# Simple Linear Models

A jack of all trades?

# Why use probabilistic models?

What questions do we want to answer?

- Statistical models should answer questions
  - What is the relation between two variables?
  - What is the difference between two groups?
  - What are the sources of variation in the data?
  - What is the expected result of an intervention?

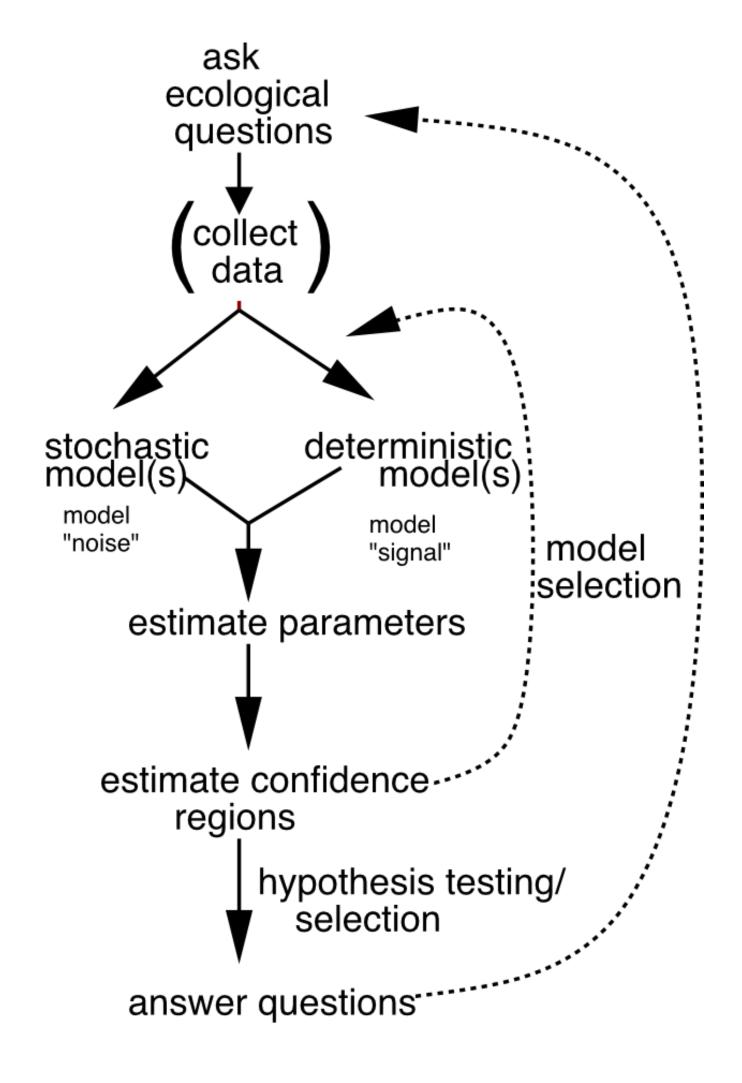
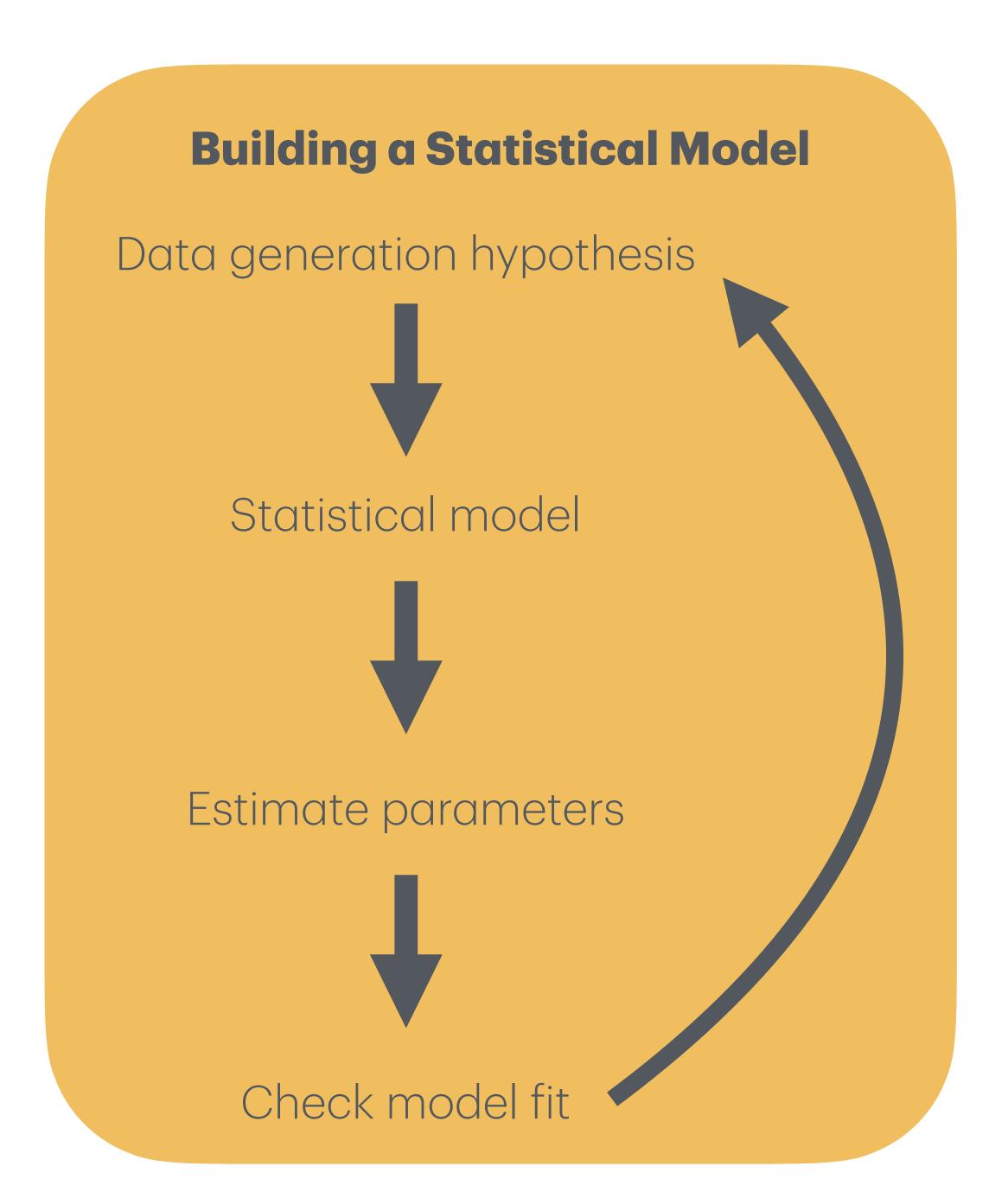


Figure 1.5 Flow of the modeling process.

## The basic model

#### Our hammer

- Our model must:
  - Define a relation between observations and parameters
  - Create a probabilistic description of the system we are studying
  - The description should capture the aspects of the data we are interested in!



## Probability definitions

#### Crash course

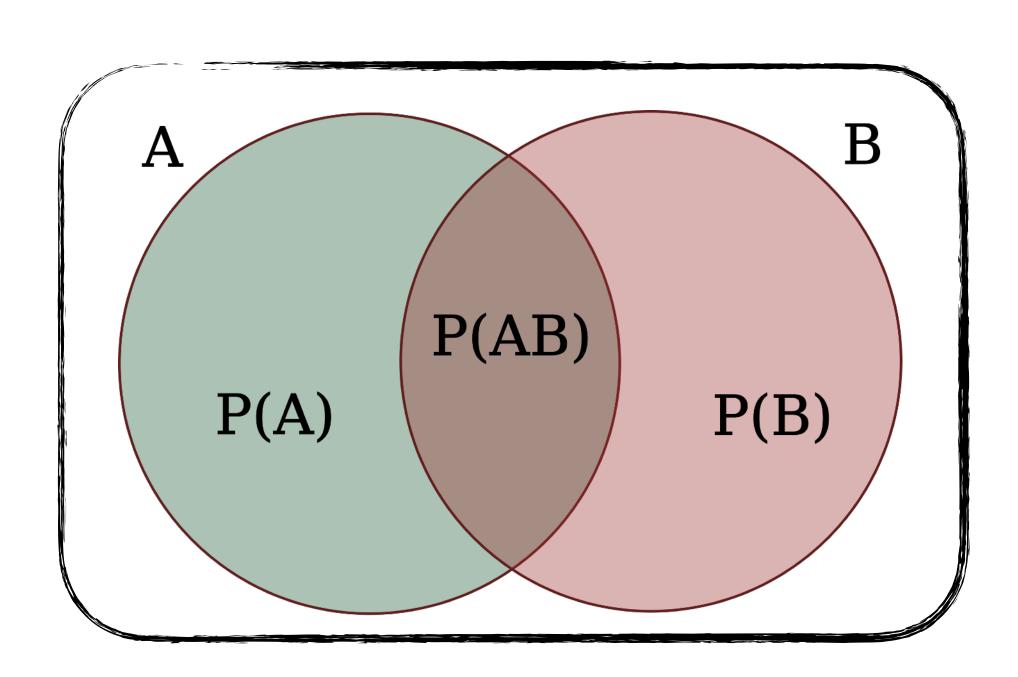
- We can express the probability of some variable as:
  - P(y): Read as the "probability of y"
- If the distribution of x depends on some parameters, we use the conditional probability:
  - P(y|x): Read as the "probability of y given x"

## Probability rules

- If two events, A and B, are independent:
  - $P(A \mid B) = P(A)$ : If A and B are independent, the probability of A given B is just the probability of A.
- More generally, we have the product rule:

