

# Even more hierarchical models

FW 891

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# Purpose

- Today we introduce a powerful extension of mixed effects models
  - Random slopes (aka varying effects models)
  - Repent for our earlier sins
- Adventures in covariance a la McElreath (2023)
  - Cover some math necessary for working with covariance matrices
- Simulate a varying effects problem
- Develop both centered and noncentered varying effects models in Stan

# Some references

- This lecture is drawing heavily on McElreath (2023), and much of the code and analyses are adapted from information in that text

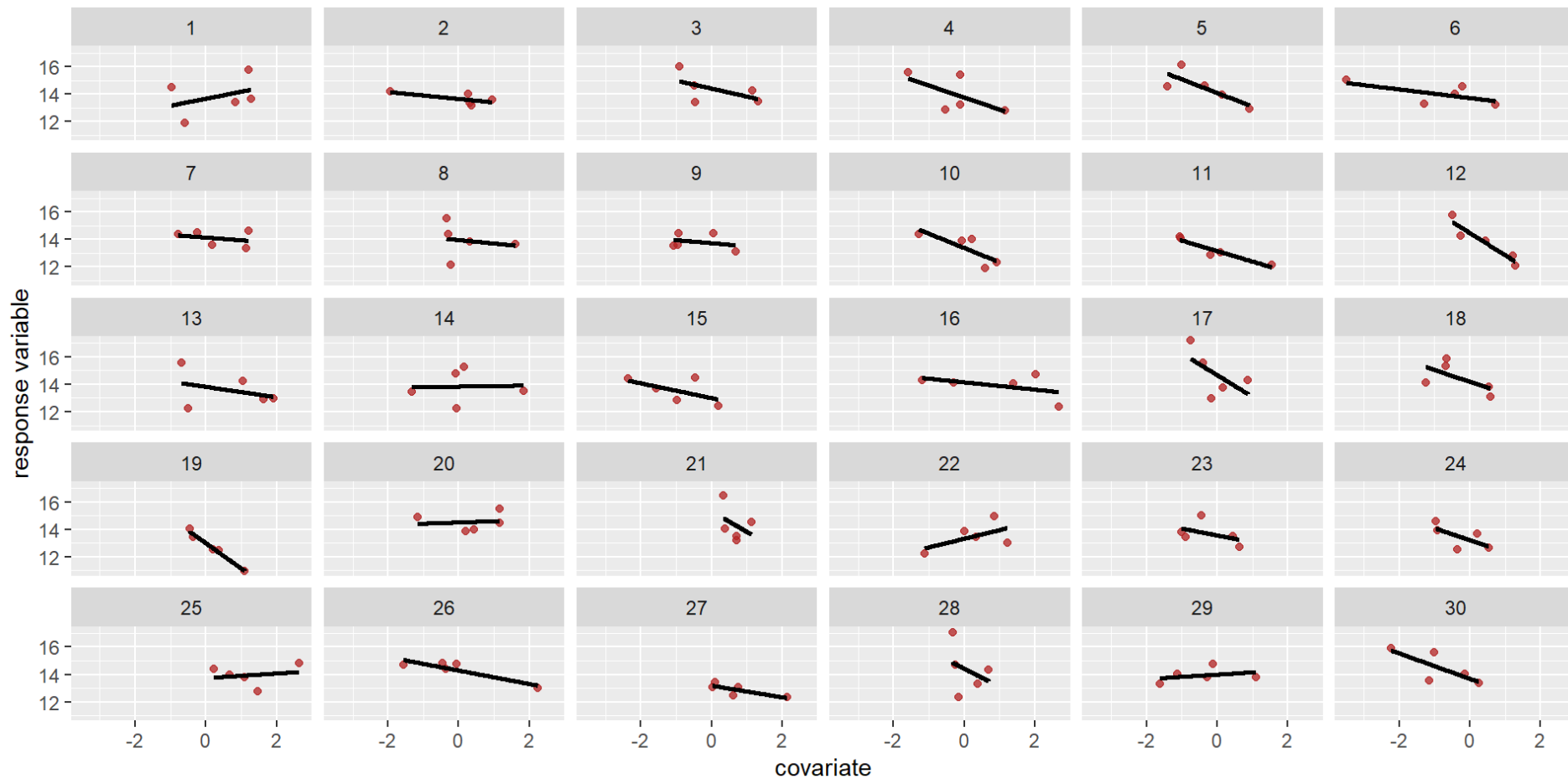
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- See also Gelman and Hill (2007)
  - Specifically chapter 13

