# EDS222 Week 8

**Hypothesis Testing** 

**November 19, 2024** 

# Agenda

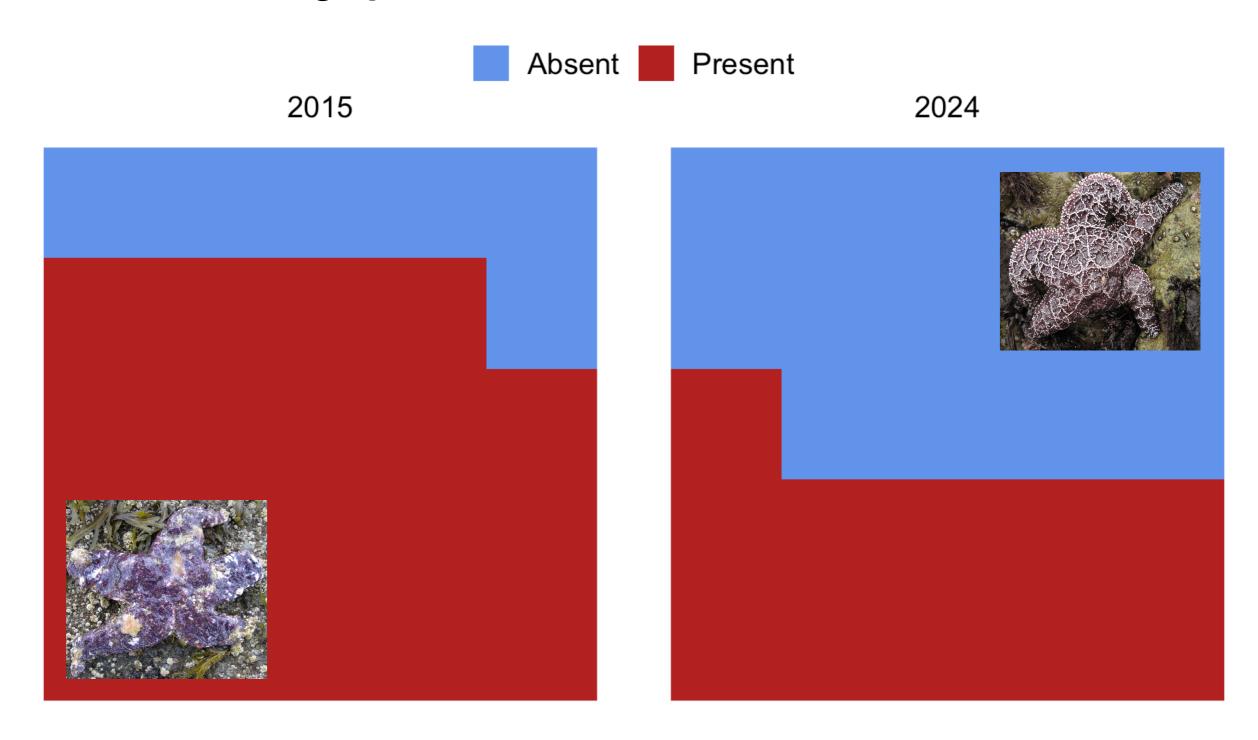
- Hypothesis testing by randomization
  - Null and alternative hypotheses
  - Sample statistics and sampling distributions
  - P-values and rejecting the null
- Hypothesis testing in practice
  - Central limit theorem
  - Standard errors
- Confidence intervals
  - Interpretation
  - Effect sizes

Sea star wasting sydrome





Sea star wasting sydrome



#### **Overview**

- Overall question
  - Did sea star wasting syndrome incidence decrease from 2015 to 2024?
- Procedure
- 1.
- 2.
- 3.
- 4.
- 5.