• 
$$P(AB) = P(A)P(B|A) = P(B)P(A|B)$$

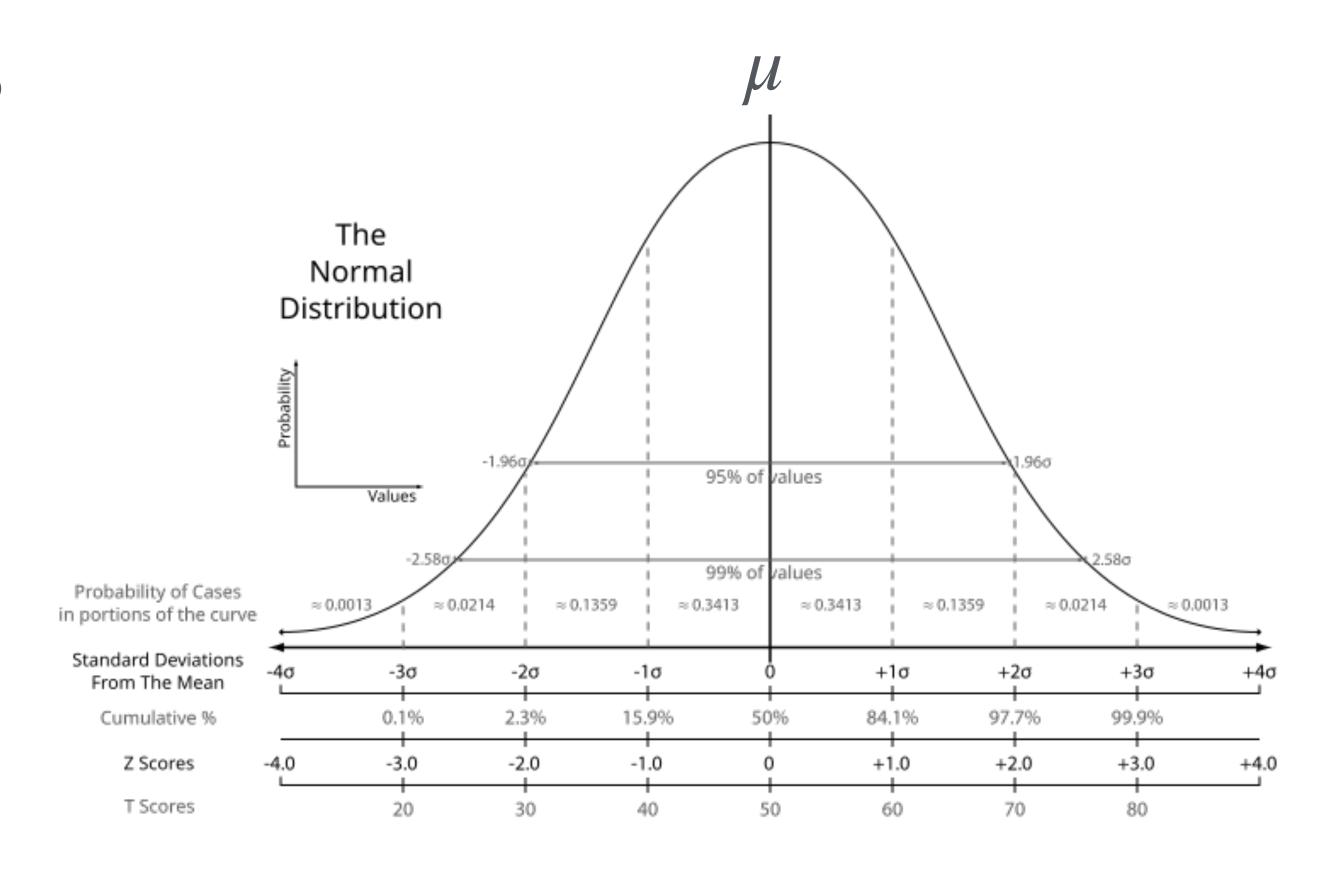
$$P(A \mid B) = \frac{P(A)P(B \mid A)}{P(B)}$$



## Probability distribution

#### Crash course

- We can use standard probability distributions to define these relations
- If a variable y follows a normal distribution:
  - $P(y) = P(y | \mu, \sigma) = Normal(y | \mu, \sigma)$
  - Where  $\mu$  and  $\sigma$  are parameters
    - $\mu$ : is the mean, a location parameter
    - $\sigma$ : sigma is the standard deviation, a scale parameter

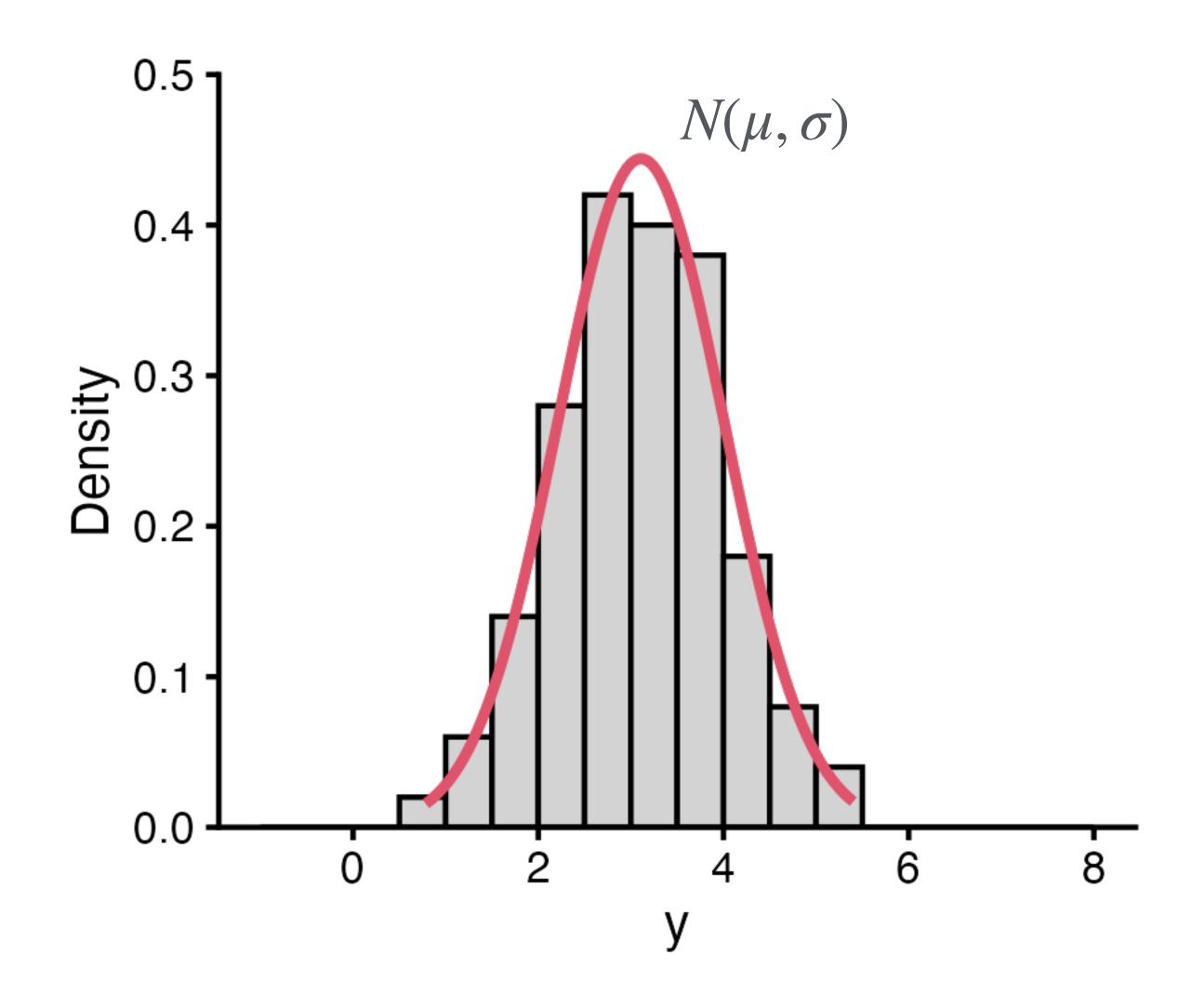


## Simplest probabilistic model

Fit parameters to a set of measurements

- Measure a set of  $y_i$  values:
- Find the "best fitting" normal distribution by choosing  $\mu$  and  $\sigma$  such that the N( $\mu$ ,  $\sigma$ ) distribution approximates the histogram of the  $y_i$  values

$$y_i \sim N(\mu, \sigma)$$



## The likelihood

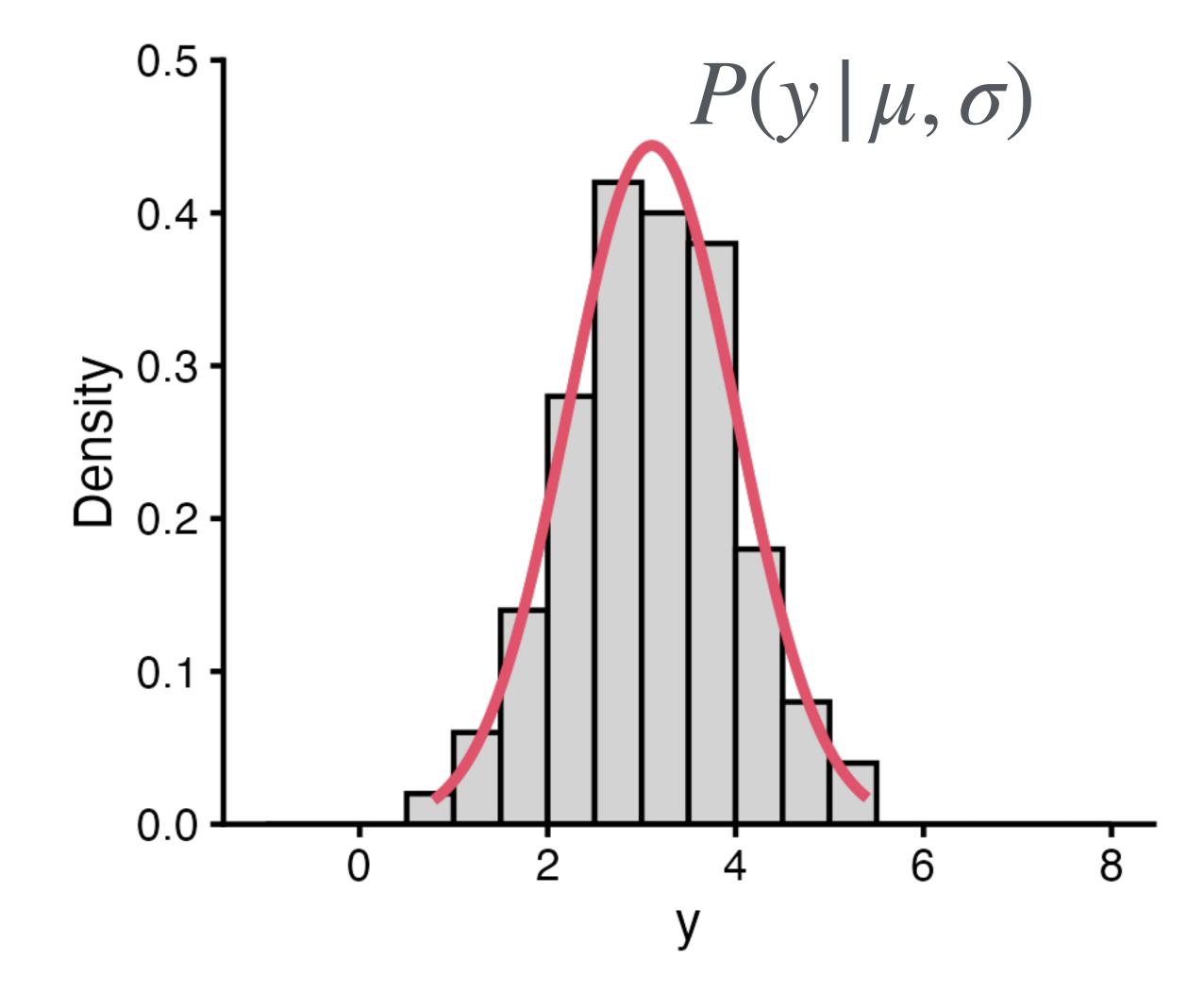
The probability of each value of y

• What does this mean?

