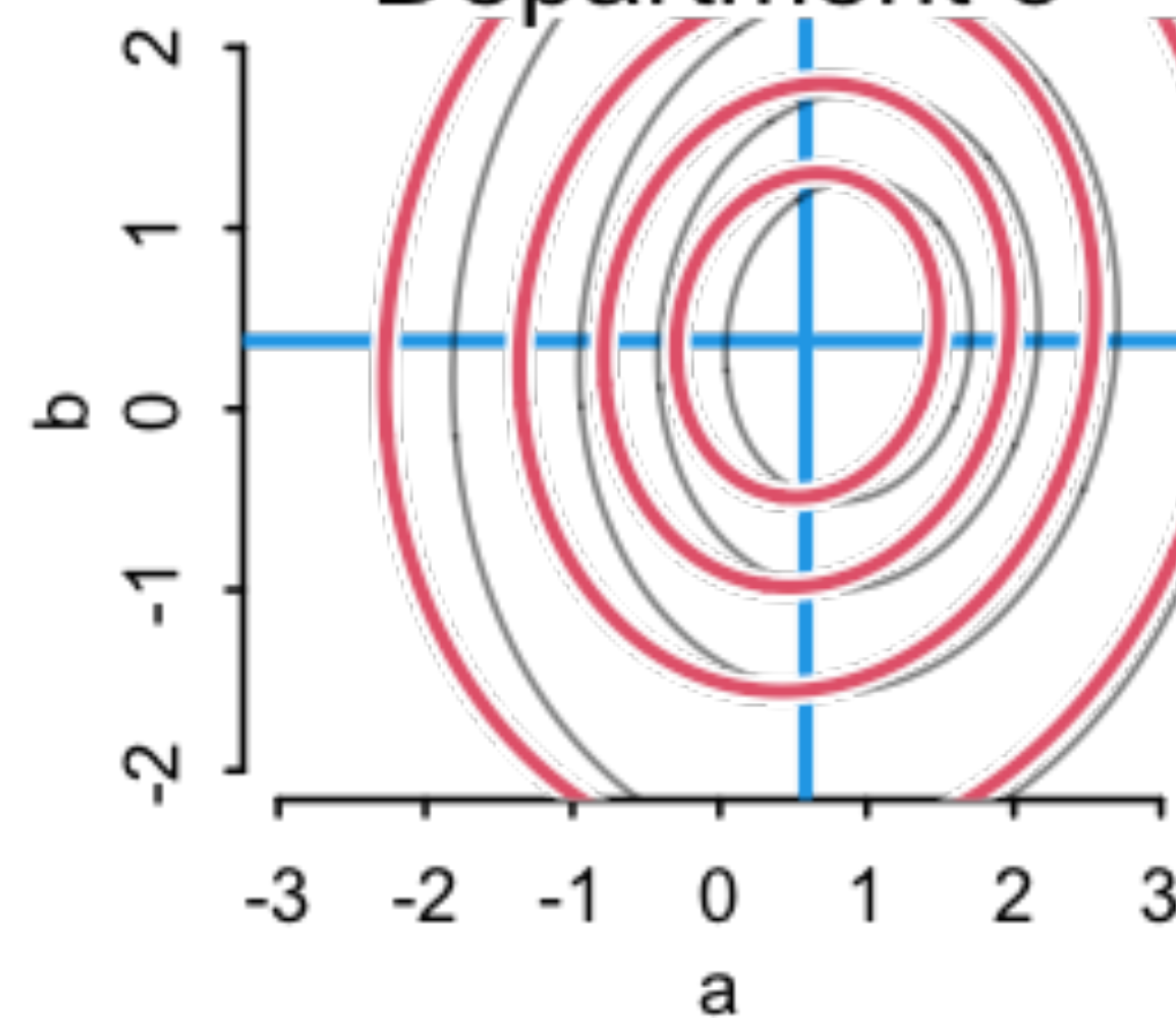
Department 3



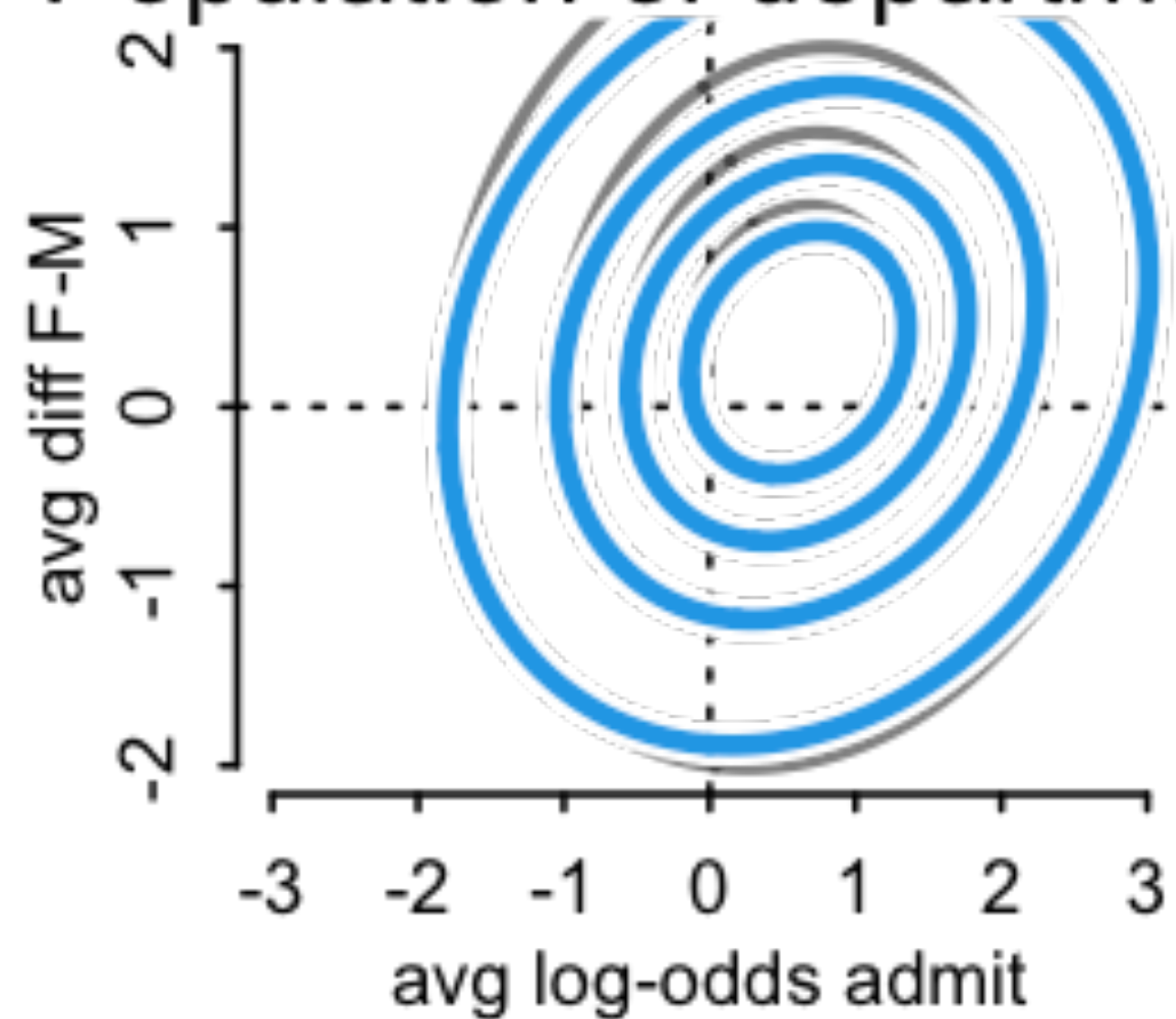
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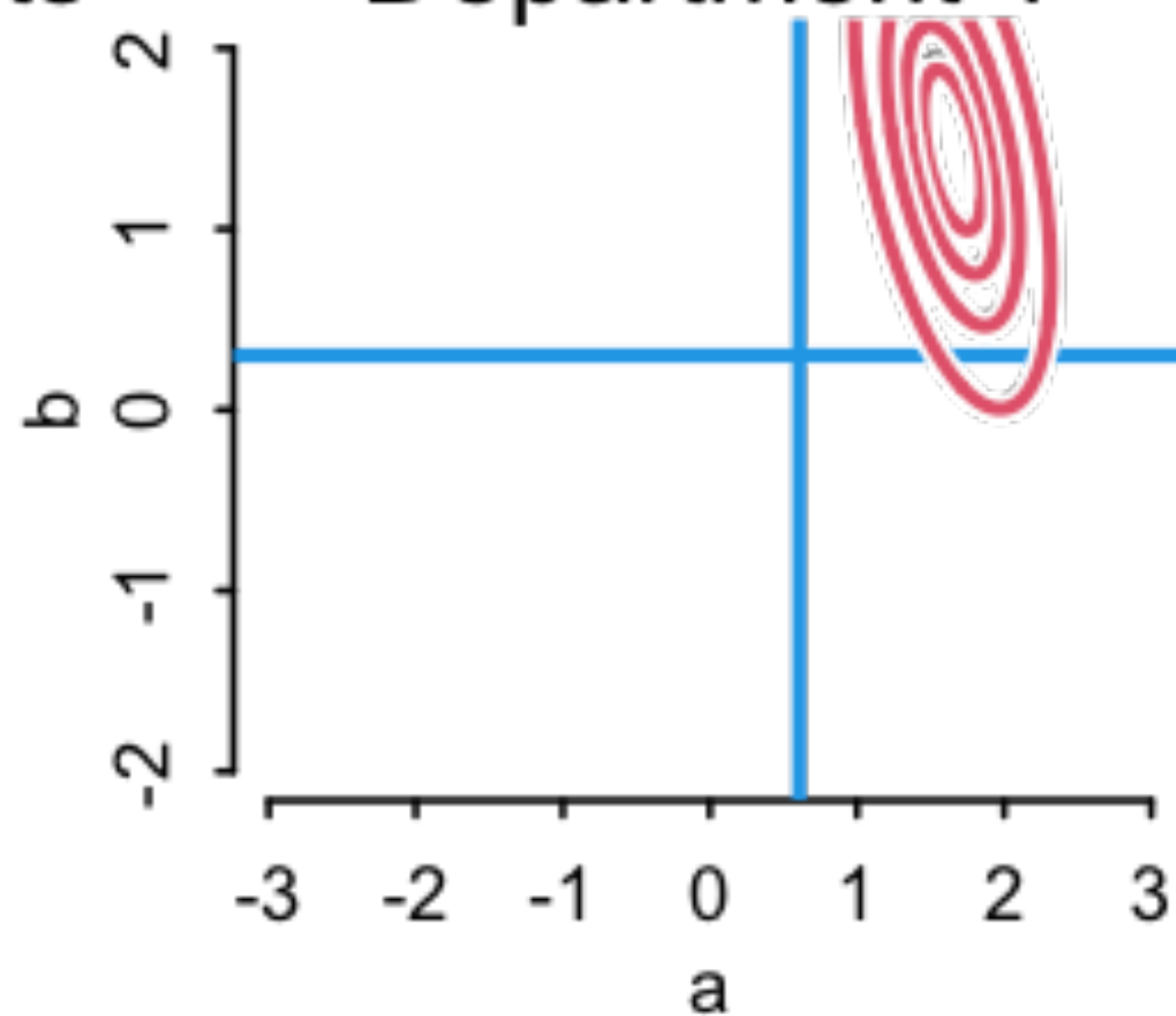
Department 5



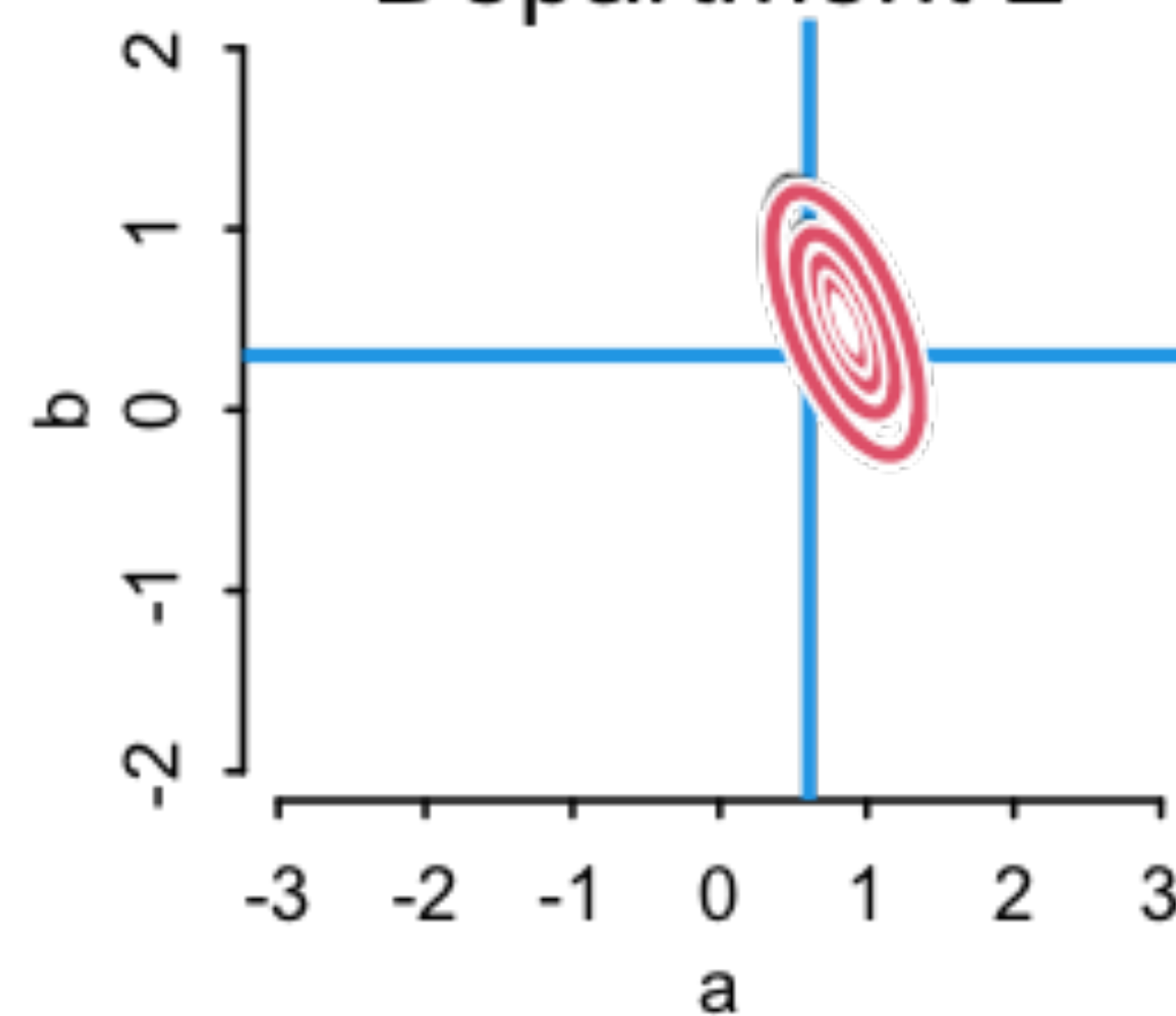
Population of departments



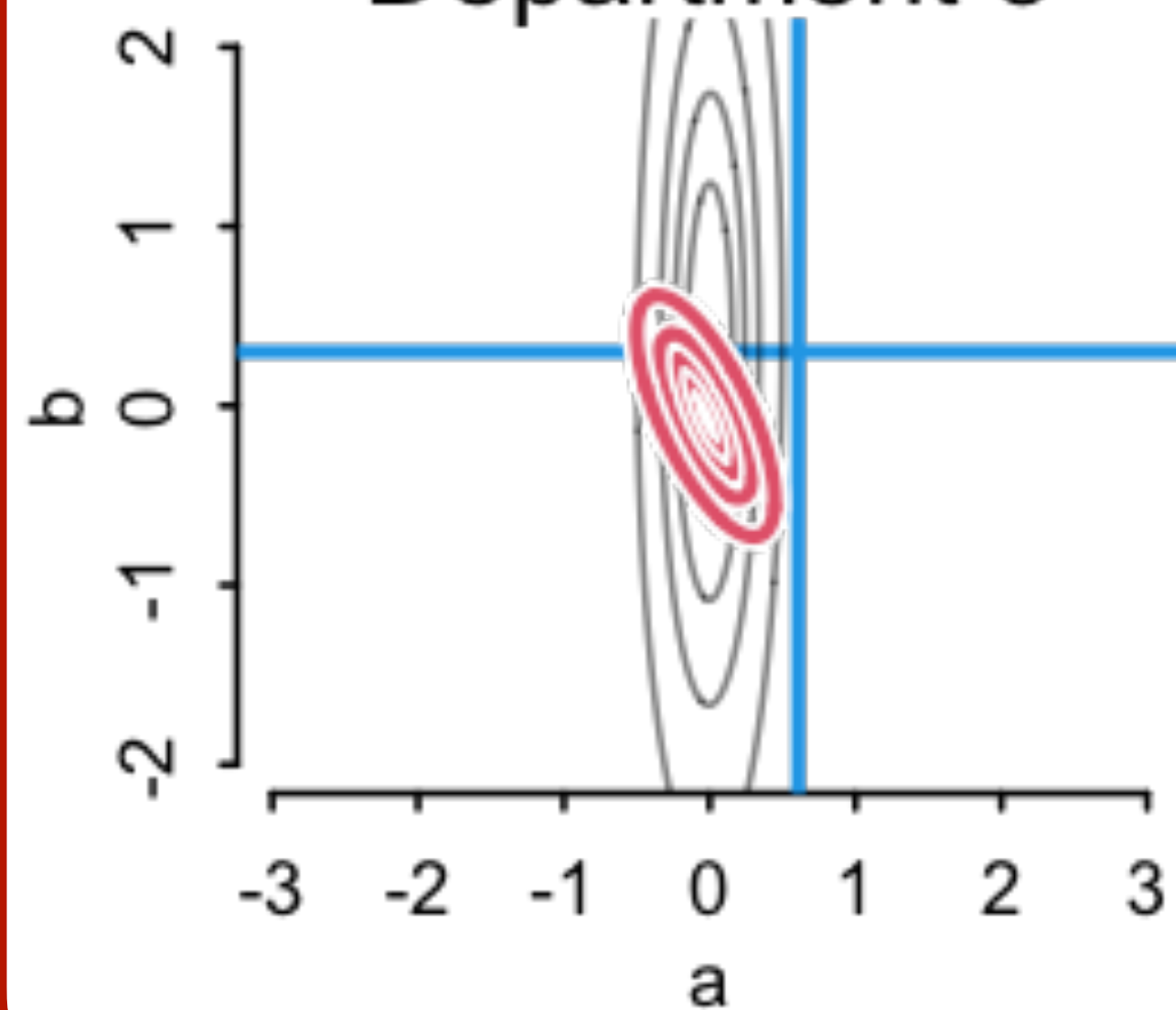
Department 1



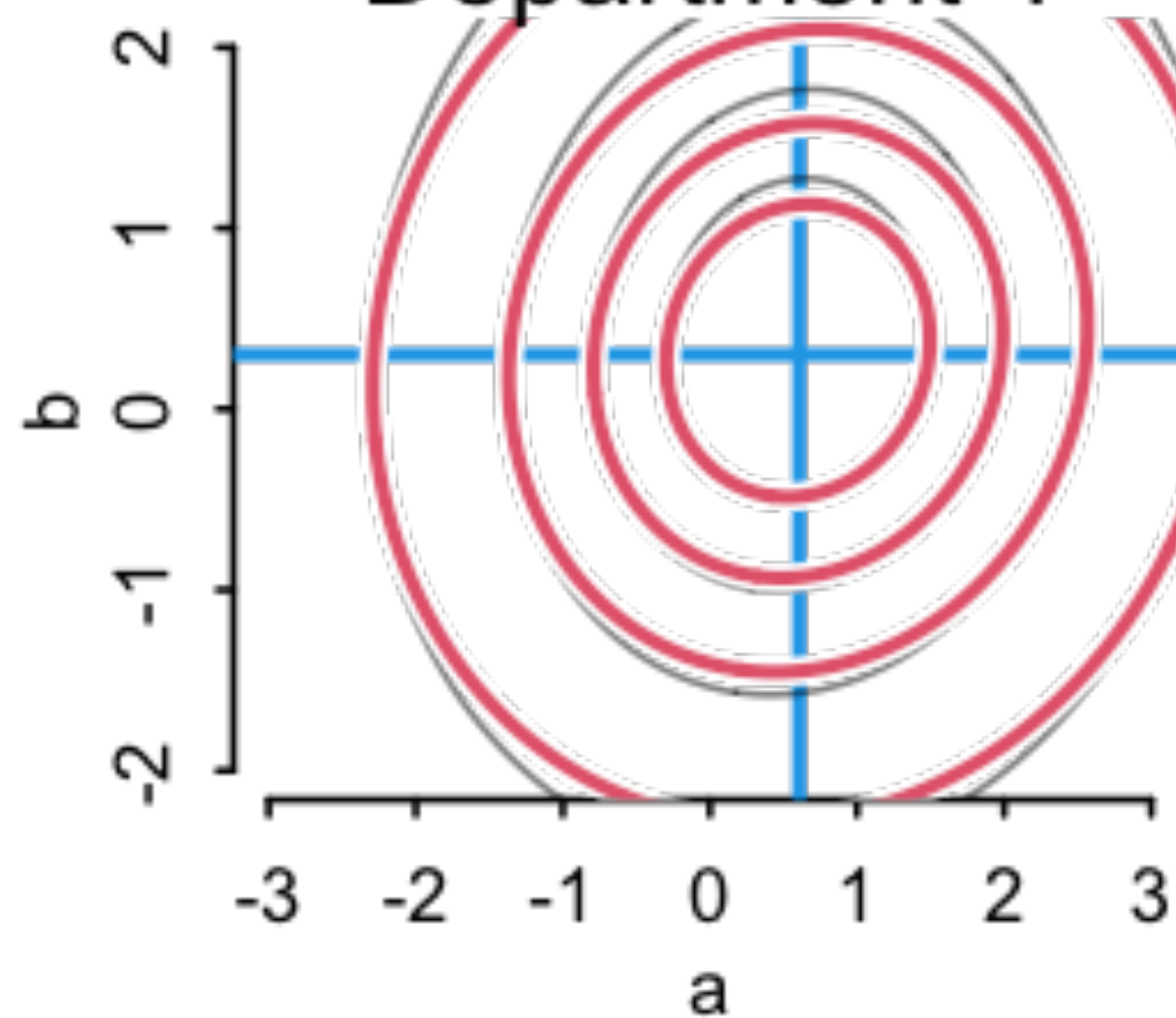
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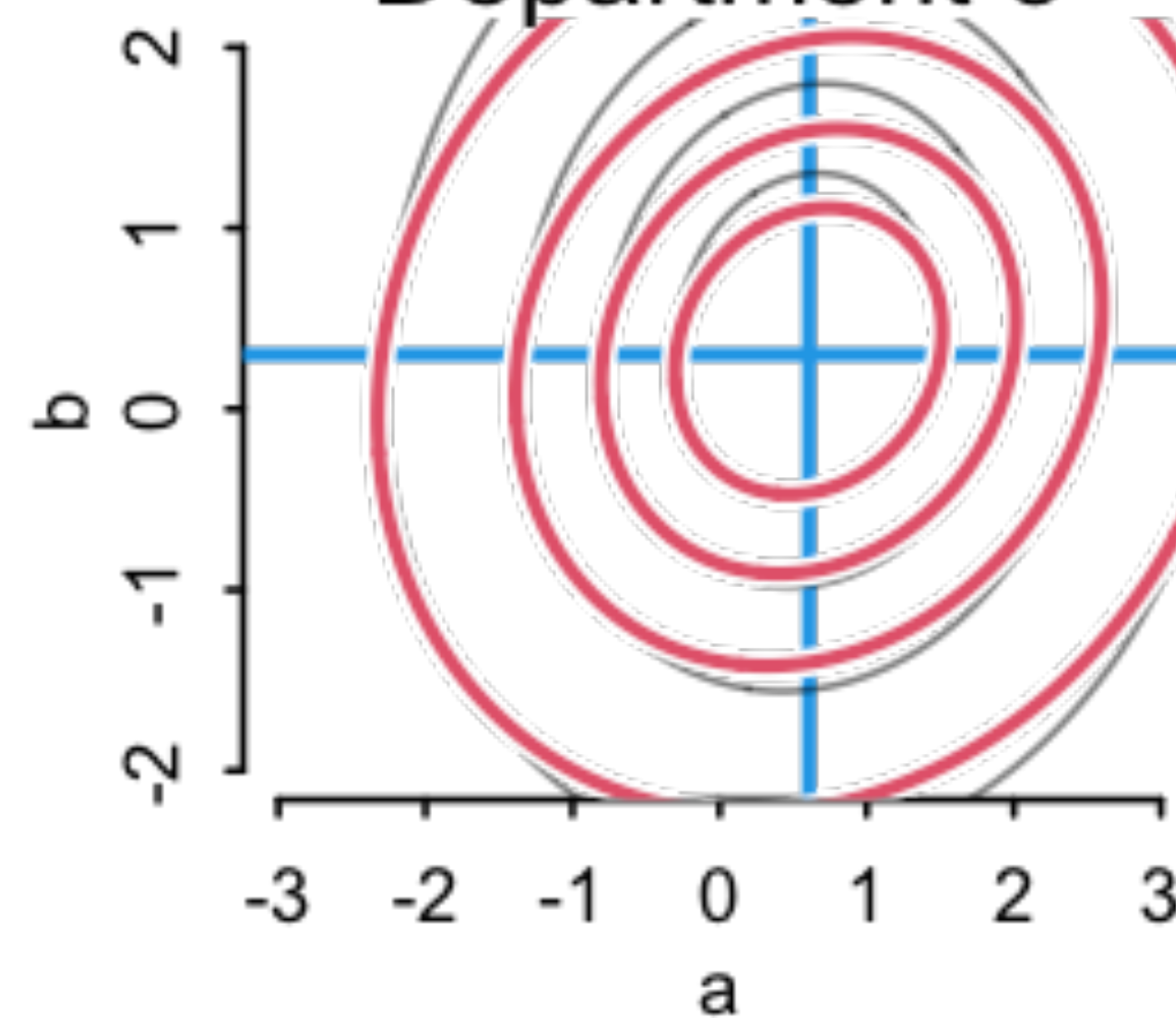
Department 3



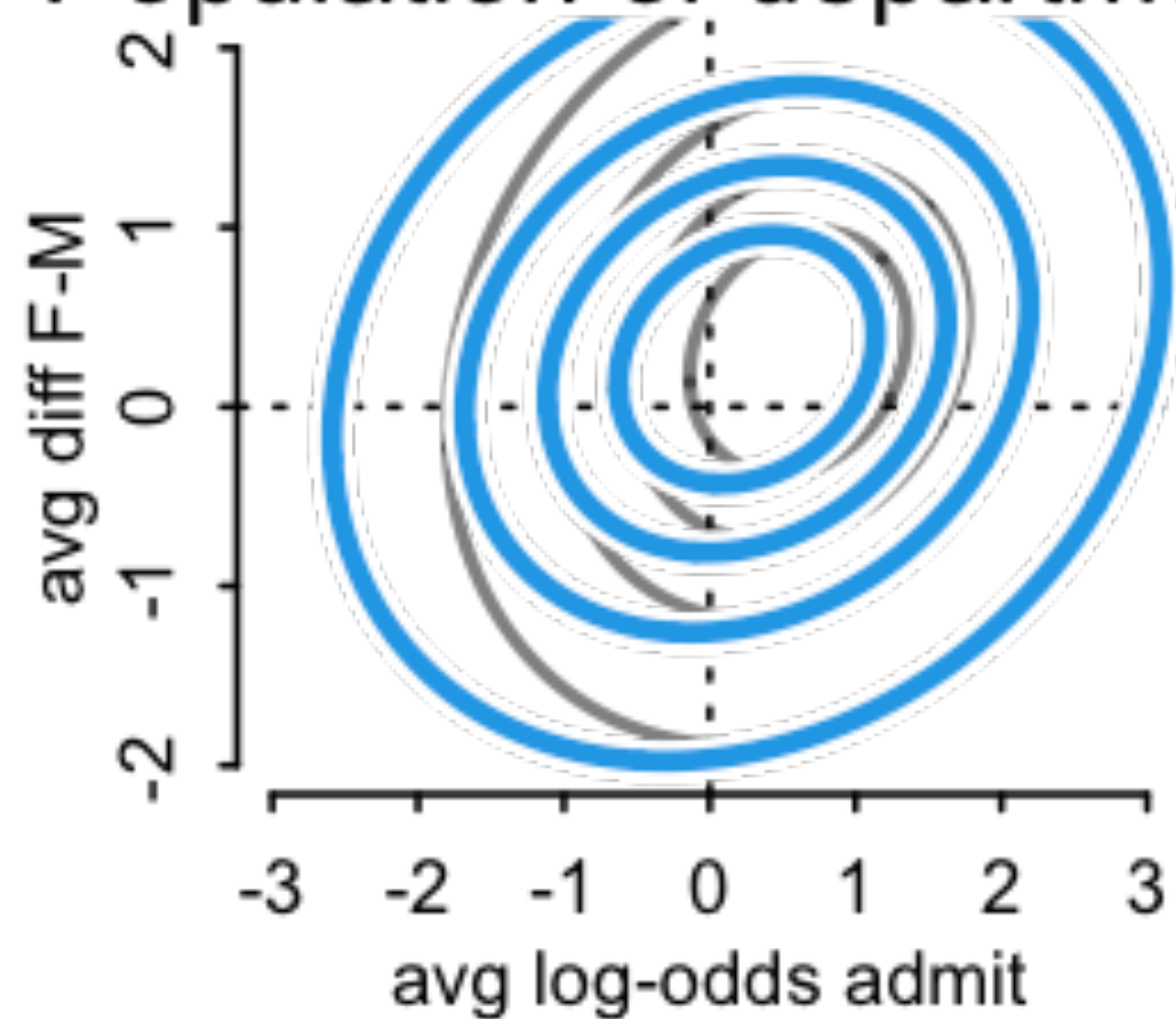
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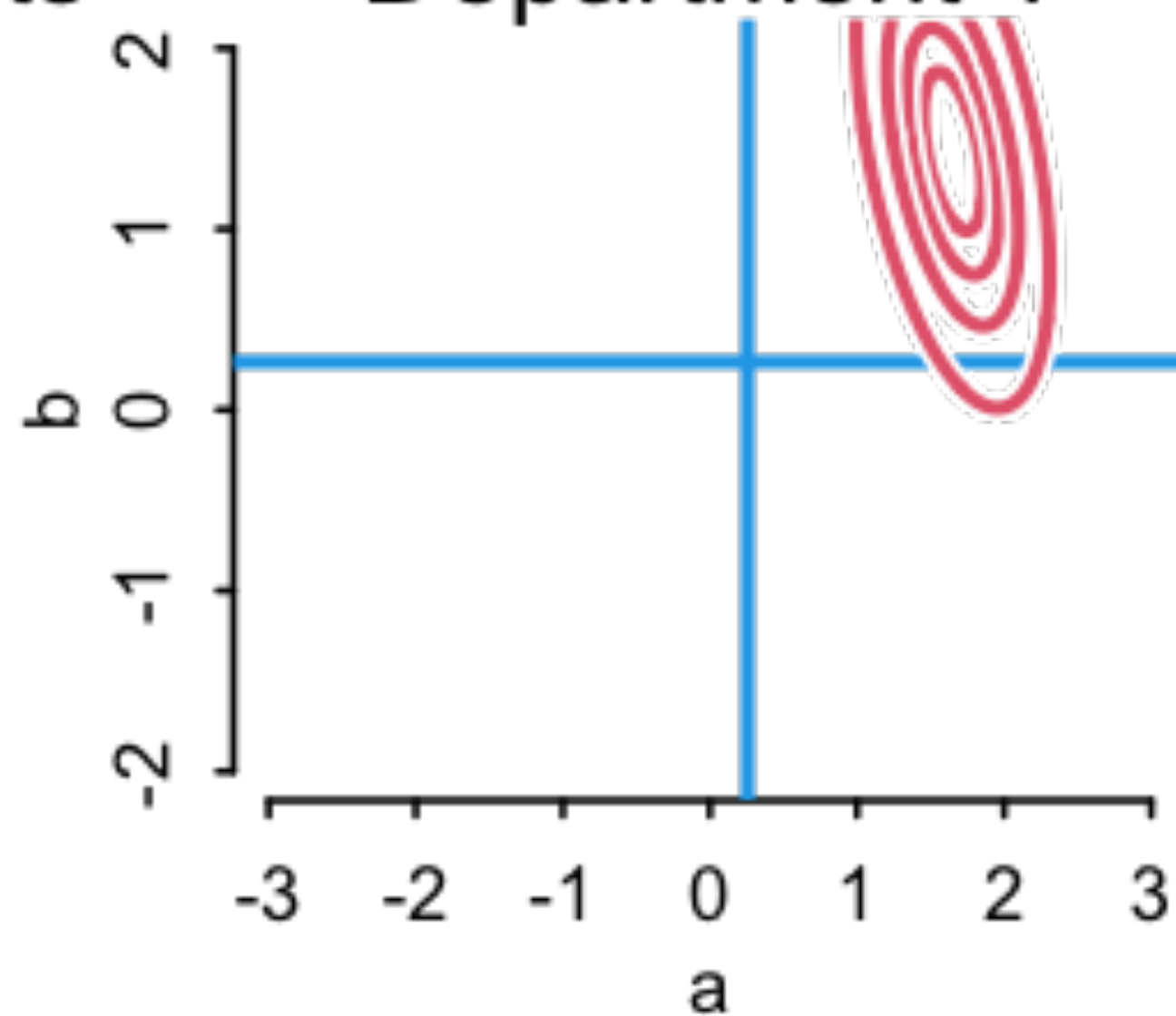
Department 5