**Key terms** 

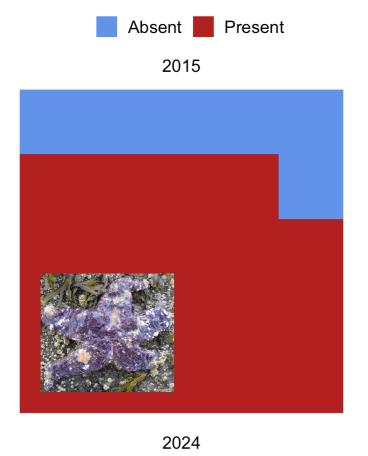
Null and alternate hypotheses

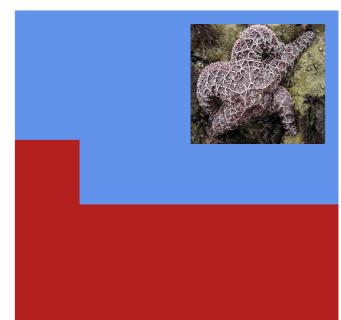
Sample statistic

Point estimate

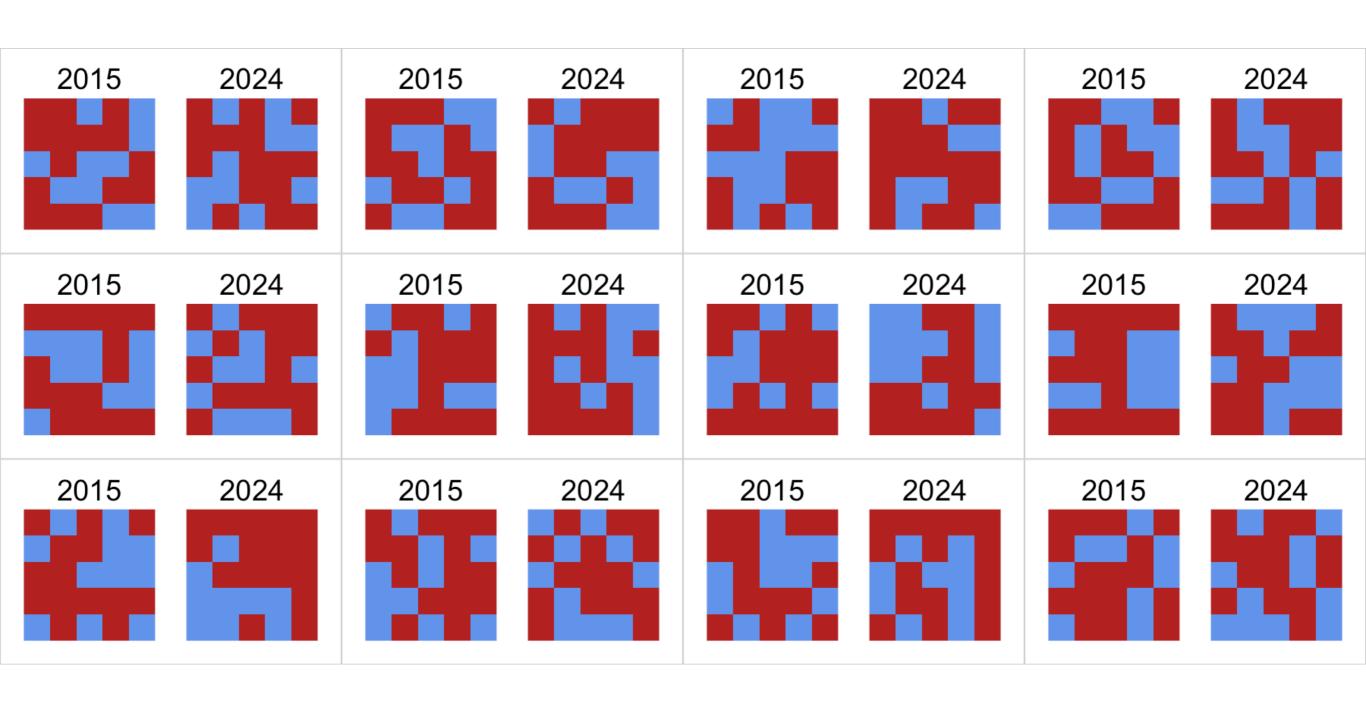
Sampling distribution

#### Sea star wasting sydrome

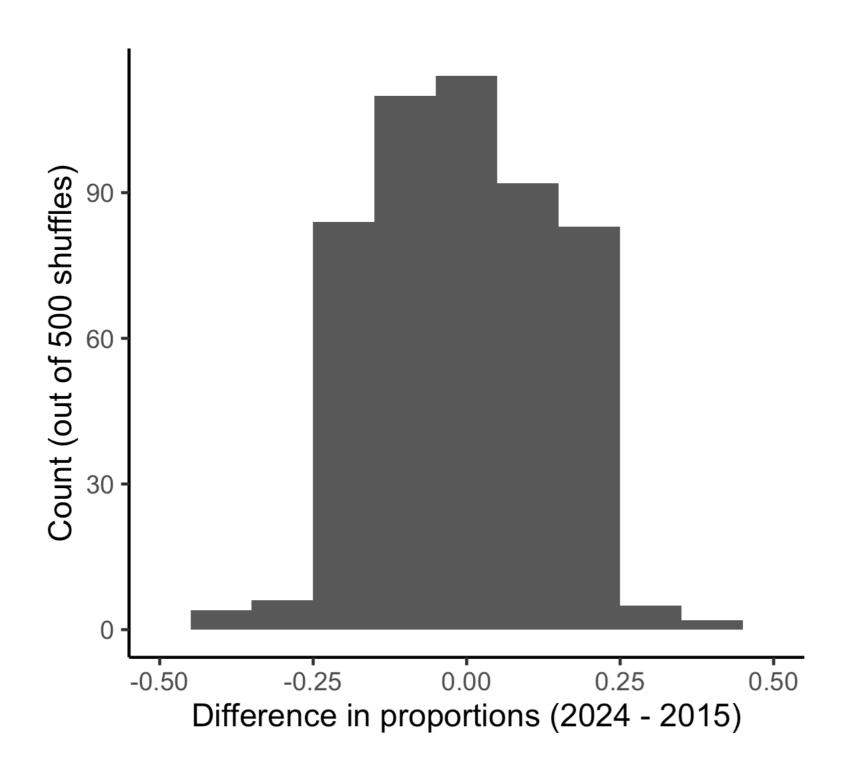




Quantify uncertainty by shuffling



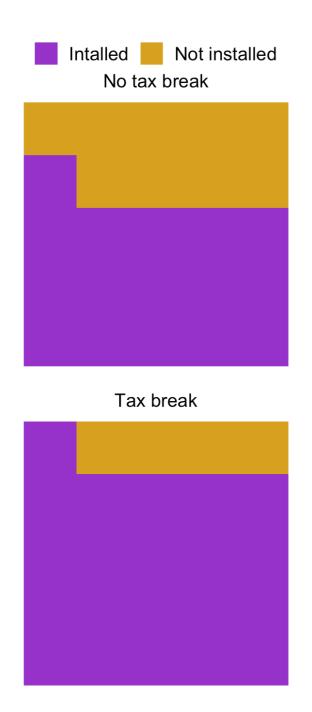
Probability of point estimate under the null



Reject or fail to reject the null?

#### Your turn

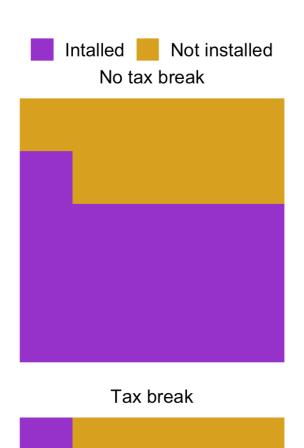
Do tax breaks incentivize solar panel installation?



- 1. Define the null and alternate hypotheses
- 2. Calculate the point estimate of the sample statistic
- 3. Quantify the uncertainty in the sampling distribution
- 4. Calculate probability of point estimate under the null
- 5. Reject or fail to reject null

#### Your turn

Do tax breaks incentivize solar panel installation?