$$y_i \sim N(\mu, \sigma)$$

• We can also write this as:

$$P(y | \mu, \sigma)$$
The likelihood of  $y$ 



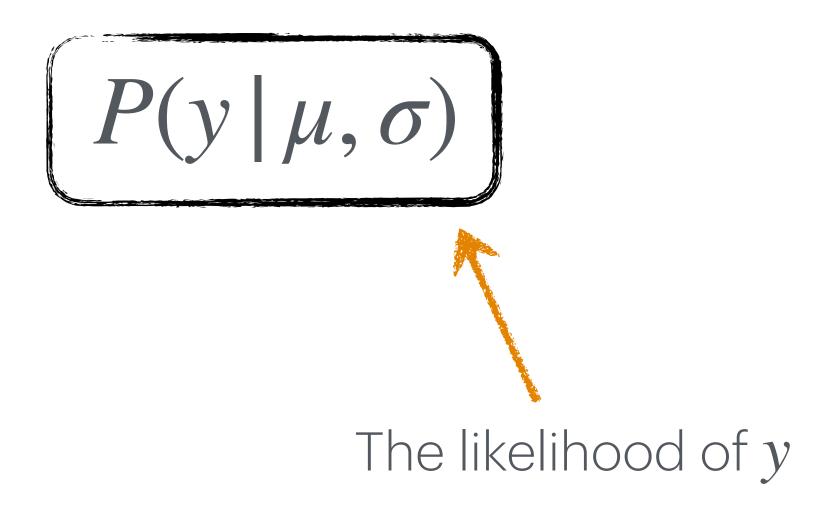
## The likelihood

The probability of each value of y

What does this mean?

$$y_i \sim N(\mu, \sigma)$$

• We can also write this as:



By the product rule:

$$P(\mu, \sigma | y) = \frac{P(y | \mu, \sigma)P(\mu, \sigma)}{P(y)}$$

- $P(\mu, \sigma) = P(\mu)P(\sigma)$ : the prior distribution
- $P(\mu, \sigma | y)$ : The posterior distribution
- P(y): A constant, the "evidence"

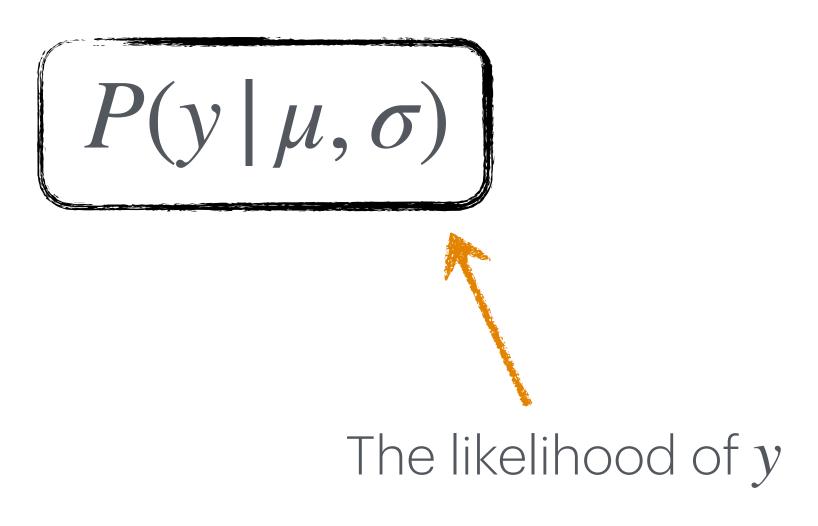
## The likelihood and friends

The probability of each value of y

What does this mean?

$$y_i \sim N(\mu, \sigma)$$

• We can also write this as:



By the product rule:

$$P(\mu, \sigma | y) \propto P(y | \mu, \sigma) P(\mu, \sigma)$$

- $P(\mu, \sigma) = P(\mu)P(\sigma)$ : the prior distribution
- $P(\mu, \sigma | y)$ : The posterior distribution
- P(y): A constant, the "evidence" Not necessary for inference, usually ignored

## The posterior distribution

### The encoding of our inference

- We can use the posterior distribution to understand what out data says about our parameter values
- By finding the posterior  $P(\mu, \sigma | y)$ , we can infer the most probable values for the parameters.
- So, we just need to define the ingredients:
  - $P(\mu), P(\sigma), P(y_i | \mu, \sigma)$

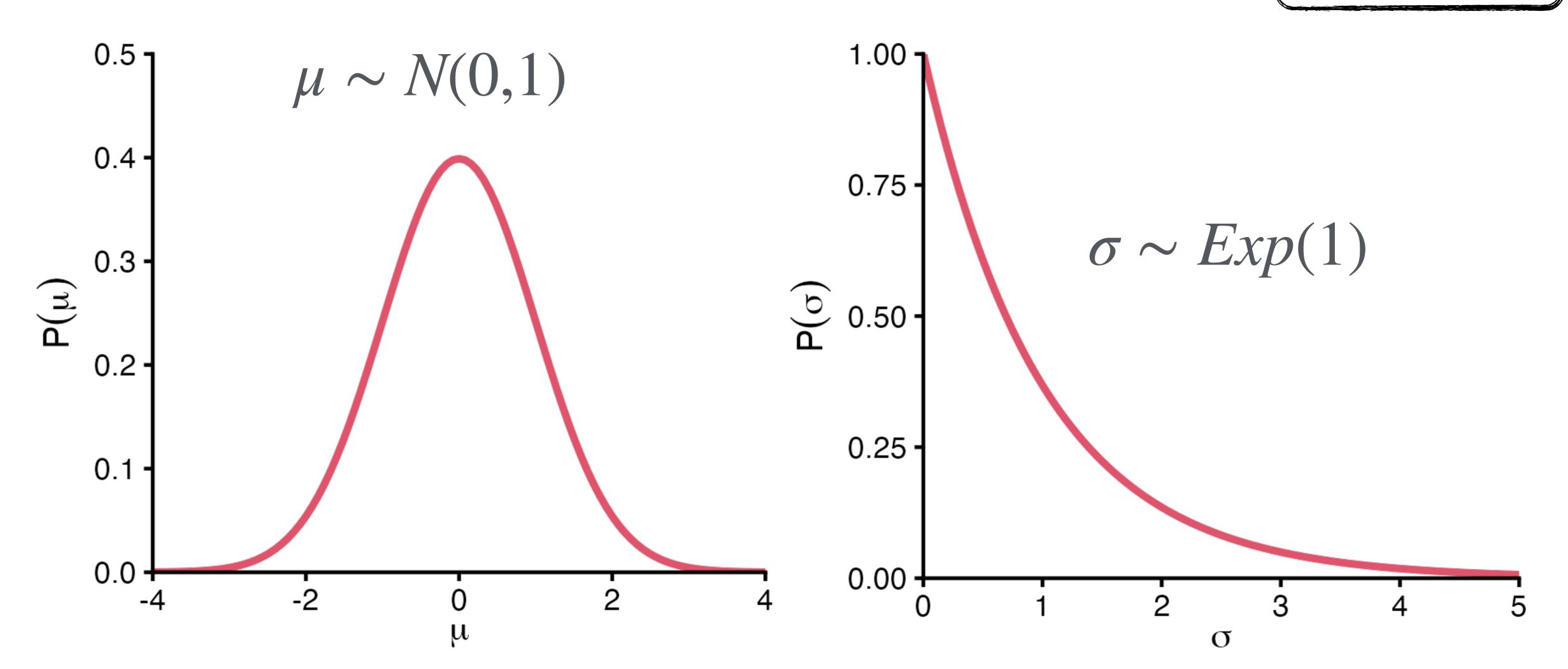
• A full model:

$$y_i \sim N(\mu, \sigma)$$
 $\mu \sim N(0, 1)$ 
 $\sigma \sim Exp(1)$ 

## The priors

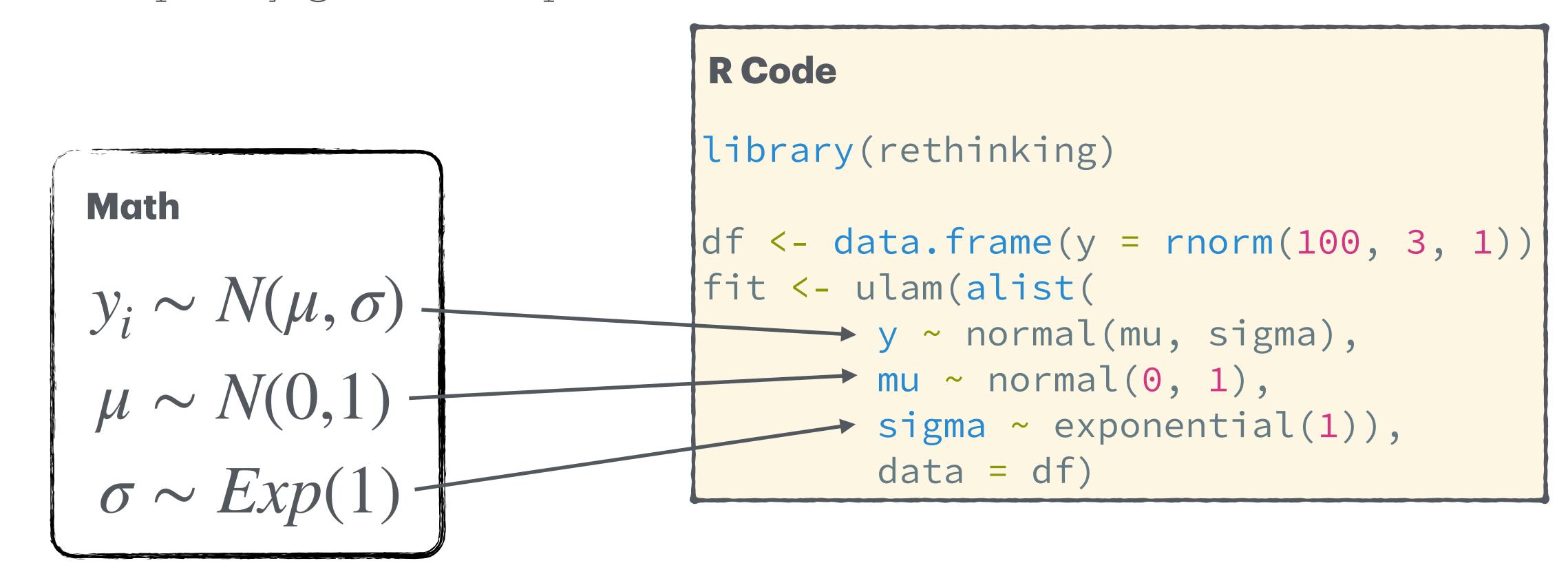
The encoding of our guesses

 $y_i \sim N(\mu, \sigma)$   $\mu \sim N(0, 1)$   $\sigma \sim Exp(1)$ 

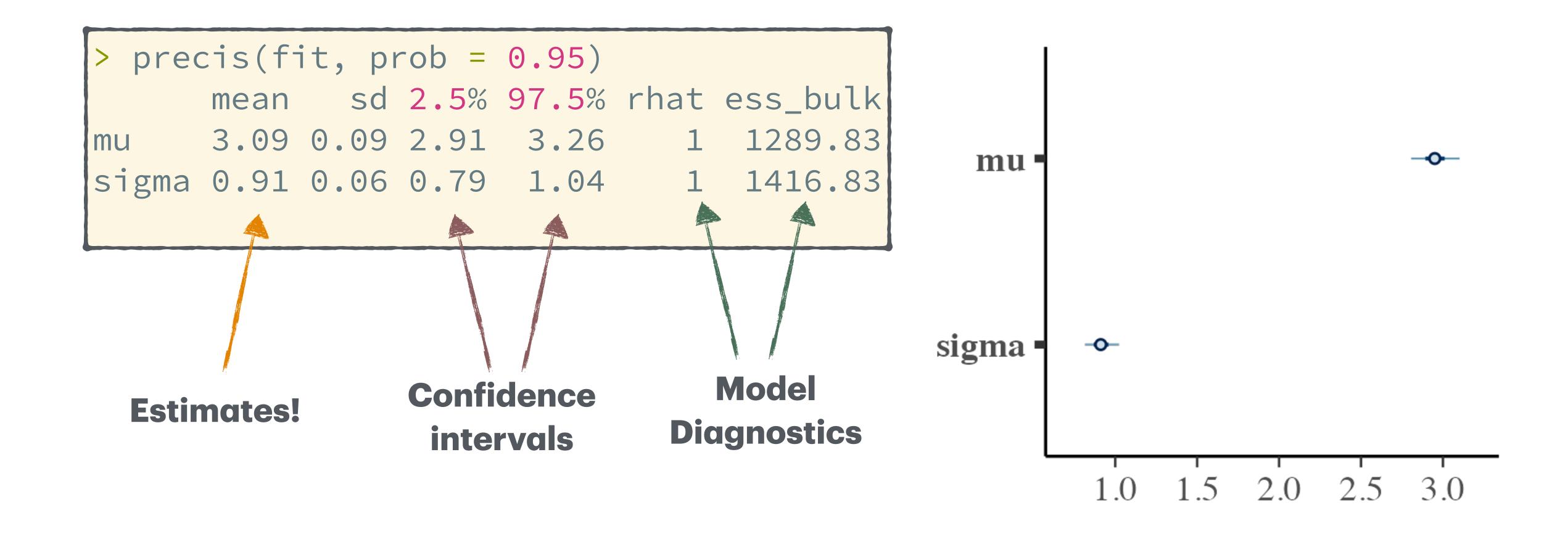


# Fitting the model

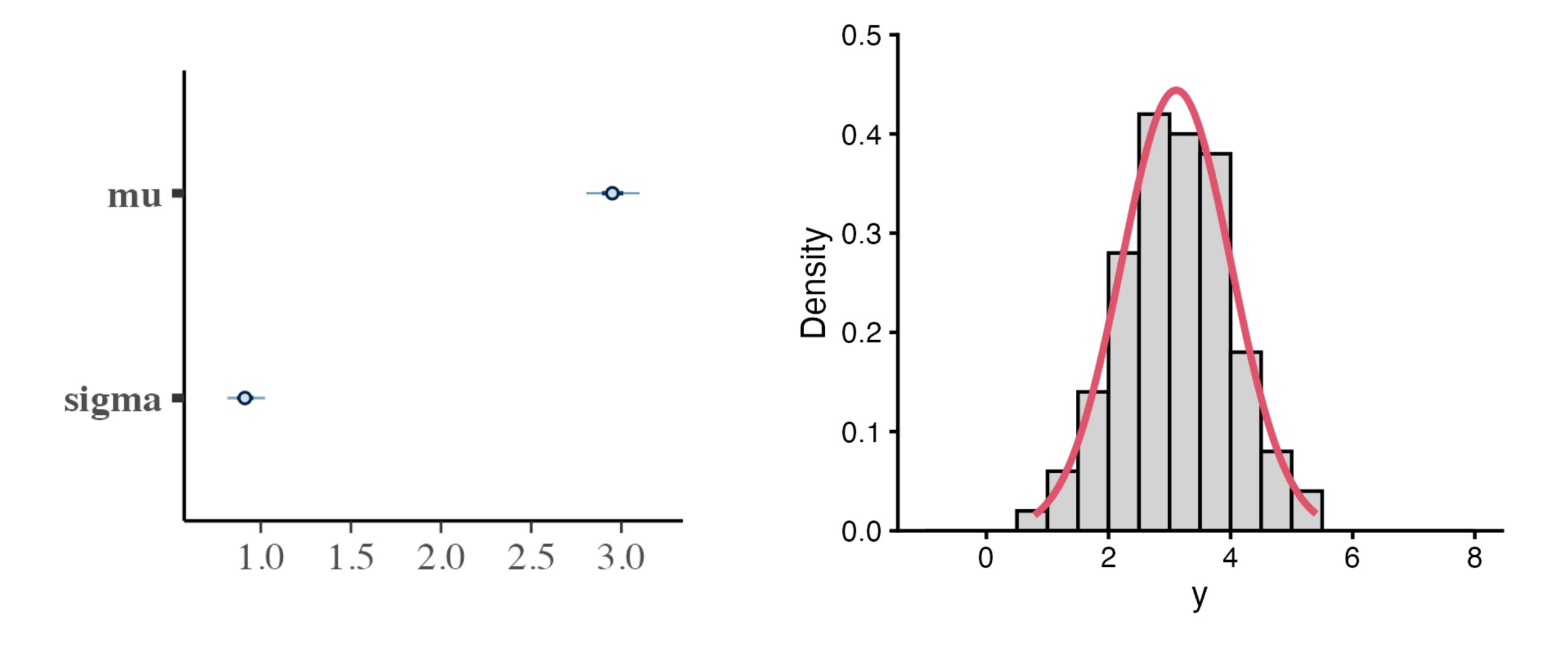
We have pretty good computers



## Model fit and parameter estimates



## Model fit and parameter estimates



## The linear model

### Adding more variables

- The main strategy for making useful probabilist models is to allow the parameters to vary
- If we have two measured variables:
  - $y_i$ : the dependent variable, the outcome, the response, the predicted
  - $x_i$ : the independent variable, the treatment, the control, the predictor

• We make the parameter  $\mu$  a linear function of the predictor variable:

$$y_i \sim N(\mu_i, \sigma)$$

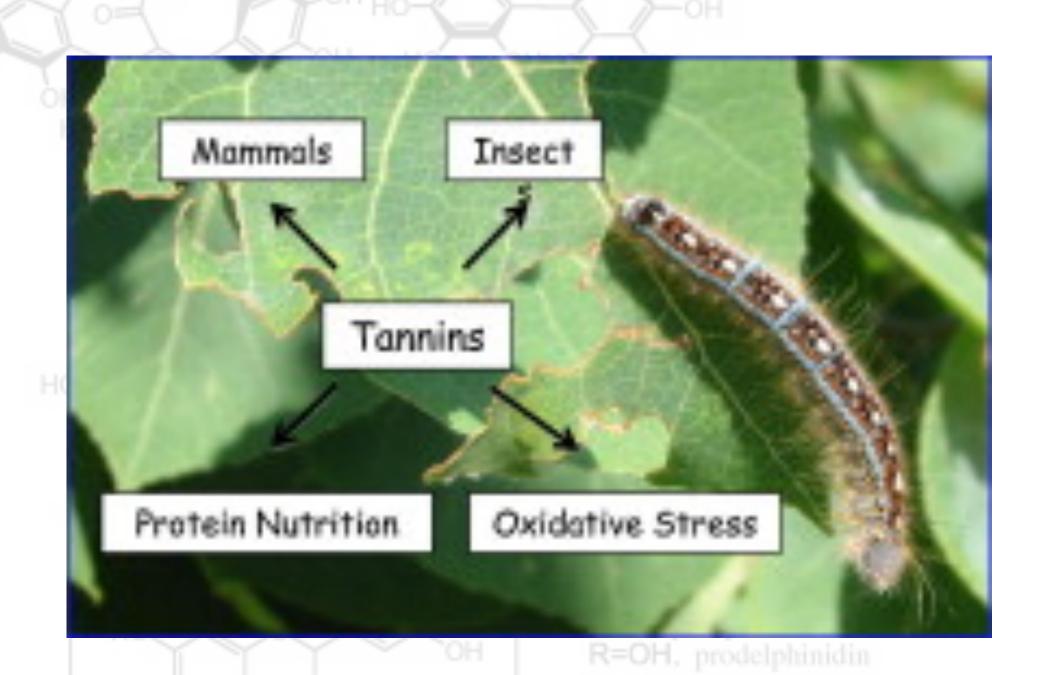
$$\mu_i = \alpha + \beta x_i$$

- This replaces  $\mu$  with two new parameters:
  - $\alpha$ : the intercept
  - $\beta$ : the slope of x

# An ecological example

Tannins in plant-herbivore interactions

- tannin are the most abundant secondary metabolites in plants
- defense mechanism: defend leaves against herbivores by deterrence or toxicity
- tannins oxidize in insects, leading to oxidative stress



## Our variables

- $y_i$ : caterpillar growth response variable
- $x_i$ : quantity of tannin in the caterpillar diet predictor variable,



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## Lets ask a question

Do leaf chemical compounds reduce the growth of caterpillars?