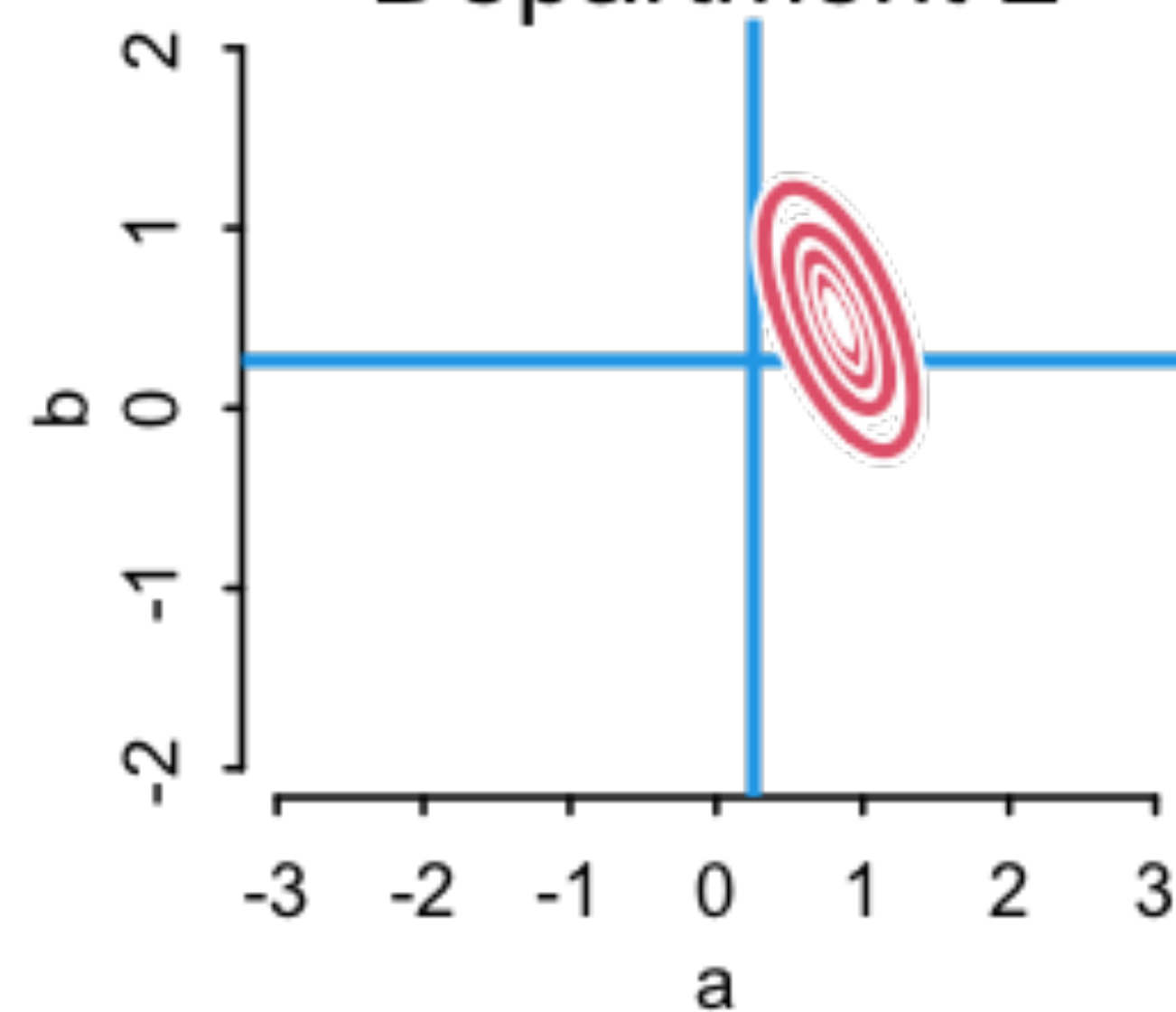
Population of departments



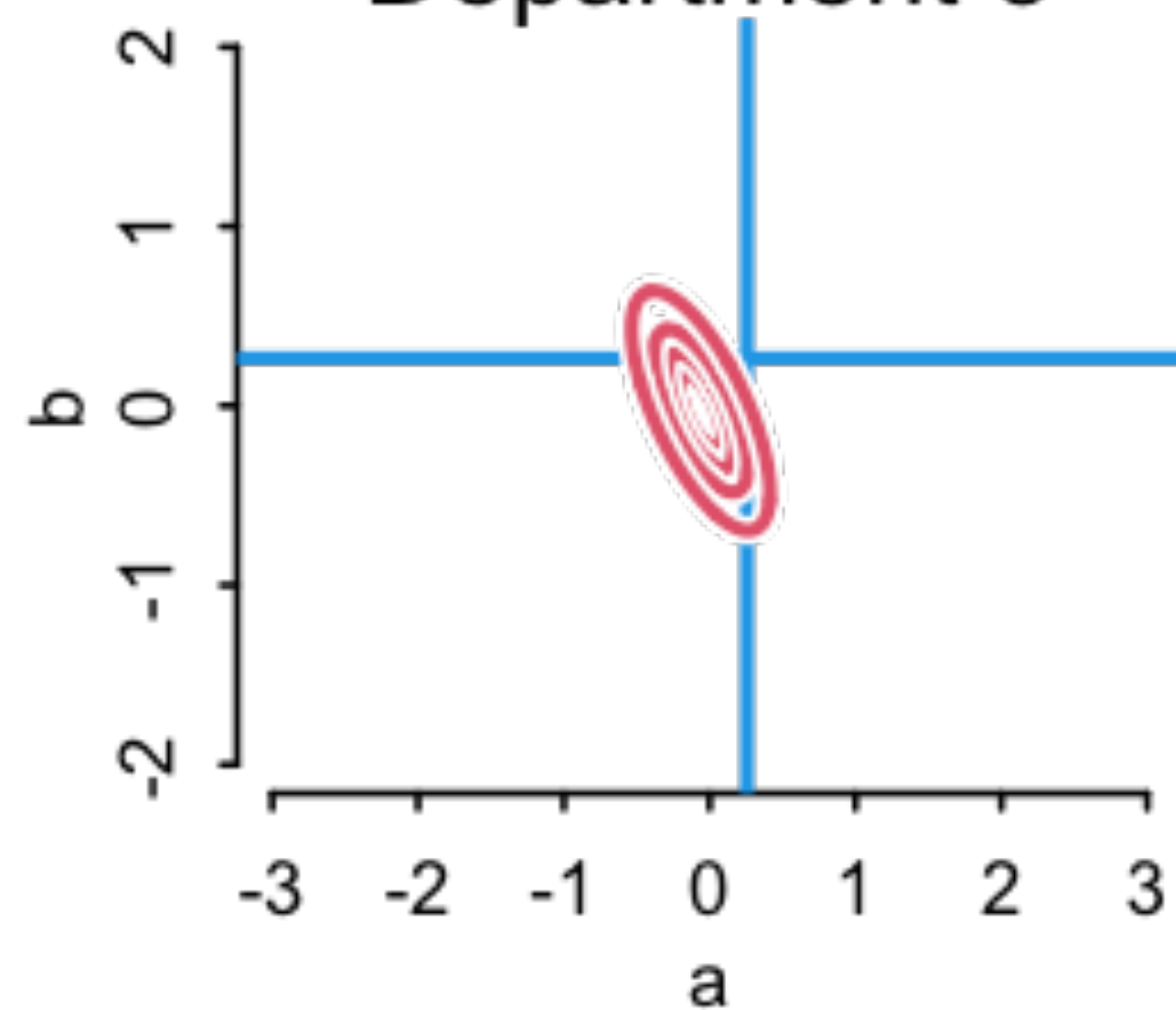
Department 1



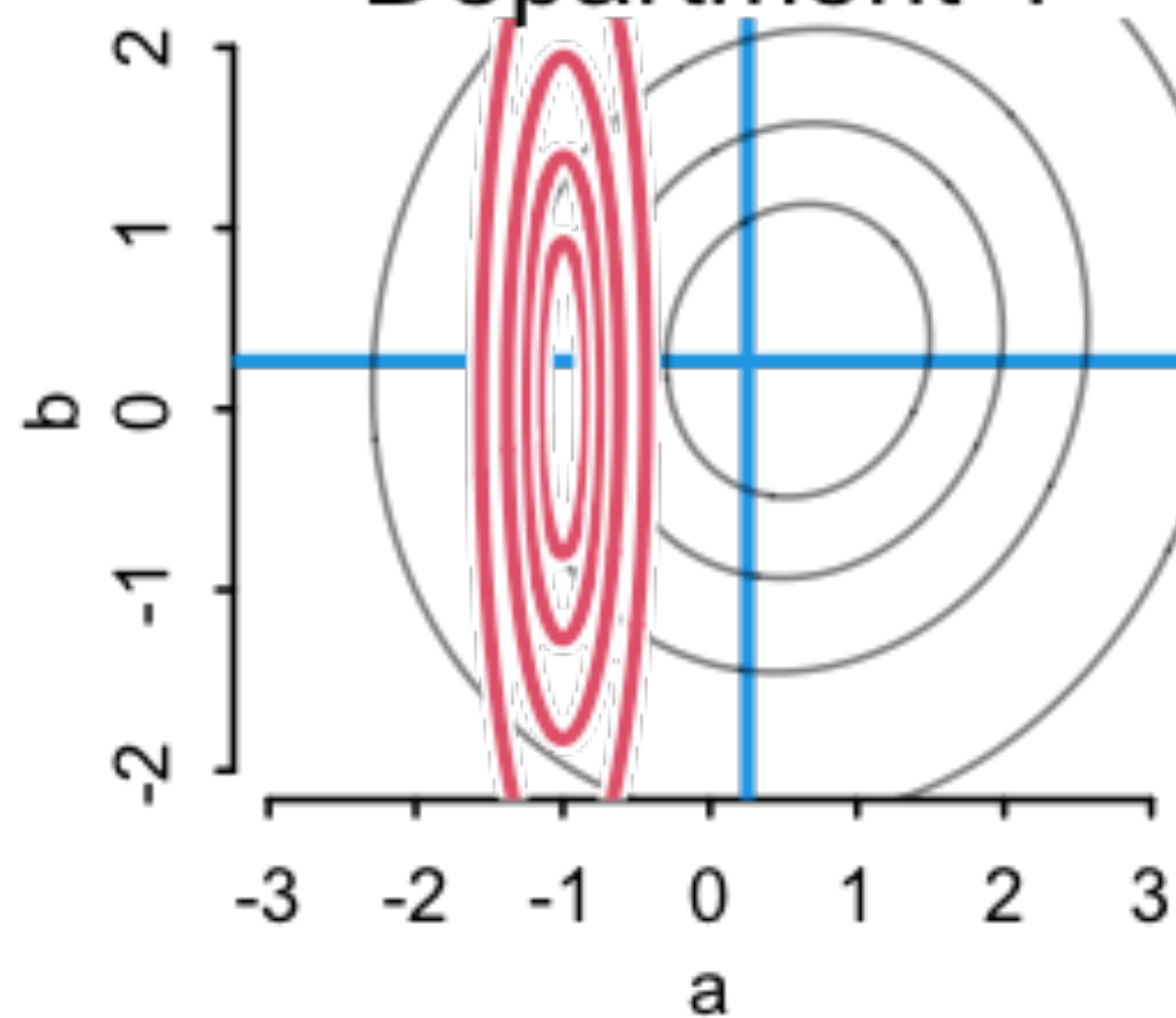
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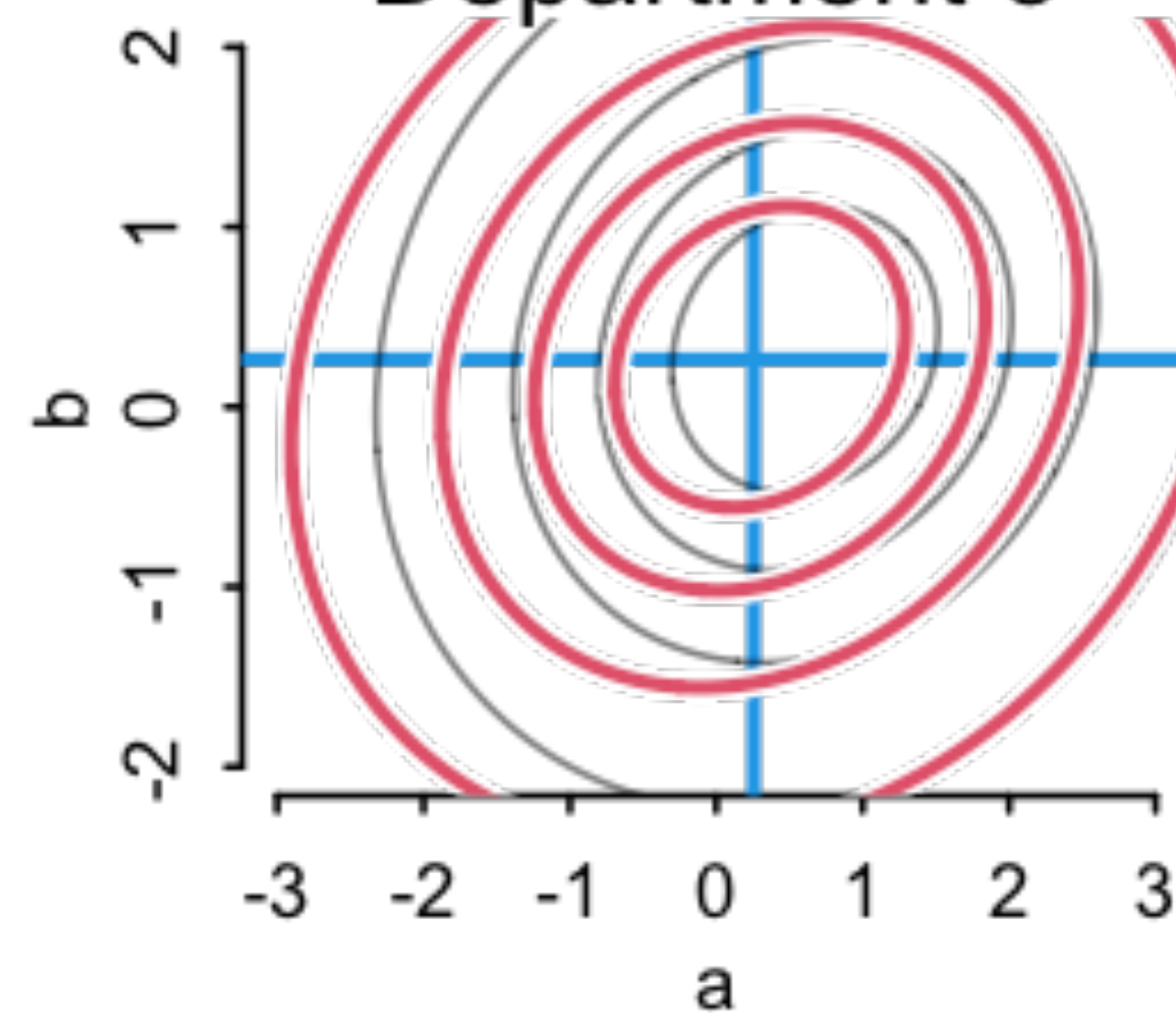
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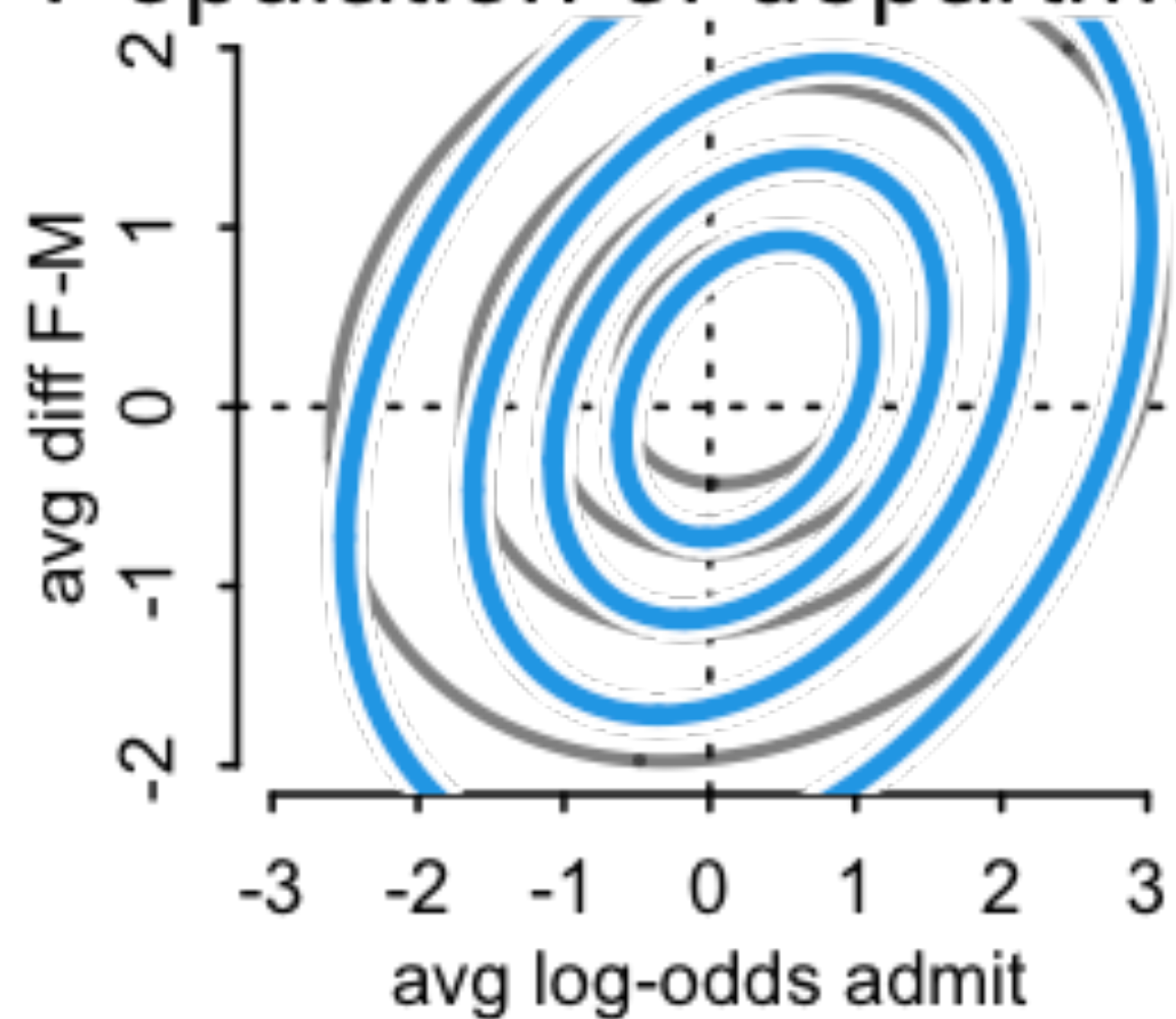
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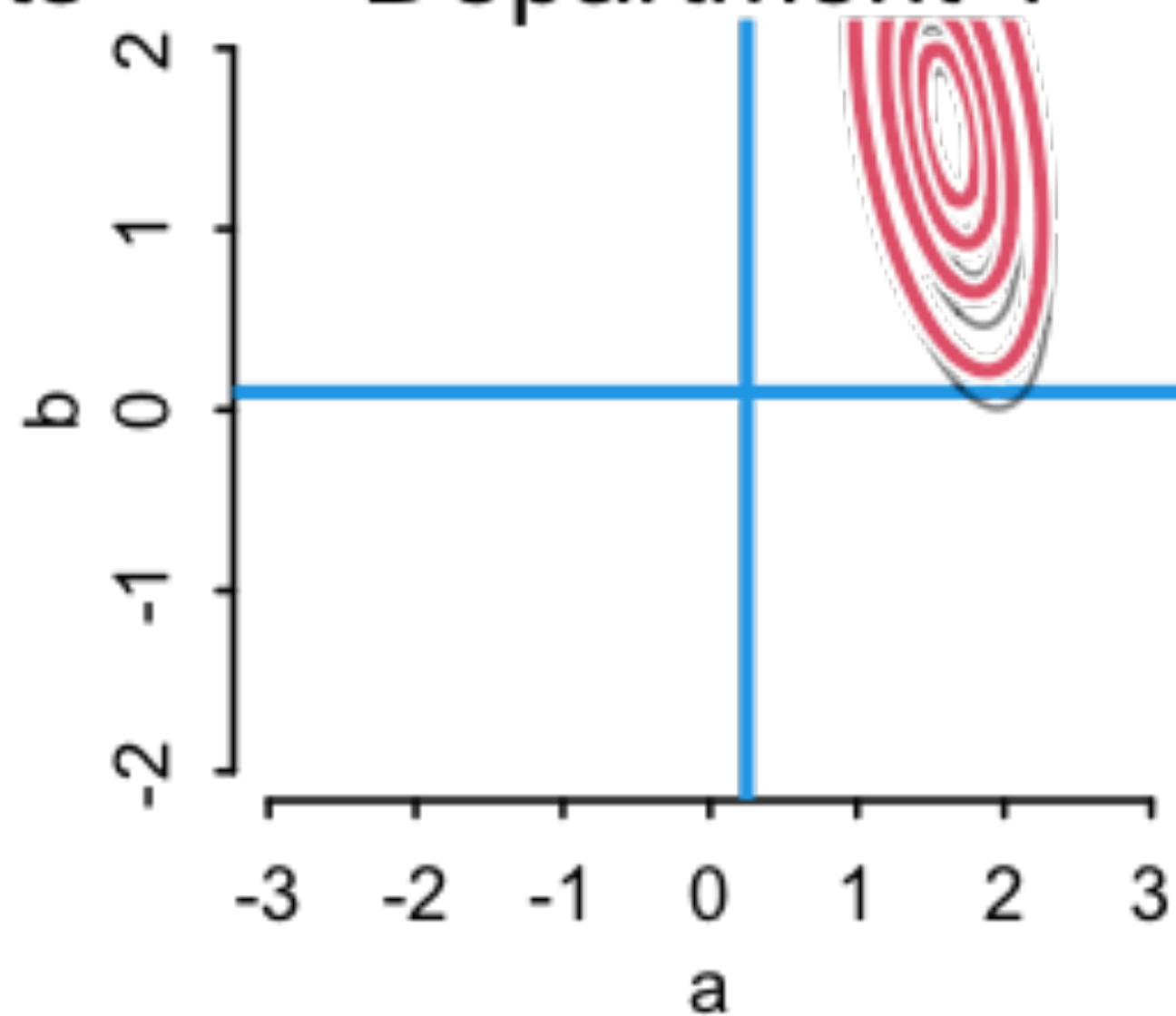
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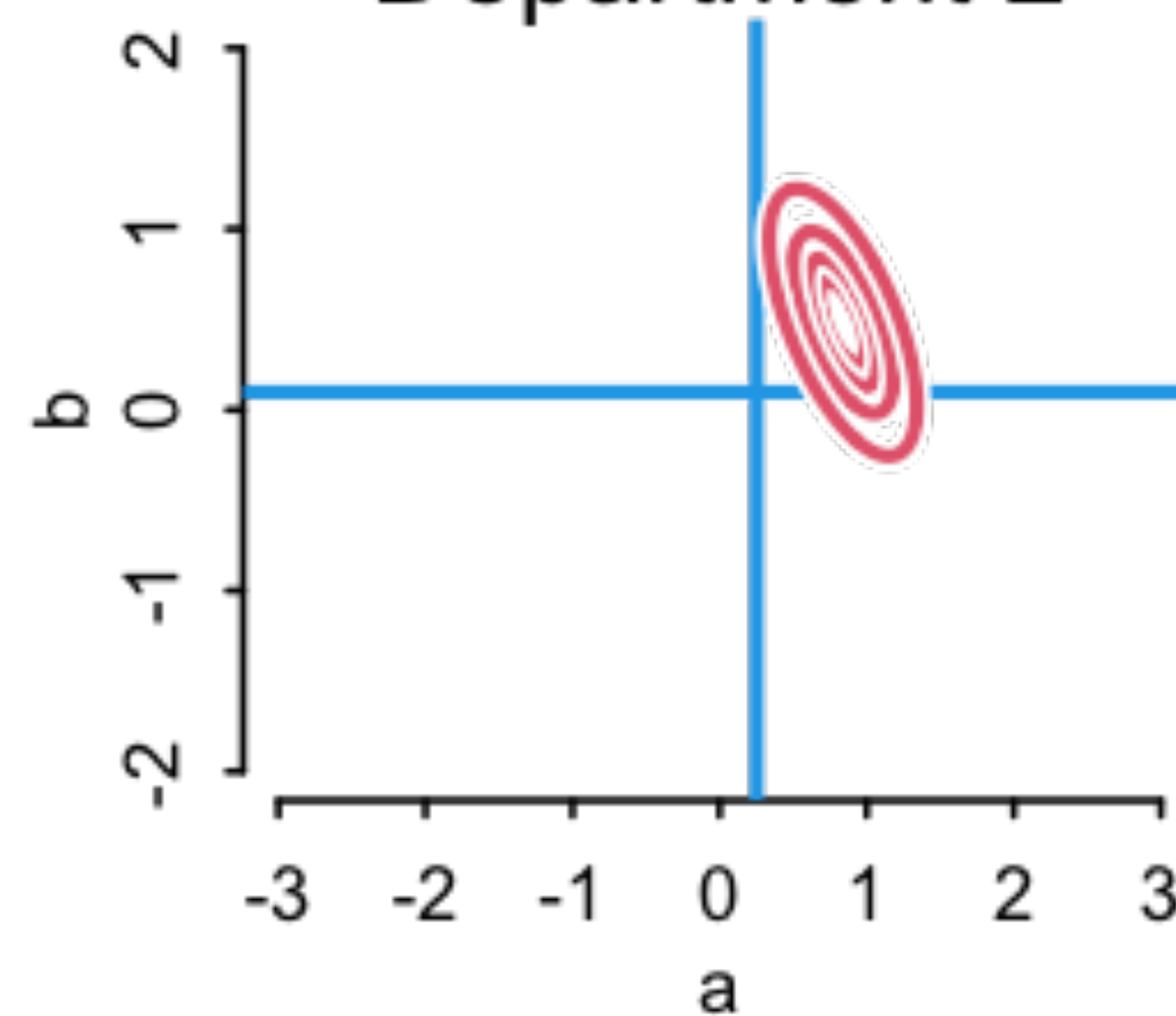
Population of departments