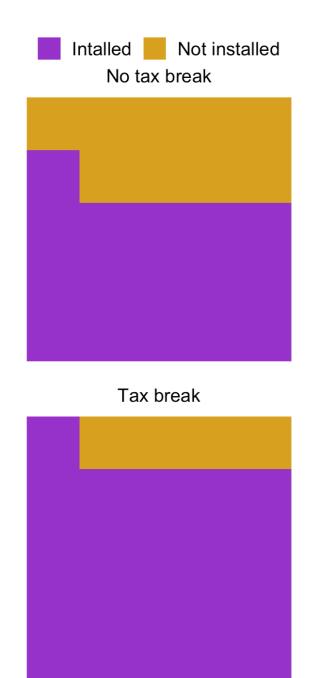


1. Define the null and alternate hypotheses

What are  $H_0$  and  $H_A$ ?

#### Your turn

Do tax breaks incentivize solar panel installation?



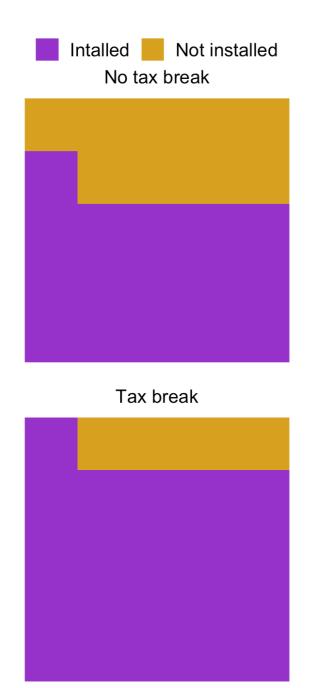
2. Calculate the point estimate of the sample statistic

Draw lines from *a,b,c,d* in the equation below to the corresponding parts of the figure on the left.

difference in proportions = 
$$\frac{a}{b} - \frac{c}{d}$$

#### Your turn

Do tax breaks incentivize solar panel installation?



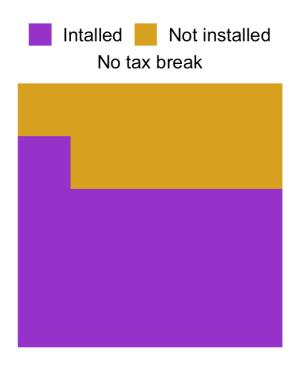
3. Quantify the uncertainty in the sampling distribution

Which R function will help?

- A) rnorm()
- B) sample()
- C) dnorm()

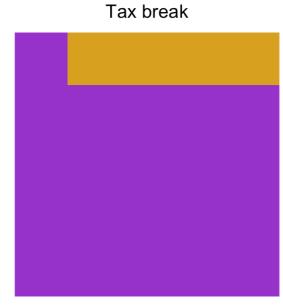
#### Your turn

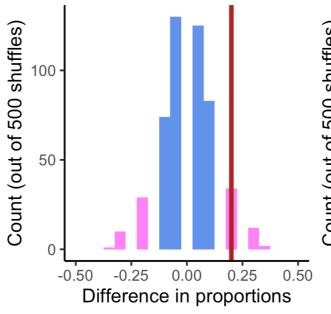
Do tax breaks incentivize solar panel installation?

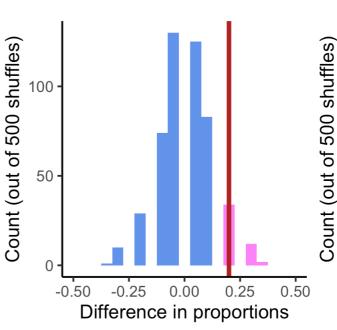


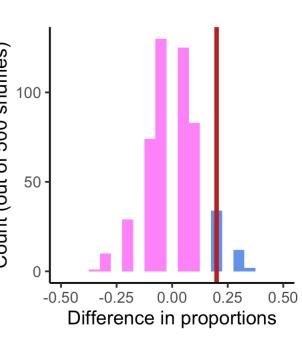
# 4. Calculate probability of point estimate under the null

The histograms below show the results of randomization and the red line is the observed difference. Which figure shows the p-value in pink?