#### Our model

$$y_i \sim N(\mu_i, \sigma)$$

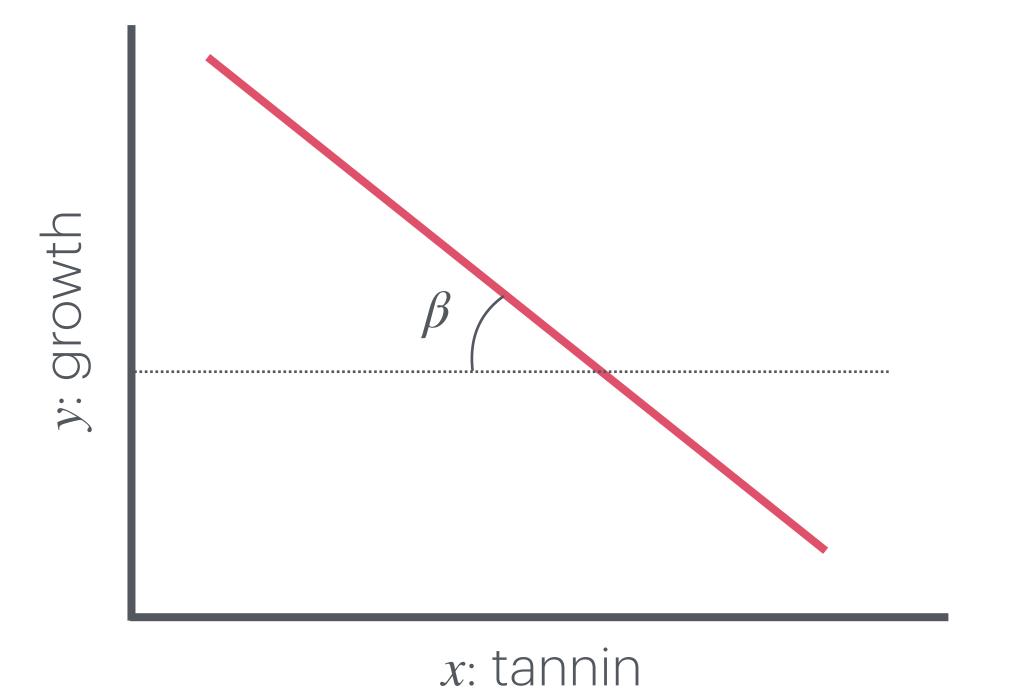
$$\mu_i = \alpha + \beta x_i$$

$$\alpha \sim N(0, 1)$$

$$\beta \sim N(0, 1)$$

$$\sigma \sim Exp(1)$$

The relation between growth and tannins is given by the slope parameter  $oldsymbol{eta}$ 

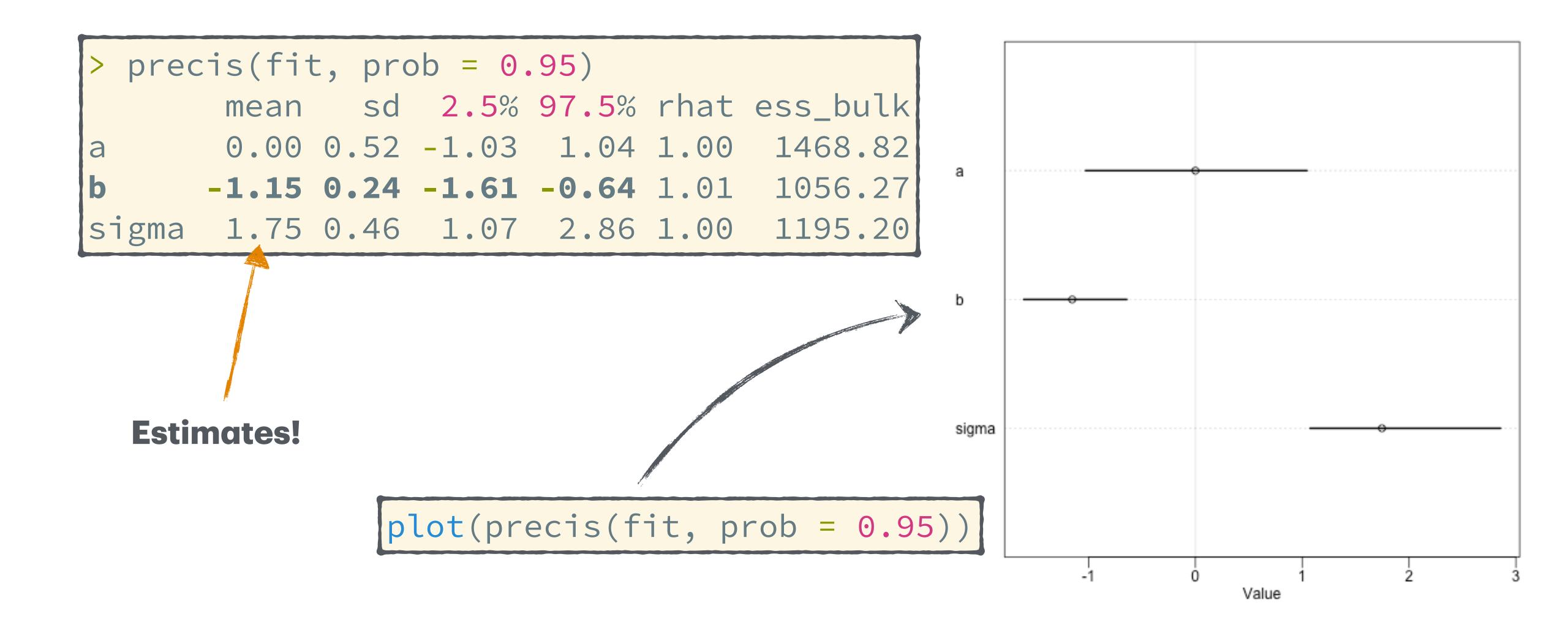


## Model in the computer

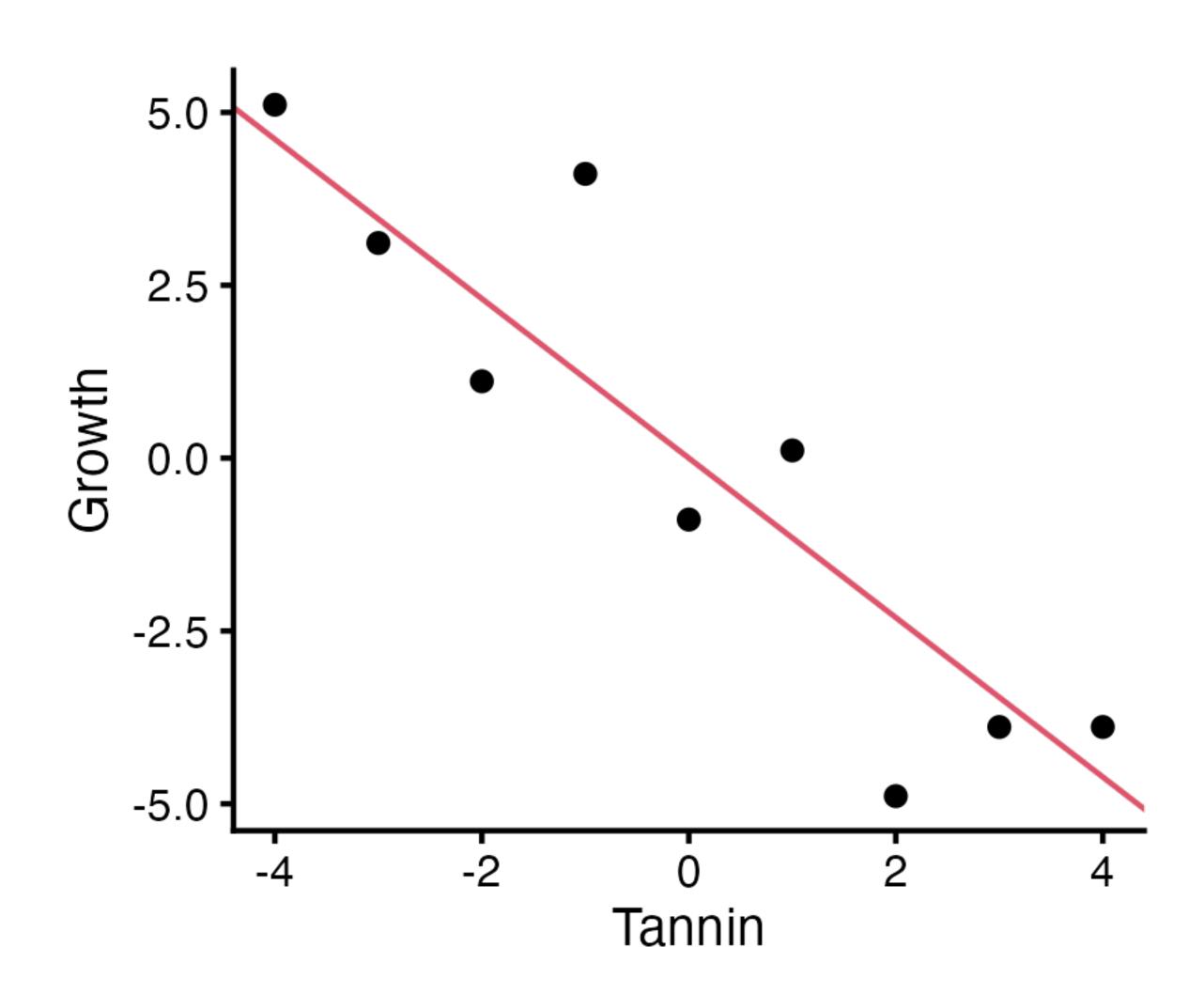
Centering both variables is always a good idea

```
df < -data.frame(growth = c(12, 10, 8, 11, 6, 7, 2, 3, 3),
                    tannin = c(0, 1, 2, 3, 4, 5, 6, 7, 8))
df$tannin = scale(df$tannin, scale = FALSE)
df$growth = scale(df$growth, scale = FALSE)
fit = ulam(alist(growth ~ normal(mu, sigma),
                                                         y_i \sim N(\mu_i, \sigma)
                     mu <- a + b*tannin,
                                                         \mu_i = \alpha + \beta x_i
                      a \sim normal(0, 1),
                                                         \alpha \sim N(0,1)
                      b \sim normal(0, 1),
                                                         \beta \sim N(0,1)
                      sigma ~ exponential(1)),
                                                        \sigma \sim Exp(1)
              data = df
```