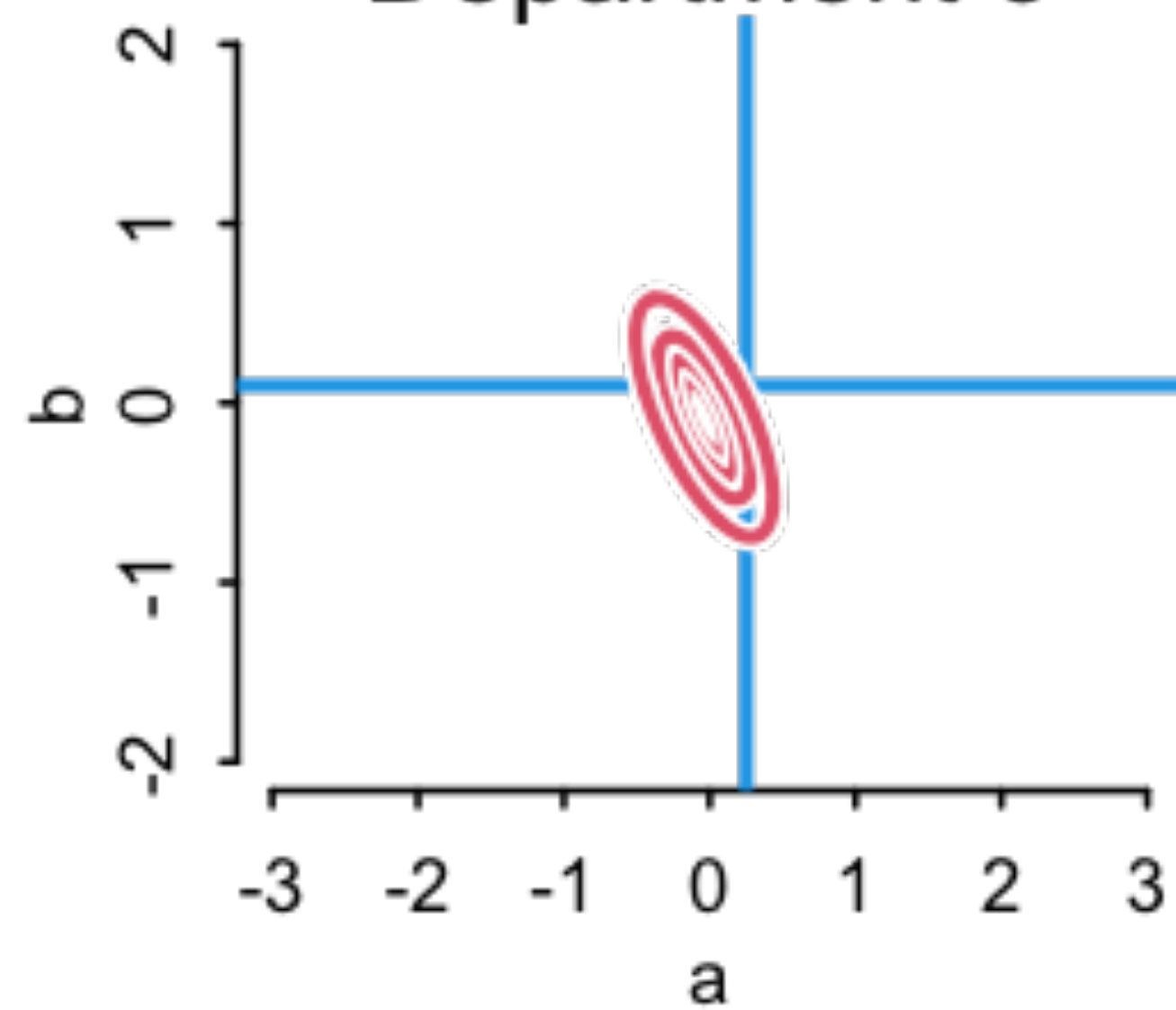
Department 1



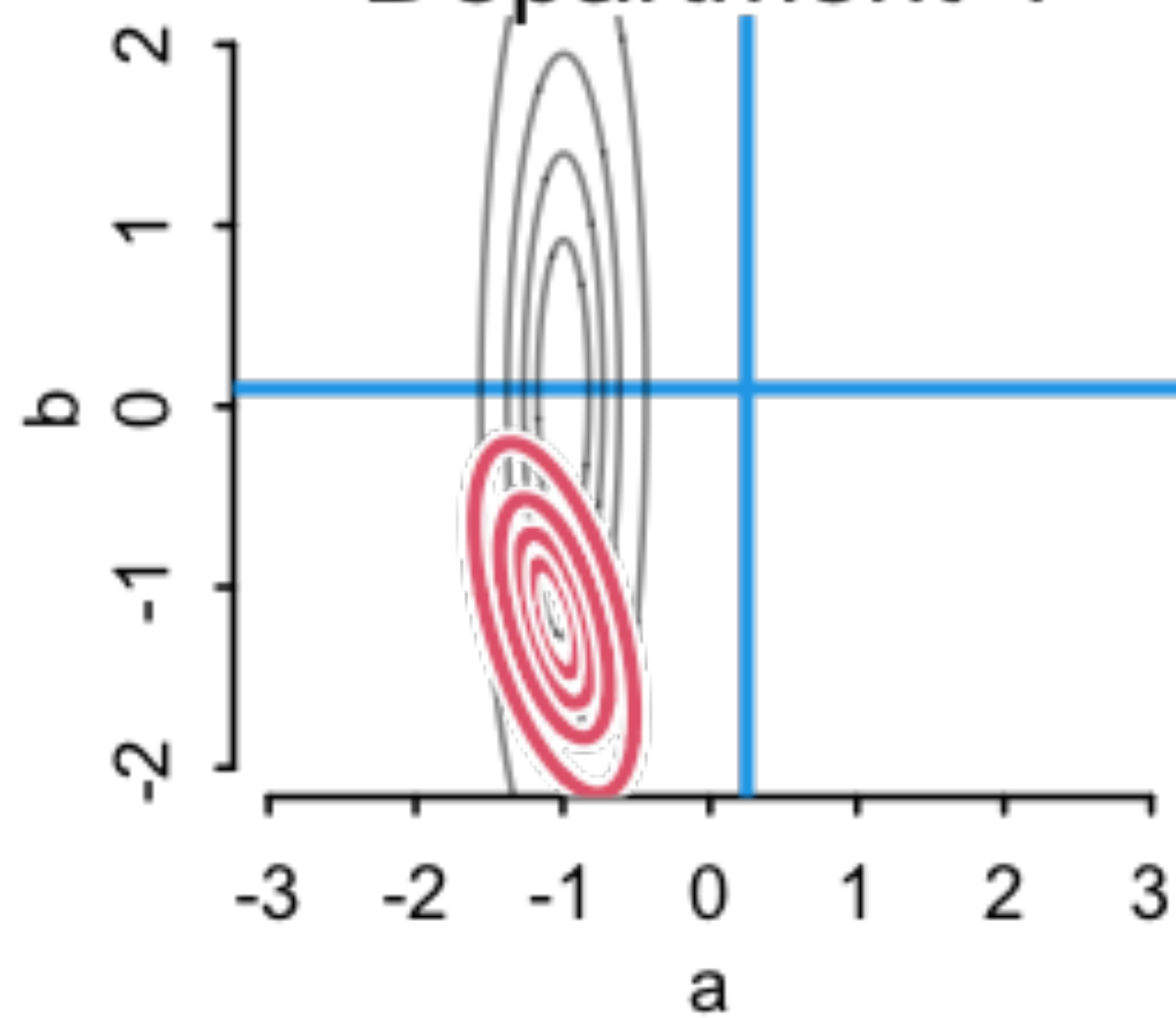
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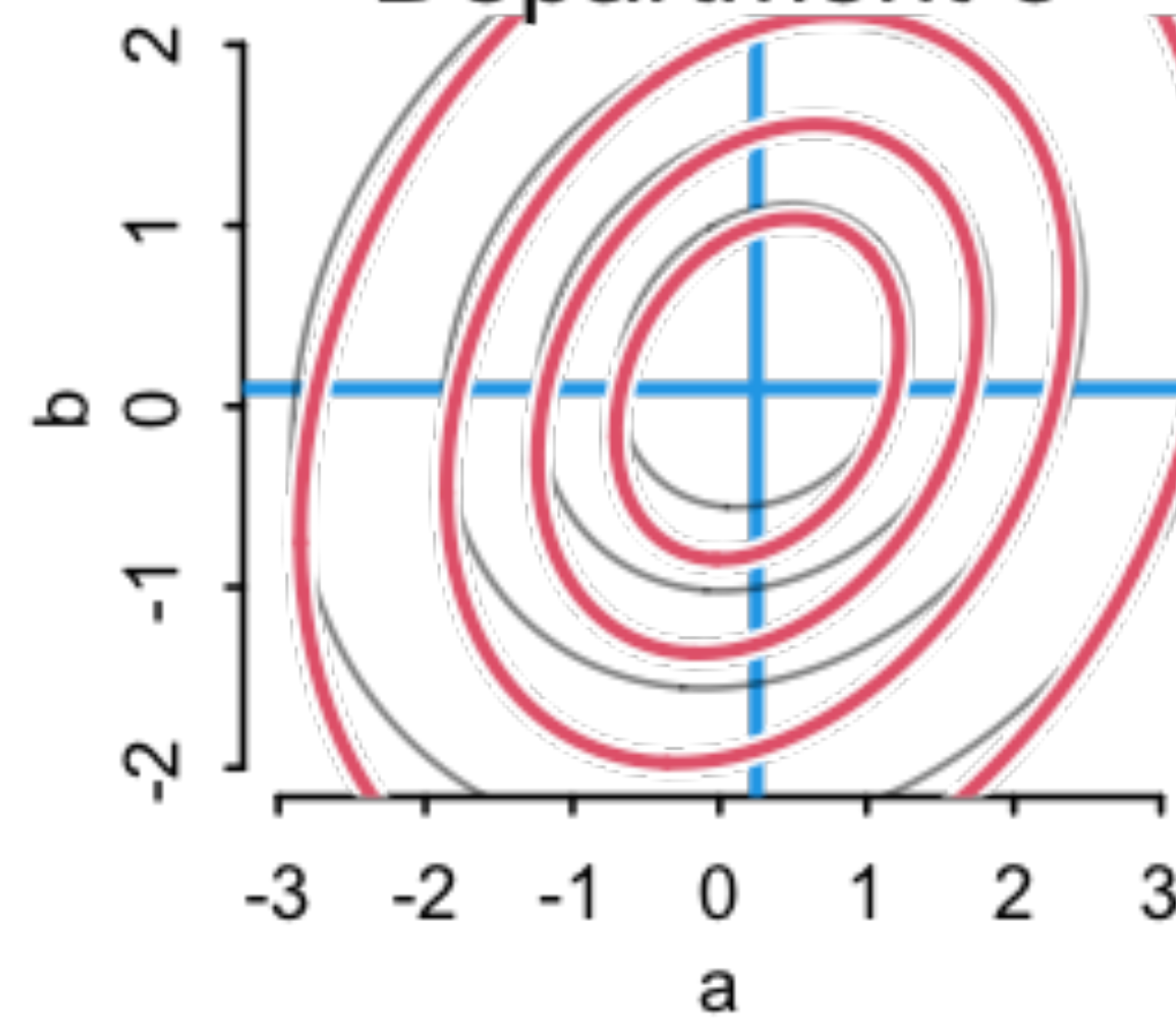
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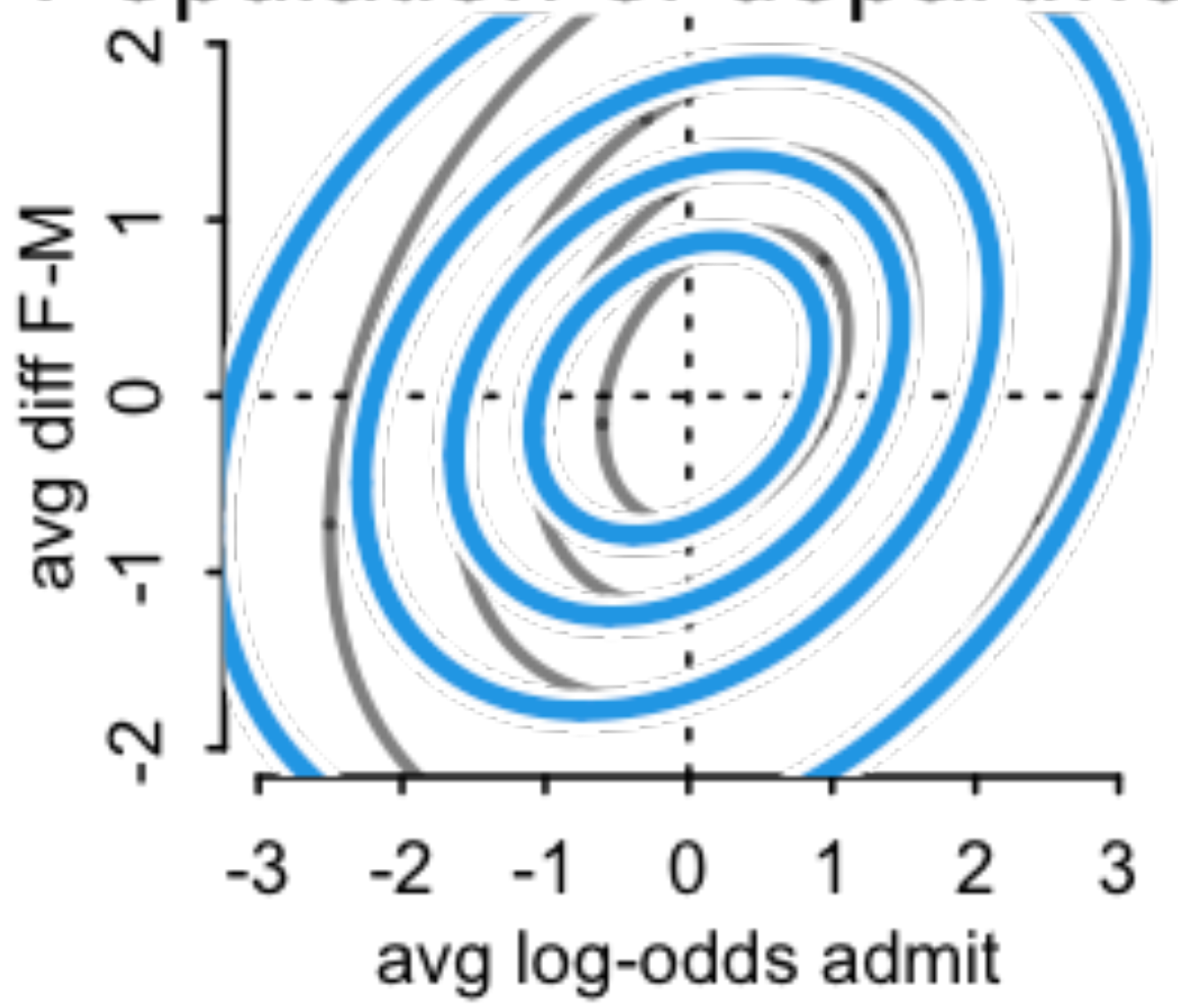
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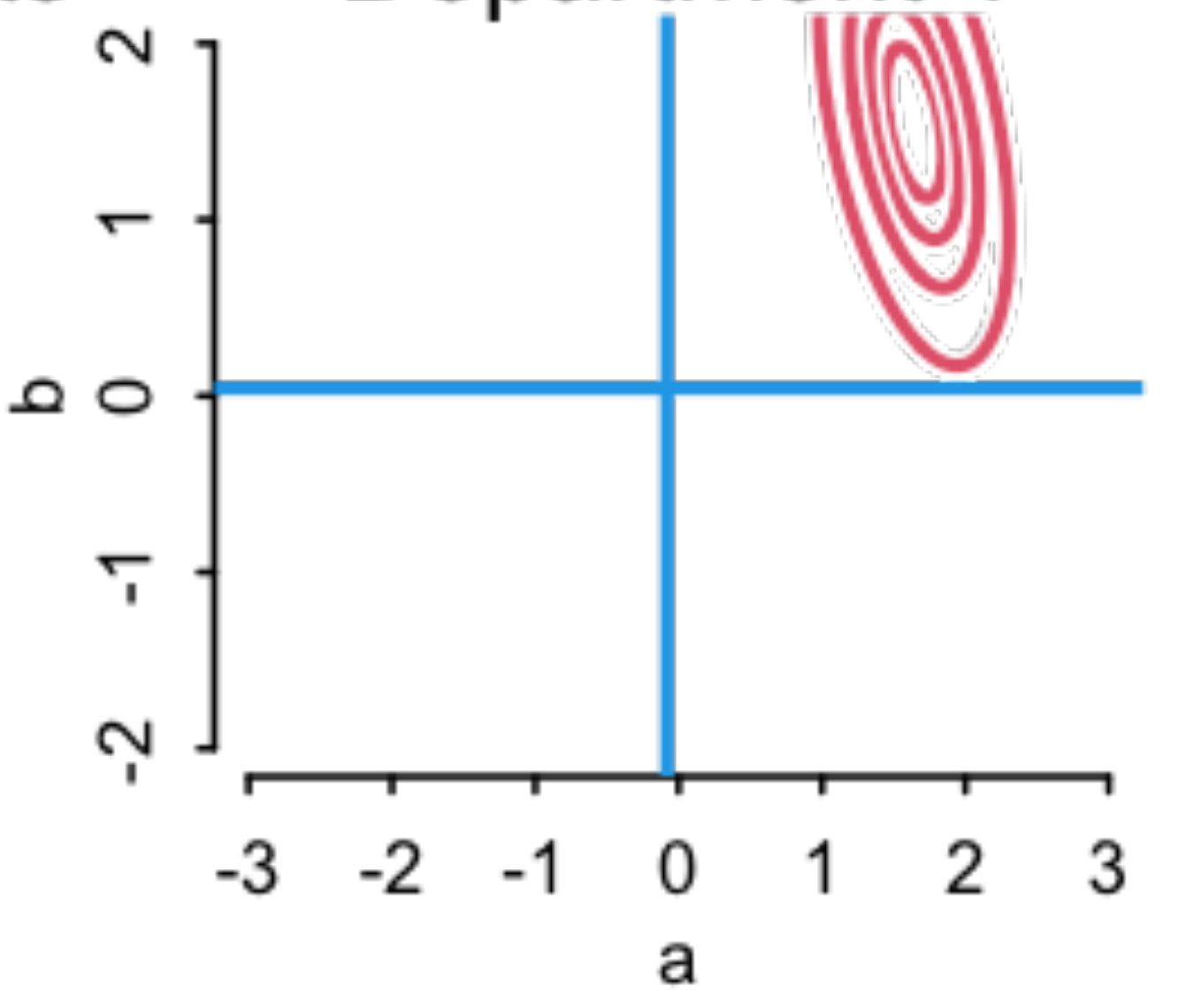
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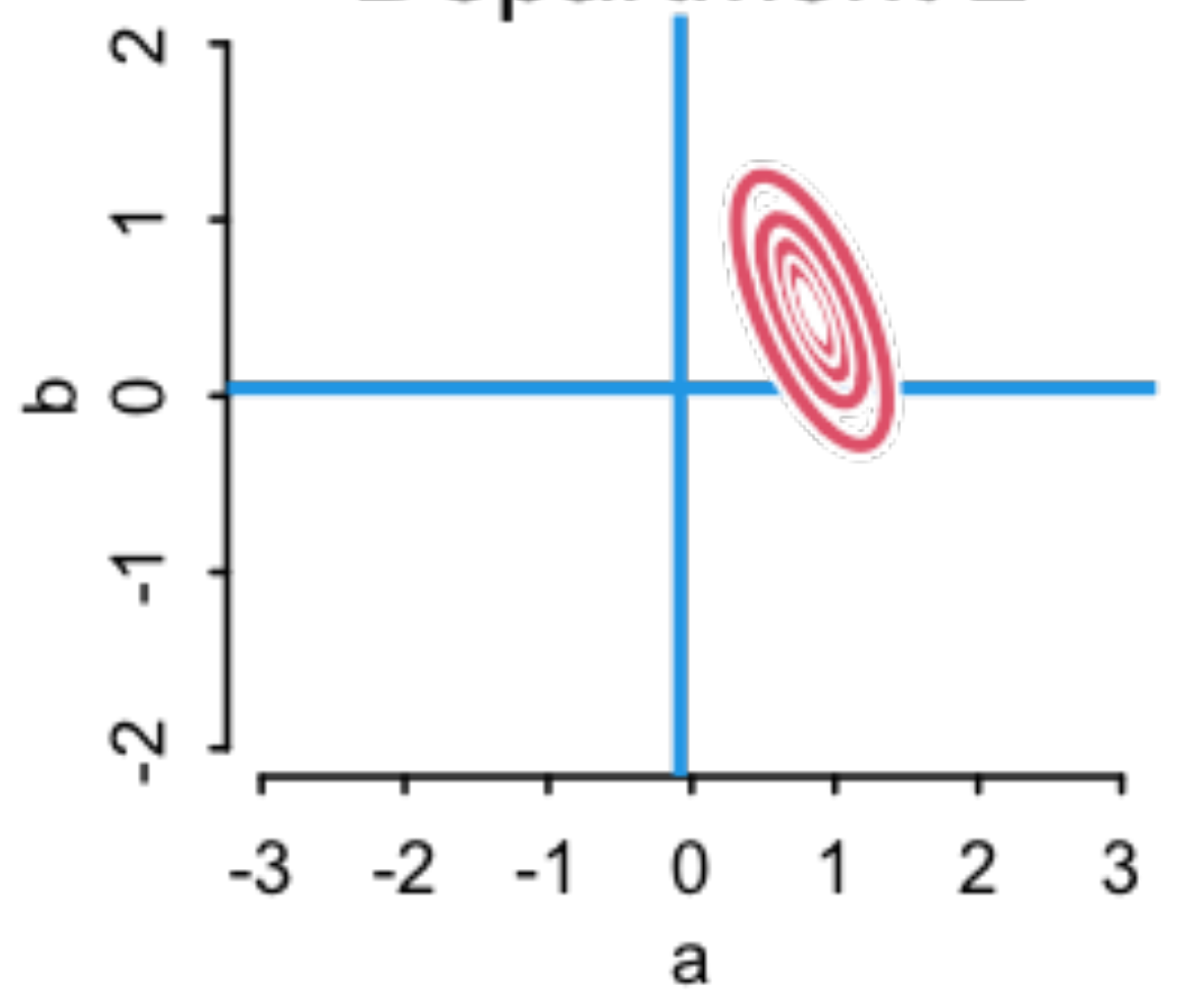
Population of departments



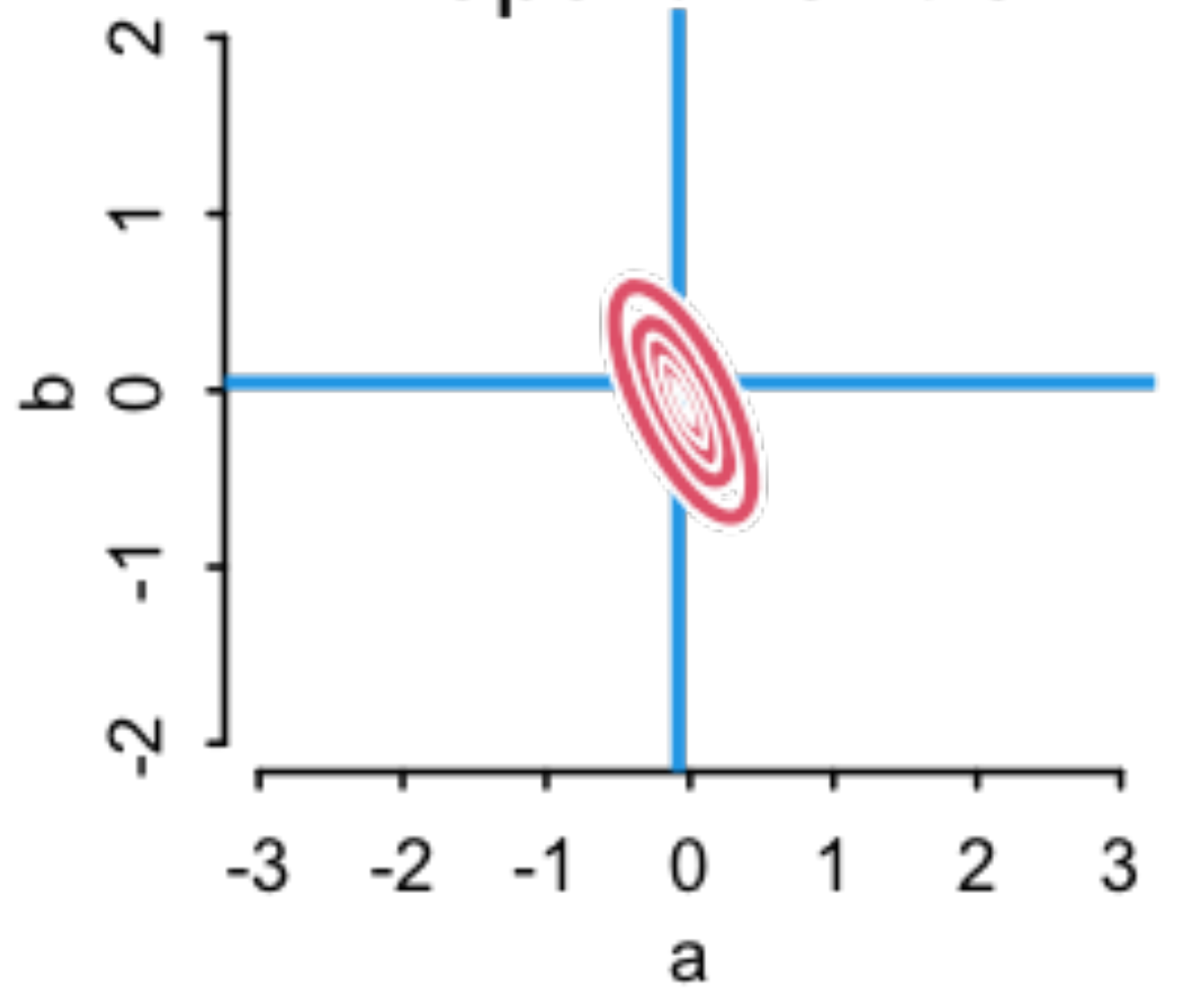
Department 1



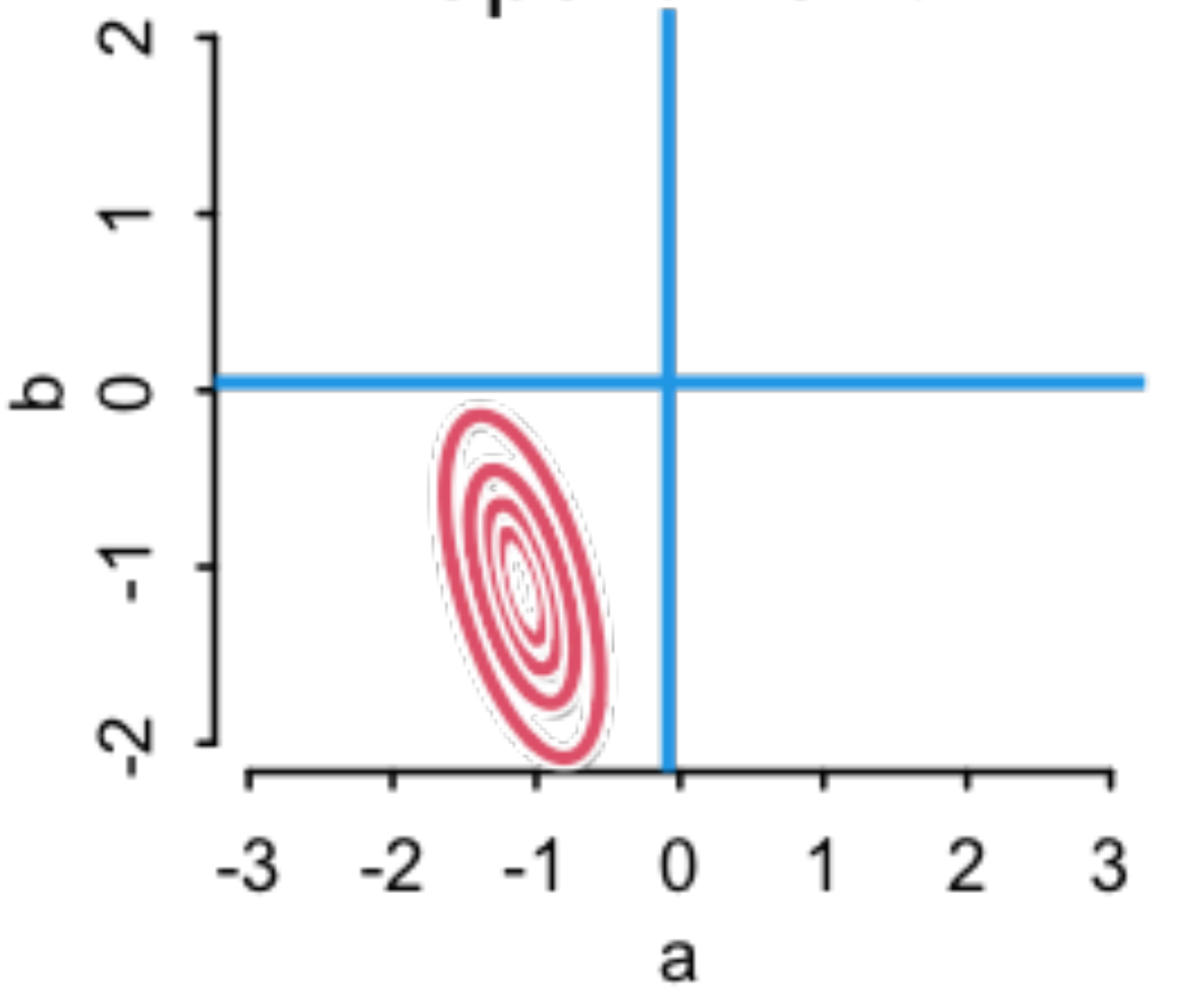
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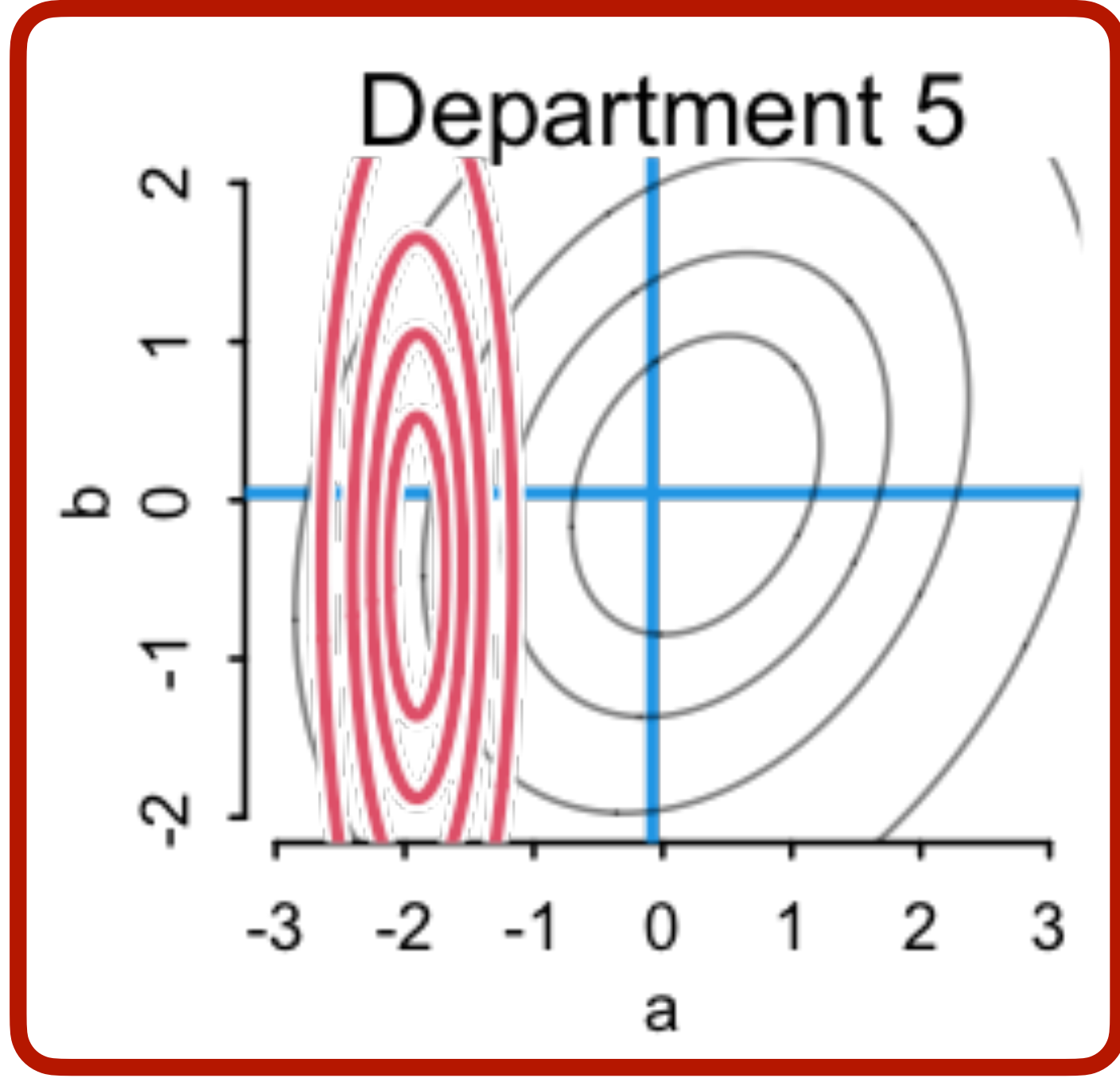
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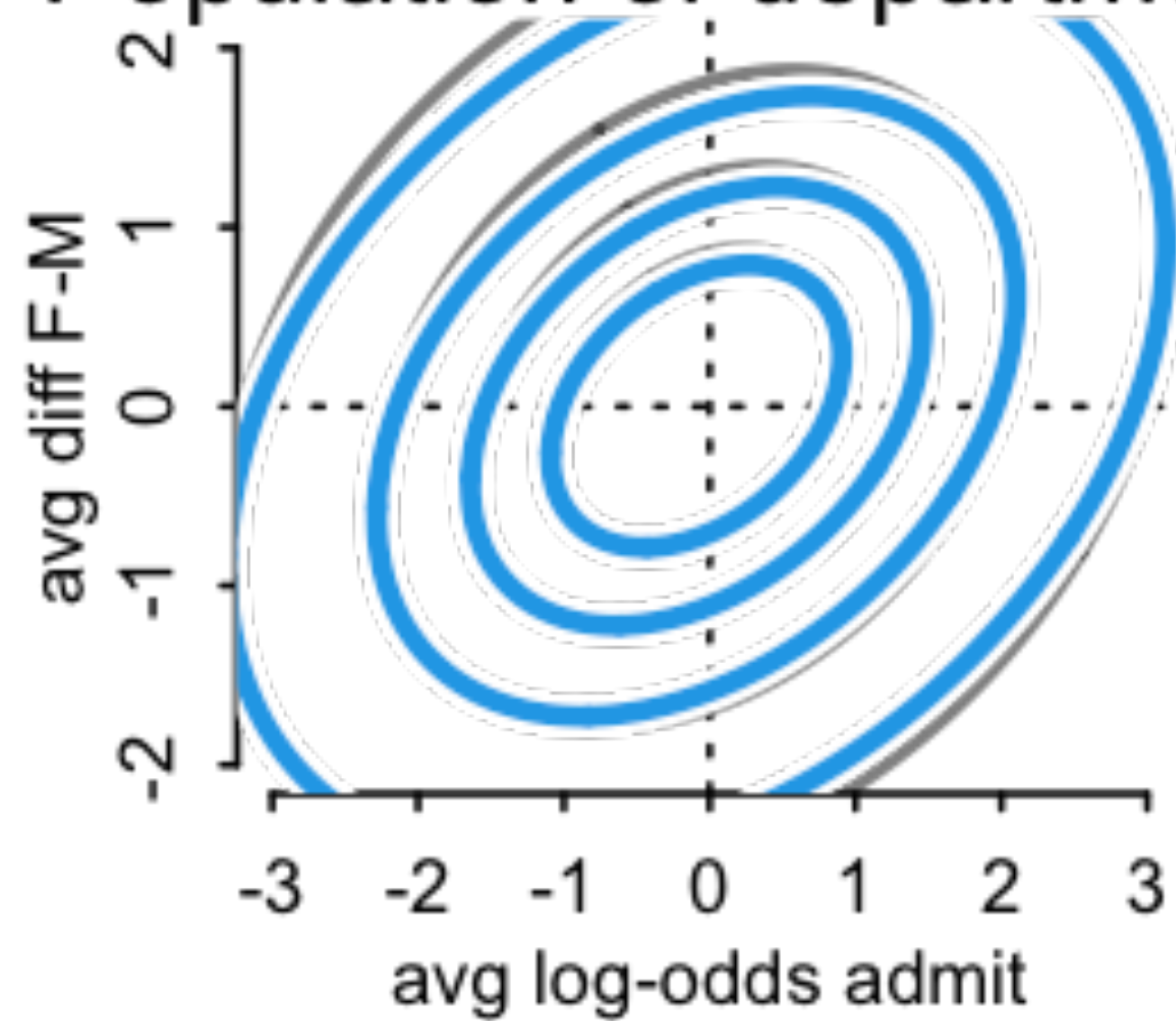
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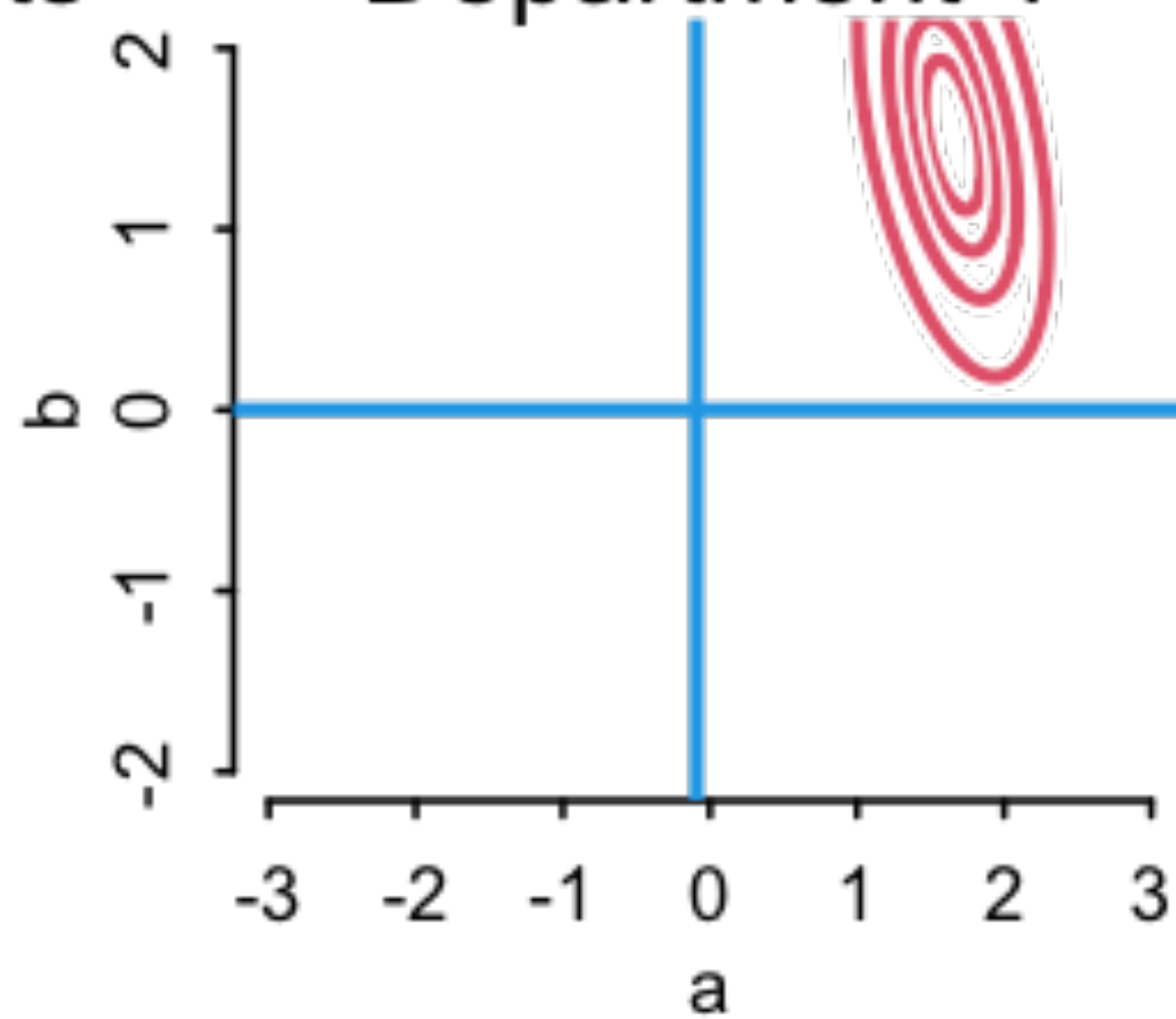
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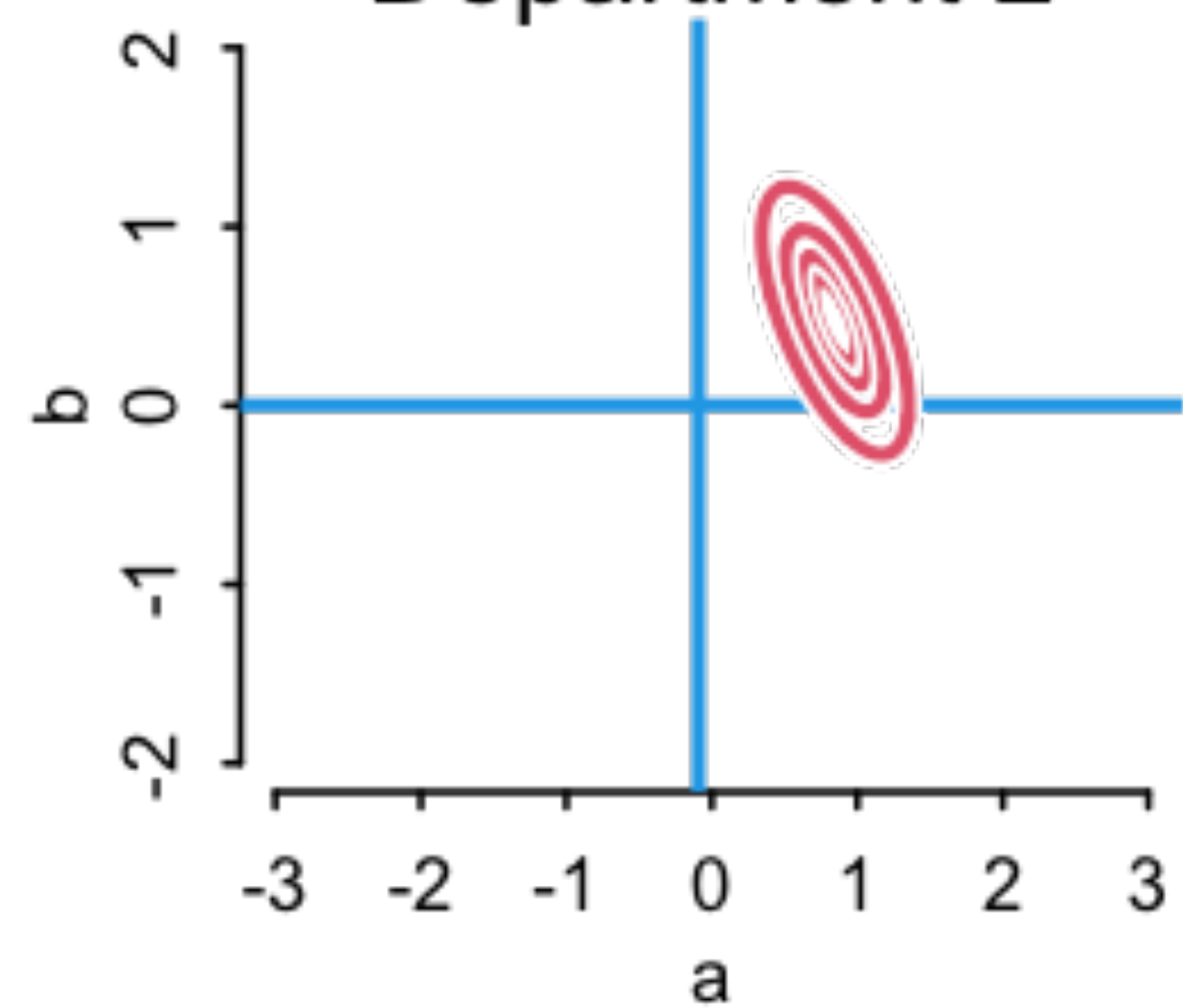
Population of departments