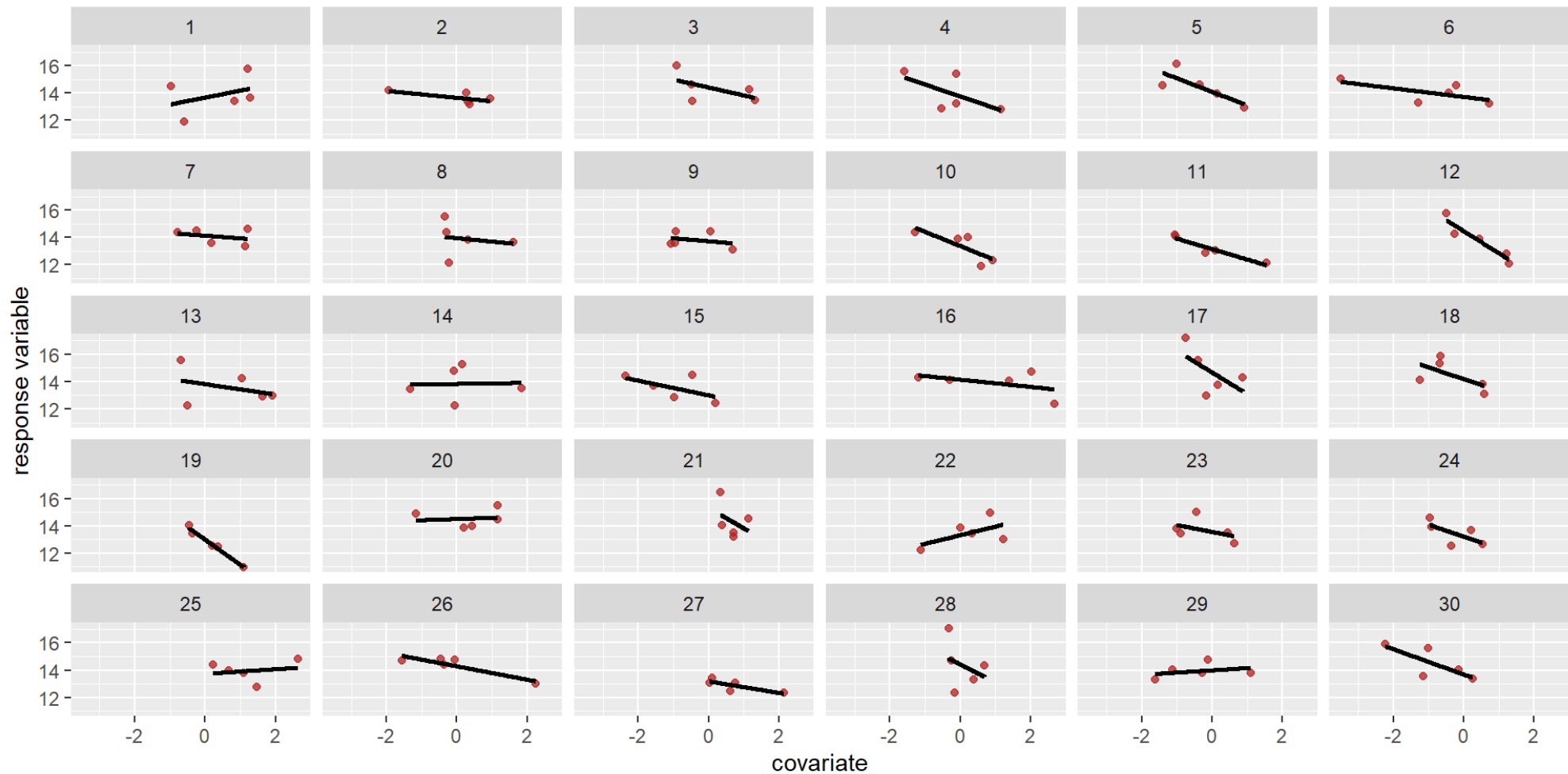
# Thinking about variability in ecological systems



# Three key takeaways

# Point #1

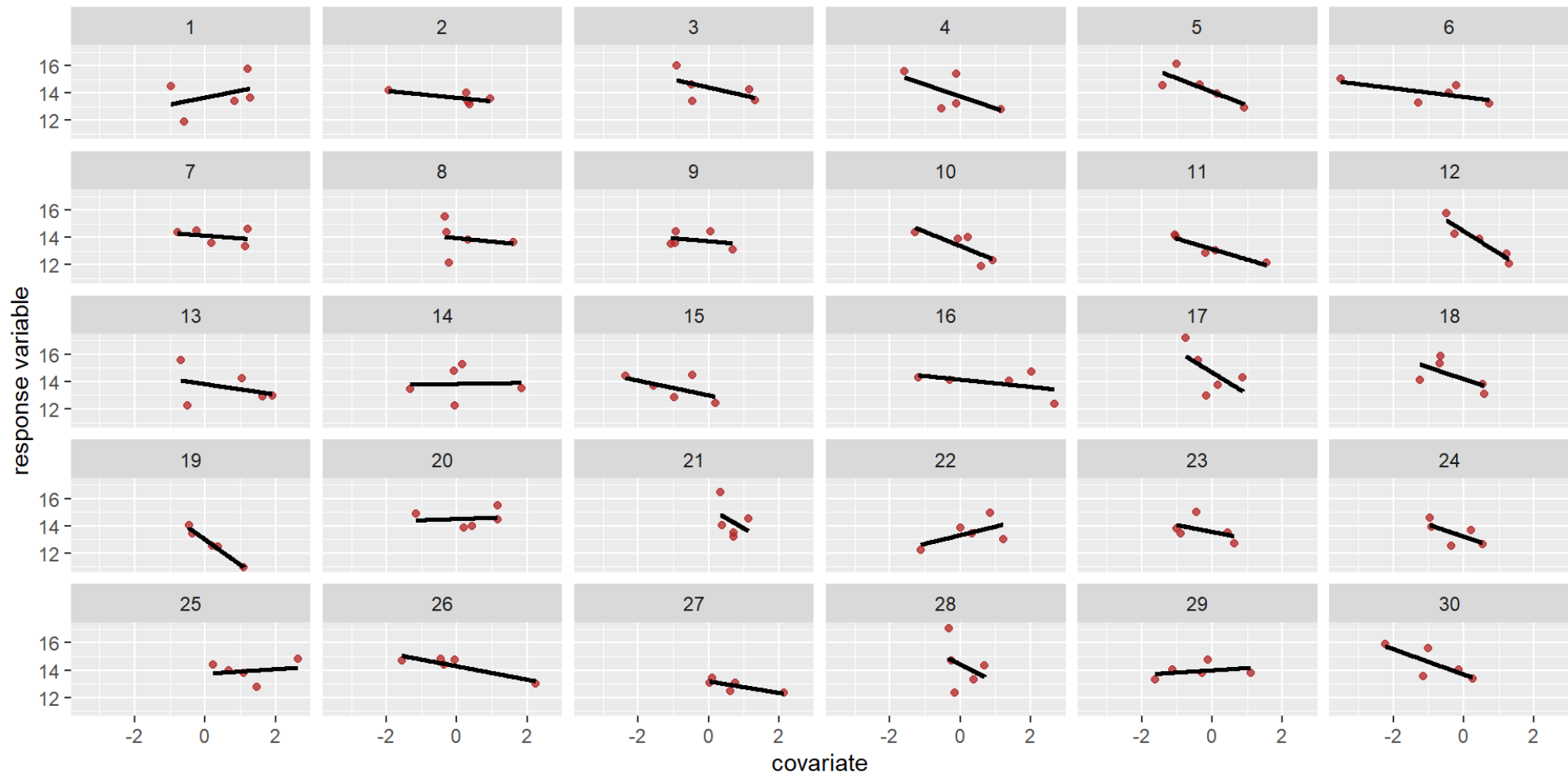
- variability in both intercepts *and* slopes among replicates