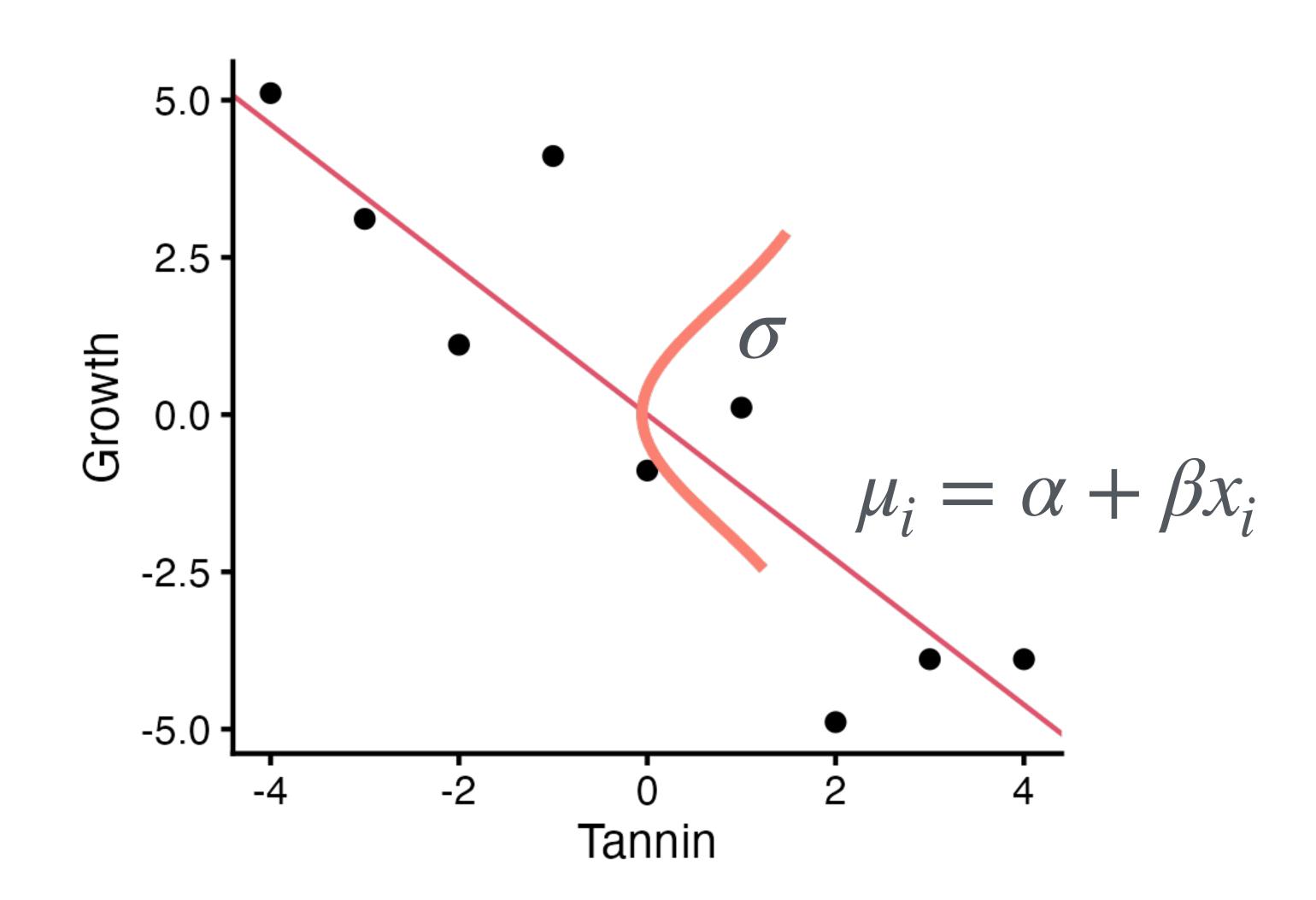
## Model fit



# Modelplot

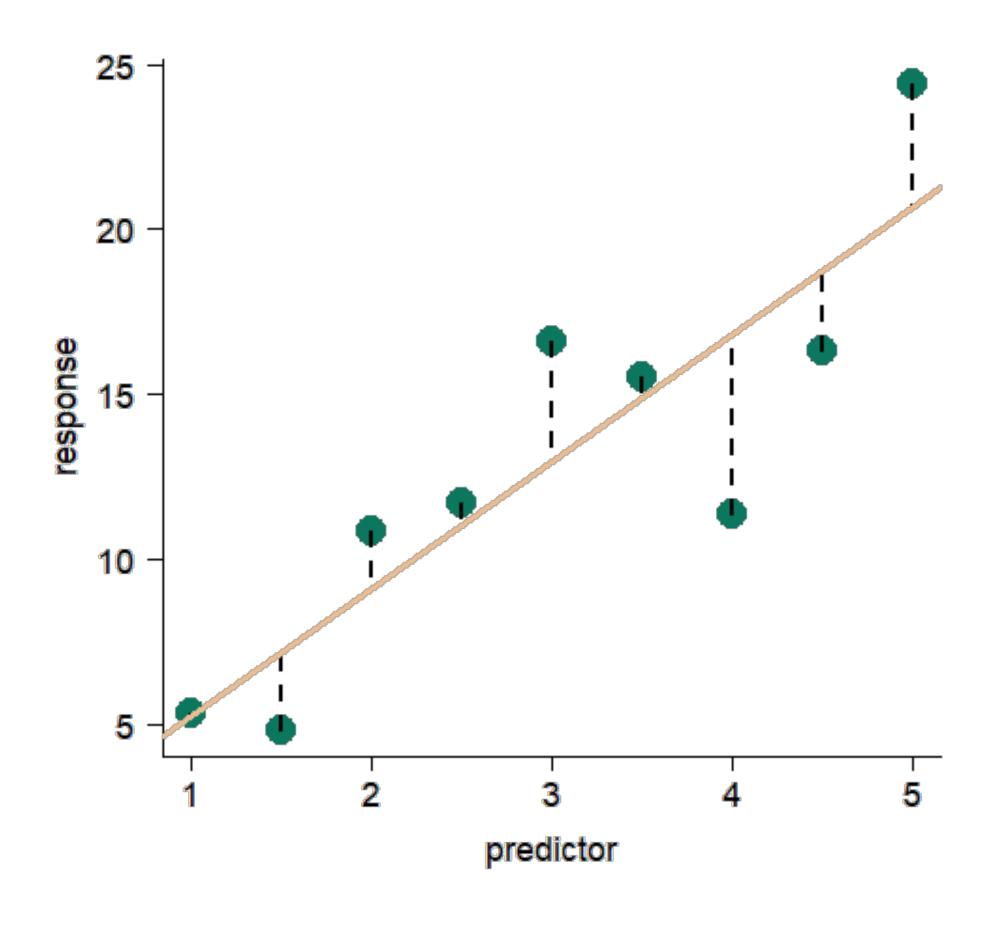


## Modelplot



# Other ways of fitting the linear model OLS and Maximum Likelihood

- The standard linear model has many flavors and justifications
- Ordinary Least Squares (OLS) is the most common introduction, and consists of minimizing the squared distance between observations and the regression line
- Maximum likelihood (ML) looks for the parameters that maximize the probability of observing  $y_i$ :
  - $P(y_i | \theta = \{\alpha, \beta, \sigma\})$
  - ML finds the same solution as OLS under a gaussian model



# Other ways of fitting the linear model

lm() function for linear models

- The lm() function in R can fit most of the models we saw with OLS using a formula notation
  - R formulas:
    - $y \sim 1$  (y as a function of a constant)
    - $y \sim x$  (y as a function of x, same as 1 + x)
- OLS assumes fixed uniform priors, so we can't change them

```
y_i \sim N(\mu_i, \sigma)
\mu_i = \alpha
```

# Other ways of fitting the linear model

lm() function for linear models

```
> df <- data.frame(growth = c(12, 10, 8, 11, 6, 7, 2, 3, 3), tannin = c(0, 1, 2, 3, 4, 5, 6, 7, 8))
y_i \sim N(\mu_i, \sigma) > \text{df$tannin} = \text{scale}(\text{df$tannin}, \text{scale} = \text{FALSE}) > \text{df$growth} = \text{scale}(\text{df$growth}, \text{scale} = \text{FALSE}) > \text{ols\_fit} = \text{lm}(\text{growth} \sim \text{tannin}, \text{data} = \text{df}) > \text{precis}(\text{ols\_fit})
                                                                       (ols_fit)
mean sd 5.5% 94.5%
                                       lpha (Intercept) 0.00 0.56 -0.90 0.90 tannin -1.22 0.22 -1.57 -0.87
```

# Other ways of fitting the linear model

stan\_glm() function for linear models and priors!

```
y_i \sim N(\mu_i, \sigma)
\mu_i = \alpha + \beta x_i
```

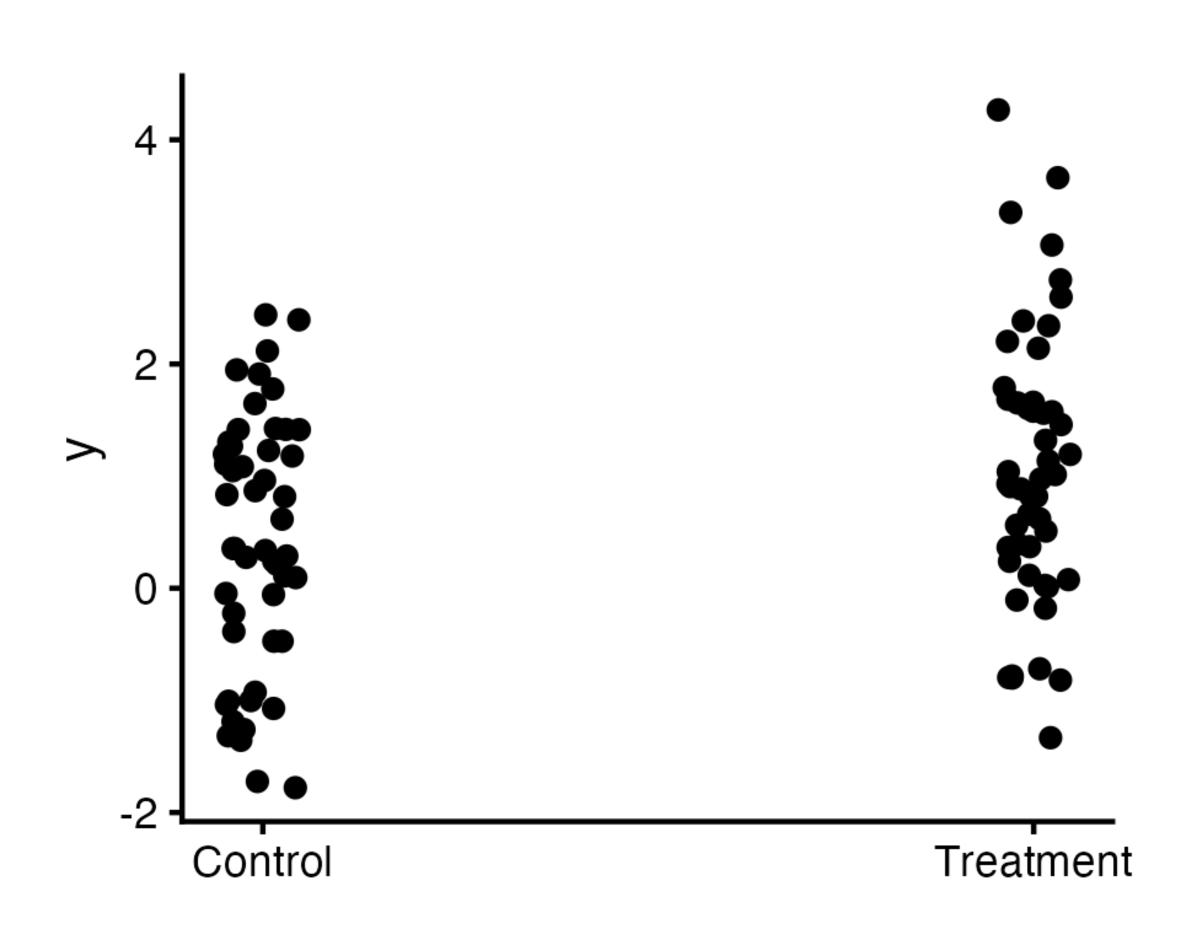
And some standard priors...

```
> sglm_fit = stan_glm(growth ~ tannin, data = df, cores = 4)
> summary(sglm_fit, probs = c(0.025, 0.975))[, 1:7]
                                                          2.5%
                                                                      97.5% n_eff
                                   mcse sd
(Intercept) lpha -0.01069275 0.015974907 0.6971944 -1.403377
                                                                  1.3696905
                                                                             1905
             eta -1.21608408 0.005609107 0.2482728 -1.716820 1.98129172 0.016244872 0.6447948 1.145132
tannin
                                                                             1959
                                                                 -0.7229236
sigma
                                                                 3.5859302
                                                                             1575
mean_PPD -0.01273309 0.020249189 0.9958192 -2.030730
                                                                  2.0076369
                                                                             2418
log-posterior -23.56539992 0.046443876 1.4480299 -27.365501 -21.9264227
                                                                              972
                   Rhat
(Intercept)
              1.000676
tannin
              1.001776
sigma
              1.000847
mean_PPD
               1.000271
log-posterior 1.003540
```

# What about categorical predictors?

Linear regression is flexible

- Our questions are frequently based on categories:
  - Is a treatment effective in improving outcomes?
  - Are two geographical regions different in some aspect?
  - Does the diet of a group of species affect their size?