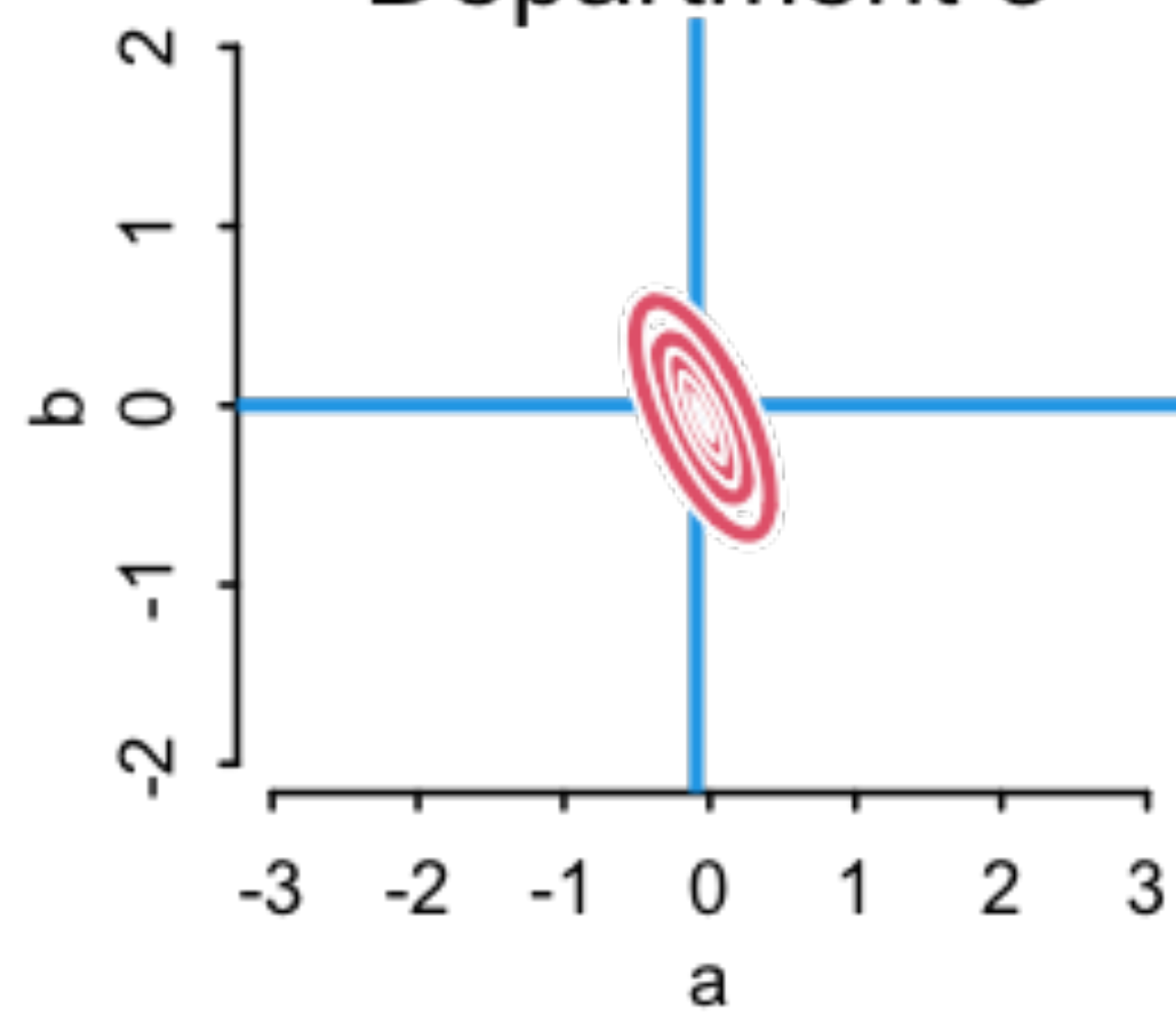
Department 1



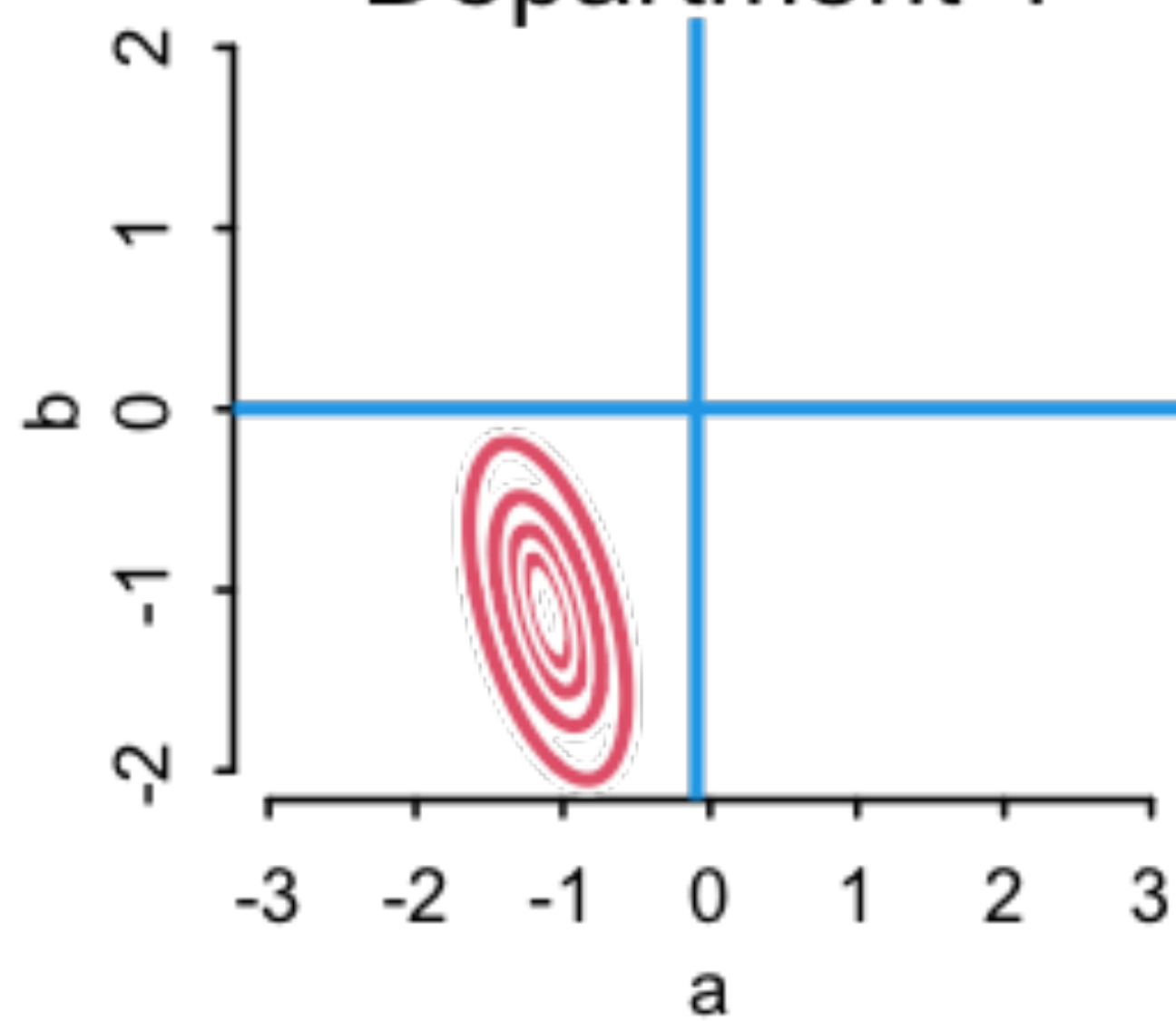
Department 2



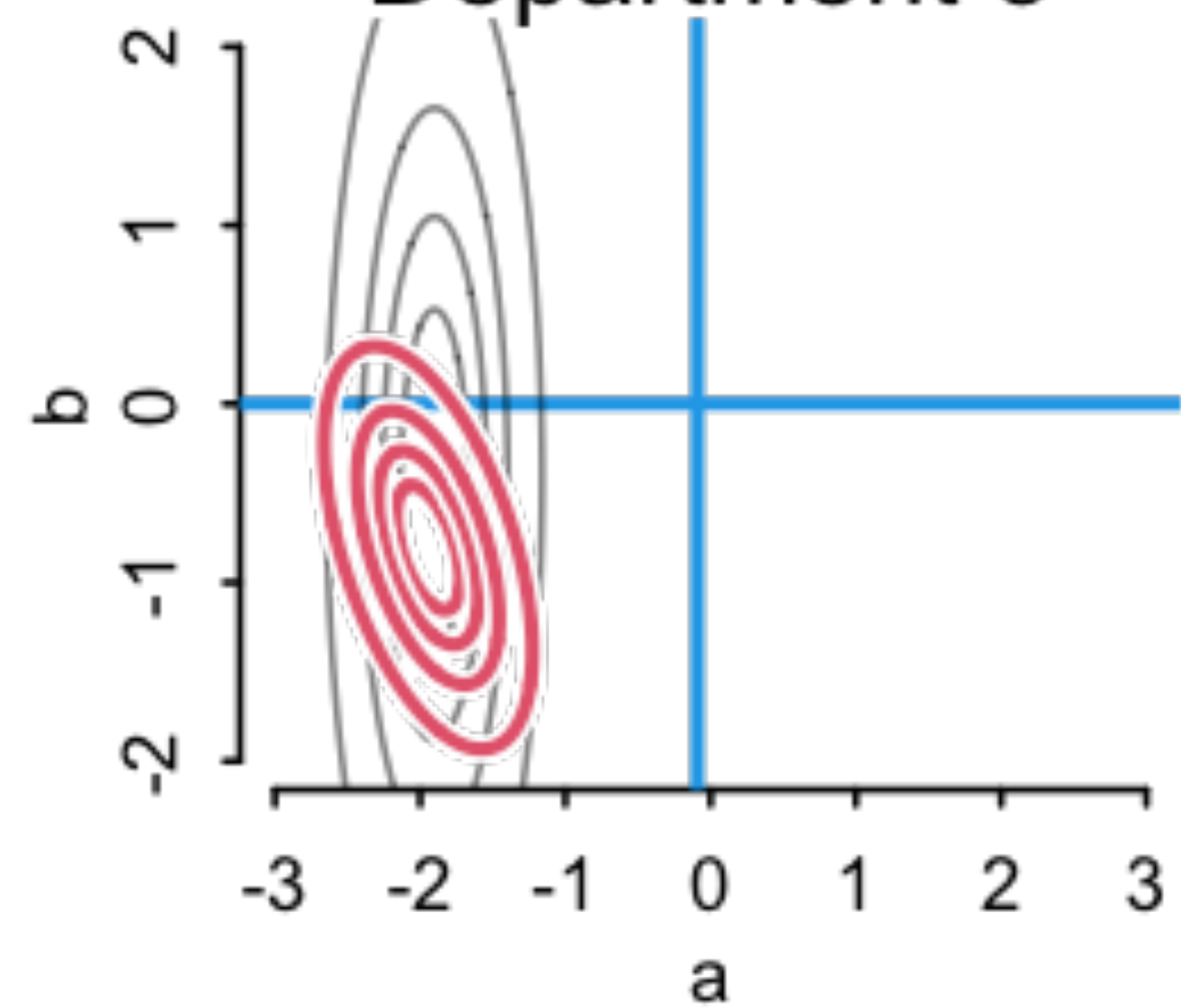
Department 3



Department 4



Department 5

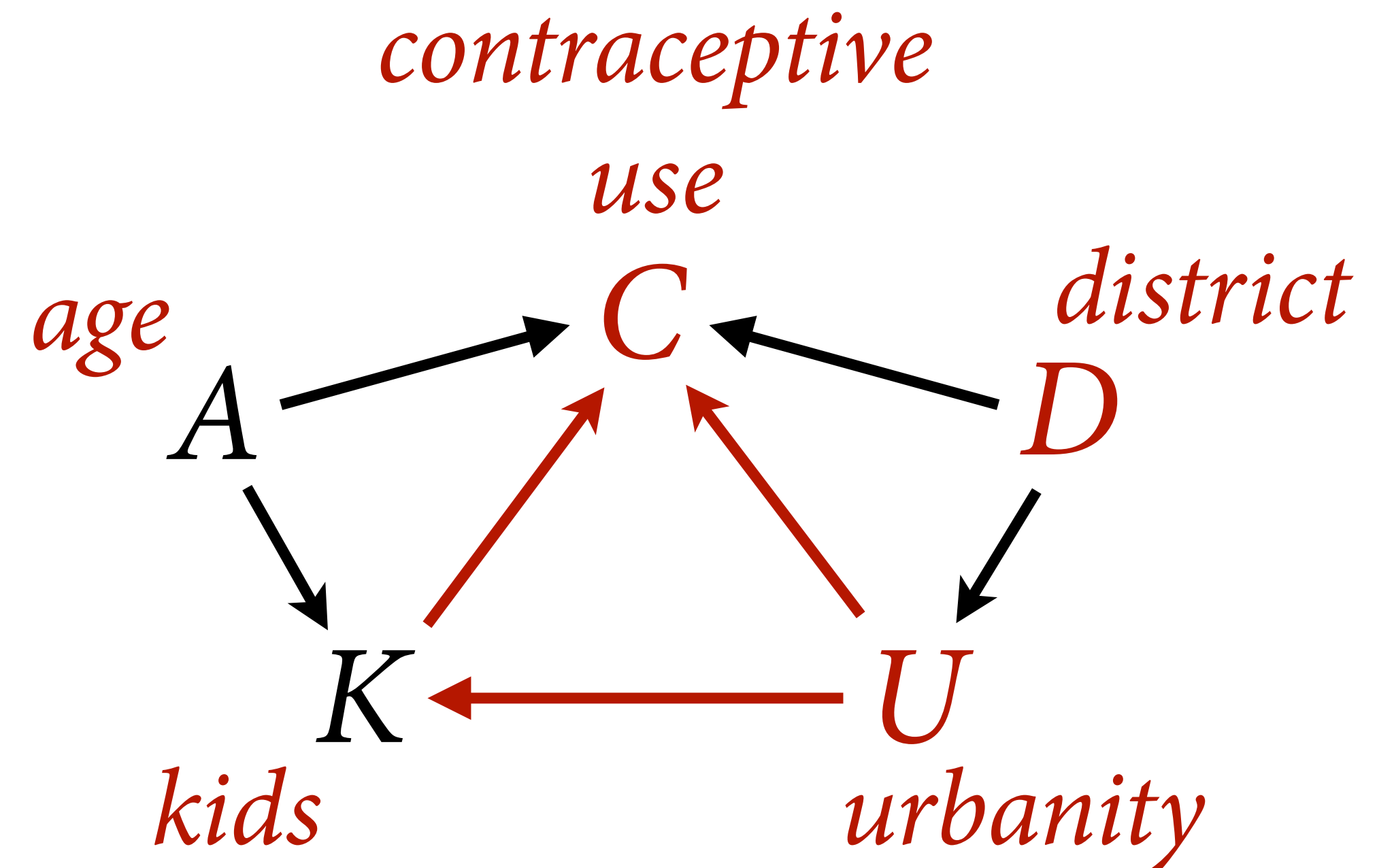


Bangladesh 1989, Party Like It's

Estimand 1: C in each district

Estimand 2: Effect of U

Estimand 3: Effects of K and A



```

mCDUnc <- ulam(
  alist(
    C ~ bernoulli(p),
    logit(p) <- a[D] + b[D]*U,
    # define effects using other parameters
    save> vector[61]:a <<- abar + za*sigma,
    save> vector[61]:b <<- bbar + zb*tau,
    # z-scored effects
    vector[61]:za ~ normal(0,1),
    vector[61]:zb ~ normal(0,1),
    # ye olde hyper-priors
    c(abar,bbar) ~ normal(0,1),
    c(sigma,tau) ~ exponential(1)
  ) , data=dat , chains=4 , cores=4 )

```

$$C_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]}U_i$$

$$\alpha_j = \bar{\alpha} + z_{\alpha,j} \times \sigma$$

$$\beta_j = \bar{\beta} + z_{\beta,j} \times \tau$$

$$z_{\alpha,j} \sim \text{Normal}(0,1)$$

$$z_{\beta,j} \sim \text{Normal}(0,1)$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0,1)$$

$$\sigma, \tau \sim \text{Exponential}(1)$$

```

dat <- list(
  C = d$use.contraception,
  D = as.integer(d$district),
  U = ifelse(d$urban==1,1,0) )

# total U
mCDU <- ulam(
  alist(
    C ~ bernoulli(p),
    logit(p) <- a[D] + b[D]*U,
    vector[61]:a ~ normal(abar,sigma),
    vector[61]:b ~ normal(bbar,tau),
    c(abar,bbar) ~ normal(0,1),
    c(sigma,tau) ~ exponential(1)
  ) , data=dat , chains=4 , cores=4 )

```

$$C_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]}U_i$$

$$\alpha_j \sim \text{Normal}(\bar{\alpha}, \sigma)$$

$$\beta_j \sim \text{Normal}(\bar{\beta}, \tau)$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0,1)$$

$$\sigma, \tau \sim \text{Exponential}(1)$$

$$C_i \sim \text{Bernoulli}(p_i)$$

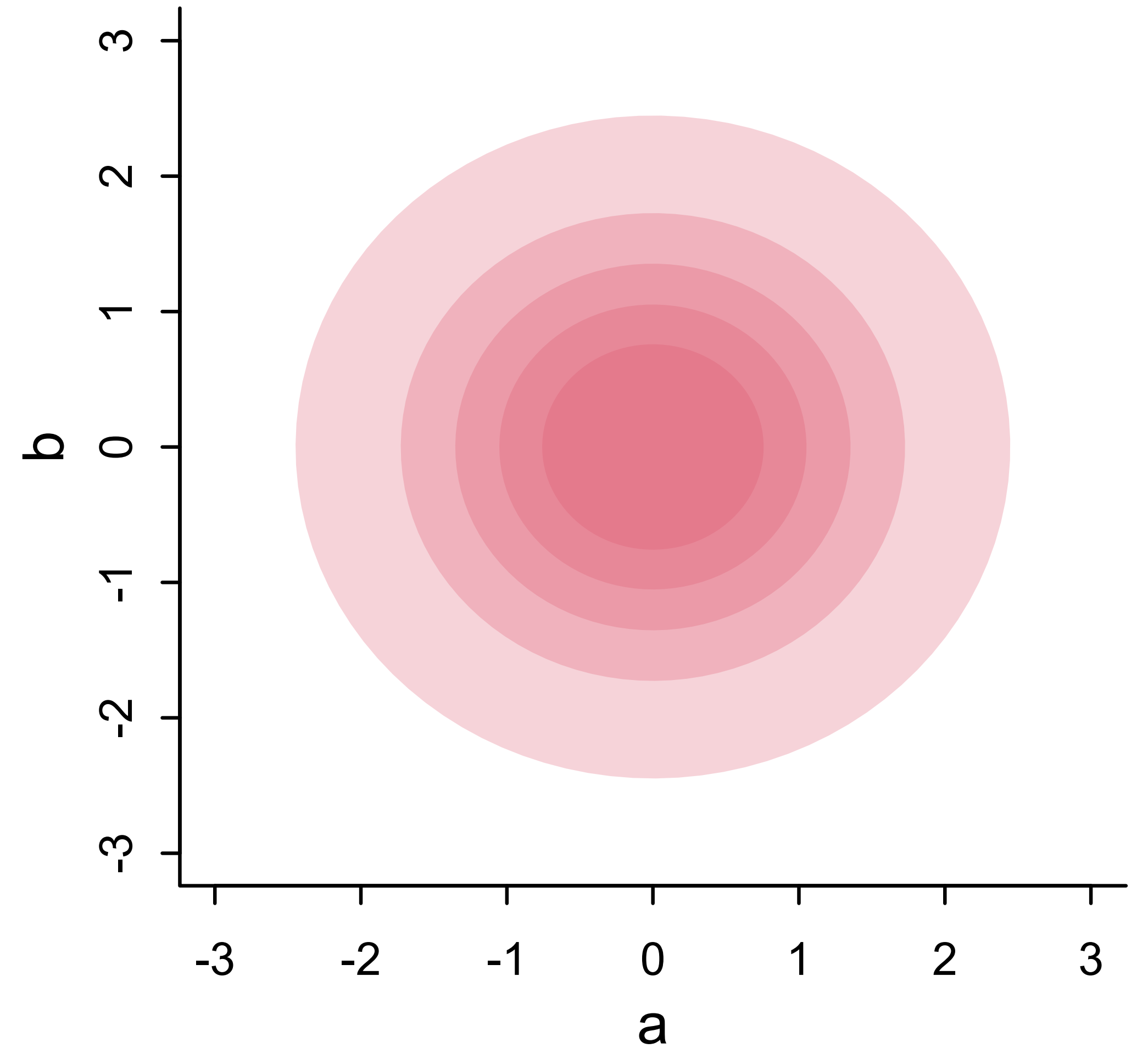
$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]} U_i$$

$$\alpha_j \sim \text{Normal}(\bar{\alpha}, \sigma)$$

$$\beta_j \sim \text{Normal}(\bar{\beta}, \tau)$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0, 1)$$

$$\sigma, \tau \sim \text{Exponential}(1)$$



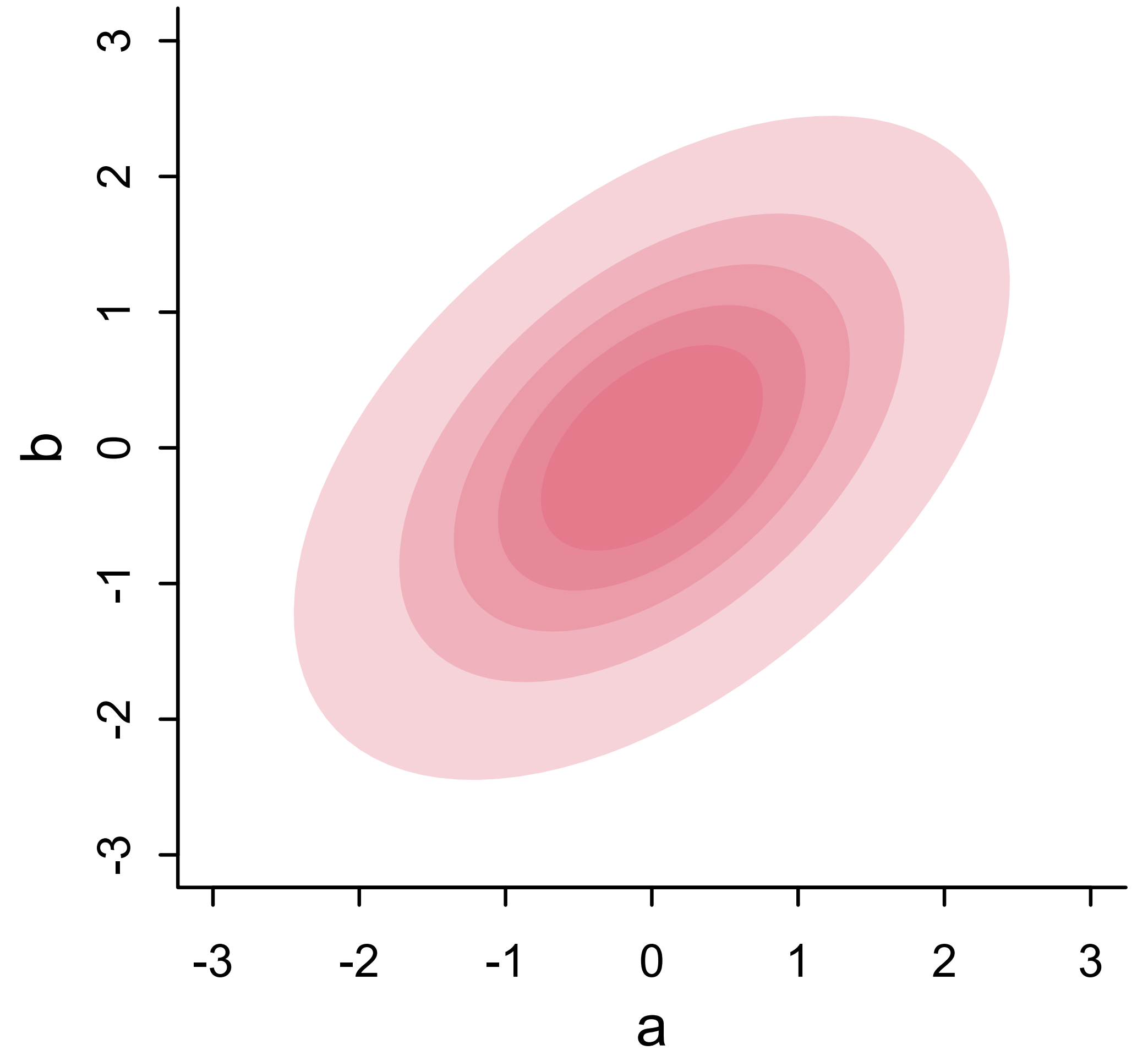
$$C_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]} U_i$$

$$[\alpha_j, \beta_j] \sim \text{MVNormal}([\bar{\alpha}, \bar{\beta}], \mathbf{R}, [\sigma, \tau])$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0,1)$$

$$\sigma, \tau \sim \text{Exponential}(1)$$



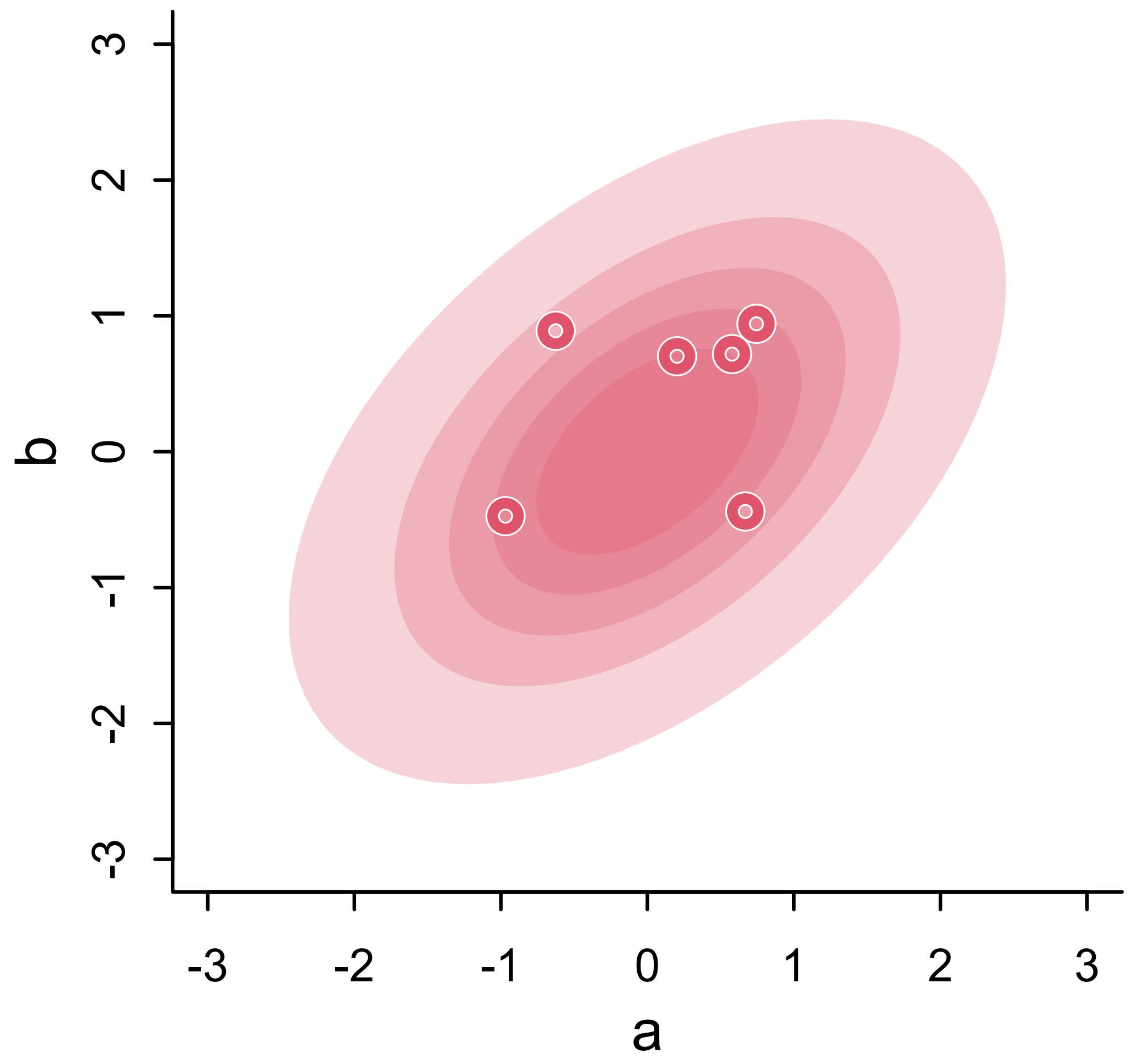
*features for
district j*

*correlation
matrix*

$$[\alpha_j, \beta_j] \sim \text{MVNormal}([\bar{\alpha}, \bar{\beta}], R, [\sigma, \tau])$$

*feature
means*

*standard
deviations*



$$C_i \sim \text{Bernoulli}(p_i)$$

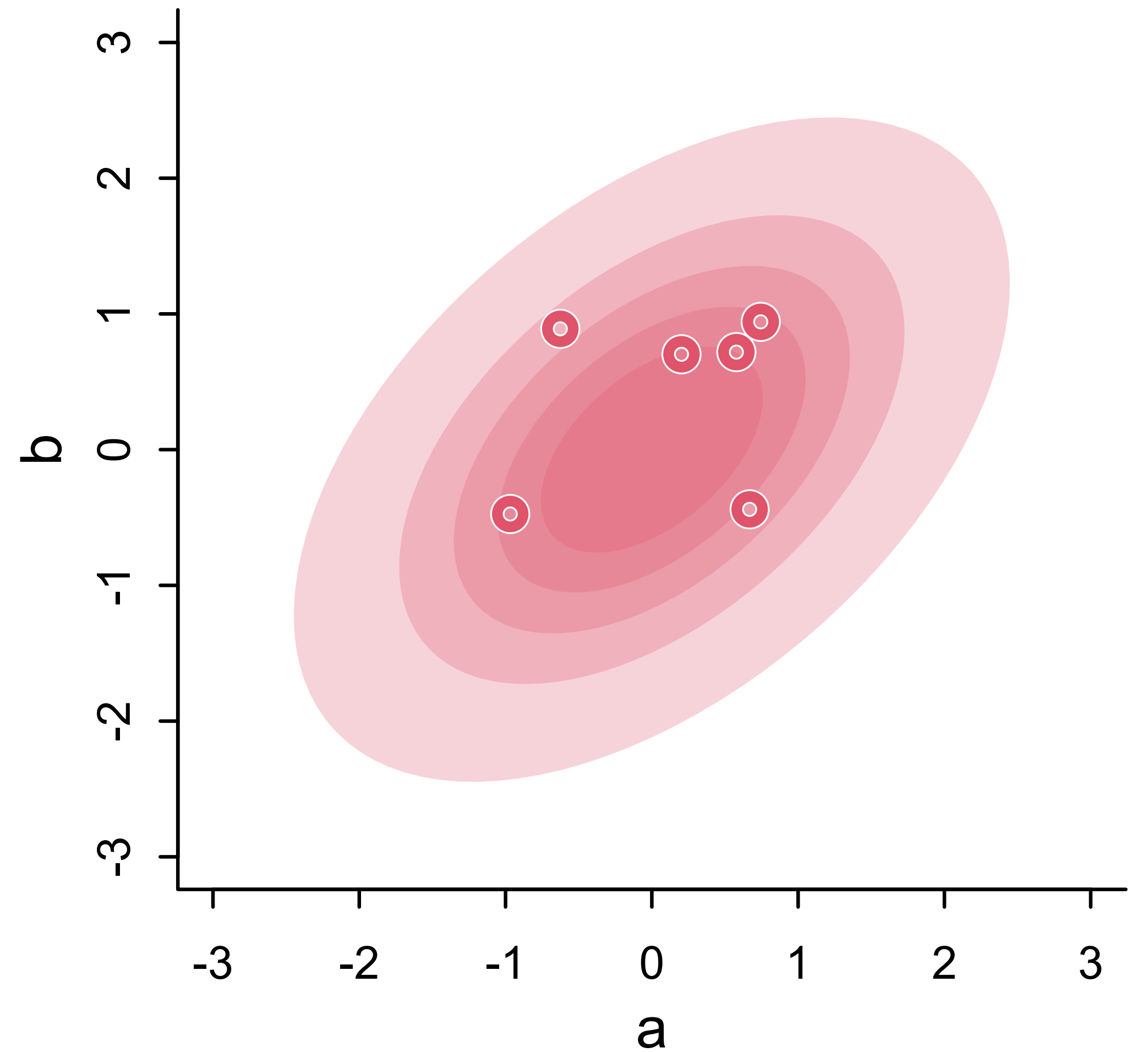
$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]} U_i$$

$$[\alpha_j, \beta_j] \sim \text{MVNormal}([\bar{\alpha}, \bar{\beta}], \mathbf{R}, [\sigma, \tau])$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0, 1)$$

$$\sigma, \tau \sim \text{Exponential}(1)$$

$$\mathbf{R} \sim ?$$



$$C_i \sim \text{Bernoulli}(p_i)$$

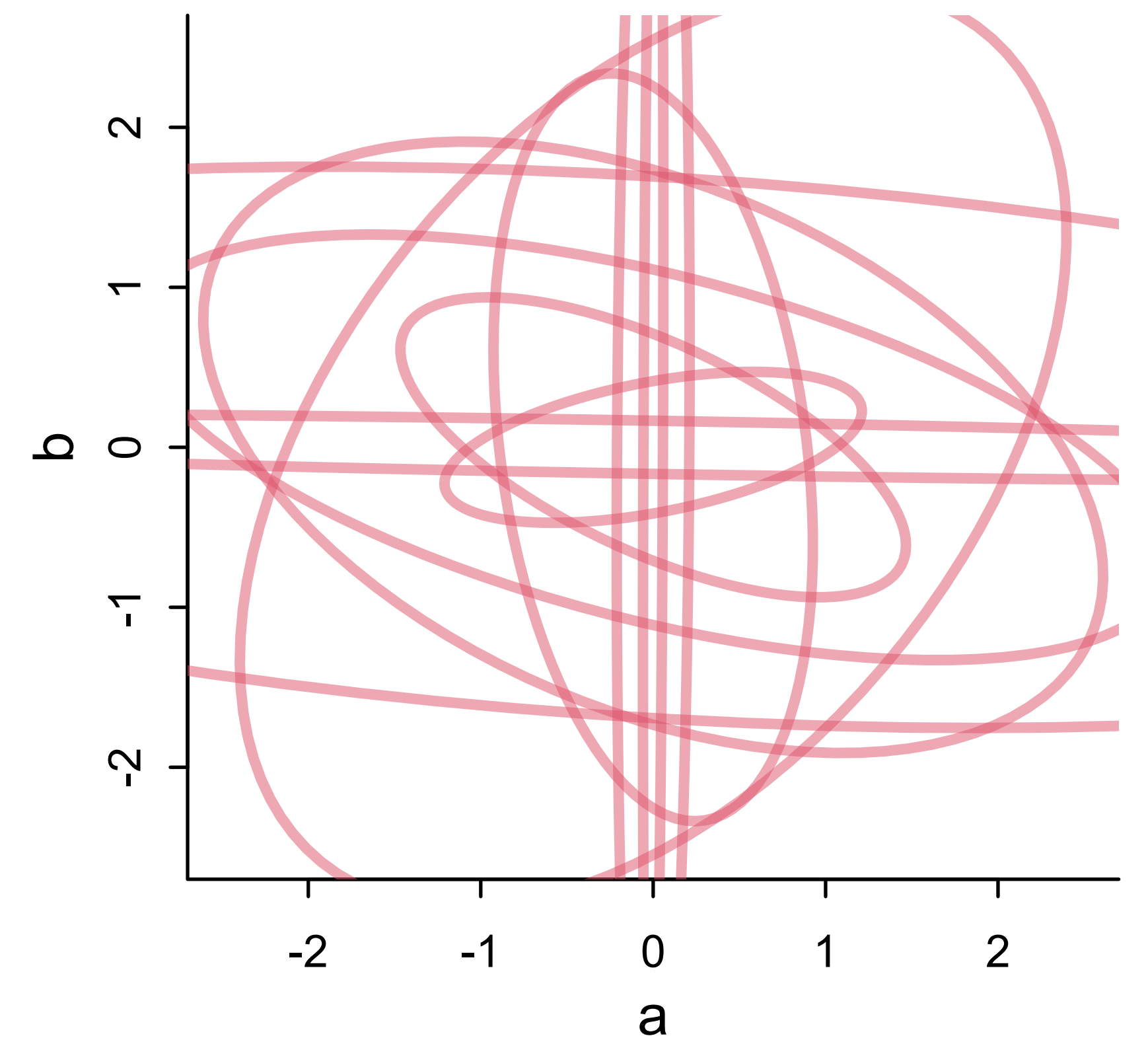
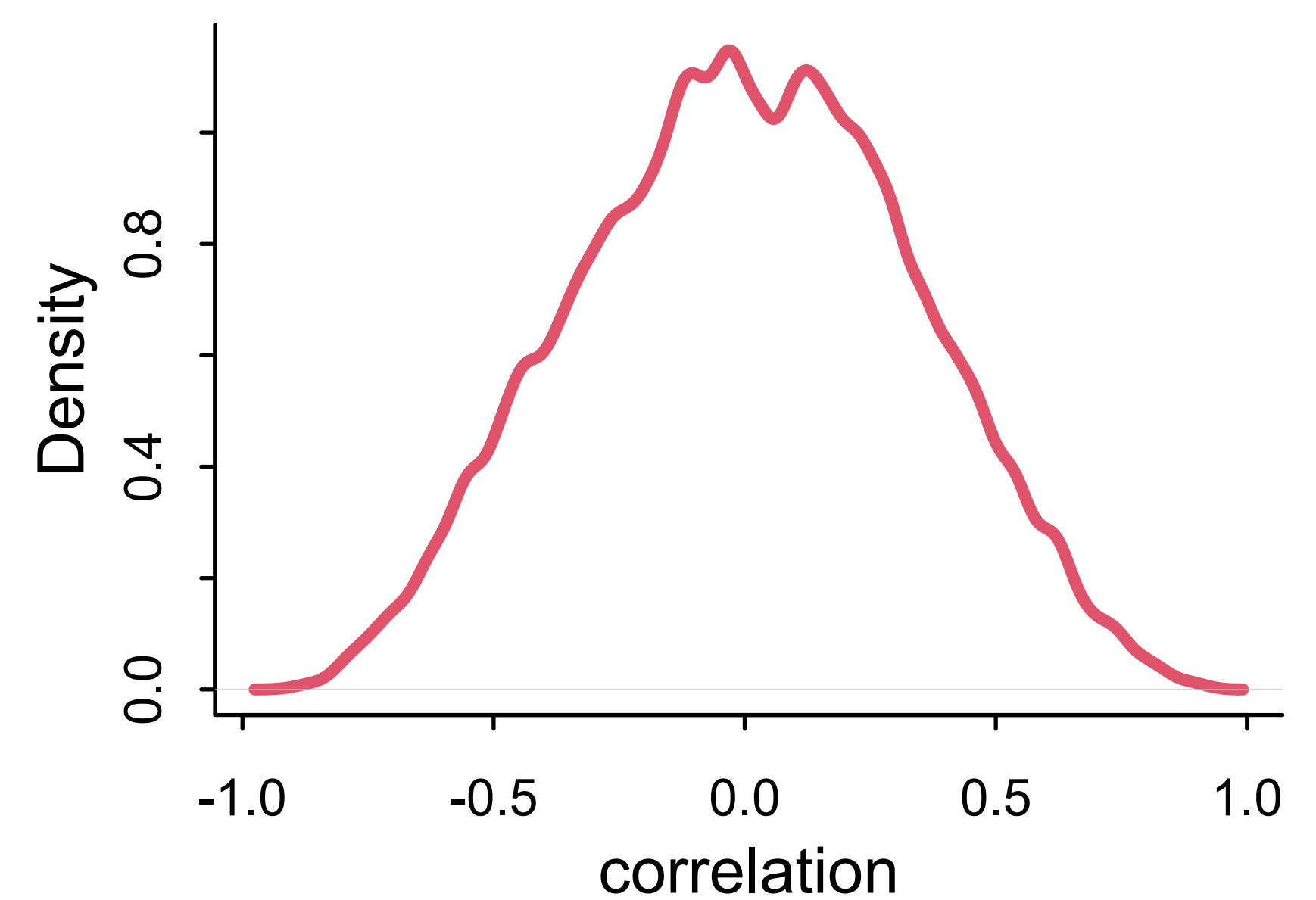
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$$\mathbf{R} \sim \text{LKJCorr}(4)$$