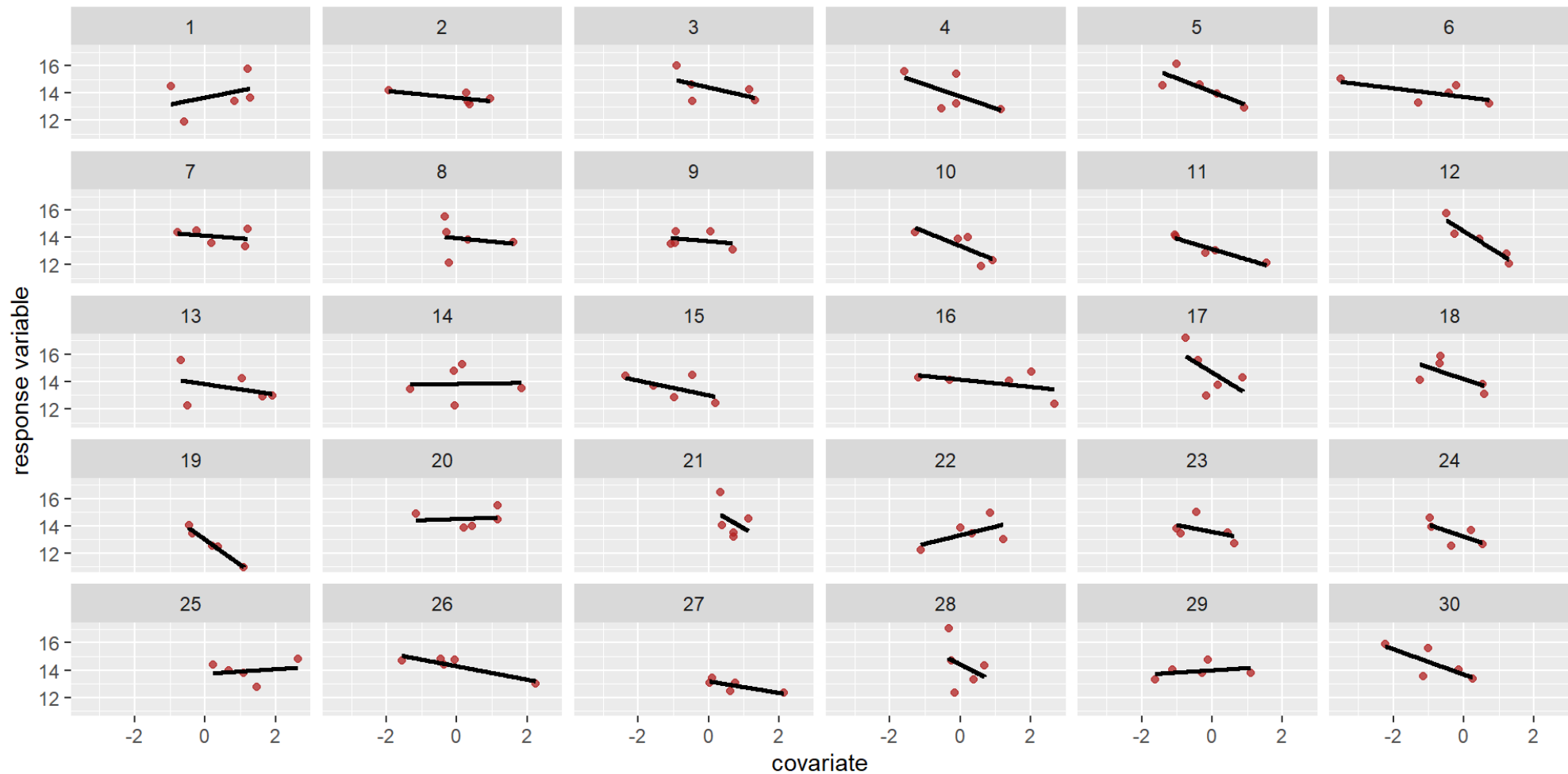
# Point #2

- slopes get steeper as intercepts get bigger



# Point #3

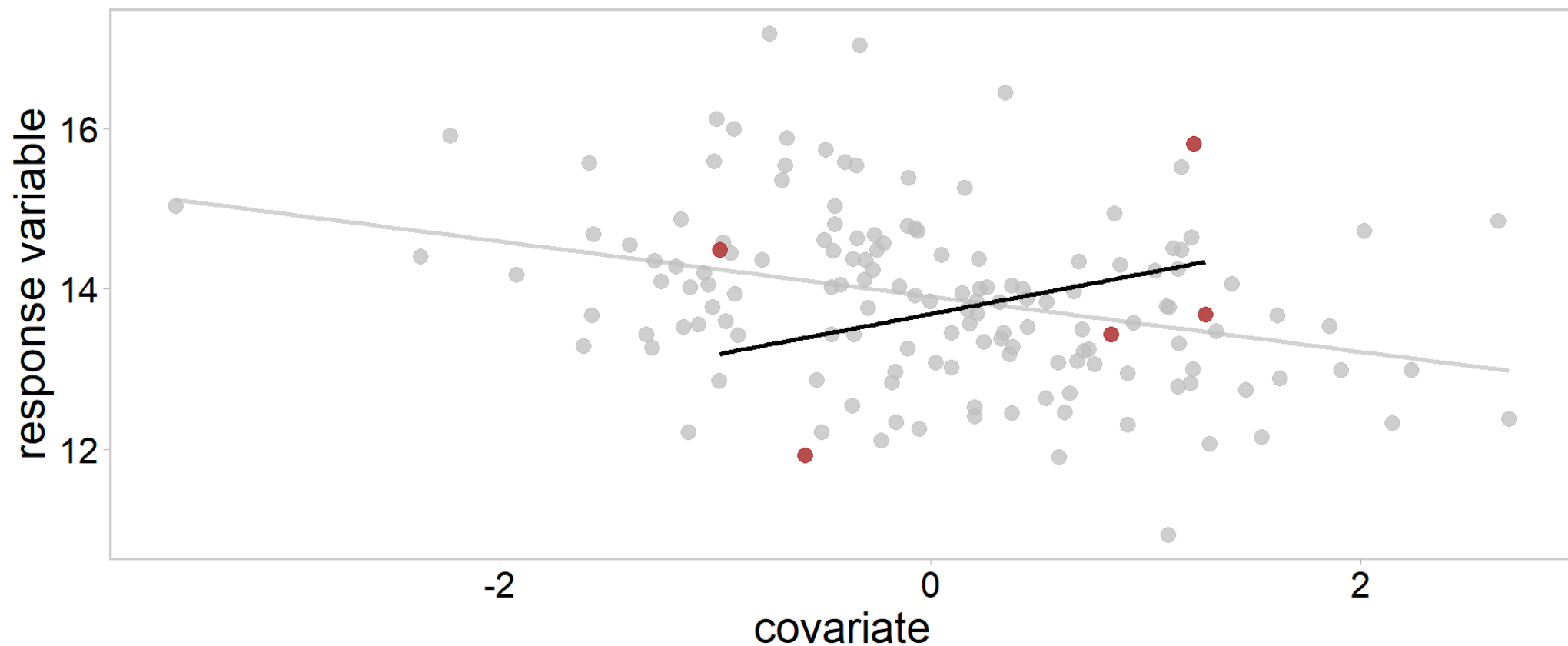
- some groups display **Simpson's paradox**