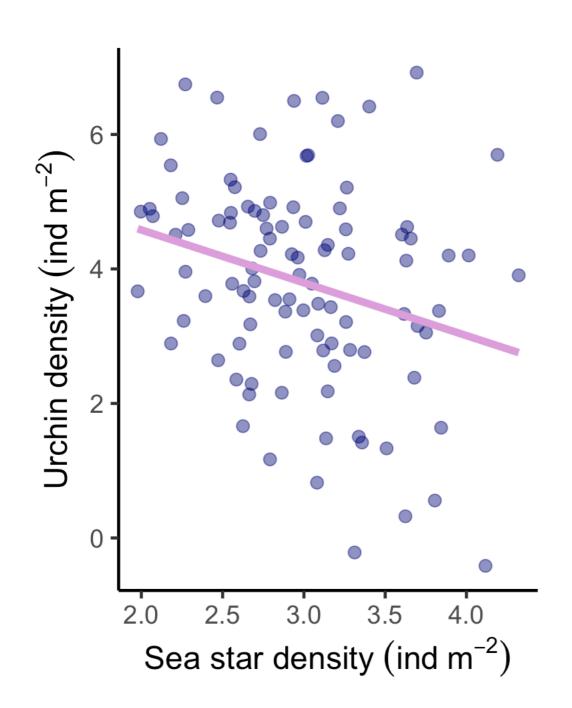






#### Applicable to regression and other models

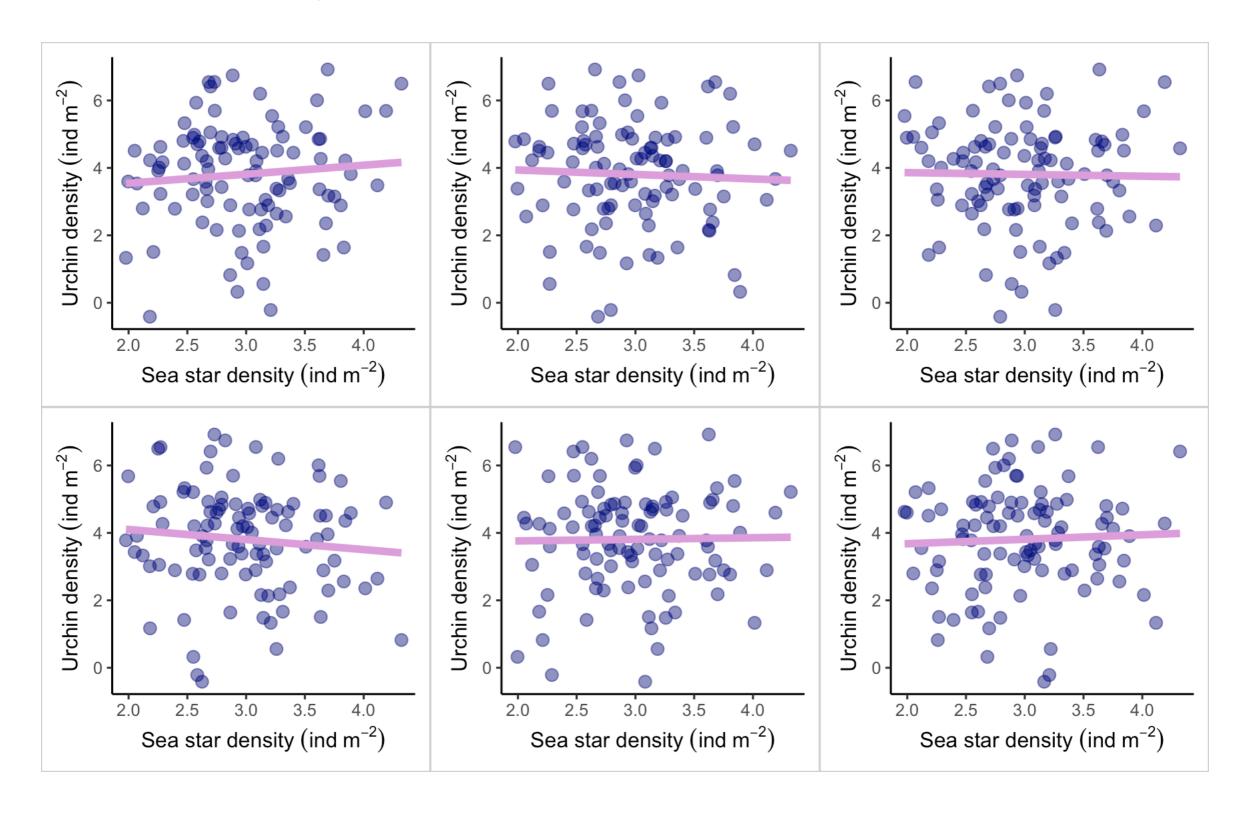
Do sea stars reduce urchin populations?