PAUSE

```

mCDUcov <- ulam(
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    matrix[61,2]:v ~ multi_normal(a_bar, Rho, sigma),
    vector[2]:a_bar ~ normal(0,1),
    corr_matrix[2]:Rho ~ lkj_corr(4),
    vector[2]:sigma ~ exponential(1)
  ), data=dat , chains=4 , cores=4 )

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    matrix[61,2]:v ~ multi_normal(abar, Rho, sigma),
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    corr_matrix[2]:Rho ~ lkj_corr(4),
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61 districts, 2 features each

```

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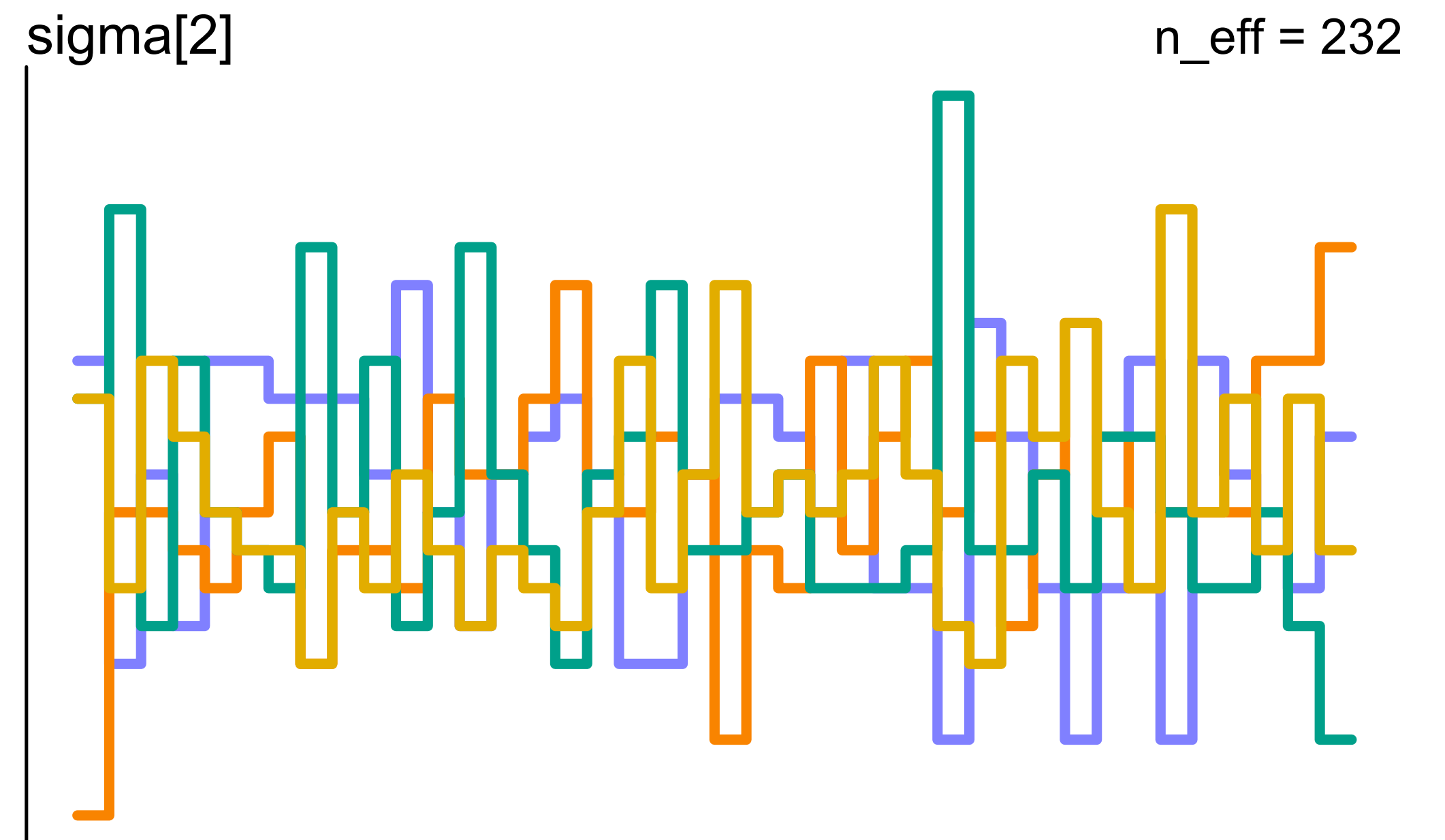
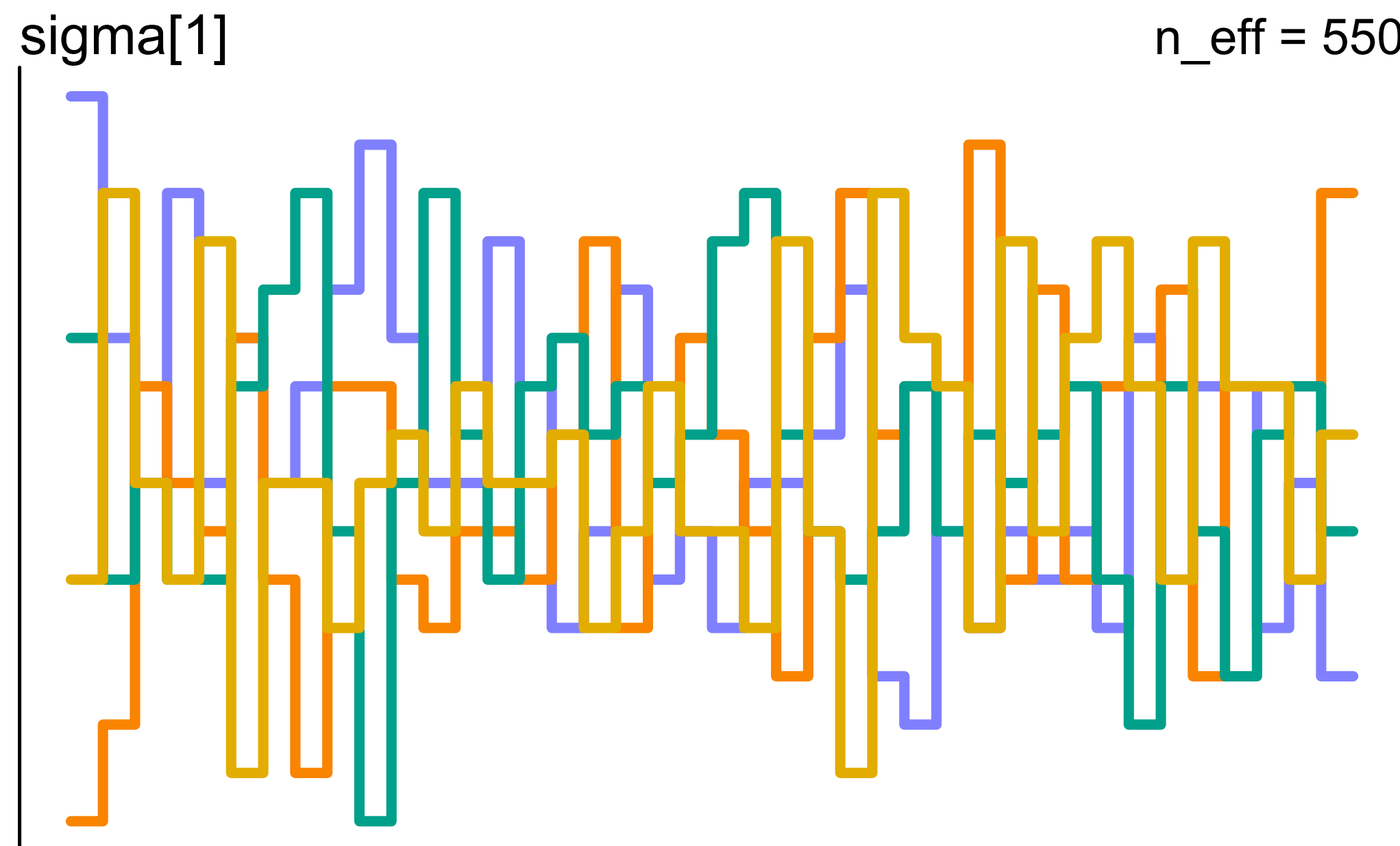
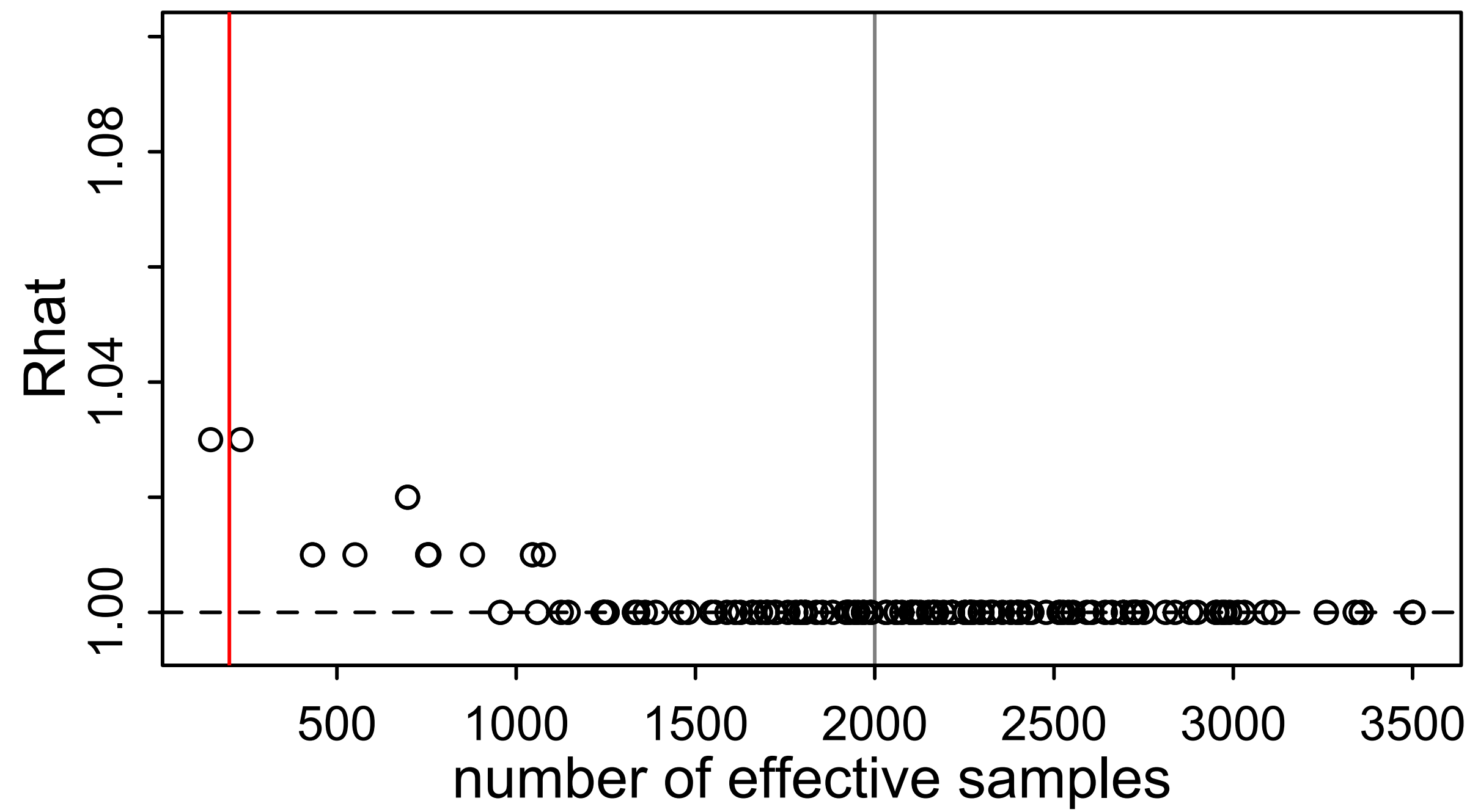
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Non-centered priors

$$C_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]} U_i$$

$$\alpha = \bar{\alpha} + v_{-,1}$$

$$\beta = \bar{\beta} + v_{-,2}$$

$$v = (\text{diag}(\sigma) \mathbf{LZ})^\top$$

$$\mathbf{Z}_{j,k} \sim \text{Normal}(0, 1)$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0, 1)$$

$$\sigma \sim \text{Exponential}(1)$$

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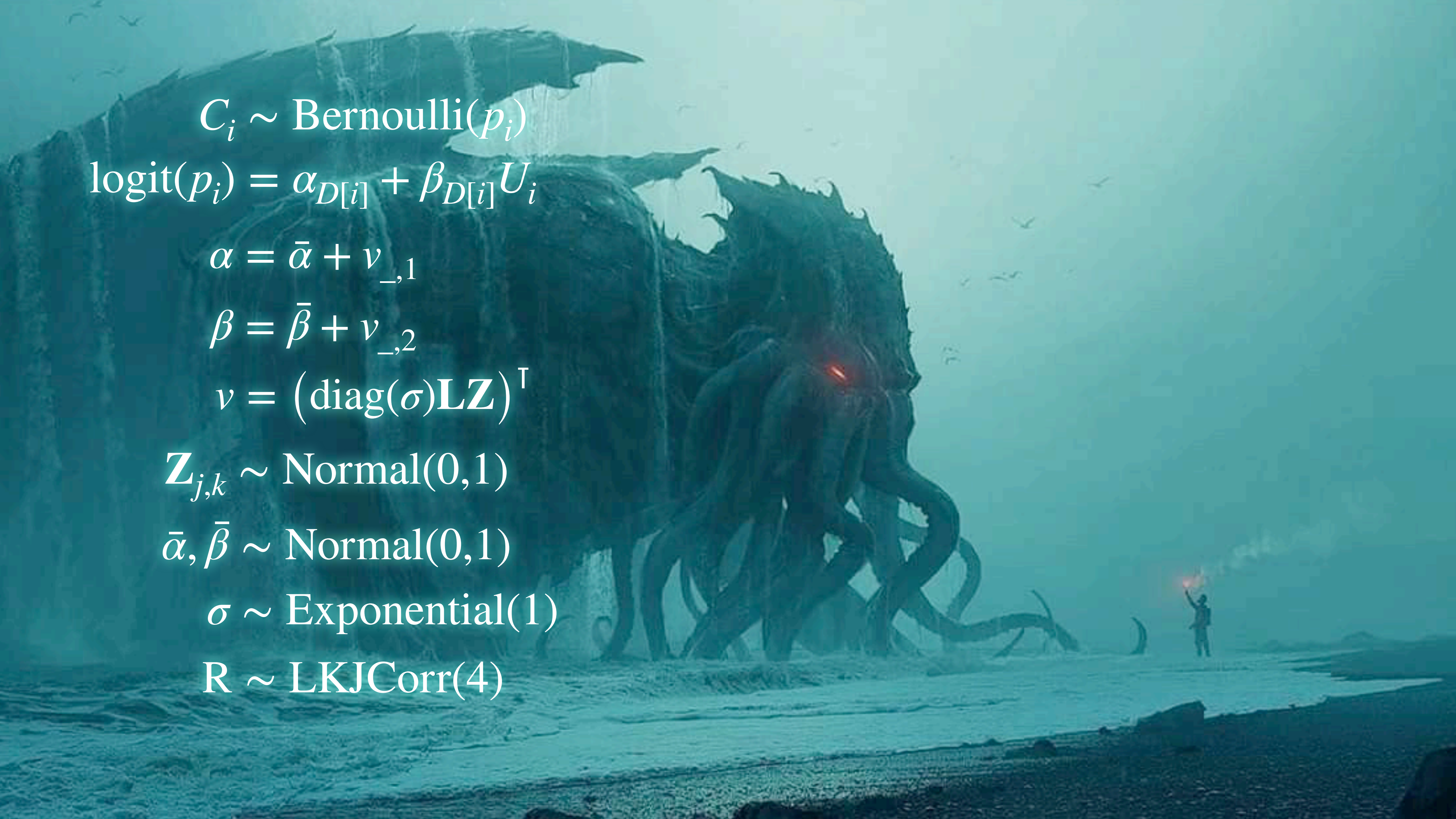
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```

# covariance - non-centered
mCDUcov_nc <- ulam(
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    # define effects using other parameters
    # this is the non-centered Cholesky machine
    transpars> vector[61]:a <<- abar[1] + v[,1],
    transpars> vector[61]:b <<- abar[2] + v[,2],
    transpars> matrix[61,2]:v <-
      compose_noncentered( sigma , L_Rho , Z ),

    # priors - note that none have parameters inside them
    # that is what makes them non-centered
    matrix[2,61]:Z ~ normal( 0 , 1 ),
    vector[2]:abar ~ normal(0,1),
    cholesky_factor_corr[2]:L_Rho ~ lkj_corr_cholesky( 4 ),
    vector[2]:sigma ~ exponential(1),

    # convert Cholesky to Corr matrix
    gq> matrix[2,2]:Rho <<- Chol_to_Corr(L_Rho)
  ) , data=dat , chains=4 , cores=4 )

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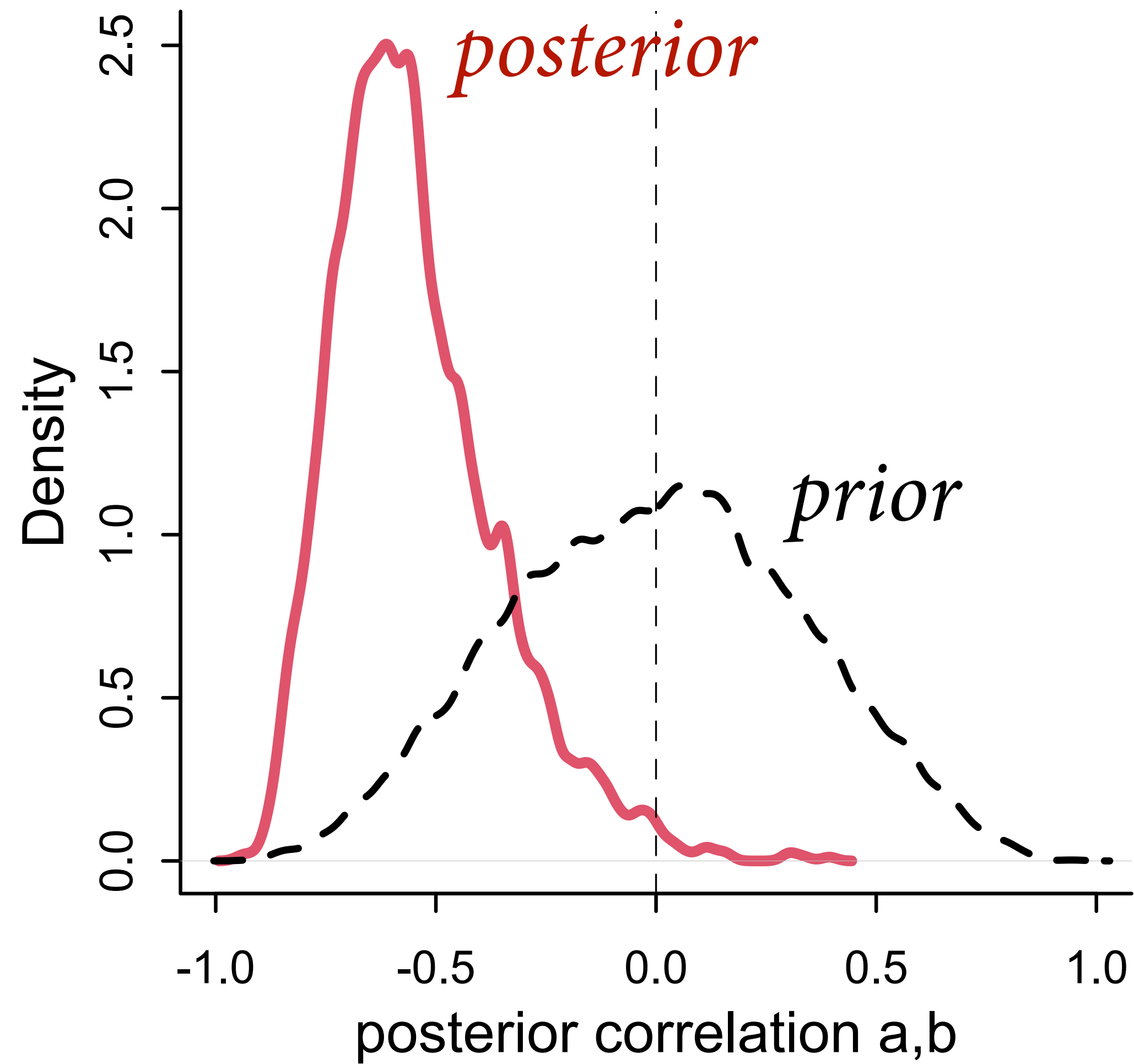
$$\mathbf{Z}_{j,k} \sim \text{Normal}(0,1)$$

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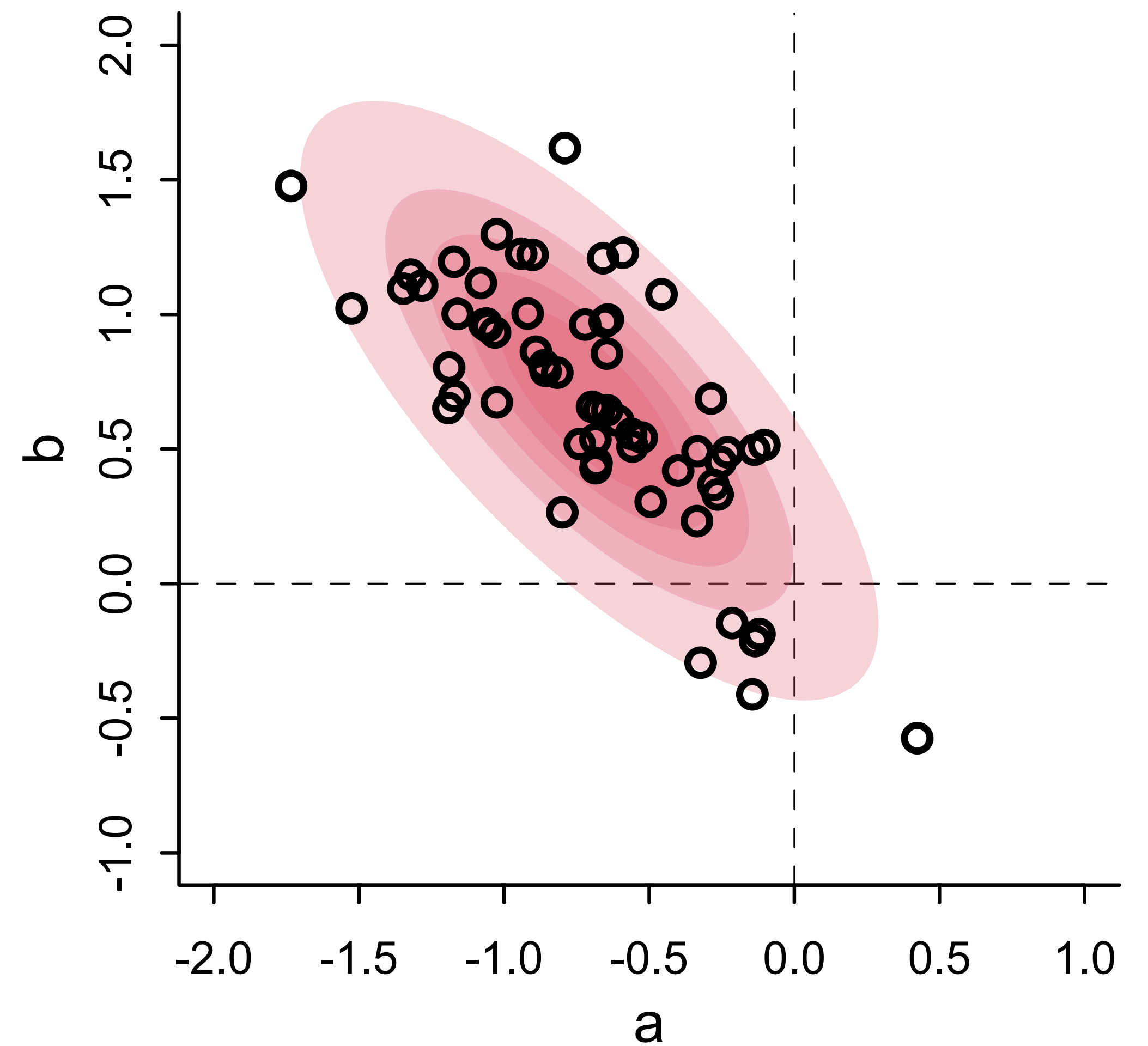
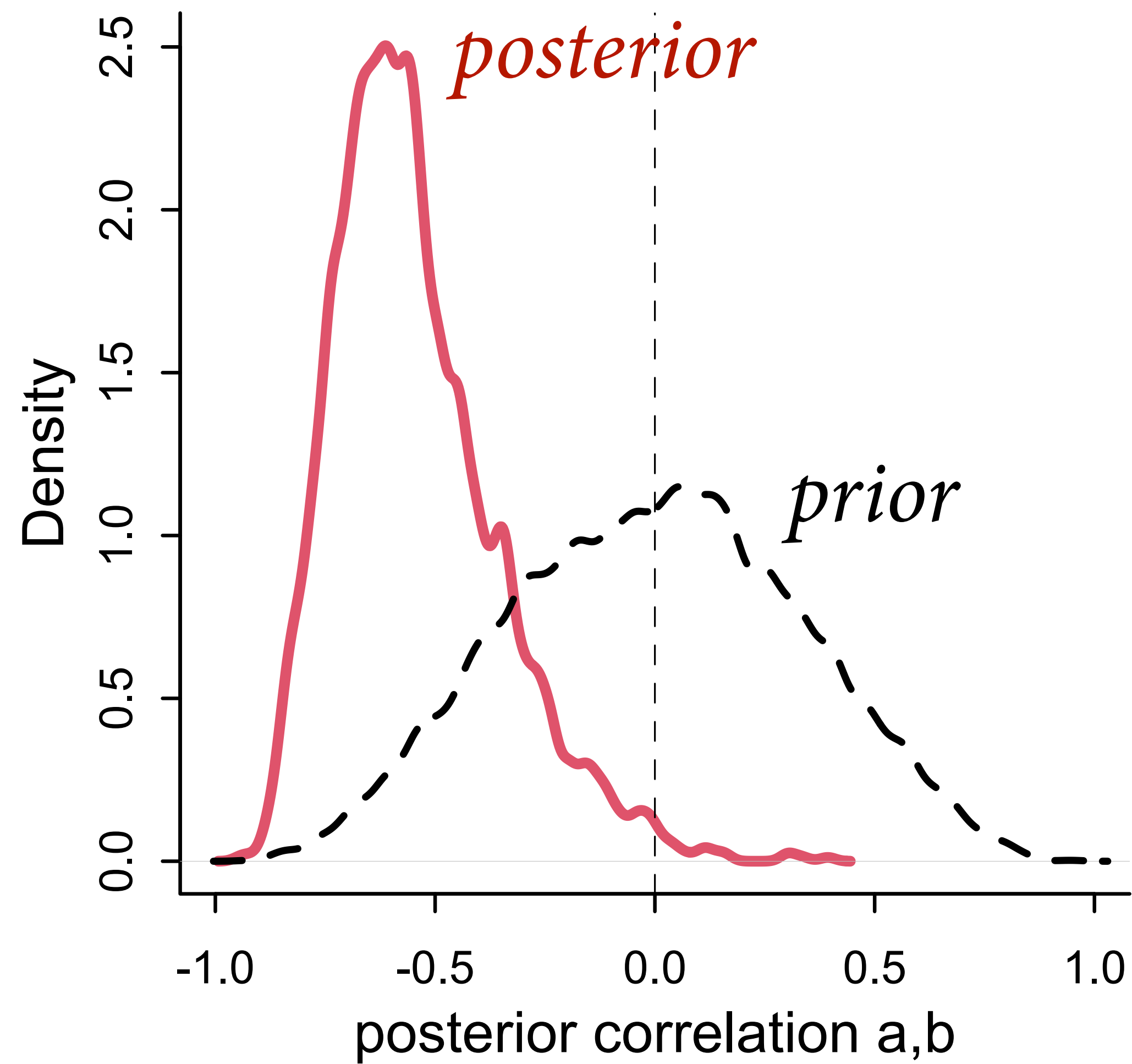
$$\sigma \sim \text{Exponential}(1)$$

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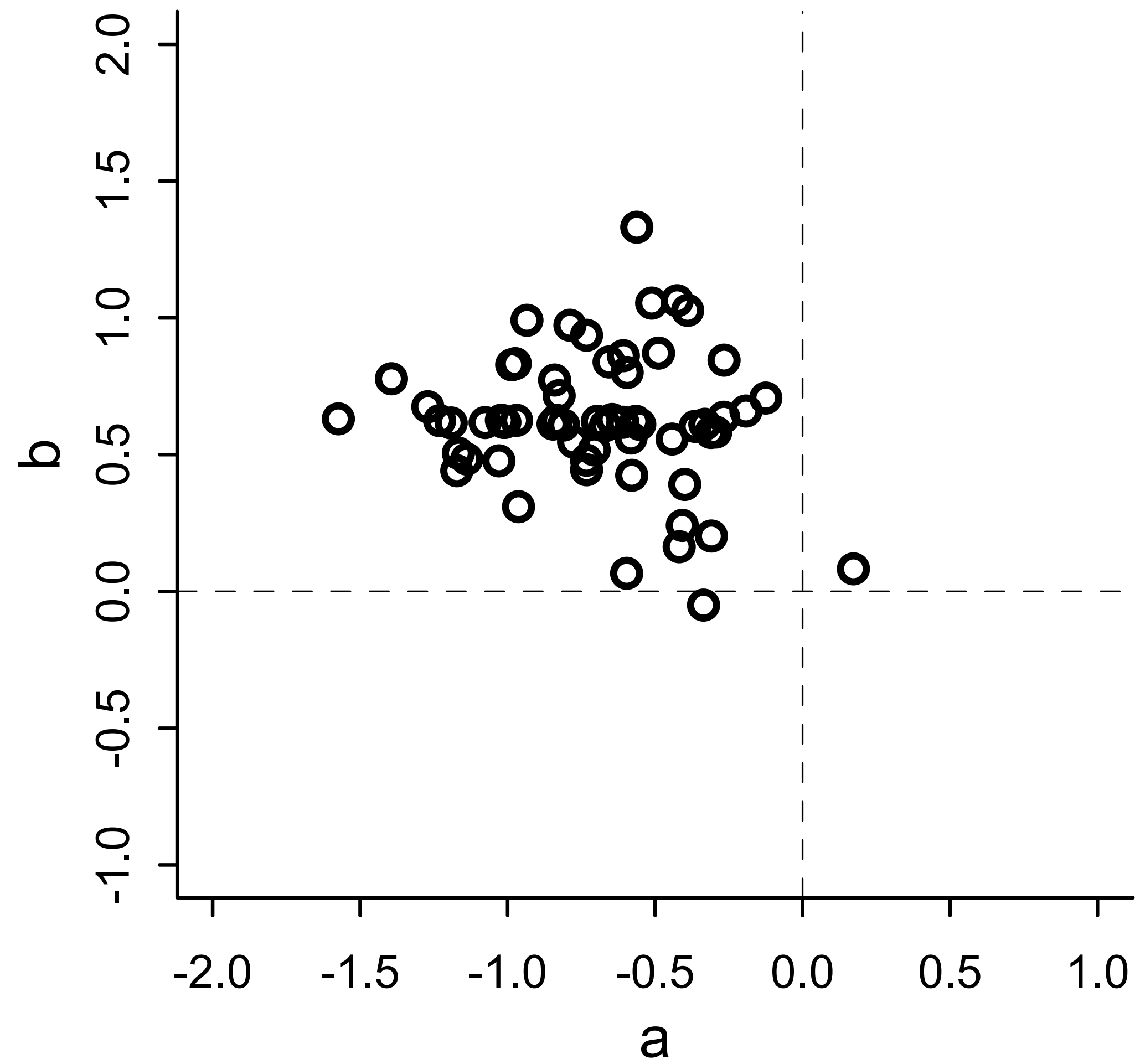
Correlated features