# Simpson's paradox

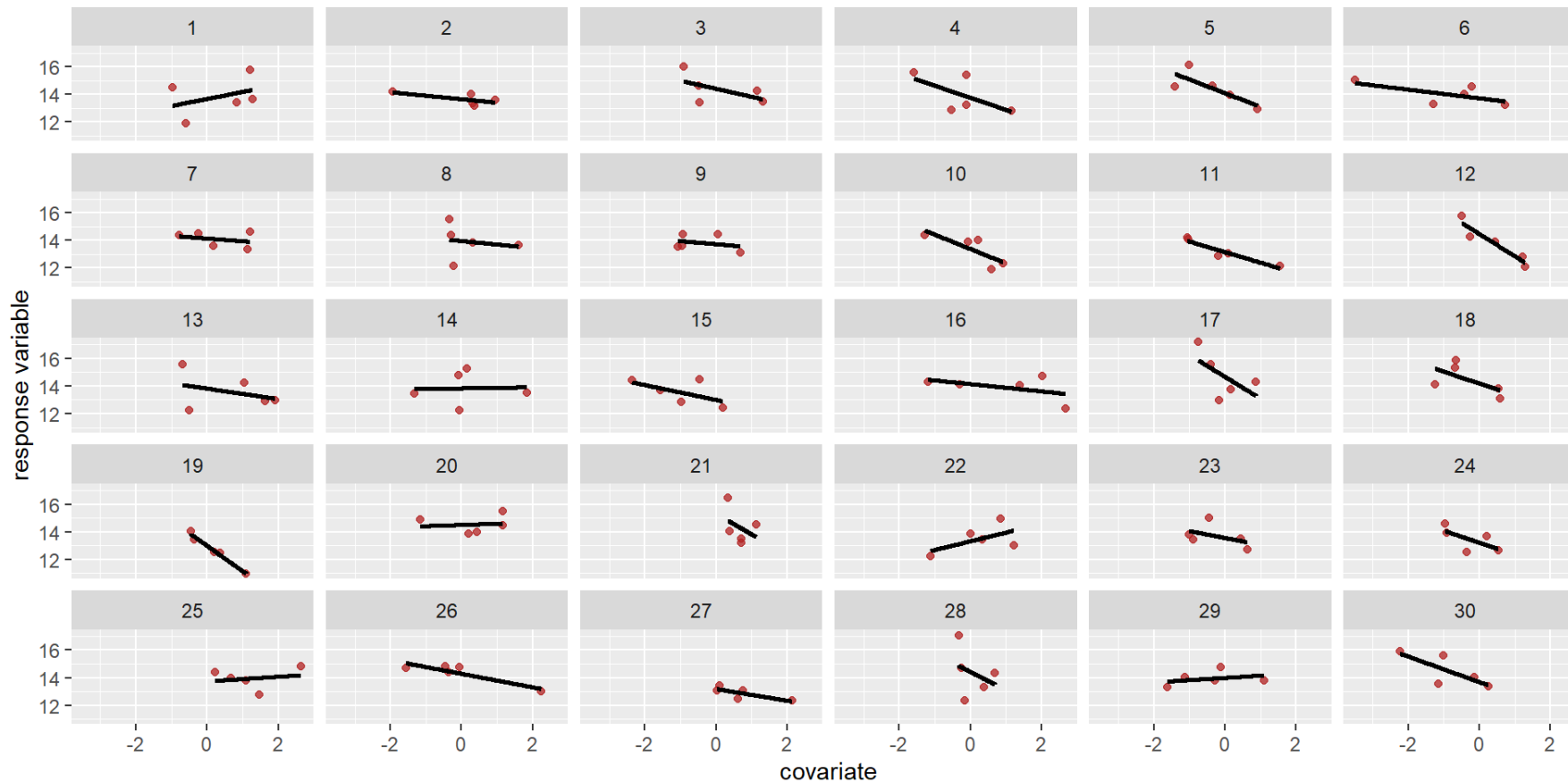
Colored points are data from replicate 1

Grey points are data from all other replicates



# Group question

- what is a question in your field of study that might show a similar pattern?



# Varying effects

- Generalization of standard multilevel regression
  - Specifically, models that allow slopes and intercepts to vary by group
- Several ways to write, here's one for the model we just visualized

# Varying effects maths

$$y_i \sim \text{N}(\mu_i, \sigma) \quad [\text{likelihood}]$$

$$\mu_i = \beta_{0[\textit{group}]} + \beta_{1[\textit{group}]} x_{1[i]} \quad [\text{linear model}]$$

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$$\begin{bmatrix} \beta_{0_{group}} \\ \beta_{1_{group}} \end{bmatrix} \sim \text{MVN} \left( \begin{bmatrix} \beta_0 \\ \beta_1 \end{bmatrix}, \Sigma \right) [\text{population of varying effects}]$$

$$\Sigma = \begin{pmatrix} \sigma_{\beta_0} & 0 \\ 0 & \sigma_{\beta_1} \end{pmatrix} \mathbf{\Omega} \begin{pmatrix} \sigma_{\beta_0} & 0 \\ 0 & \sigma_{\beta_1} \end{pmatrix} [\text{construct covariance matrix}]$$

adapted from McElreath 2023



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# Let's break that down

- Math on the previous slide says that each group has a  $\beta_{0[group]}$  and  $\beta_{1[group]}$  with a prior distribution defined by the two dimensional Gaussian distribution with means  $\beta_0$  and  $\beta_1$  and covariance matrix  $\Sigma$