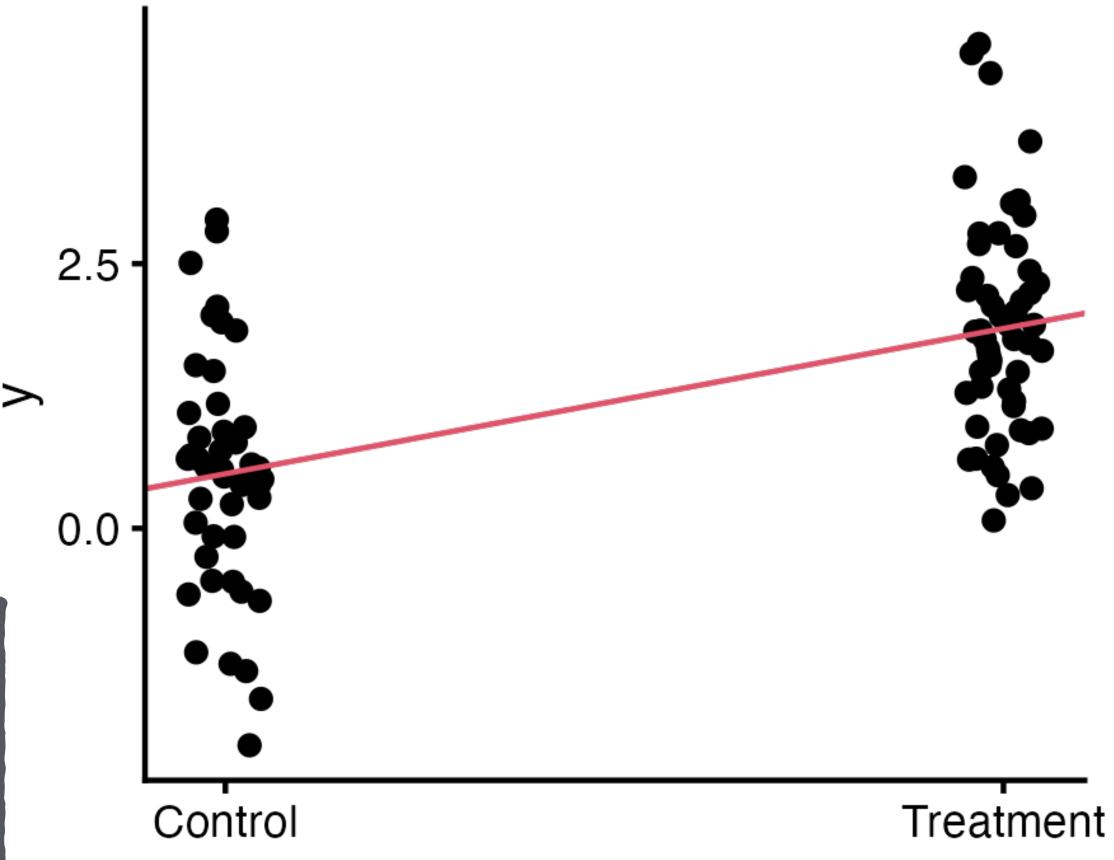
## Binary predictor, same model

Control-treatment, two categories...

$$y_i \sim N(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta x_i$$

- $x_i$ : O for control, 1 for treatment
- $\alpha$ : the intercept is the **mean of** the Control group
- $\beta$ : the slope is the difference in the means across the groups



- There are a few ways of modeling predictors with many categories:
  - Contrasts: Each category is compared to a baseline, and the coefficients are comparisons between baseline and levels
  - One-hot: coefficients are means of each level of the predictor
  - Residuals: an overall mean is measured, and coefficients are differences between each level and the global mean

```
> x = sample(LETTERS[1:3], 9, replace = TRUE)
> y = 1 + ifelse(x == "A", 0,
          ifelse(x == "B", 1, 2)) + rnorm(9)
 df = tibble(y, x)
> df
# A tibble: 9 × 2
     <dbl> <chr>
   0.870
  -0.00180 A
  2.03
   2.55
   3.92
   4.66
```

Contrasts, the default in most lm() functions

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Q: How do we get estimates for the mean in each class?