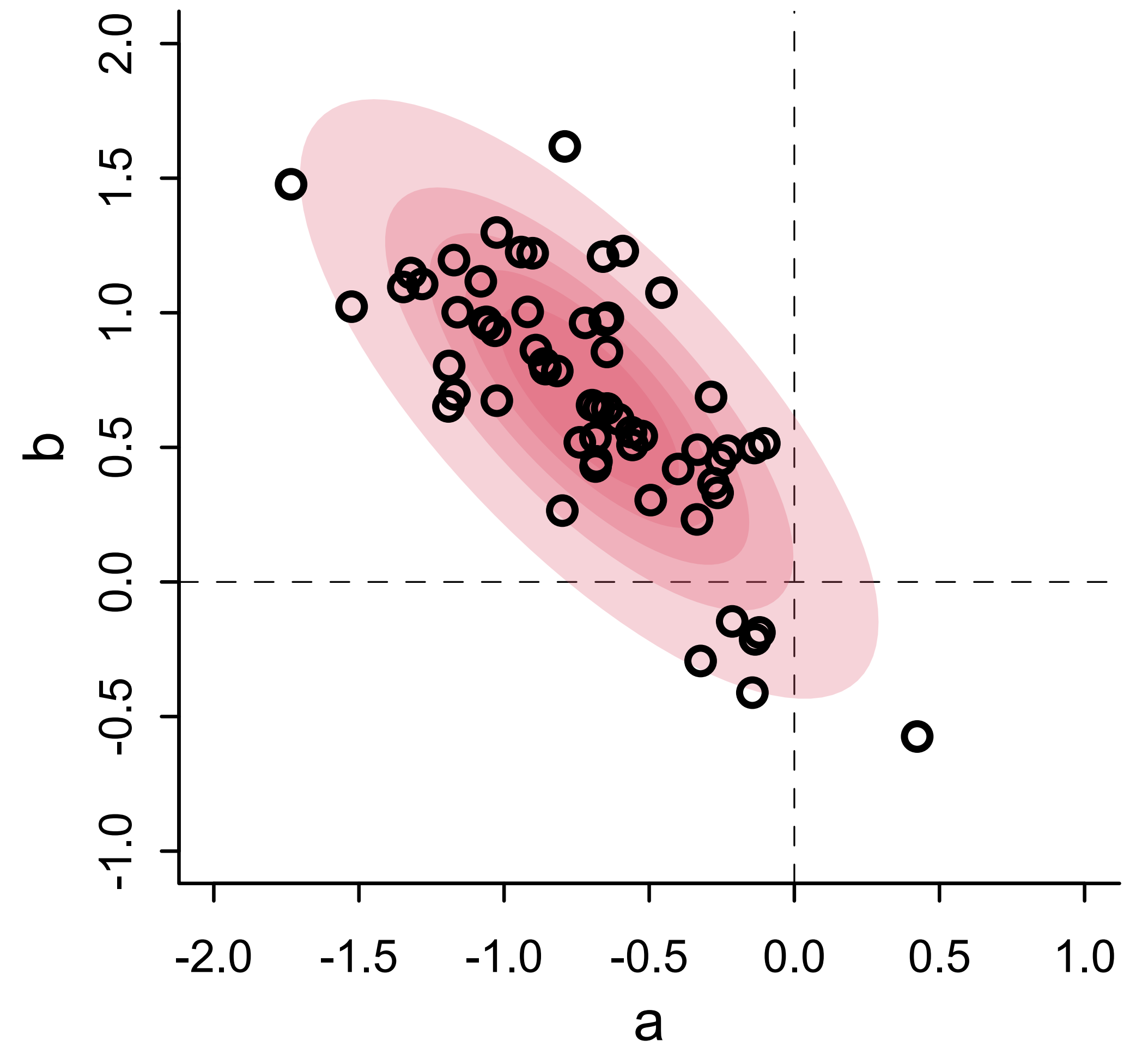
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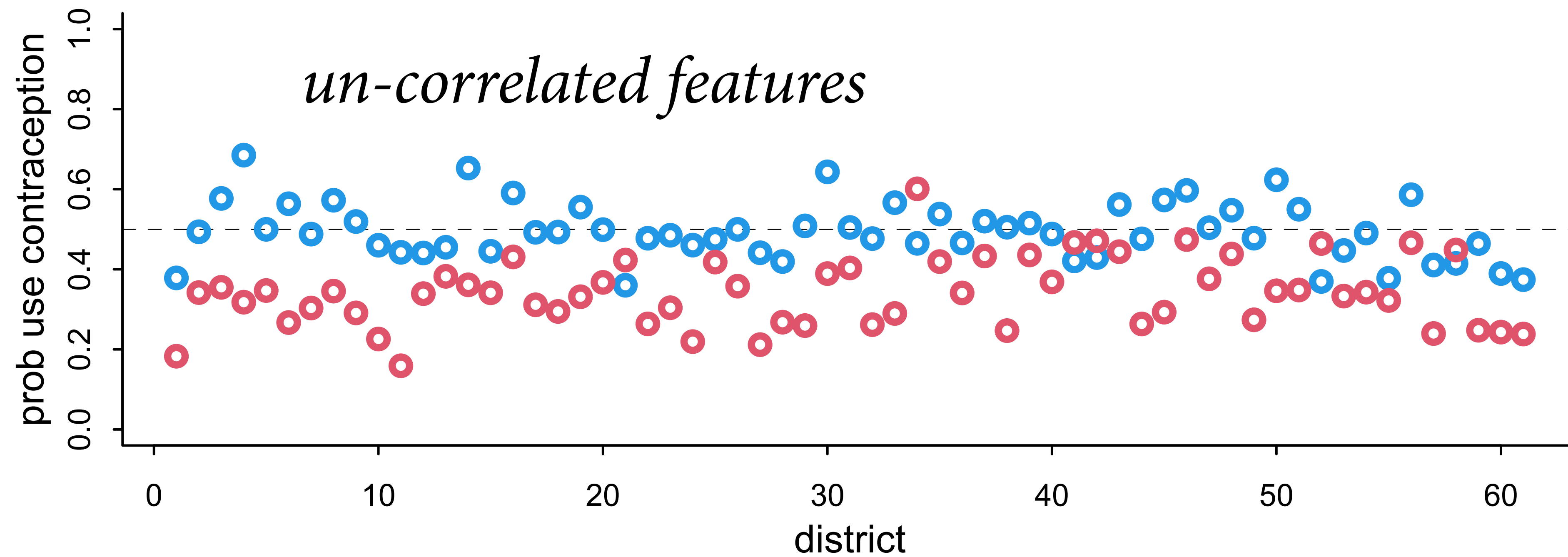
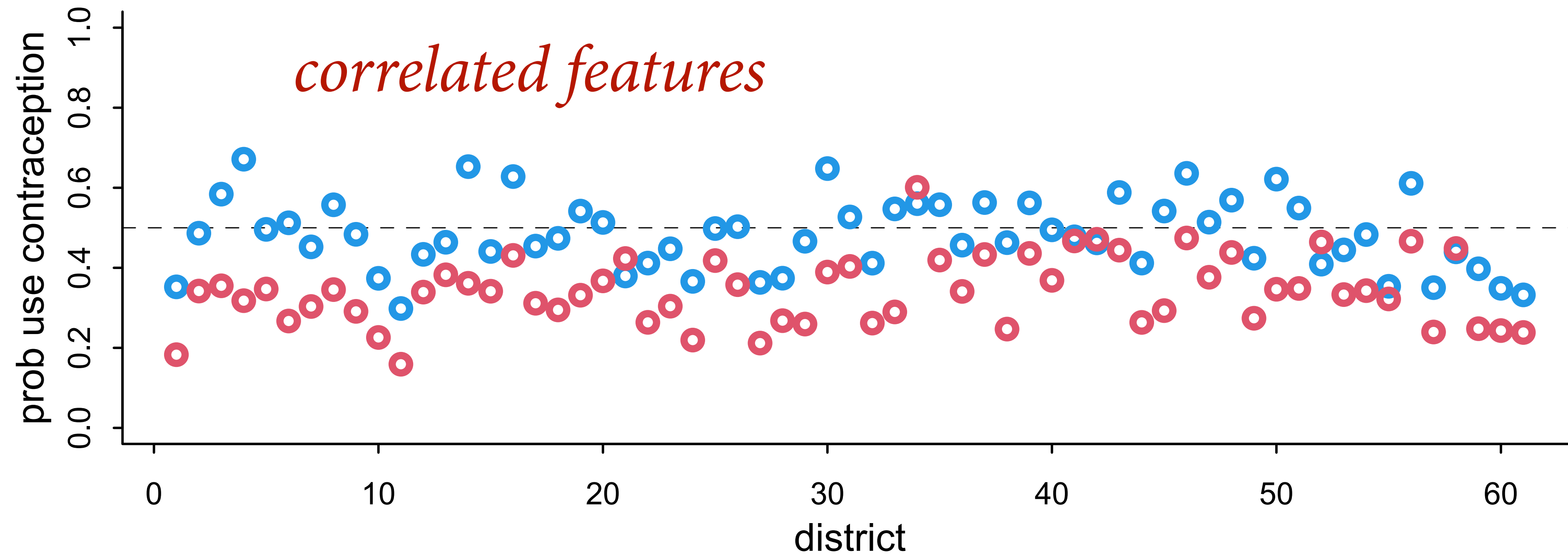


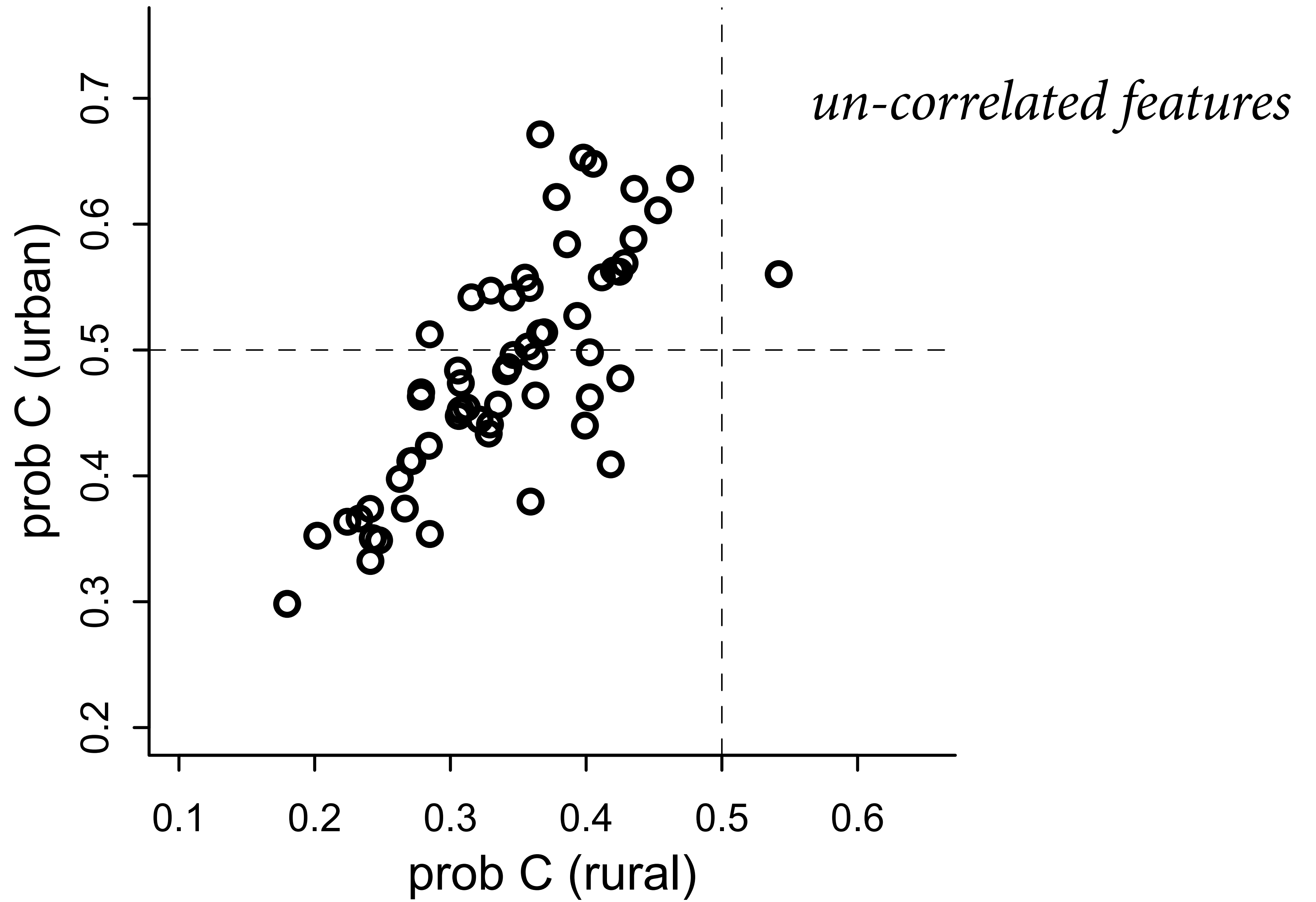
un-correlated features

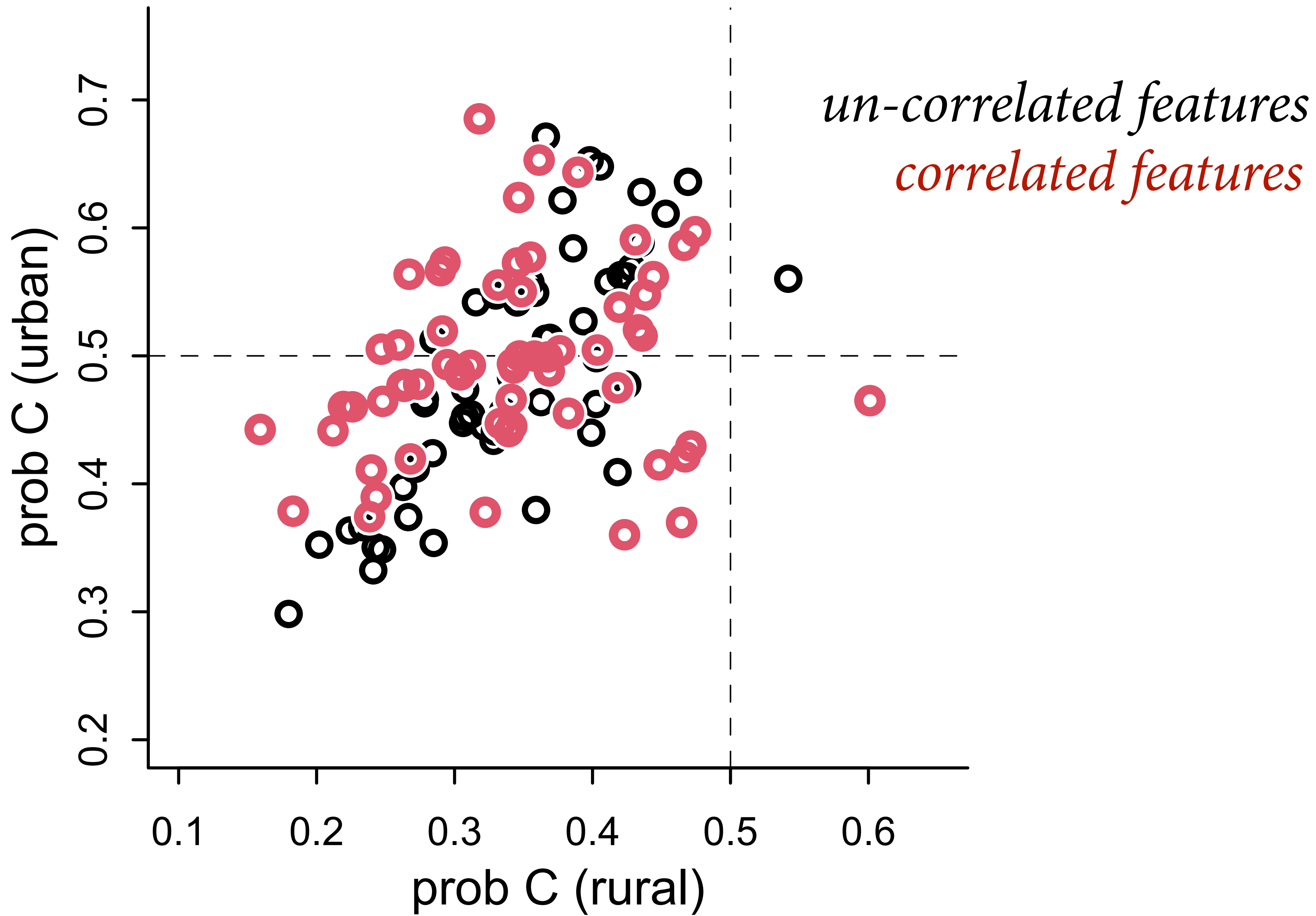


correlated features







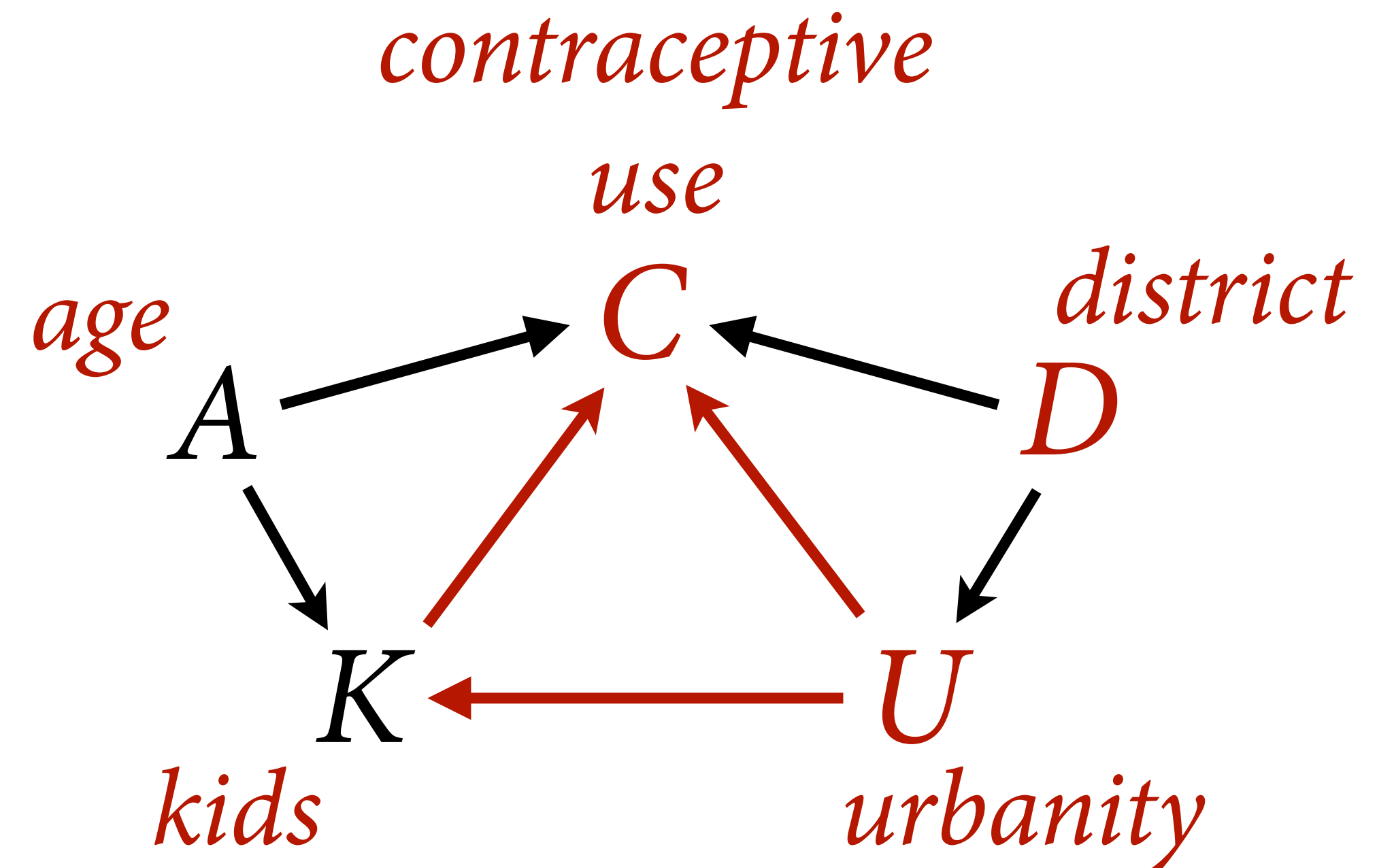


Bangladesh 1989, Party Like It's

Estimand 1: C in each district

Estimand 2: Effect of U

Estimand 3: Effects of K and A

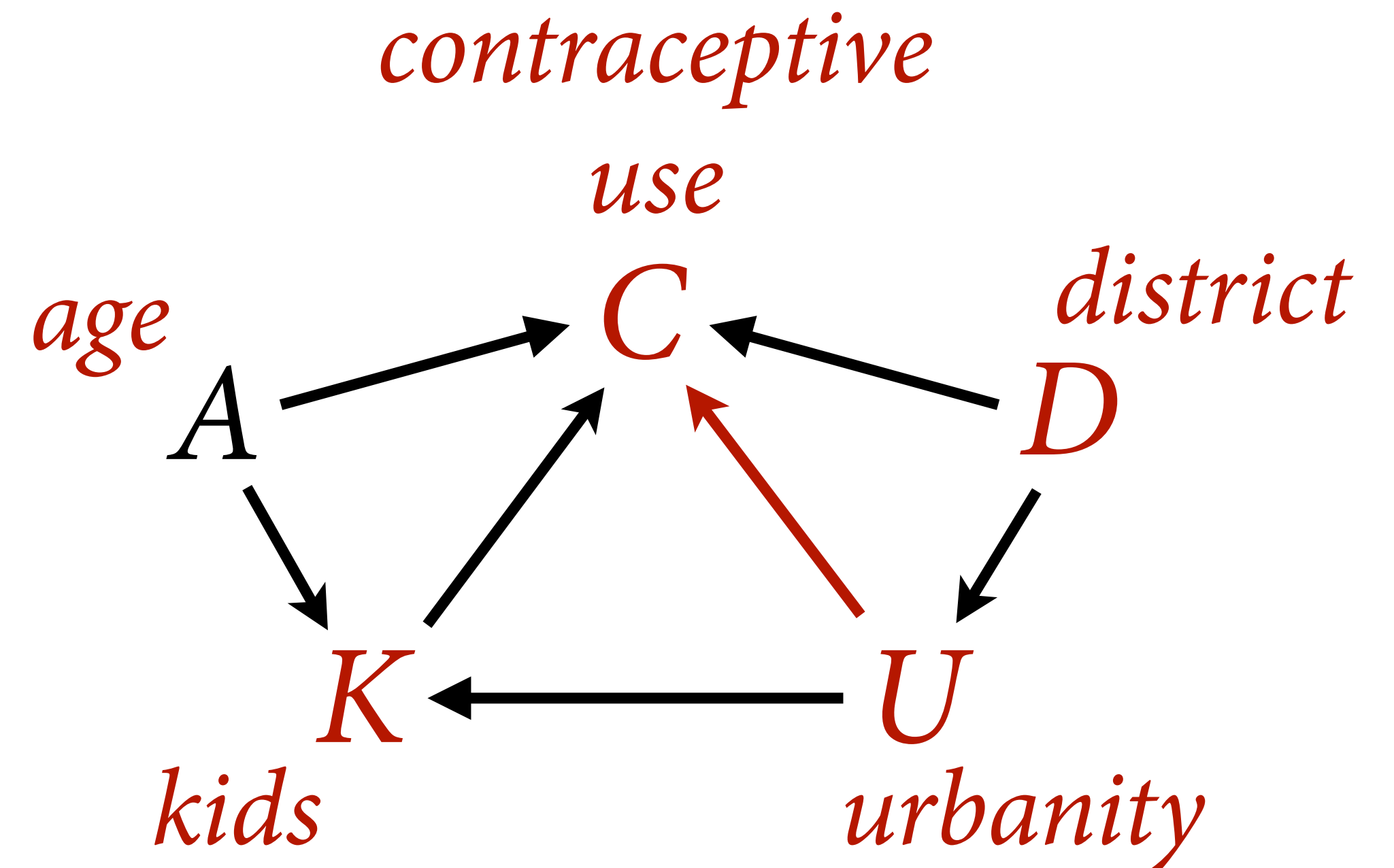


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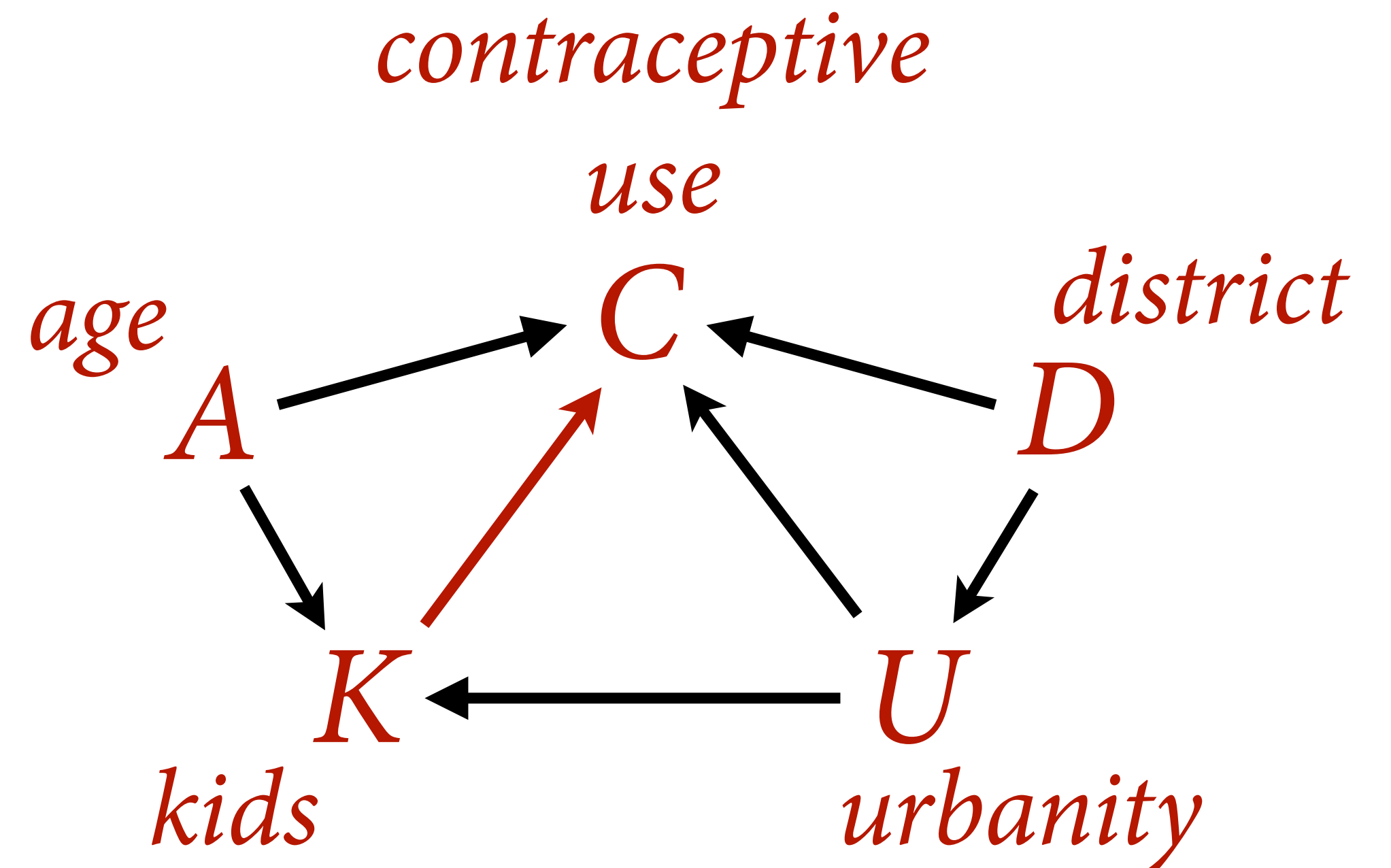


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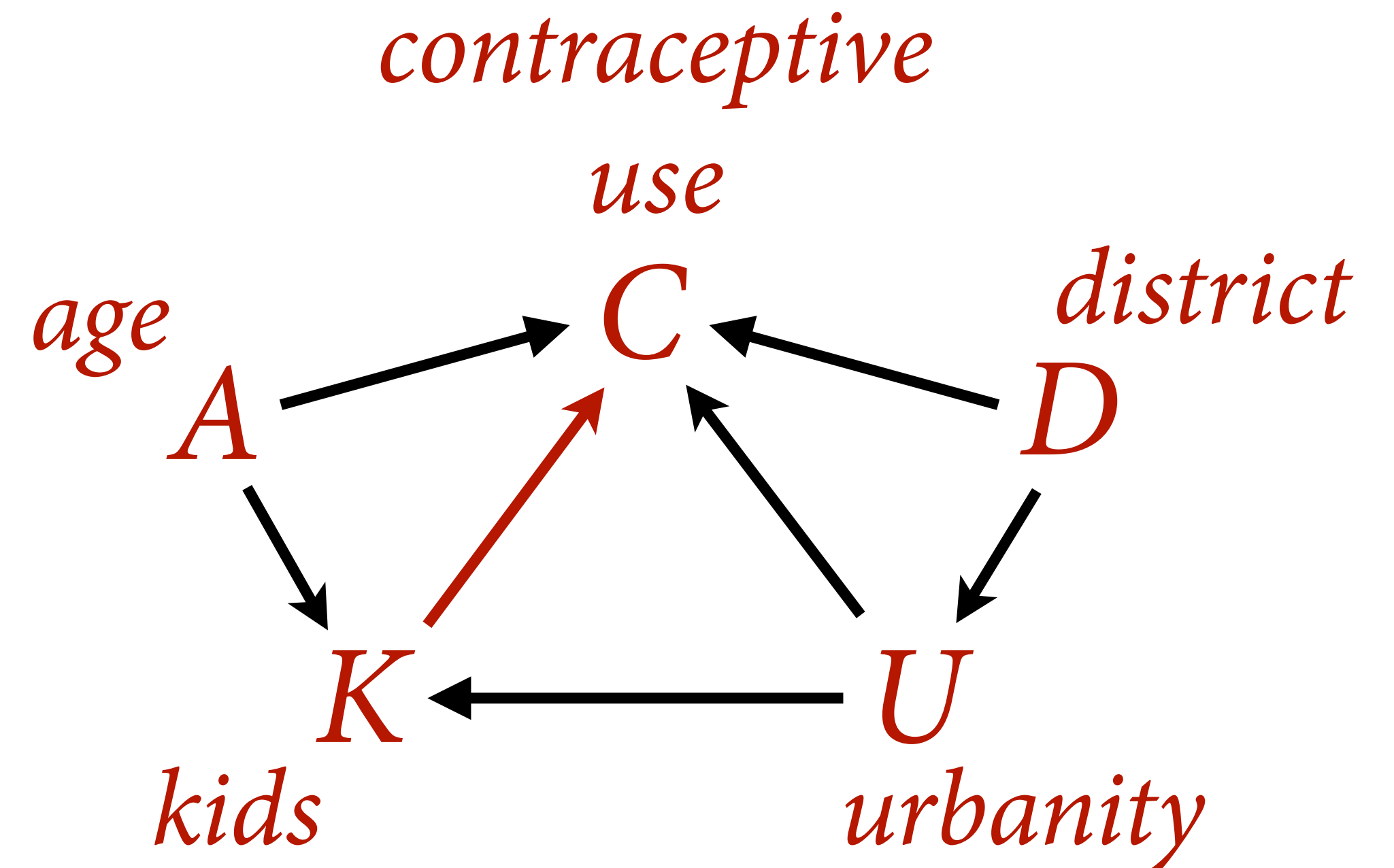
Bangladesh 1989, Party Like It's

Estimand 1: C in each district

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Estimand 3: Effects of K and A

Synthetic data & testing



Correlated Varying Effects

Priors that **learn correlation**

structure:

- (1) partial pooling across features
- (2) learn correlations

$$\alpha_j \sim \text{Normal}(0, 1)$$

$$\beta_j \sim \text{Normal}(0, 1)$$

$$\alpha_j \sim \text{Normal}(\bar{\alpha}, \sigma)$$

$$\beta_j \sim \text{Normal}(\bar{\beta}, \tau)$$

$$\begin{bmatrix} \alpha_j \\ \beta_j \end{bmatrix} \sim \text{MVN} \left(\begin{bmatrix} \bar{\alpha} \\ \bar{\beta} \end{bmatrix}, \Sigma \right)$$



Correlated Varying Effects

Priors that **learn correlation** structure:

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Varying effects can be correlated even if the prior doesn't learn the correlations

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Correlated Varying Effects

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Ethical obligation to do our best

$$\alpha_j \sim \text{Normal}(0, 1)$$

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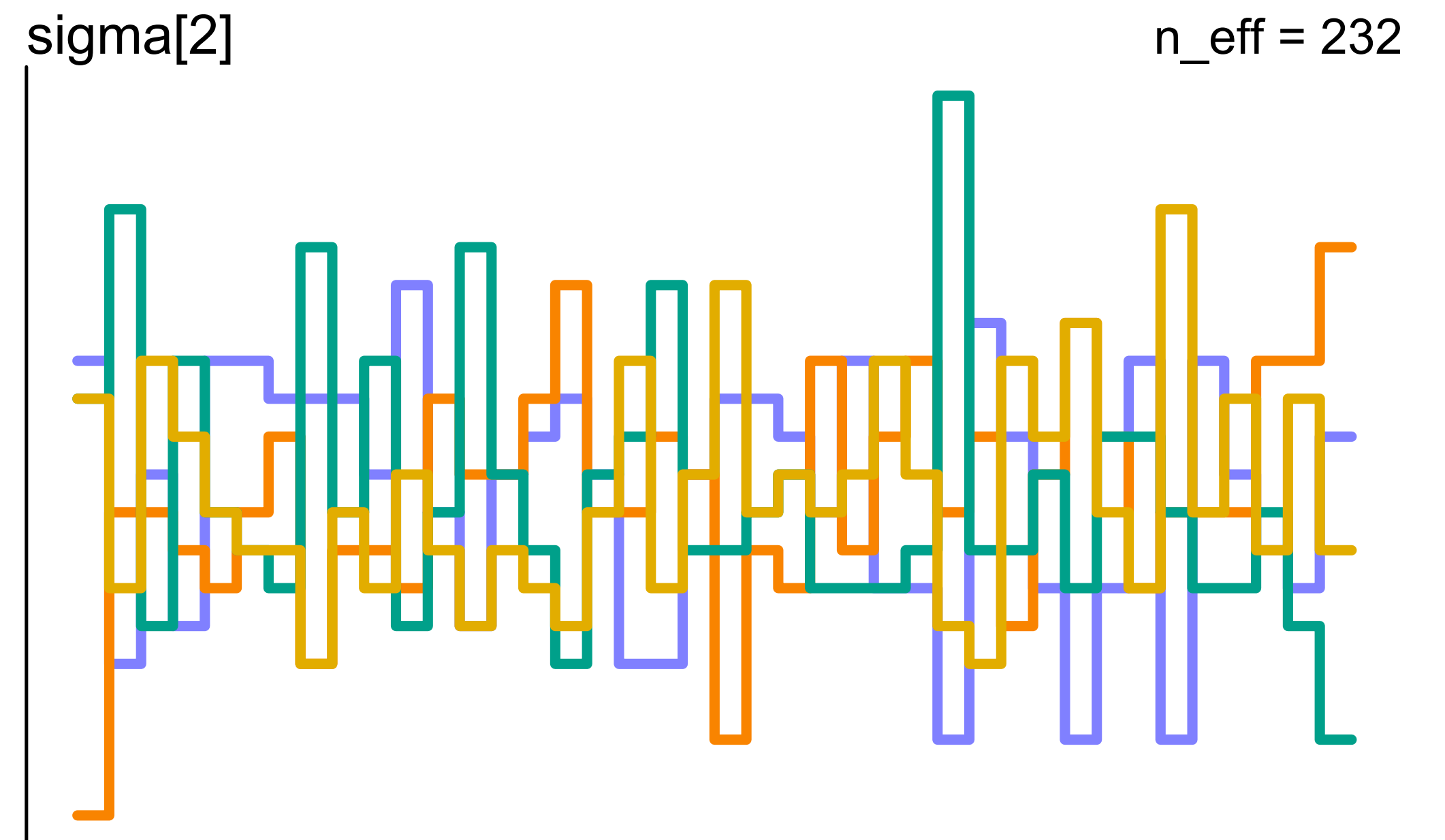
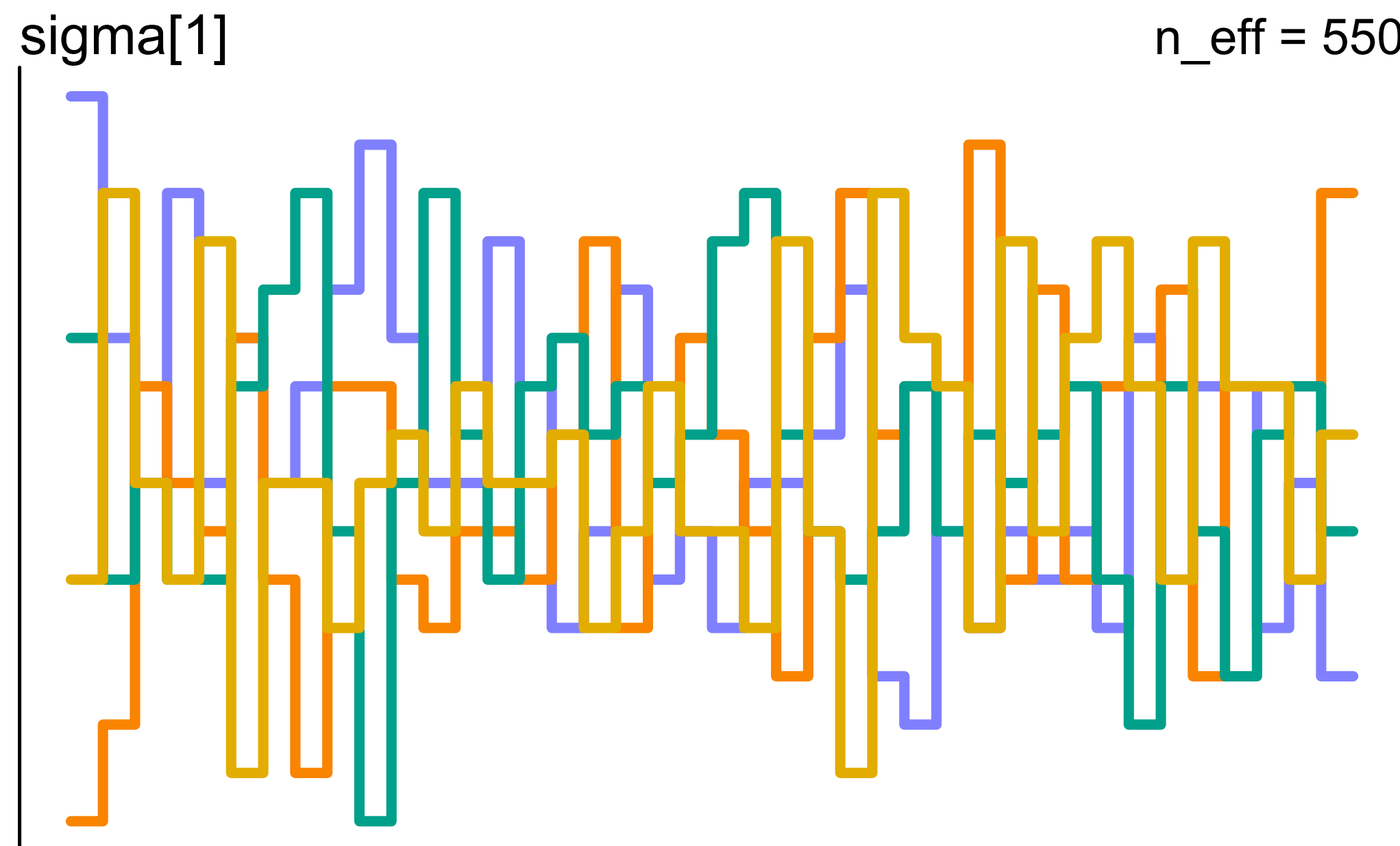
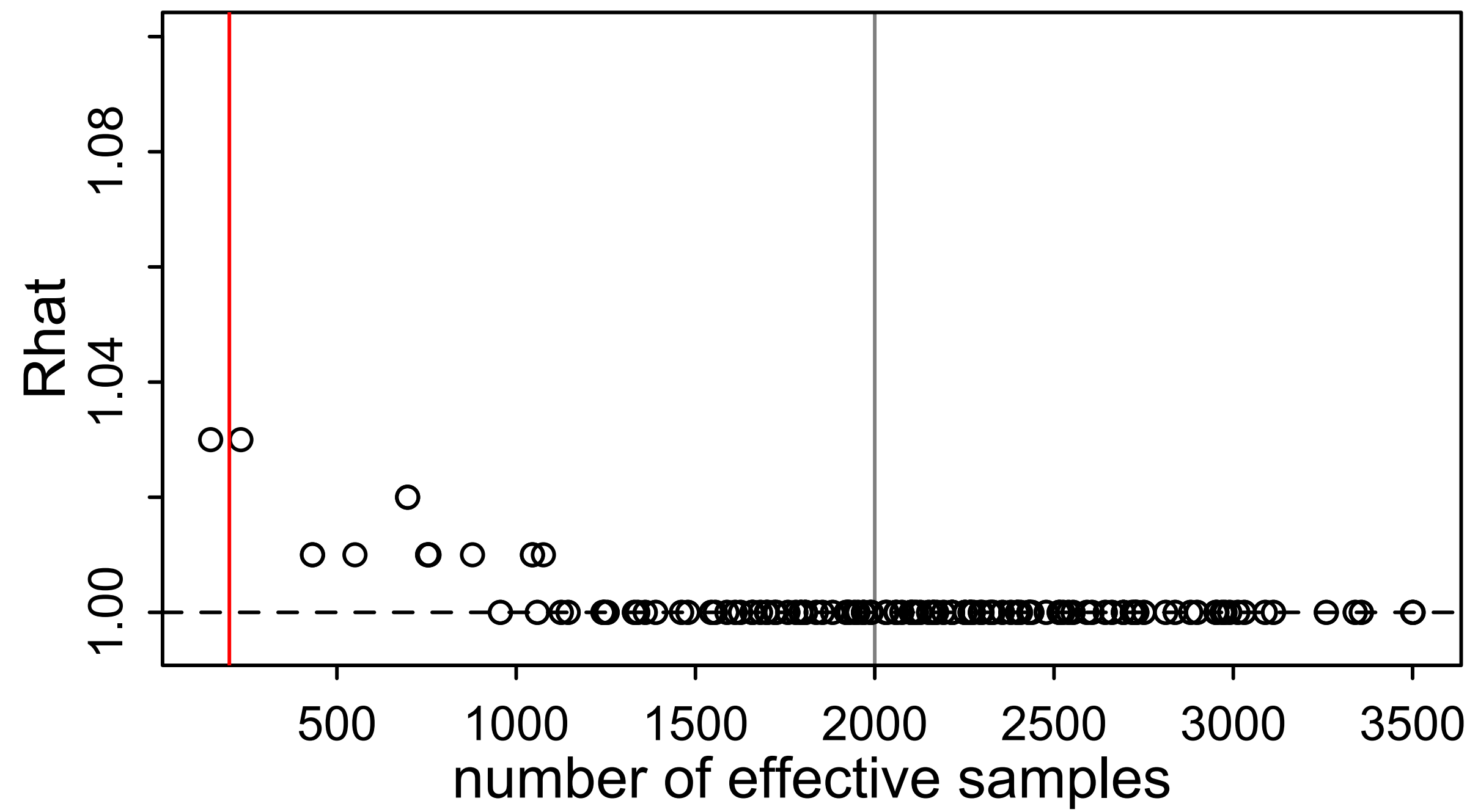