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  - Several ways to construct  $\Sigma$ , but splitting it into standard deviations,  $\sigma_{\beta_0}$  and  $\sigma_{\beta_1}$ , and a correlation matrix  $\Omega$  helps with learning
- Compare this with a standard normal distribution which takes a mean and a standard deviation

# The correlation matrix

- For this simple example, the correlation matrix looks like

$$\mathbf{\Omega} = \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix}$$

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- where  $\rho$  is the correlation between  $\beta_0$  and  $\beta_1$
- $\mathbf{\Omega}$  gets more complicated for models with more varying slopes

# Cholesky decomposition

- Note that we can take any arbitrary symmetric, positive-definite matrix  $\mathbf{A}$ , and factor or decompose it into

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- where  $\mathbf{L}$  is a lower triangular matrix with real and positive diagonal entries and  $\mathbf{L}^T$  is a transpose of  $\mathbf{L}$

# Cholesky Decomposition

- If we visualize a Cholesky decomposition

$$\begin{bmatrix} A_{00} & A_{01} & A_{02} \\ A_{10} & A_{11} & A_{12} \\ A_{20} & A_{21} & A_{22} \end{bmatrix} = \begin{bmatrix} L_{00} & 0 & 0 \\ L_{10} & L_{11} & 0 \\ L_{20} & L_{21} & L_{22} \end{bmatrix} \begin{bmatrix} L_{00} & L_{10} & L_{20} \\ 0 & L_{11} & L_{21} \\ 0 & 0 & L_{22} \end{bmatrix}$$

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- This is helpful from a numerical perspective, particularly with noncentered parameterizations

# Cholesky factors continued

- Note that there is a lot of convenient linear algebra that can be done with Cholesky factors of covariance matrices  $\mathbf{L}$  or of correlation matrices  $\mathbf{L}_{\text{corr}}$ 
  - For example,

$$\mathbf{L} = \begin{pmatrix} \sigma_{\beta_0} & 0 \\ 0 & \sigma_{\beta_1} \end{pmatrix} \mathbf{L}_{\text{corr}}$$

- See this [link](#) for a useful review

# Cholesky factors

```
1 # create a correlation matrix and declare sigmas
2 OMEGA <- matrix(c(1, 0.7, 0.7, 1), nrow = 2)
3 sigmas <- c(1, 2) # sd_b0, sd_b1
4
5 OMEGA
```

```
      [,1] [,2]
[1,]  1.0  0.7
[2,]  0.7  1.0
```

```
1 sigmas
```

```
[1] 1 2
```

```
1 # note also
2 diag(sigmas) # diagonal matrix
```

```
      [,1] [,2]
[1,]    1    0
[2,]    0    2
```

# Cholesky factors

```
1 # calculate covariance matrix:
2 SIGMA <- diag(sigmas) %*% OMEGA %*% diag(sigmas)
3 SIGMA
```

```
      [,1] [,2]
[1,]  1.0  1.4
[2,]  1.4  4.0
```



# Cholesky factors

```
1 # convert Cholesky factor of correlation matrix
2 # to covariance Cholesky factor
3 L_corr <- t(chol(OMEGA)) # note chol() returns upper triangular matrix
4 diag(sigmas) %*% L_corr
```

```
      [,1]      [,2]
[1,]  1.0  0.000000
[2,]  1.4  1.428286
```

```
1 t(chol(SIGMA)) # L of SIGMA
```

```
      [,1]      [,2]
[1,]  1.0  0.000000
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```

# Cholesky factors

```
1 # Cholesky factor of correlation matrix to
2 # covariance matrix Cholesky factor:
3 L_corr <- t(chol(OMEGA)) # note chol() returns upper tri
4 Lambda <- diag(sigmas) %*% L_corr
5
6 t(chol(SIGMA)) # L of SIGMA
```

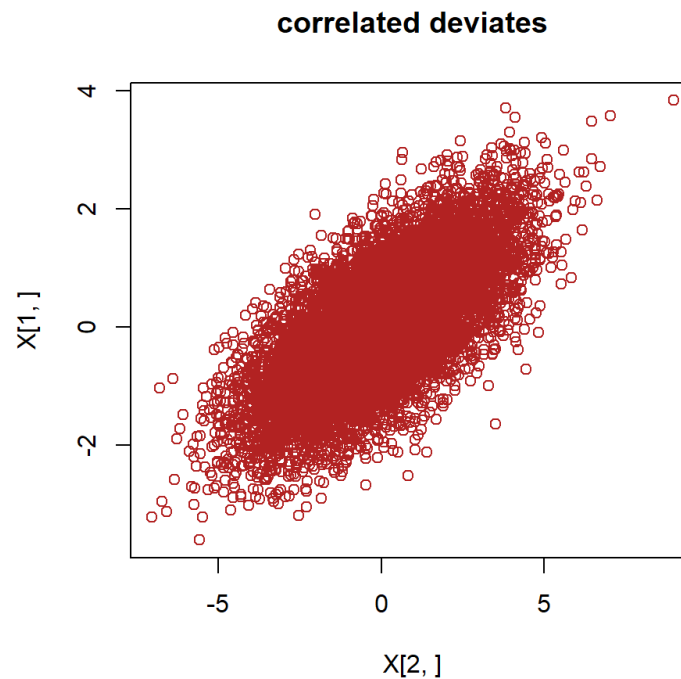
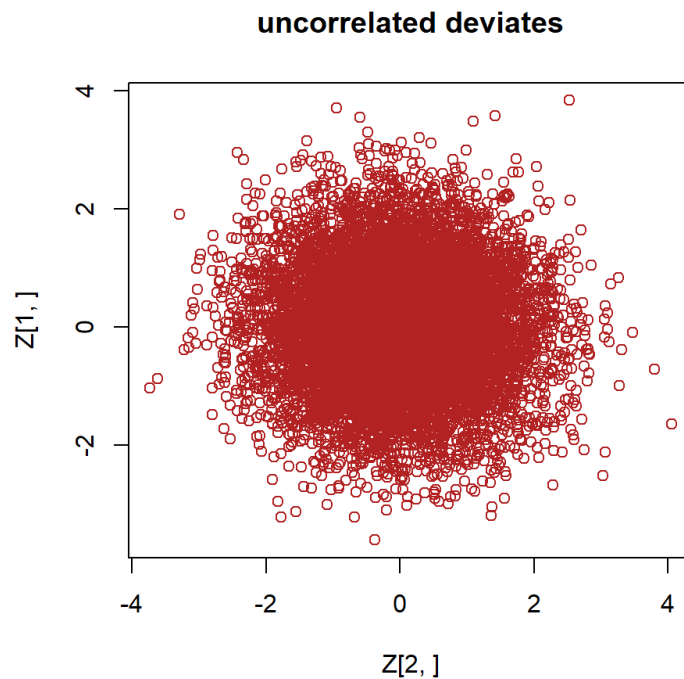
```
      [,1]      [,2]
[1,]  1.0 0.000000
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```

```
1 Lambda
```

```
      [,1]      [,2]
[1,]  1.0 0.000000
[2,]  1.4 1.428286
```