#### One-hot

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```
"B" "A" "A" "B" "C" "B" "B" "C" "B"
onehot
xA xB xC
```

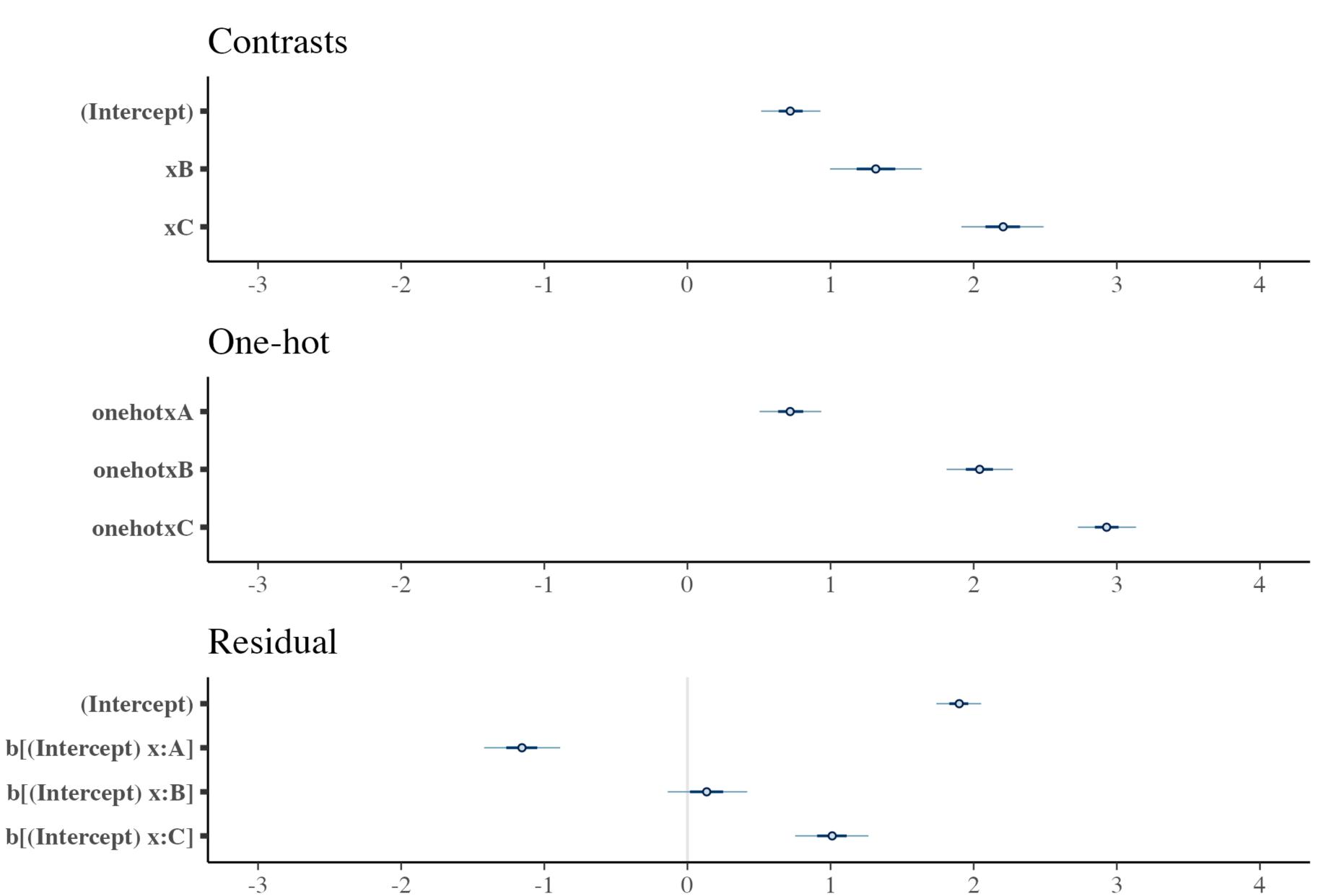
#### One-hot

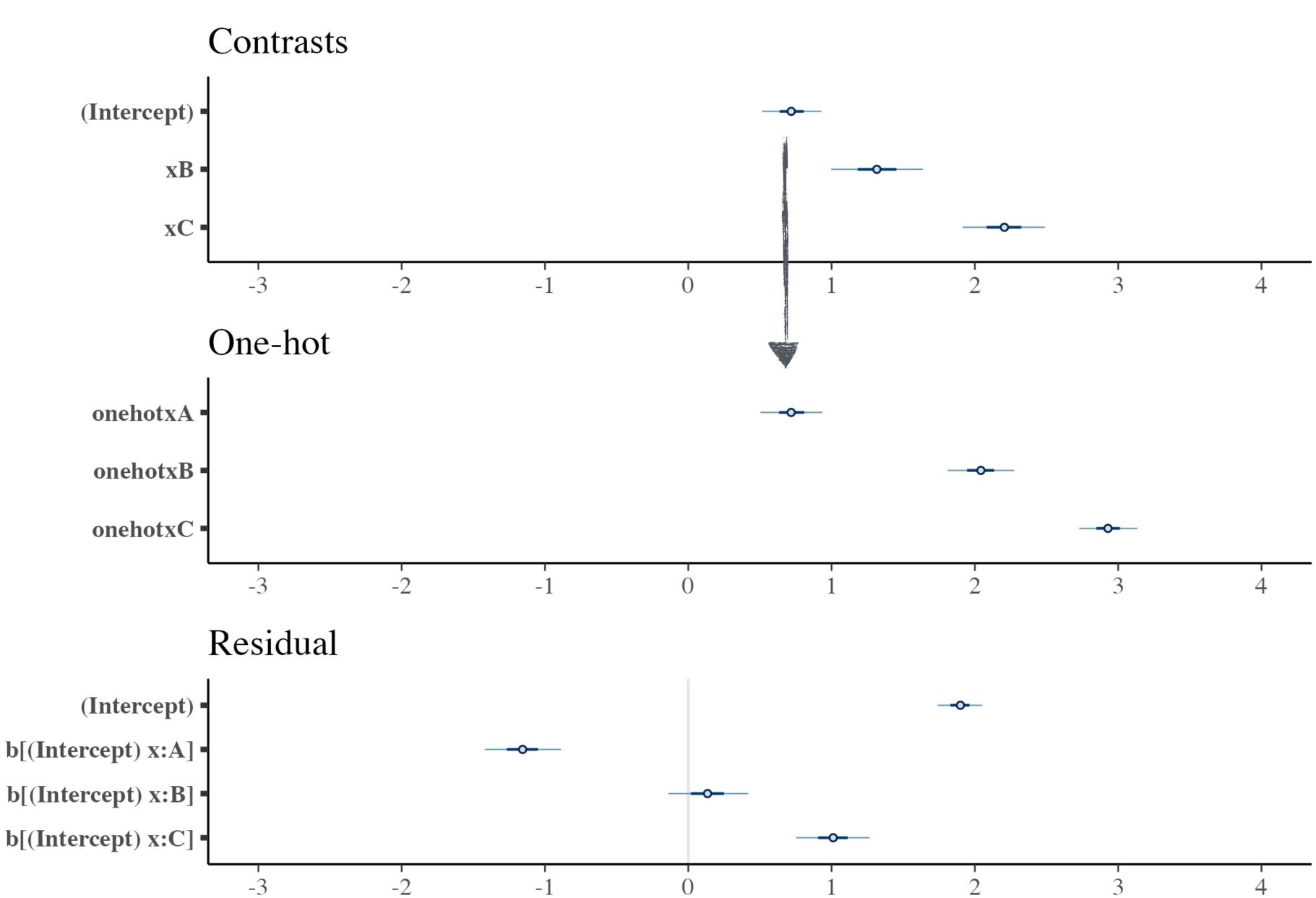
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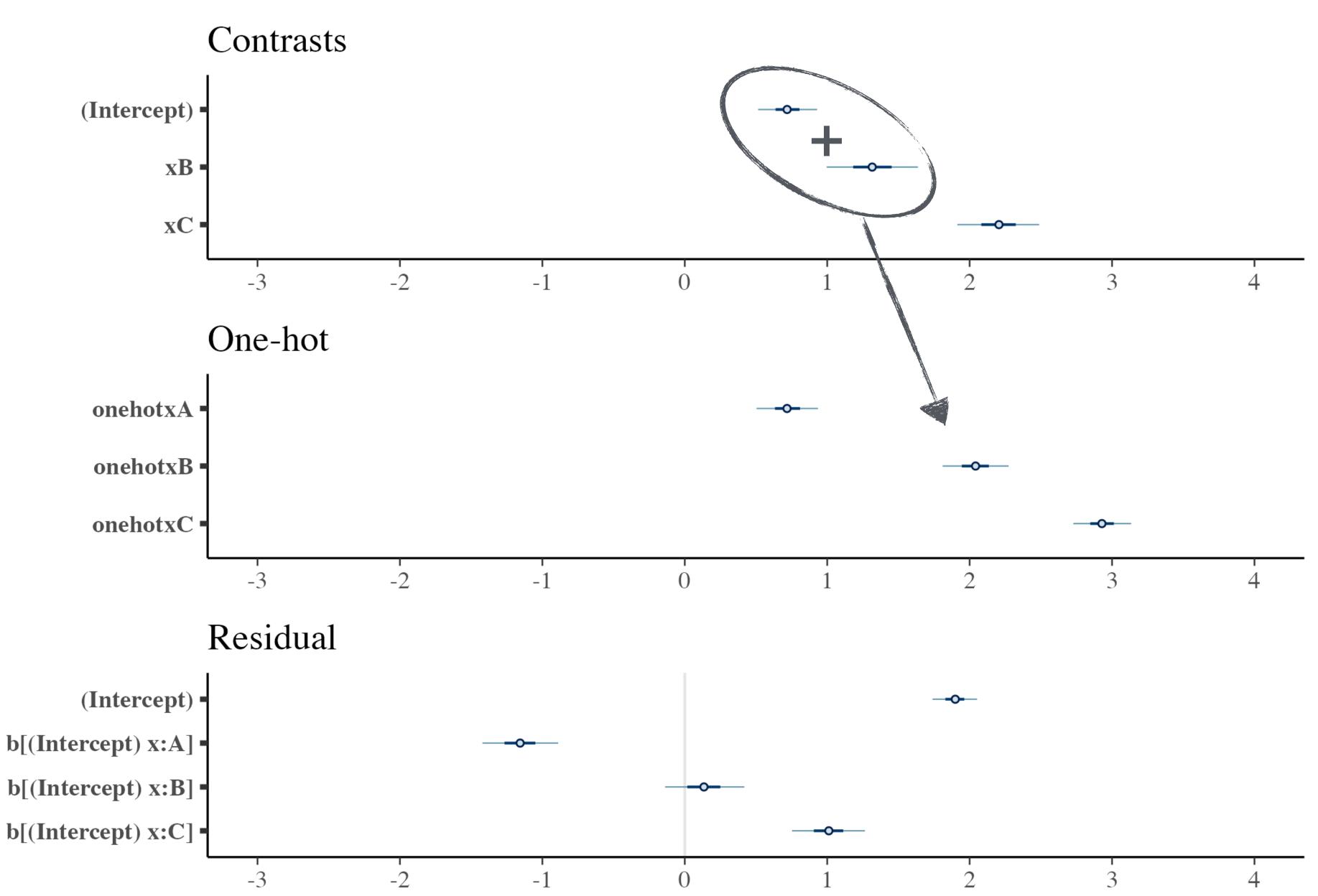
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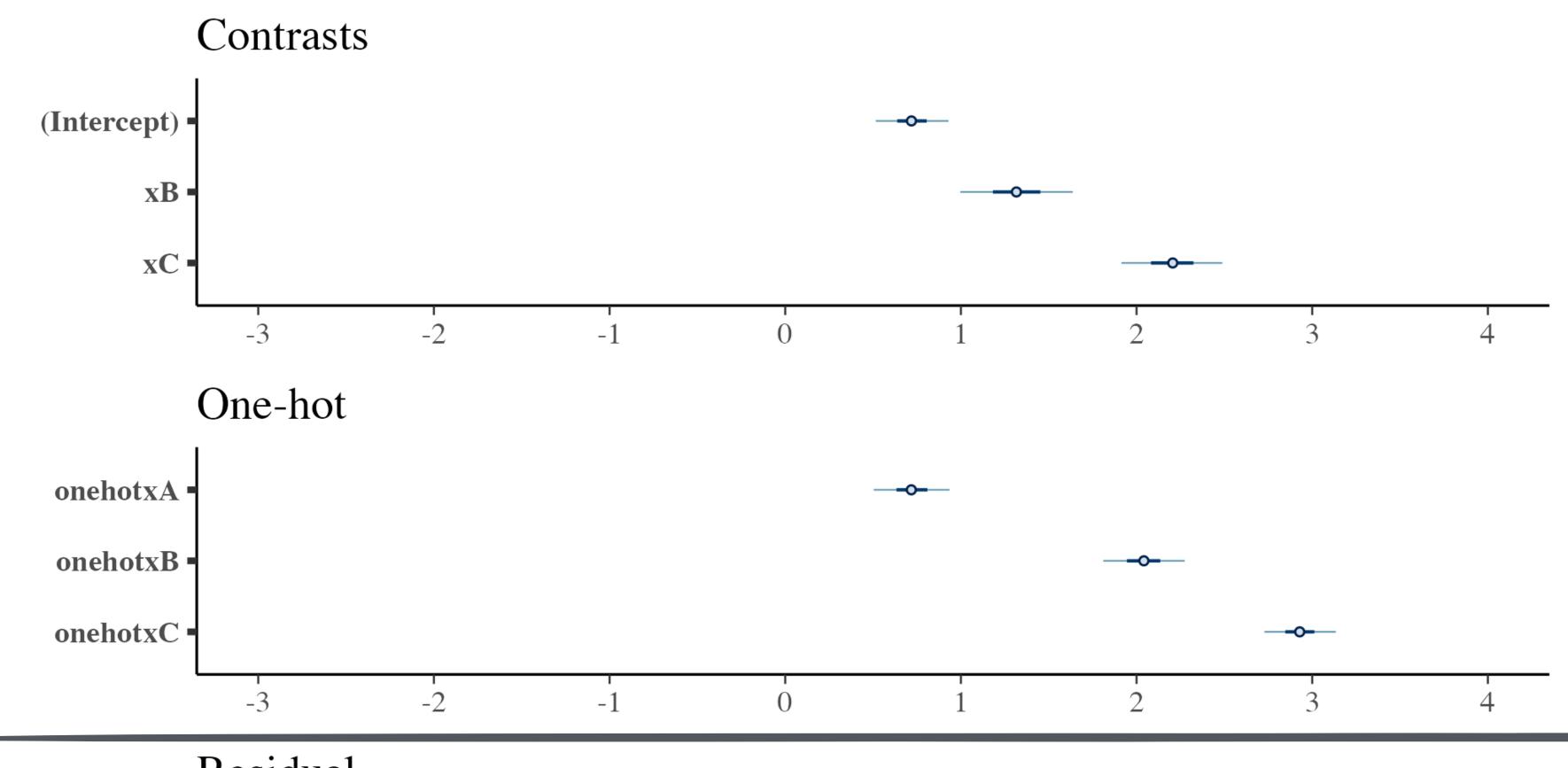
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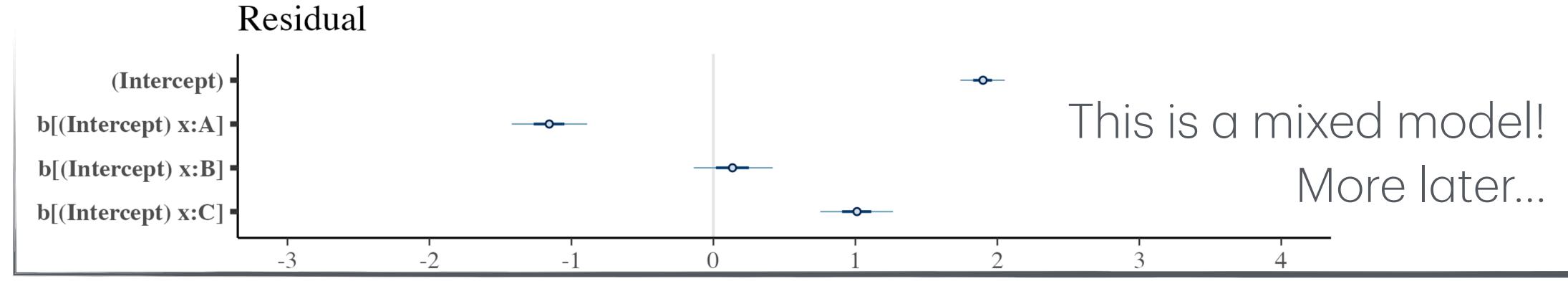
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## Summary

- Linear models help us answer questions about differences across groups
- The basic strategy is to formulate a
   probabilistic description of our data, and
   to establish a relation between data and
   parameters in the statistics model
- In linear models, coefficients are comparisons across categories, and we can interpret their estimates

- There are many ways to fit the models
- We like to encode the information the data brings about parameters using the posterior distribution, and this requires the definition of priors
- We can also fit the models using OLS and Maximum Likelihood, but if we do, we are stuck with default priors that don't bring useful information