Course Schedule

Week 1	Bayesian inference	Chapters 1, 2, 3
Week 2	Linear models & Causal Inference	Chapter 4
Week 3	Causes, Confounds & Colliders	Chapters 5 & 6
Week 4	Overfitting / MCMC	Chapters 7, 8, 9
Week 5	Generalized Linear Models	Chapters 10, 11
Week 6	Ordered categories & Multilevel models	Chapters 12 & 13
Week 7	More Multilevel models	Chapters 13 & 14
Week 8	Multilevel Tactics & Gaussian Processes	Chapter 14
Week 9	Measurement & Missingness	Chapter 15
Week 10	Generalized Linear Madness	Chapter 16

https://github.com/rmcelreath/stat_rethinking_2023

BONUS



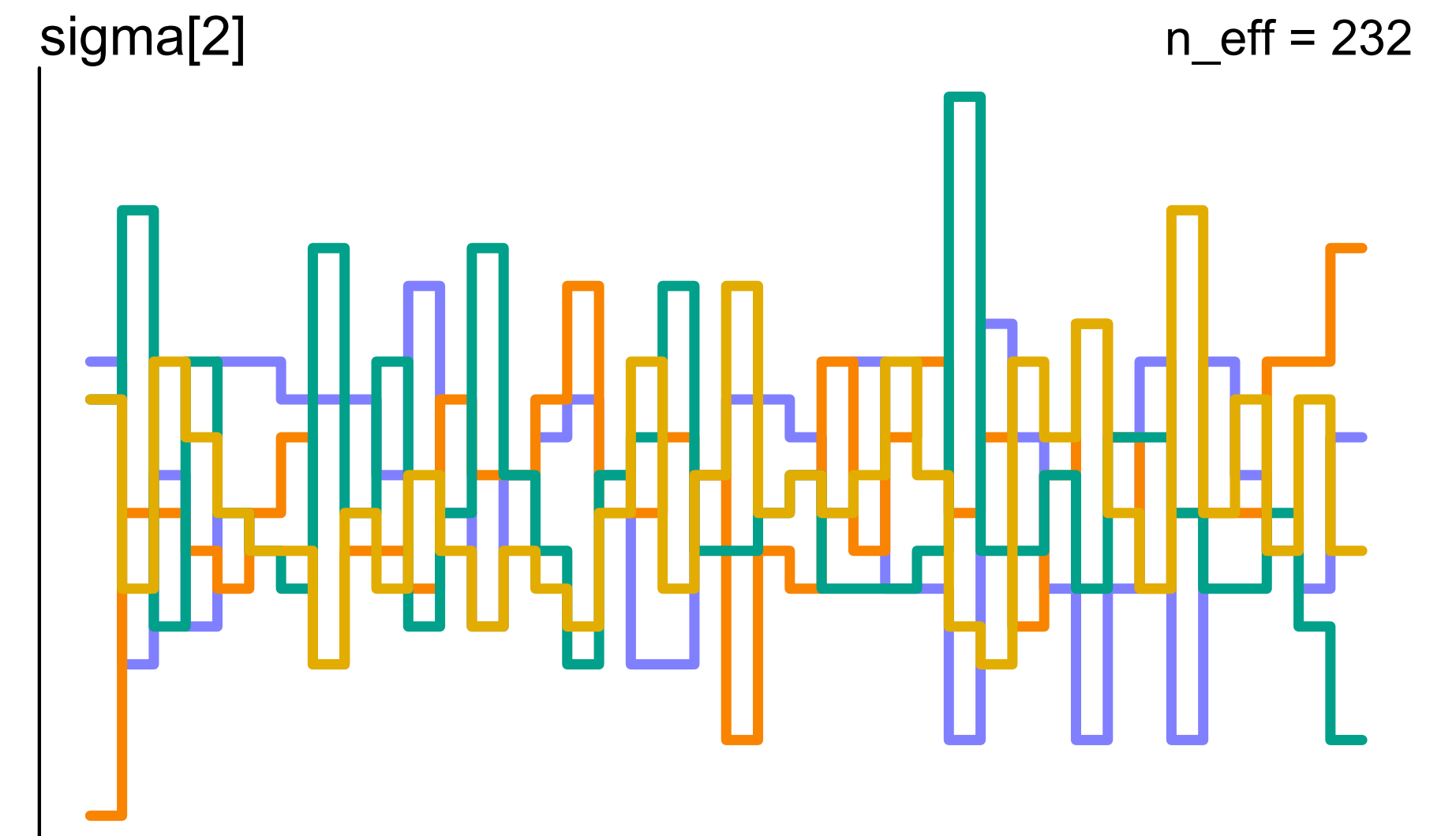
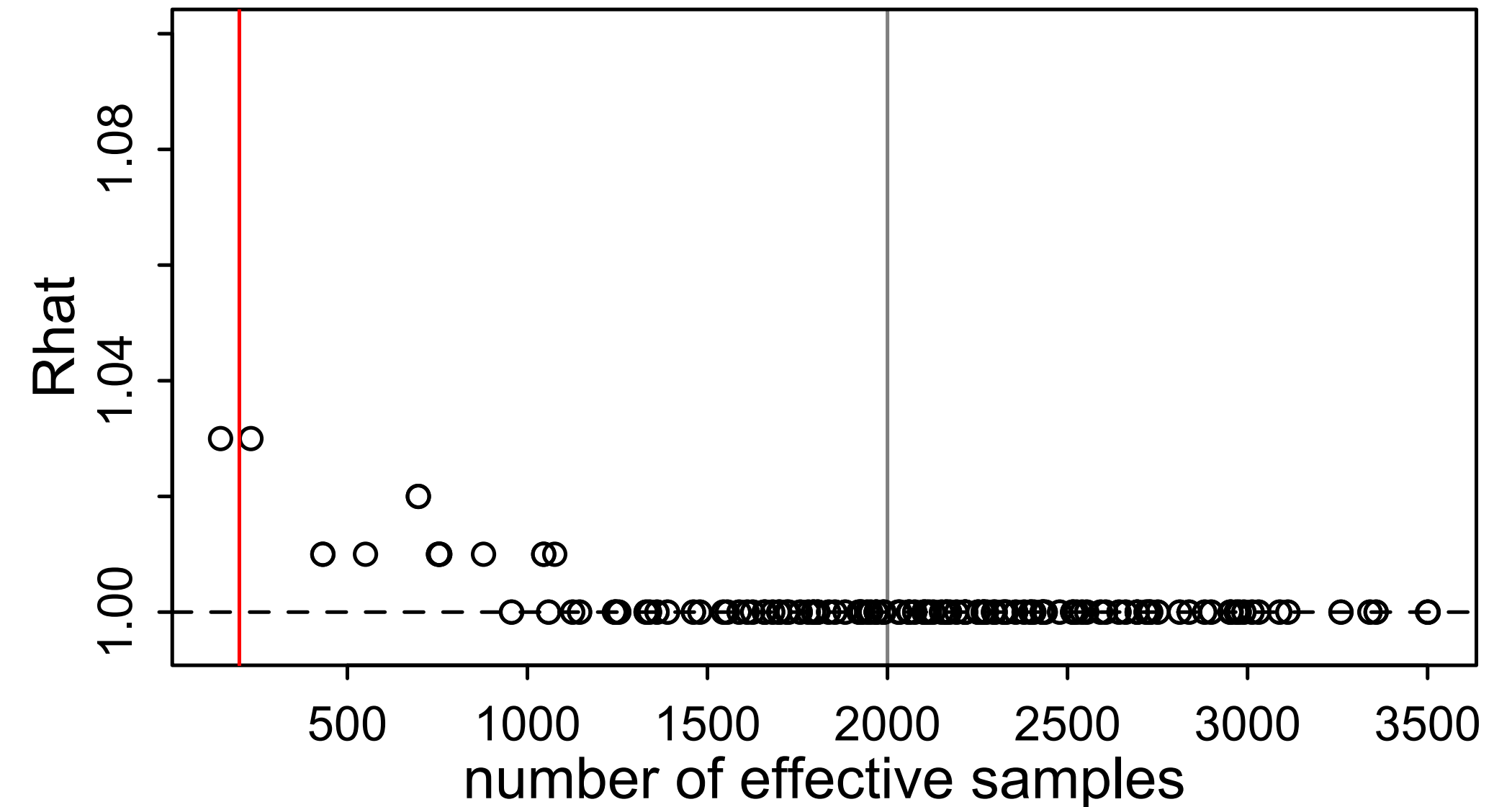
Inconvenient Posteriors

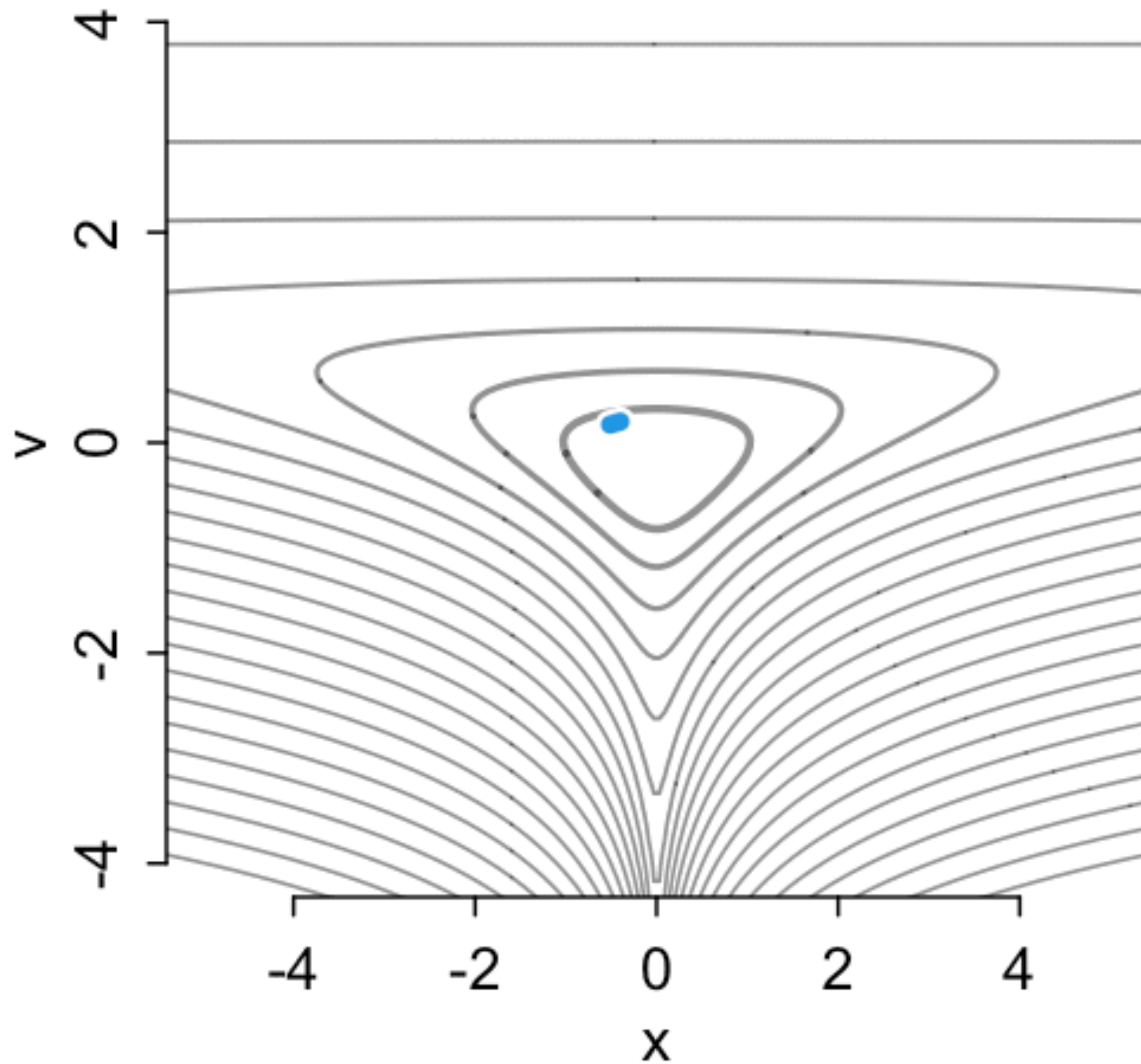
Inefficient MCMC can be caused by steep curvature

Hamiltonian simulation has trouble exploring surface

“Divergent transitions”

Transforming priors can help

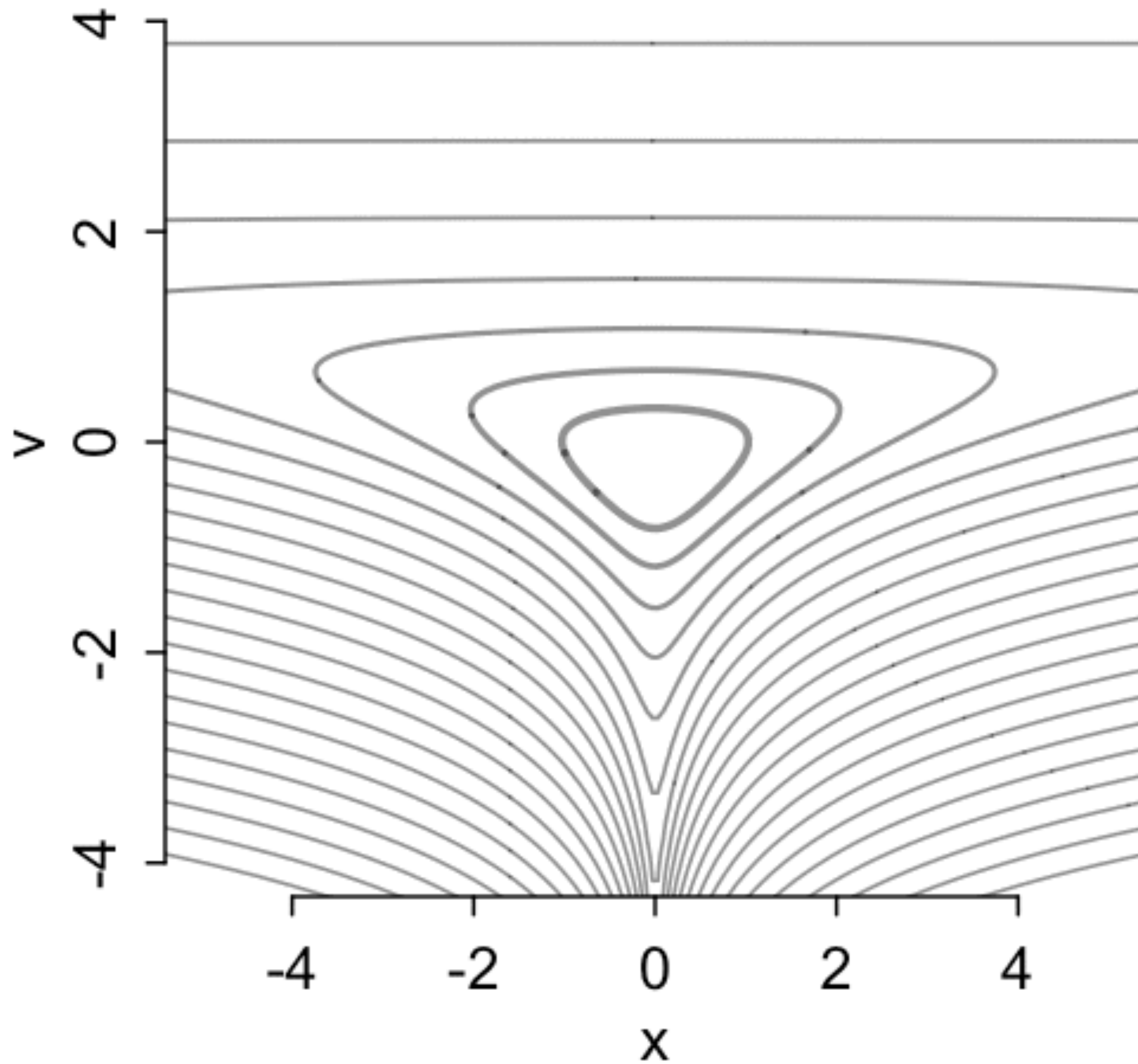




$$v \sim \text{Normal}(0, 0.5)$$

$$x \sim \text{Normal}(0, \exp(v))$$

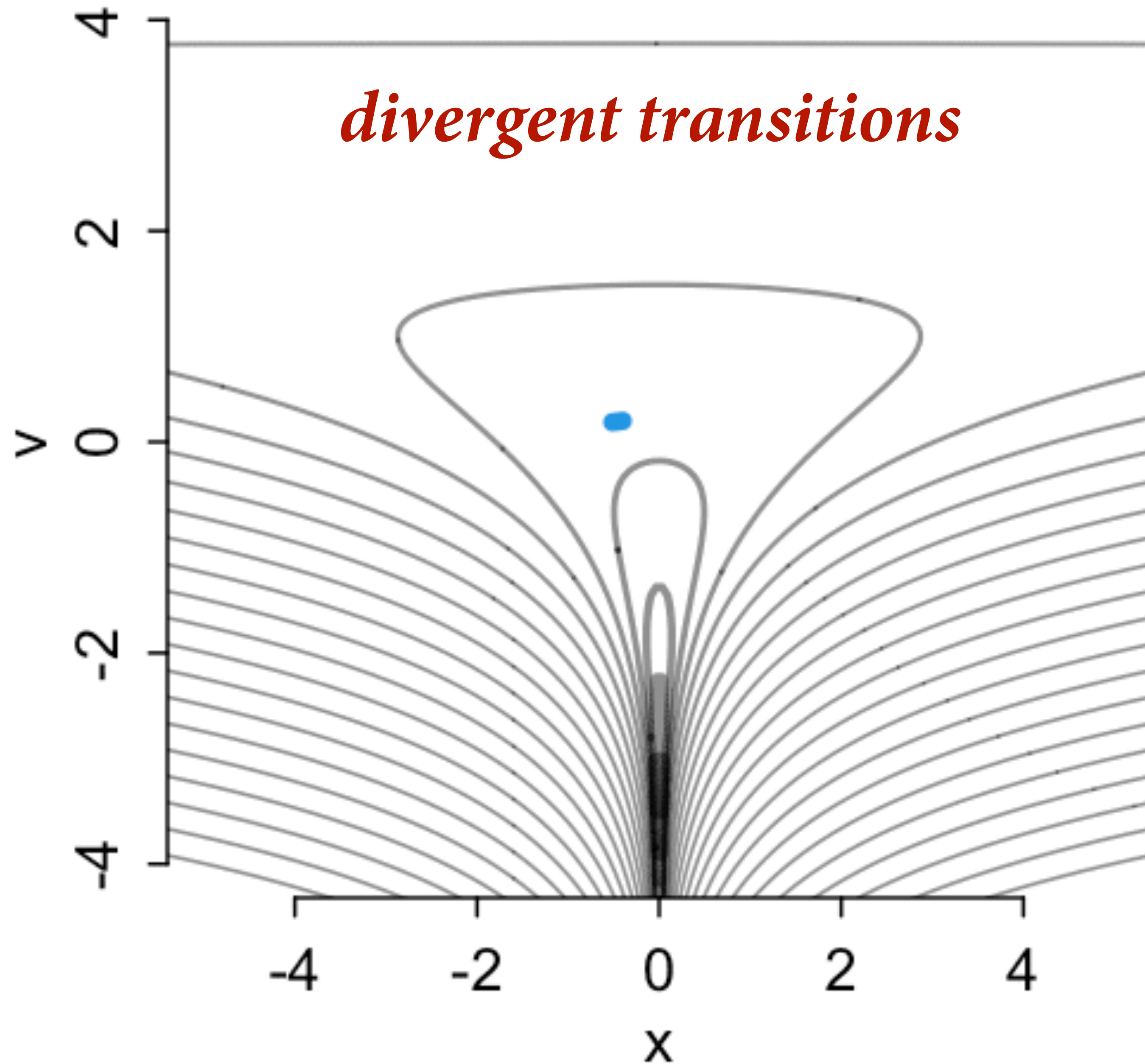
$v \sim \text{normal}(0, 0.5)$



$v \sim \text{Normal}(0, _)$

$x \sim \text{Normal}(0, \exp(v))$

divergent transitions



$$v \sim \text{Normal}(0, 3)$$

$$x \sim \text{Normal}(0, \exp(v))$$



Divergent transitions

Why? Same step size not optimal everywhere

High curvature = simulation cannot follow surface

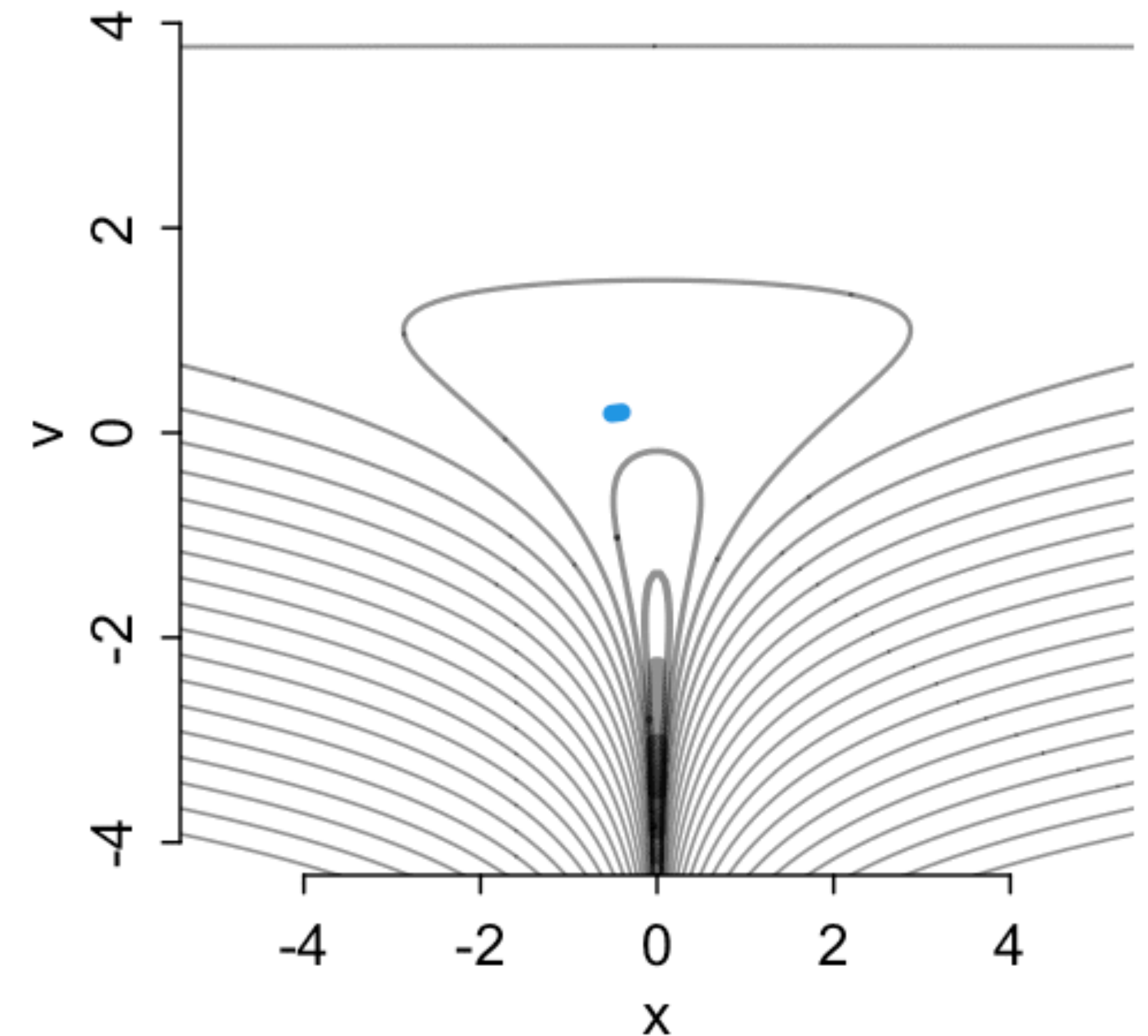
What can we do?

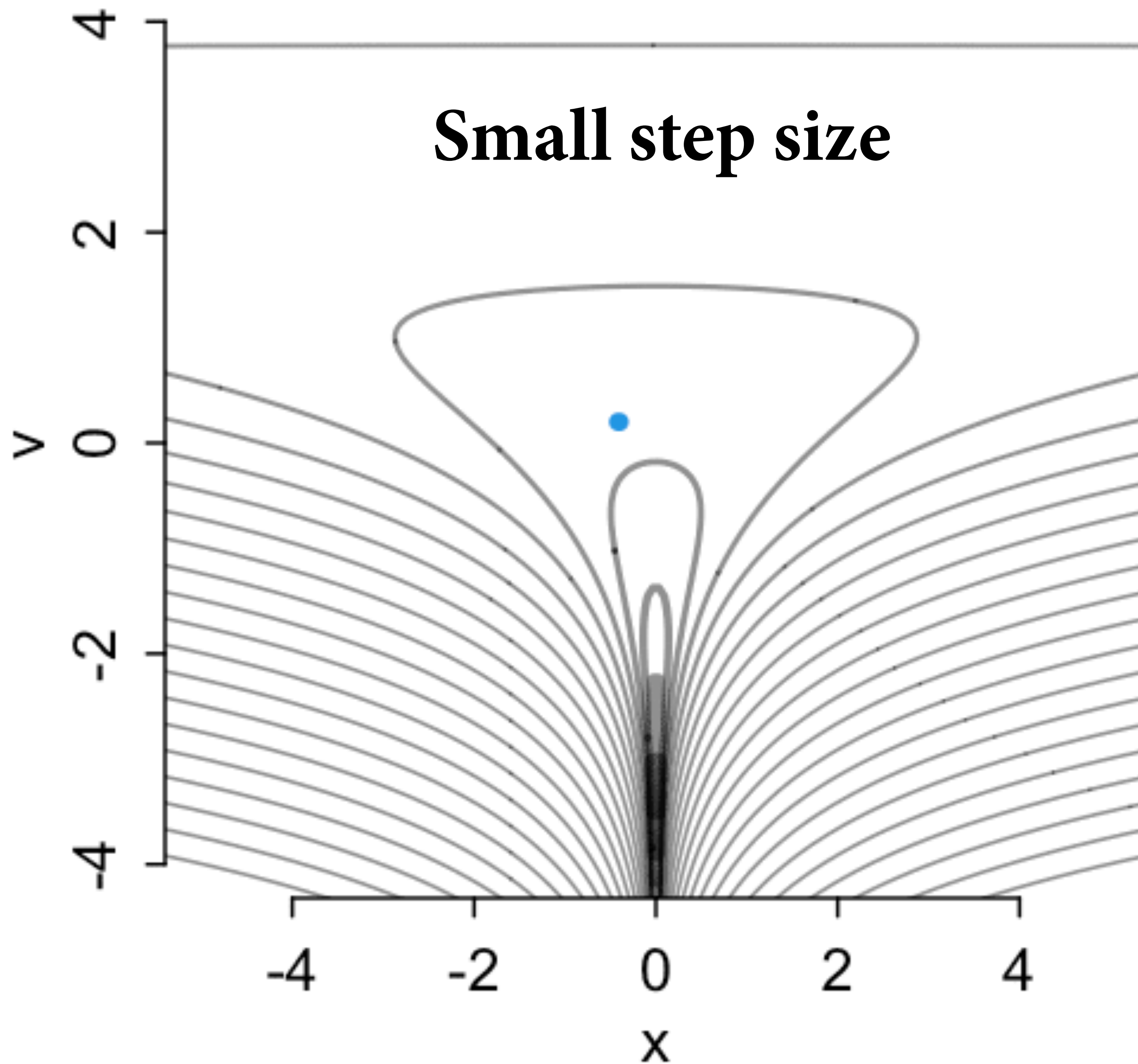
(1) use a smaller step size

(2) reparameterize!

$$v \sim \text{Normal}(0, 3)$$

$$x \sim \text{Normal}(0, \exp(v))$$





$$v \sim \text{Normal}(0, 3)$$

$$x \sim \text{Normal}(0, \exp(v))$$

Small step size helps, but
makes exploration slow

“Centered”

$$v \sim \text{Normal}(0, 3)$$

$$x \sim \text{Normal}(0, \exp(v))$$

“Centered”

$$v \sim \text{Normal}(0,3)$$

$$x \sim \text{Normal}(0, \exp(v))$$

“Non-centered”

$$v \sim \text{Normal}(0,3)$$

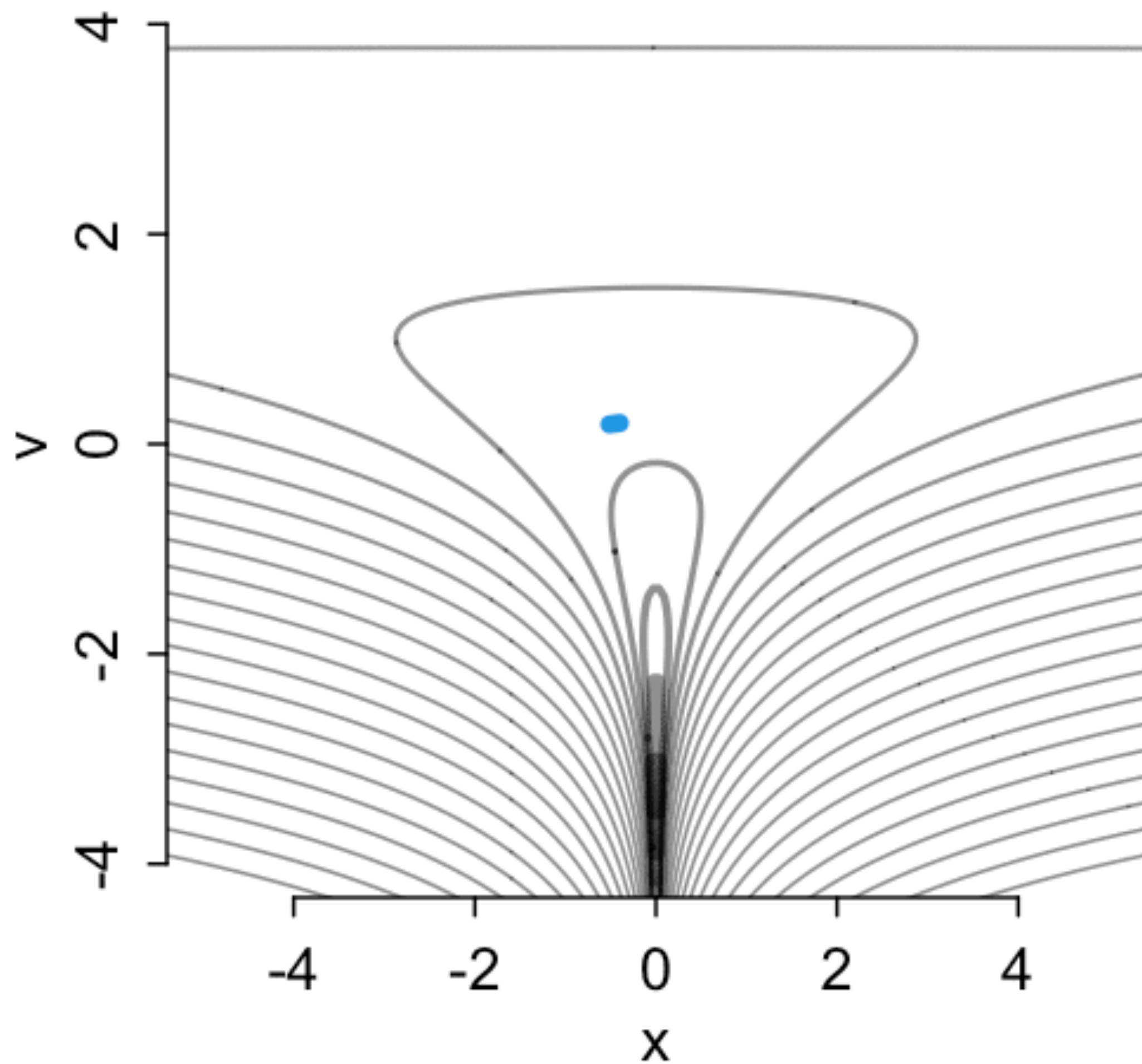
$$z \sim \text{Normal}(0,1)$$

$$x = z \exp(v)$$

“Centered”

$$v \sim \text{Normal}(0,3)$$

$$x \sim \text{Normal}(0, \exp(v))$$

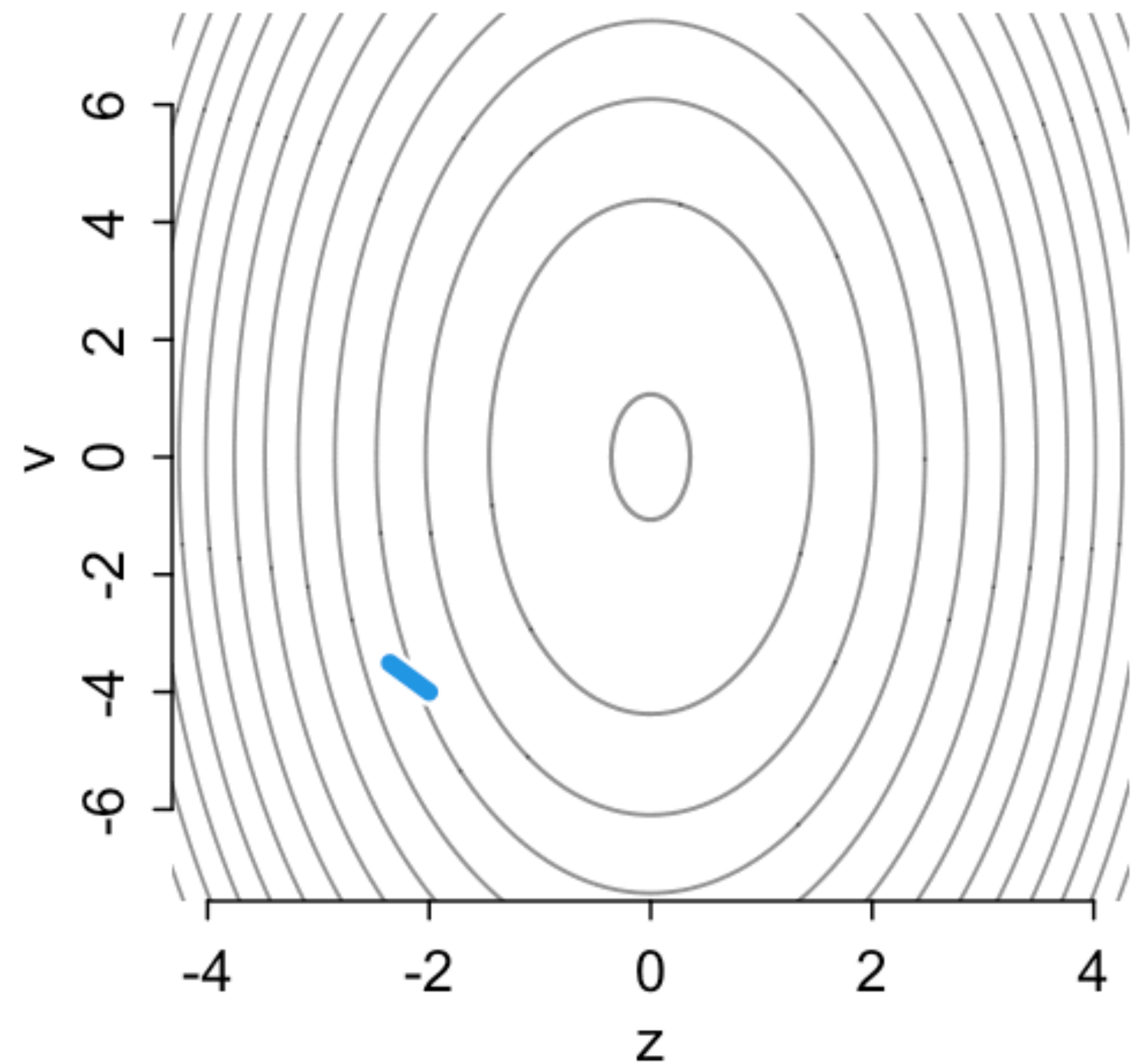


“Non-centered”

$$v \sim \text{Normal}(0,3)$$

$$z \sim \text{Normal}(0,1)$$

$$x = z \exp(v)$$



```
m13.7 <- ulam(  
  alist(  
    v ~ normal(0,3),  
    x ~ normal(0,exp(v))  
  ), data=list(N=1) , chains=4 )
```

```
m13.7nc <- ulam(  
  alist(  
    v ~ normal(0,3),  
    z ~ normal(0,1),  
    gq> real[1]:x <<- z*exp(v)  
  ), data=list(N=1) , chains=4 )
```