# Cholesky factors

```
1 # generate random values with desired covariance
2 Z <- rbind(rnorm(1e4), rnorm(1e4))
3 X <- Lambda %*% Z
4 par(mfrow=c(1,2))
5 plot(Z[,1]~Z[,2], main = "uncorrelated deviates", col = "firebrick")
6 plot(X[,1]~X[,2], main = "correlated deviates", col = "firebrick")
```



# A problem

- People want to know the extent to which juvenile walleye growth rate is density dependent
  - Has implications for both basic ecology and management
- DNR Biologists go to a collection of lakes and measure length of age-0 walleye in fall as a proxy of juvenile growth rate
- Each year, the biologists attempt to go to 30 lakes in total (weather pending)
  - They also conduct surveys to get an estimate of juvenile density
- Let's simulate some fake data representing this problem, and then build some Stan models to recover
- go to the `varying_effects.r` script

# Hyperpriors for varying effects model

$\beta_0 \sim \text{Normal}(0, 25)$  [prior for average intercept]

$\beta_1 \sim \text{Normal}(0, 25)$  [prior for average slope]

$\sigma \sim \text{Exponential}(0.01)$  [prior for stddev within group]

$\sigma_{\beta_0} \sim \text{Exponential}(0.01)$  [prior stddev among intercepts]

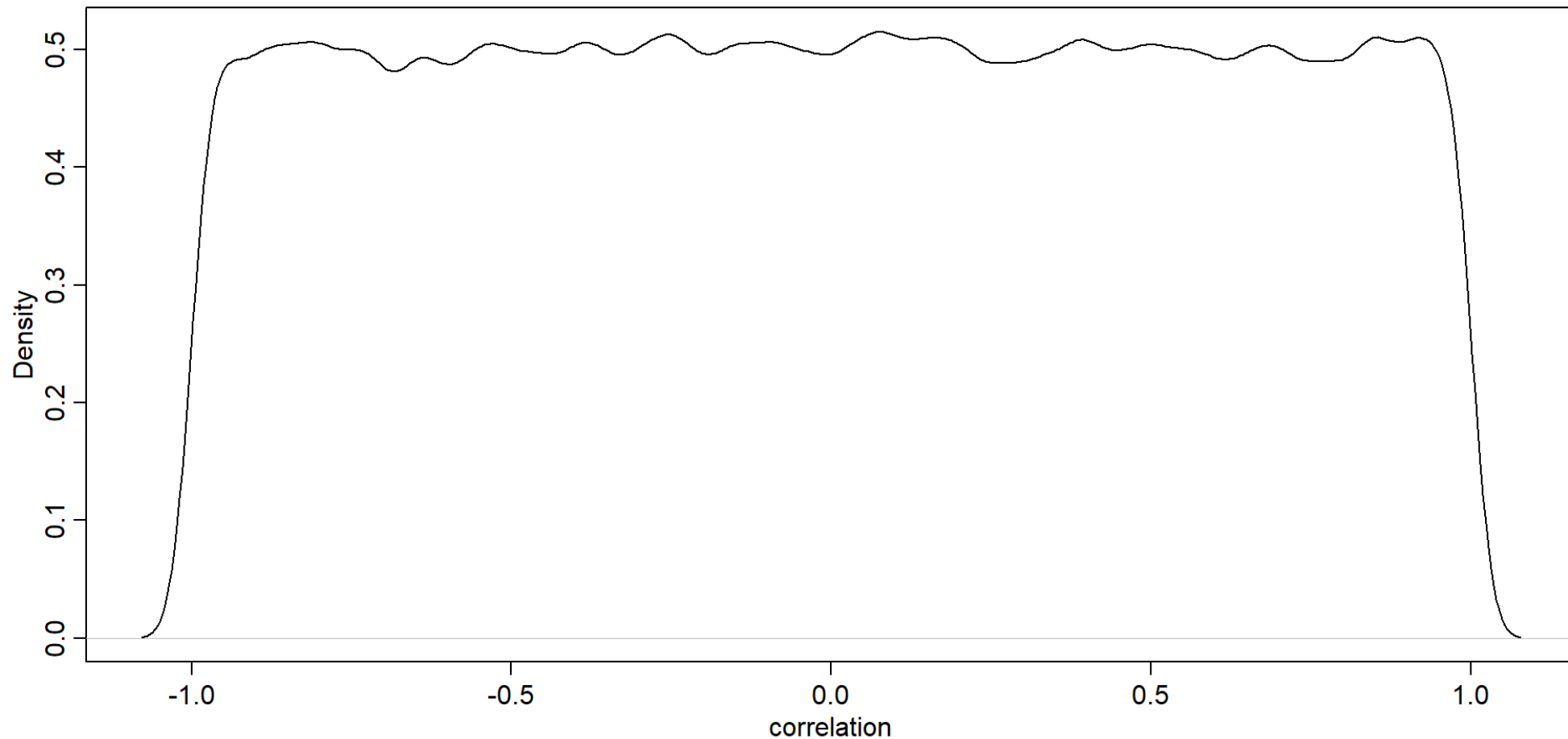
$\sigma_{\beta_1} \sim \text{Exponential}(0.01)$  [prior stddev among intercepts]

$\mathbf{\Omega} \sim \text{LKJcorr}(2)$  [prior for correlation matrix]

- LKJcorr(2) defines a weakly informative prior on  $\rho$  that is skeptical of extreme correlations near -1 or 1

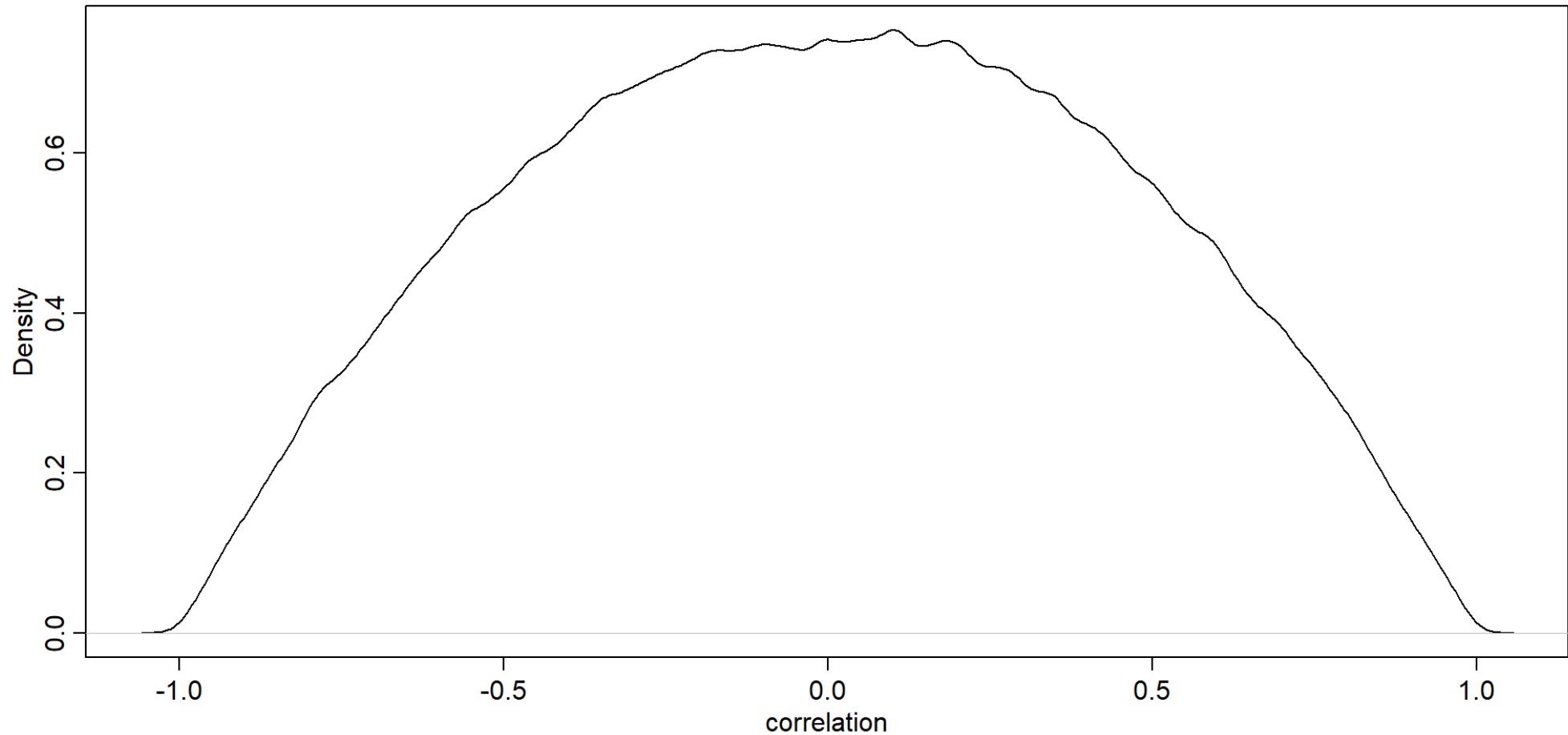
# Visualizing the LKJcorr prior

- LKJcorr(1)