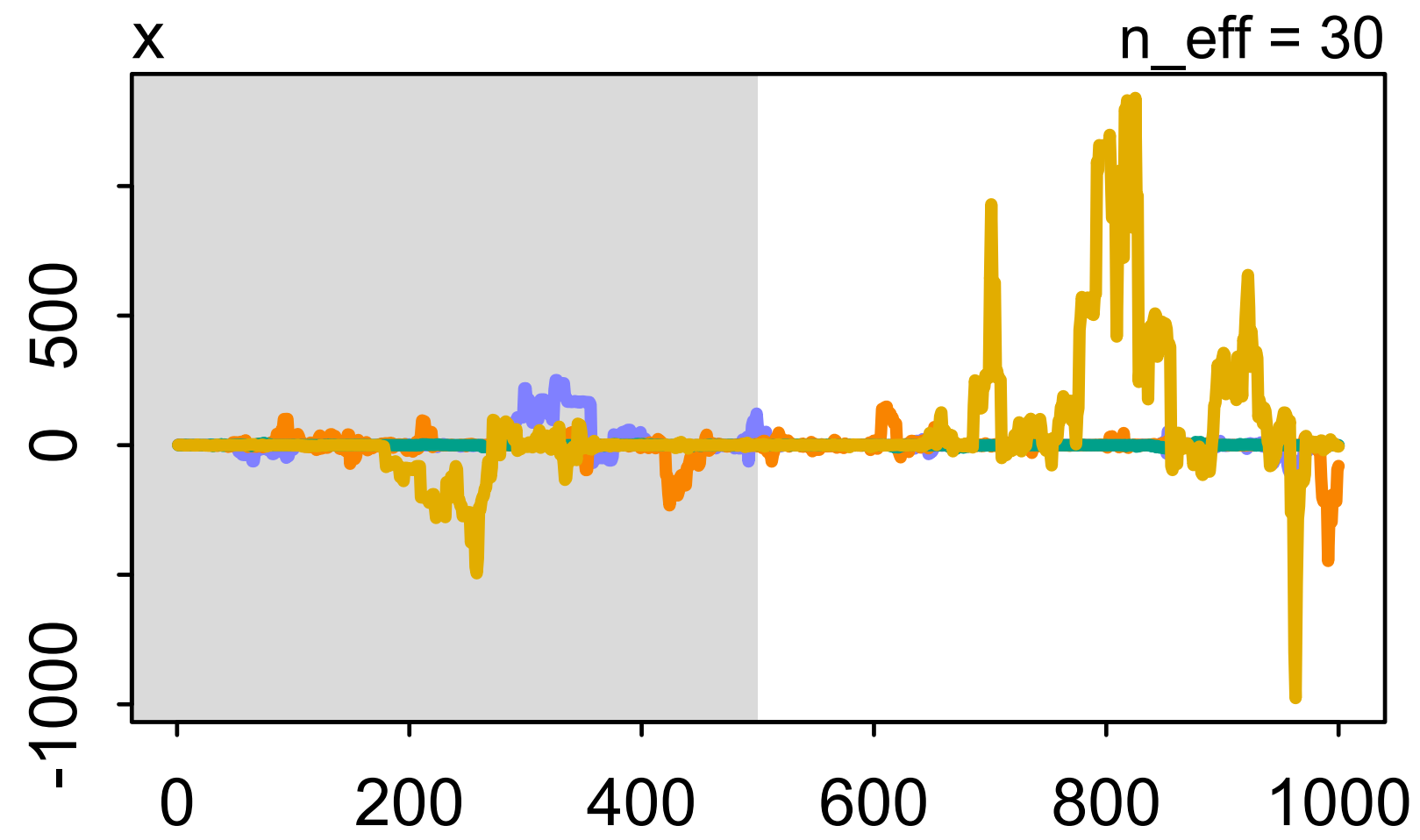
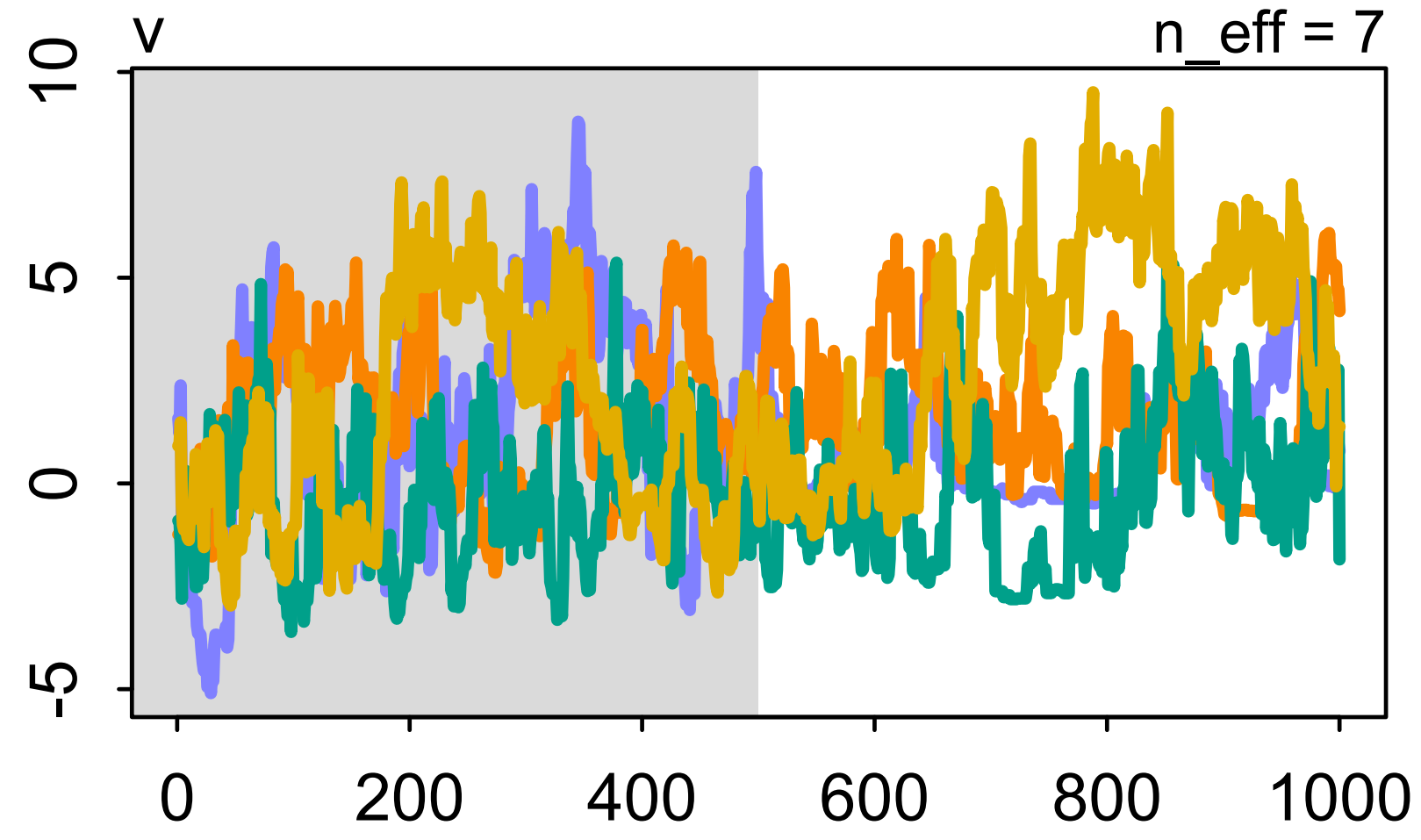
```
m13.7 <- ulam(
  alist(
    v ~ normal(0,3),
    x ~ normal(0,exp(v))
  ), data=list(N=1) , chains=4 )
```

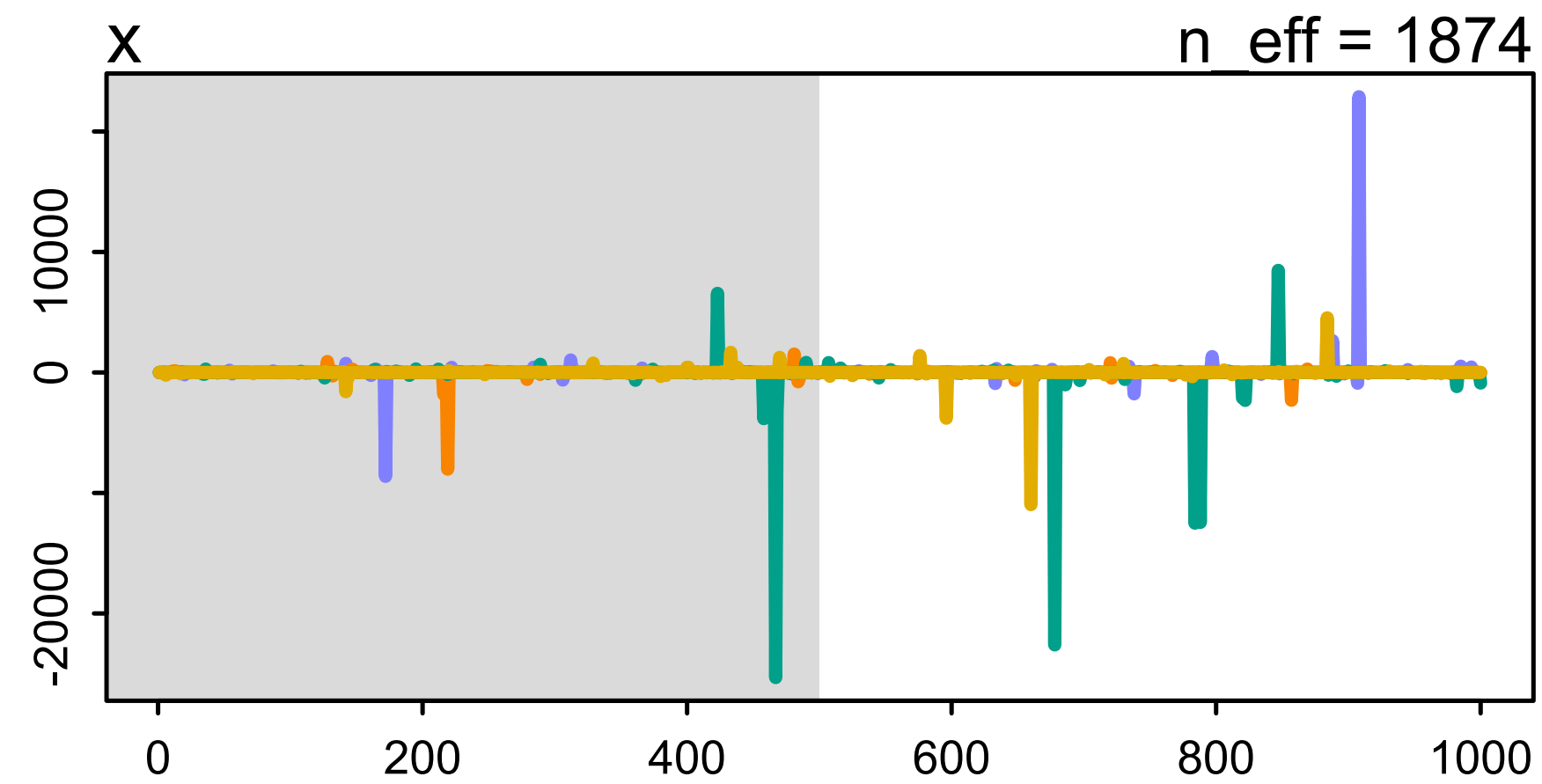
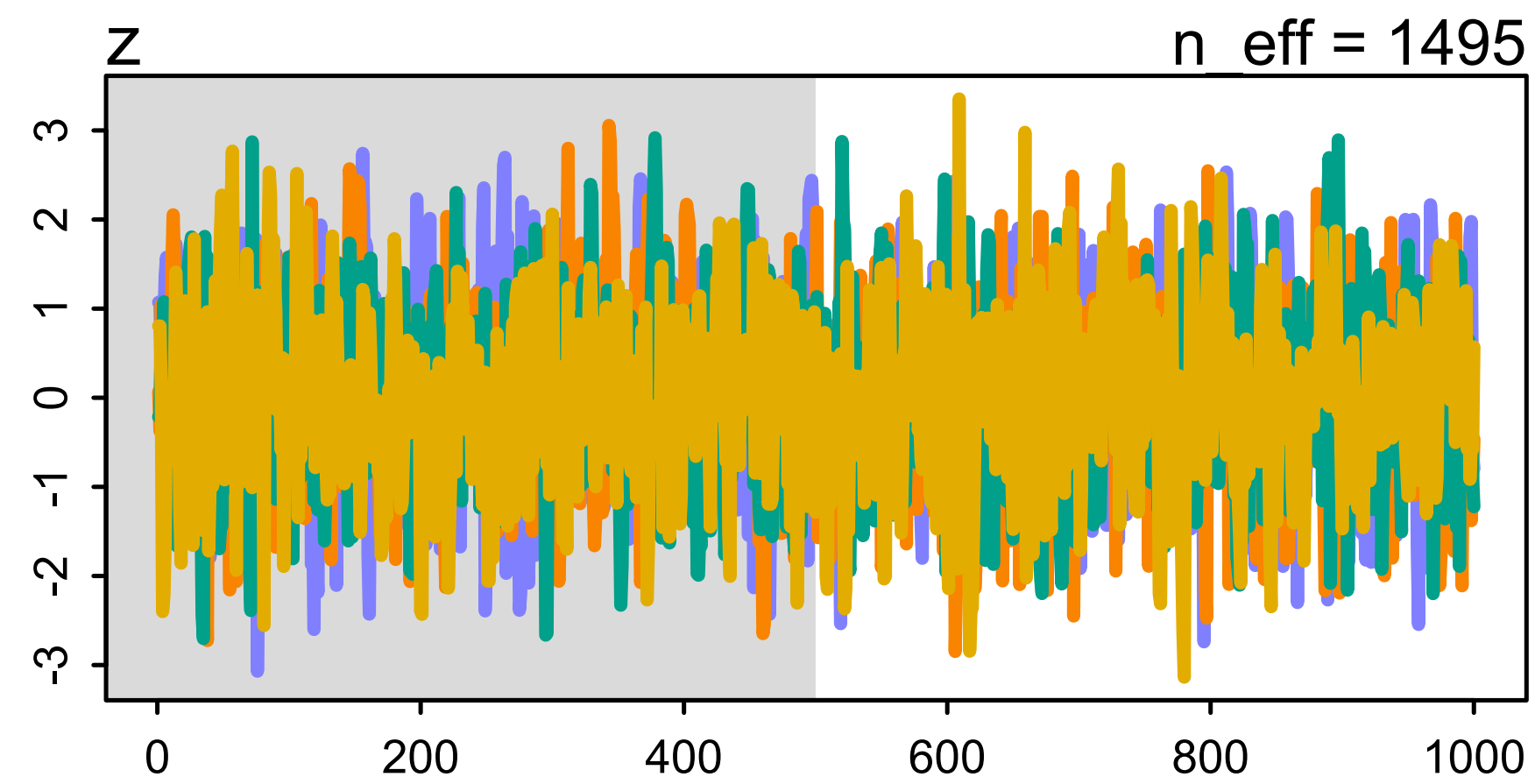
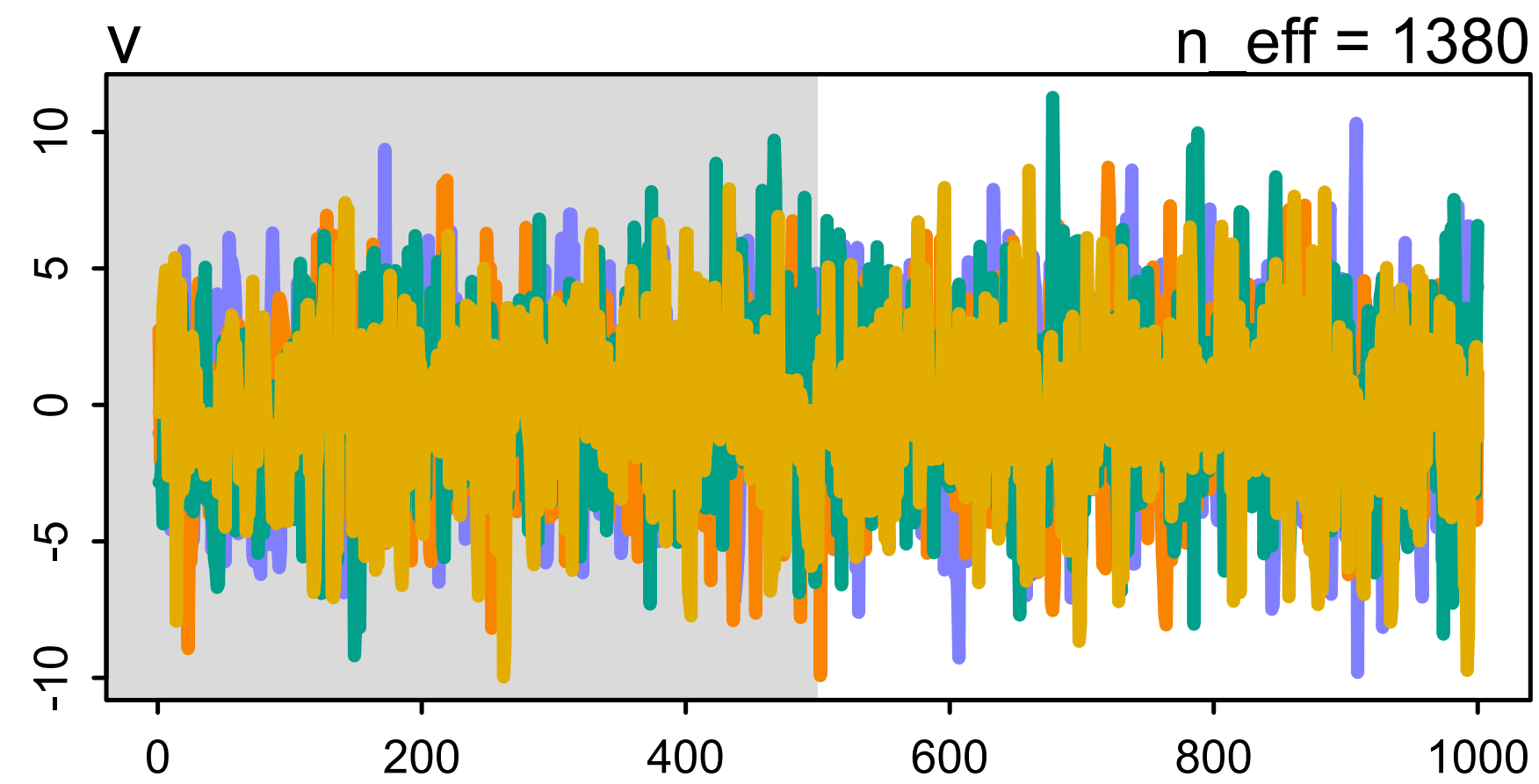
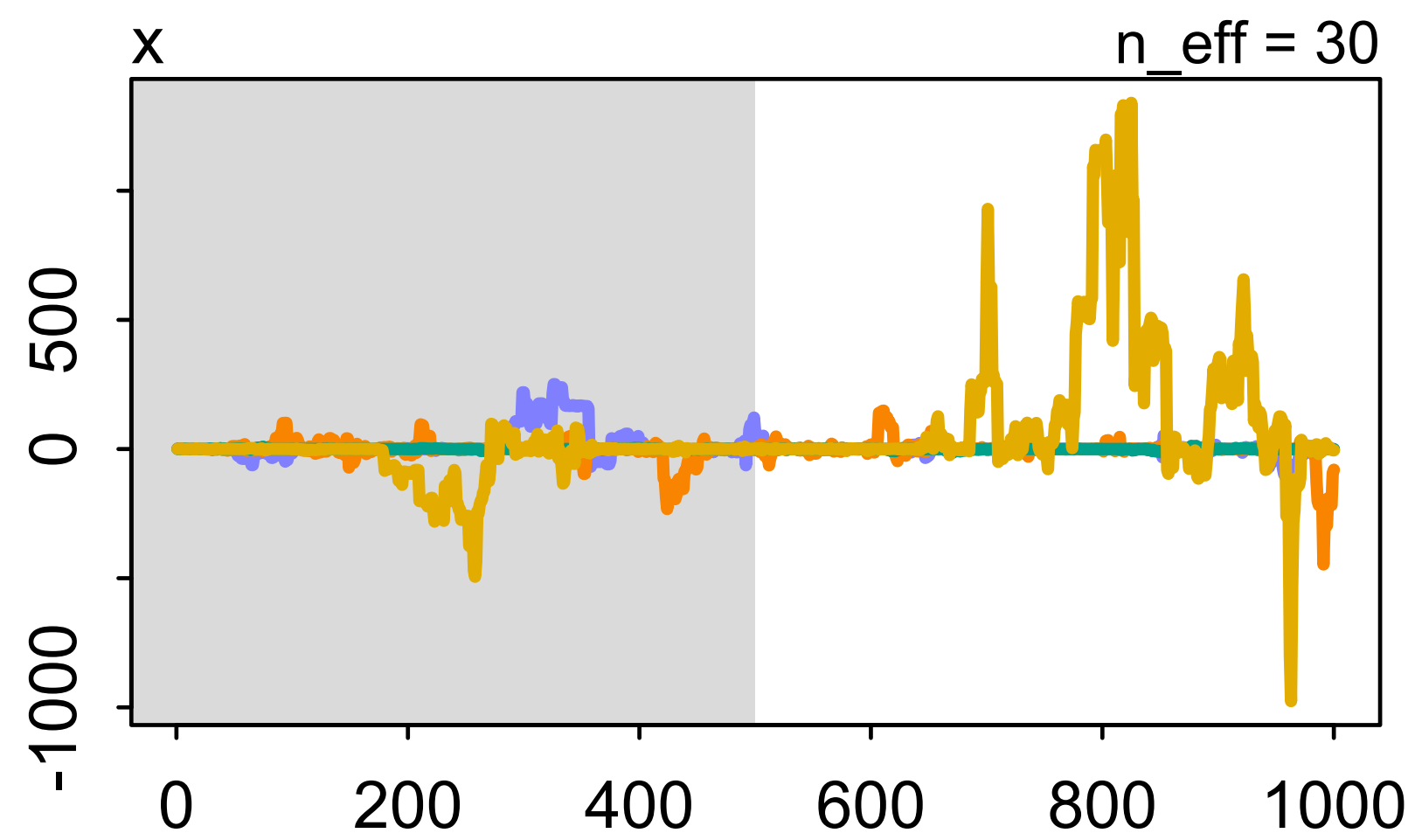
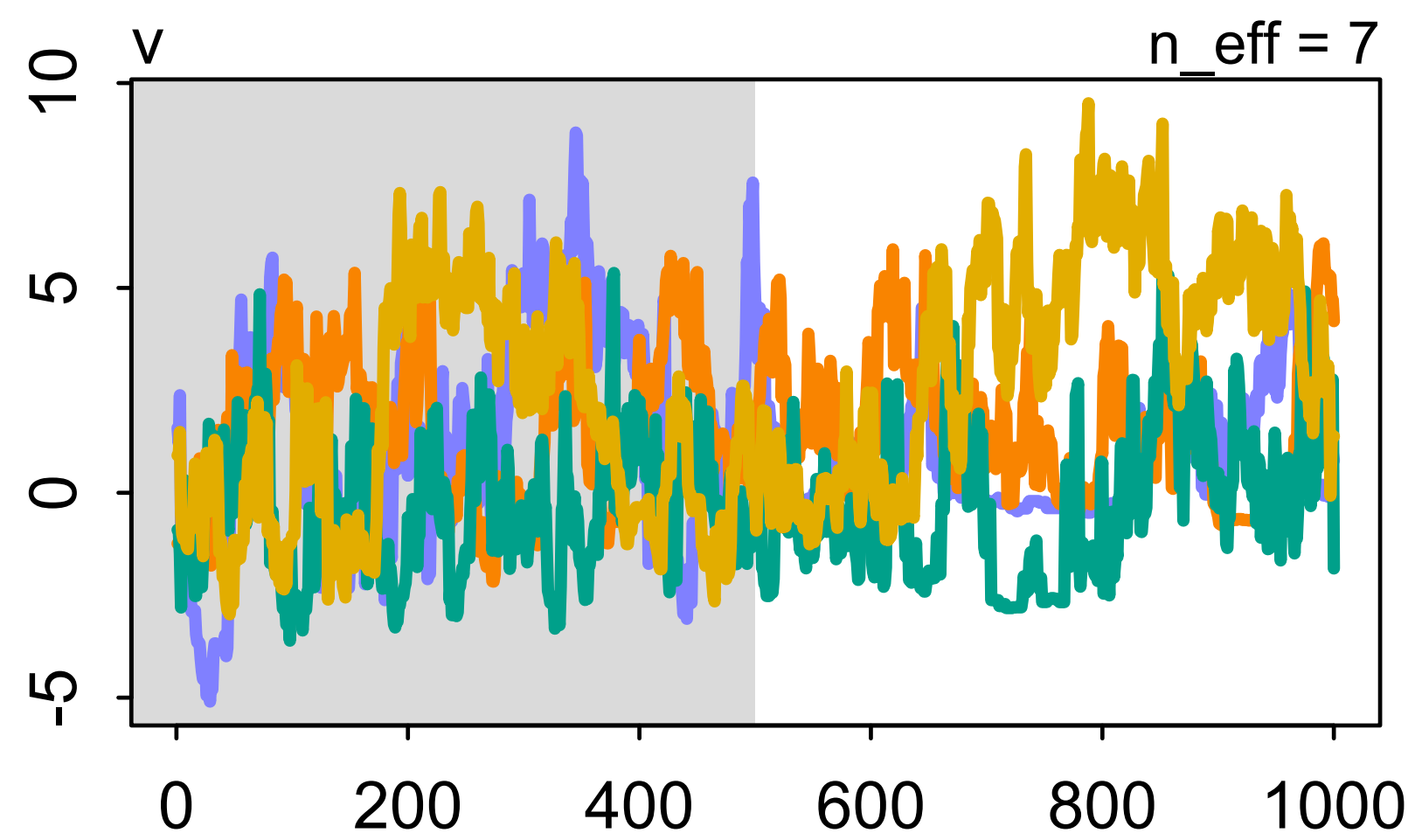
```
m13.7nc <- ulam(
  alist(
    v ~ normal(0,3),
    z ~ normal(0,1),
    gq> real[1]:x <<- z*exp(v)
  ), data=list(N=1) , chains=4 )
```

Warning: 112 of 2000 (6.0%) transitions ended with a divergence.

```
> precis( m13.7 )
      mean      sd  5.5%  94.5% n_eff Rhat4
v  1.41    2.37  -1.84   5.93    7   1.46
x 35.93 168.42 -21.15 258.86   30   1.19
```

```
> precis( m13.7nc )
      mean      sd  5.5%  94.5% n_eff Rhat4
v  -0.04    3.12  -5.17   4.84 1380    1
z  -0.01    0.96  -1.60   1.51 1495    1
x -19.34 899.98 -30.81 24.86 1874    1
```





Centered priors

$$C_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]} U_i$$

$$[\alpha_j, \beta_j] \sim \text{MVNormal}([\bar{\alpha}, \bar{\beta}], \mathbf{R}, [\sigma, \tau])$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0, 1)$$

$$\sigma, \tau \sim \text{Exponential}(1)$$

$$\mathbf{R} \sim \text{LKJCorr}(4)$$

Non-centered priors

$$C_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]} U_i$$

$$\alpha = \bar{\alpha} + v_{-,1}$$

$$\beta = \bar{\beta} + v_{-,2}$$

$$v = (\text{diag}(\sigma) \mathbf{LZ})^\top$$

$$\mathbf{Z}_{j,k} \sim \text{Normal}(0, 1)$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0, 1)$$

$$\sigma \sim \text{Exponential}(1)$$

$$\mathbf{R} \sim \text{LKJCorr}(4)$$

$$C_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha_{D[i]} + \beta_{D[i]} U_i$$

$$\alpha = \bar{\alpha} + v_{-,1}$$

$$\beta = \bar{\beta} + v_{-,2}$$

$$v = (\text{diag}(\sigma)\mathbf{LZ})^\top$$

$$\mathbf{Z}_{j,k} \sim \text{Normal}(0,1)$$

$$\bar{\alpha}, \bar{\beta} \sim \text{Normal}(0,1)$$

$$\sigma \sim \text{Exponential}(1)$$

$$\mathbf{R} \sim \text{LKJCorr}(4)$$

$$v = \left(\begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix} \mathbf{LZ} \right)^\top$$

Matrix v is back on non-standard, correlated scale

a 61-by-2 matrix $\mathbf{v} = \left(\begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix} \mathbf{LZ} \right)^\top$

a 61-by-2 matrix $\mathbf{v} = \left(\overbrace{\begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix}}^{\text{diagonal matrix of standard deviations}} \mathbf{LZ} \right)^T$

a 61-by-2 matrix $\mathbf{v} = \left(\begin{array}{c} \overbrace{\begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix}}^{\text{diagonal matrix of standard deviations}} \mathbf{LZ} \end{array} \right)^T$

Cholesky factor of correlation matrix across features

a 61-by-2 matrix $v = \left(\begin{array}{c} \overbrace{\begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix}}^{\text{diagonal matrix of standard deviations}} \mathbf{LZ} \end{array} \right)^T$

Cholesky factor of correlation matrix across features

2-by-61 matrix of feature z-scores

a 61-by-2 matrix $v =$

diagonal matrix of standard deviations

$$\left(\begin{array}{c} \overline{\left[\begin{array}{cc} \sigma_1 & 0 \\ 0 & \sigma_2 \end{array} \right]} \mathbf{LZ} \end{array} \right)^T$$

transpose! flips rows and columns

2-by-61 matrix of feature z-scores

Cholesky factor of correlation matrix across features

I. — NOTICES SCIENTIFIQUES

Commandant BENOIT¹.

NOTE SUR UNE MÉTHODE DE RÉOLUTION DES ÉQUATIONS NORMALES PROVENANT DE L'APPLICATION DE LA MÉTHODE DES MOINDRES CARRÉS A UN SYSTÈME D'ÉQUATIONS LINÉAIRES EN NOMBRE INFÉRIEUR A CELUI DES INCONNUES. — APPLICATION DE LA MÉTHODE A LA RÉOLUTION D'UN SYSTÈME DEFINI D'ÉQUATIONS LINÉAIRES.

(Procédé du Commandant CHOLESKY².)

Le Commandant d'Artillerie Cholesky, du Service géographique de l'Armée, tué pendant la grande guerre, a imaginé, au cours de recherches sur la compensation des réseaux géodésiques, un procédé très ingénieux de résolution des équations dites *normales*, obtenues par application de la méthode des moindres carrés à des équations linéaires en nombre inférieur à celui des inconnues. Il en a conclu une méthode générale de résolution des équations linéaires.

Nous suivrons, pour la démonstration de cette méthode, la progression même qui a servi au Commandant Cholesky pour l'imaginer.



André-Louis Cholesky (1875–1918)

I. — NOTICES SCIENTIFIQUES

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André-Louis Cholesky (1875–1918)

I. — NOTICES SCIENTIFIQUES

Commandant BENOIT¹.

Note on a method for solving normal equations resulting from the application of the method of least squares to a system of linear equations of smaller number than the unknowns.

(Procédé du Commandant CHOLESKY².)

The artillery commander Cholesky, of the Geographical Service of the army, killed during the Great War, imagined, during research on the compensation of the geodesic networks, a very ingenious process of solving the equations known as normal, obtained by application of the method of the least squares to linear equations in lower number than that of the unknowns.



André-Louis Cholesky (1875–1918)

ques, un procédé très ingénieux de résolution des équations dites *normales*, obtenues par application de la méthode des

```
# define 2D Gaussian with correlation 0.6
N <- 1e4
sigma1 <- 2
sigma2 <- 0.5
rho <- 0.6

# independent z-scores
z1 <- rnorm( N )
z2 <- rnorm( N )

# use Cholesky to blend in correlation
a1 <- z1 * sigma1
a2 <- ( rho*z1 + sqrt( 1-rho^2 )*z2 )*sigma2
```

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# use Cholesky to blend in correlation
a1 <- z1 * sigma1
a2 <- ( rho*z1 + sqrt( 1-rho^2 )*z2 ) * sigma2
```

```
> cor(z1,z2)
[1] -0.0005542644
> cor(a1,a2)
[1] 0.5999334
> sd(a1)
[1] 1.997036
> sd(a2)
[1] 0.4989456
```

$$v = \left(\begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix} \mathbf{LZ} \right)^T$$

Transformed Priors

Both centered and non-centered better in different contexts

Centered: Lots of data in each cluster (data probability dominant)

Non-centered: Many clusters, sparse evidence (prior dominant)