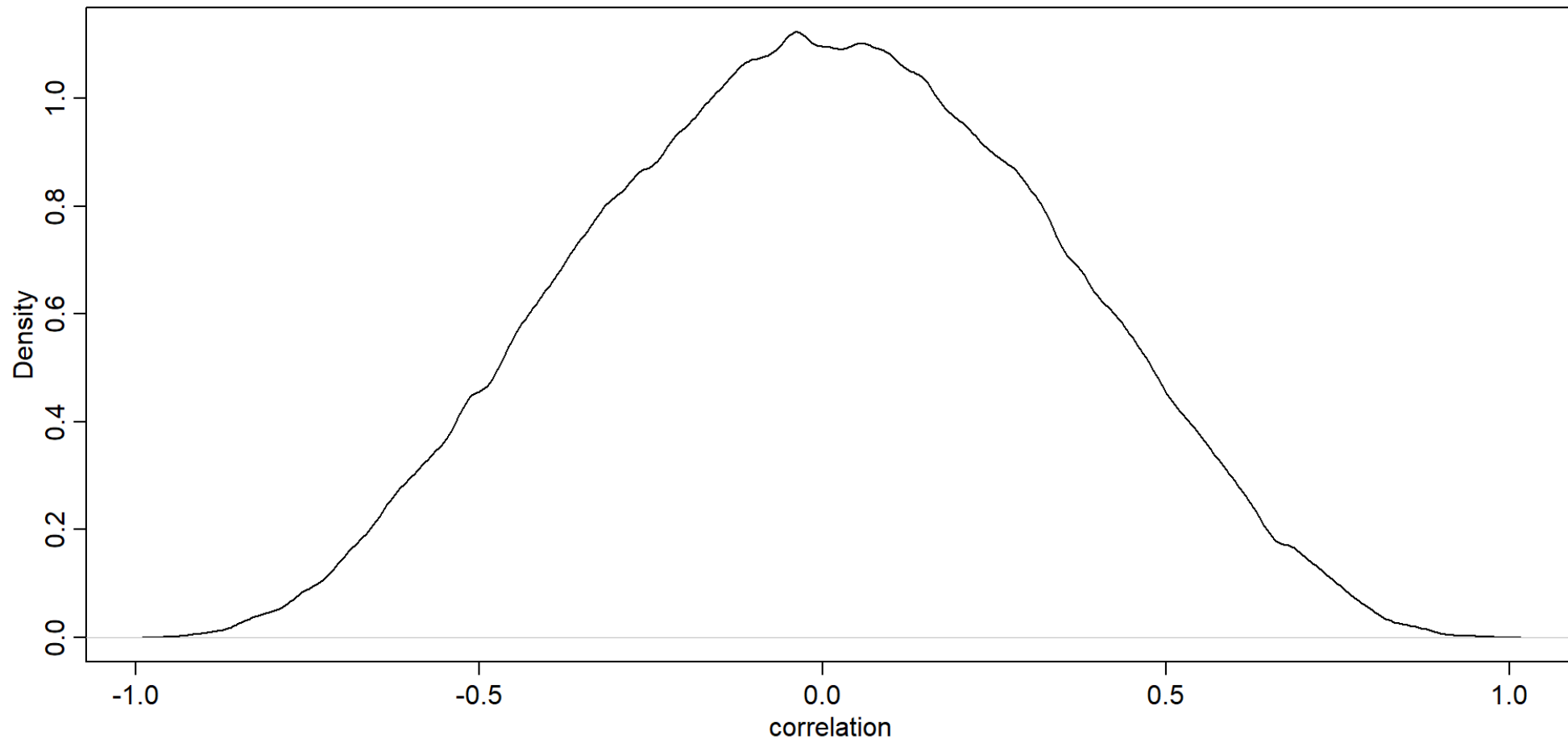
# Visualizing the LKJcorr prior

- LKJcorr(2)



# Visualizing the LKJcorr prior

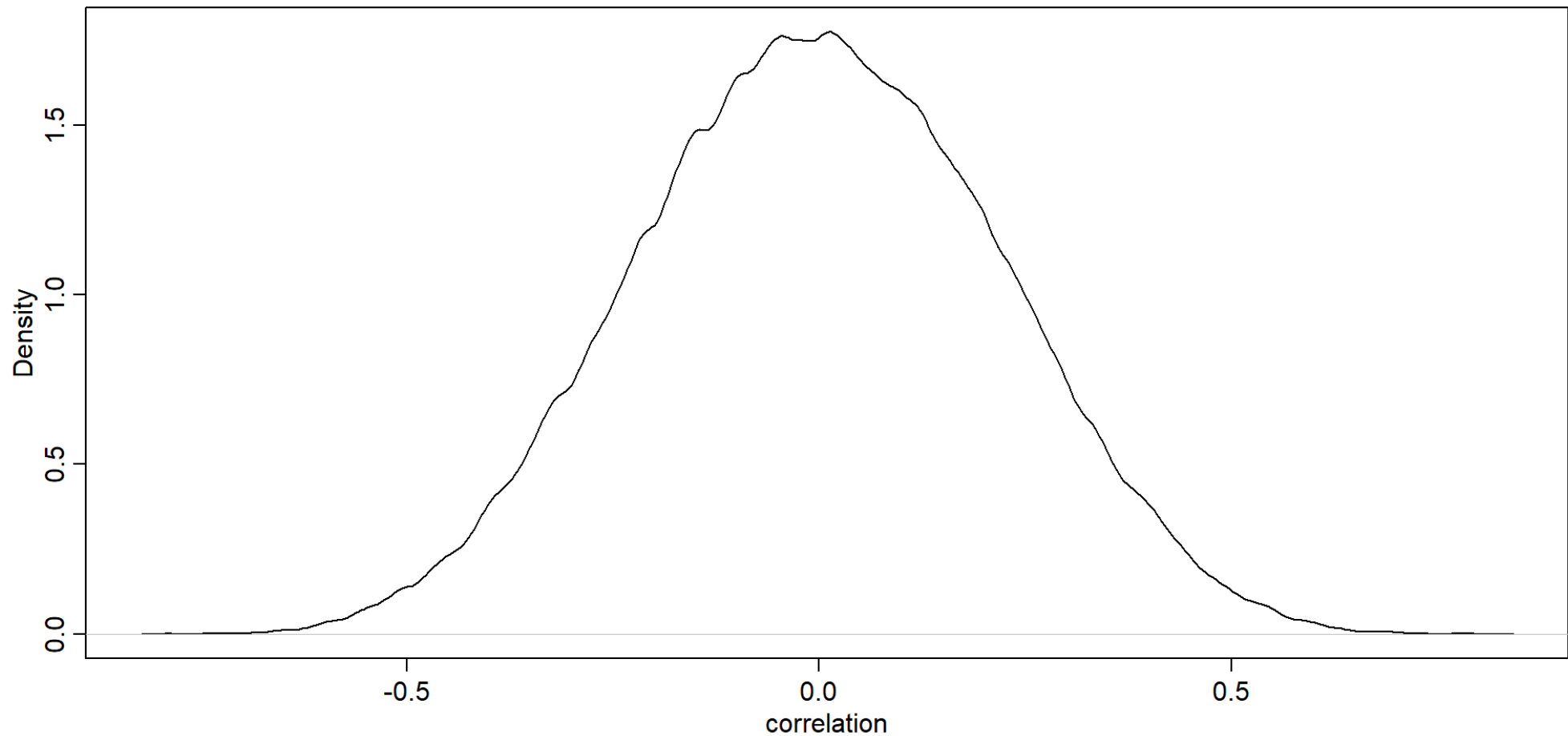
- LKJcorr(4)





# Visualizing the LKJcorr prior

- LKJcorr(10)



# Why go through all of that



# Wrap up

- Introduced a powerful extension to mixed effects models: varying effects
- Went through a bunch of math to show how to play with multivariate normal distributions
- Simulated an example, conducted prior predictive checks, estimated the model
- Showed a non-centered version of this model as well
- Up next, spatial random effects

# References

Cahill et al. 2020. A spatial-temporal approach to modeling somatic growth across inland fisheries landscapes. CJFAS.

Gelman, A. and J. Hill. 2007. Data analysis using regression and multilevel/hierarchical models

McElreath 2023. Statistical Rethinking.

Simpson's paradox Wikipedia:

[https://en.wikipedia.org/wiki/Simpson%27s\\_paradox](https://en.wikipedia.org/wiki/Simpson%27s_paradox)

<https://mlisi.xyz/post/simulating-correlated-variables-with-the-cholesky-factorization/>