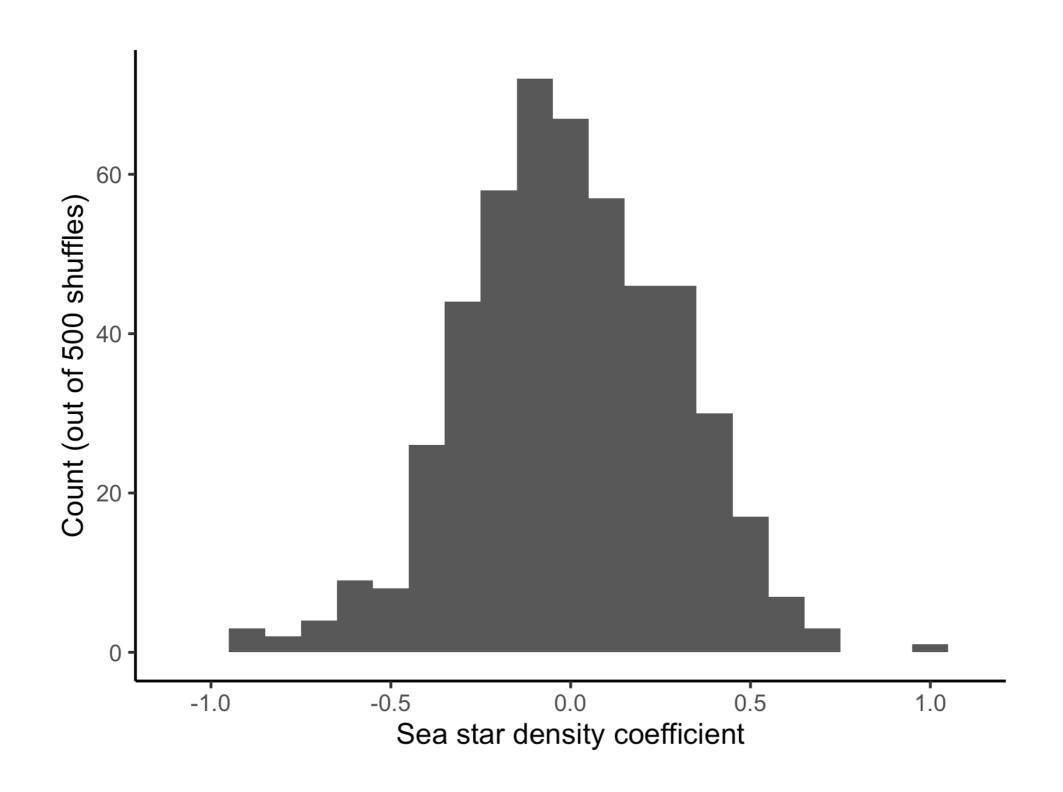
1. What are the null and alternate hypotheses?

2. What's the relevant sample statistic?

#### Applicable to regression and other models



Applicable to regression and other models

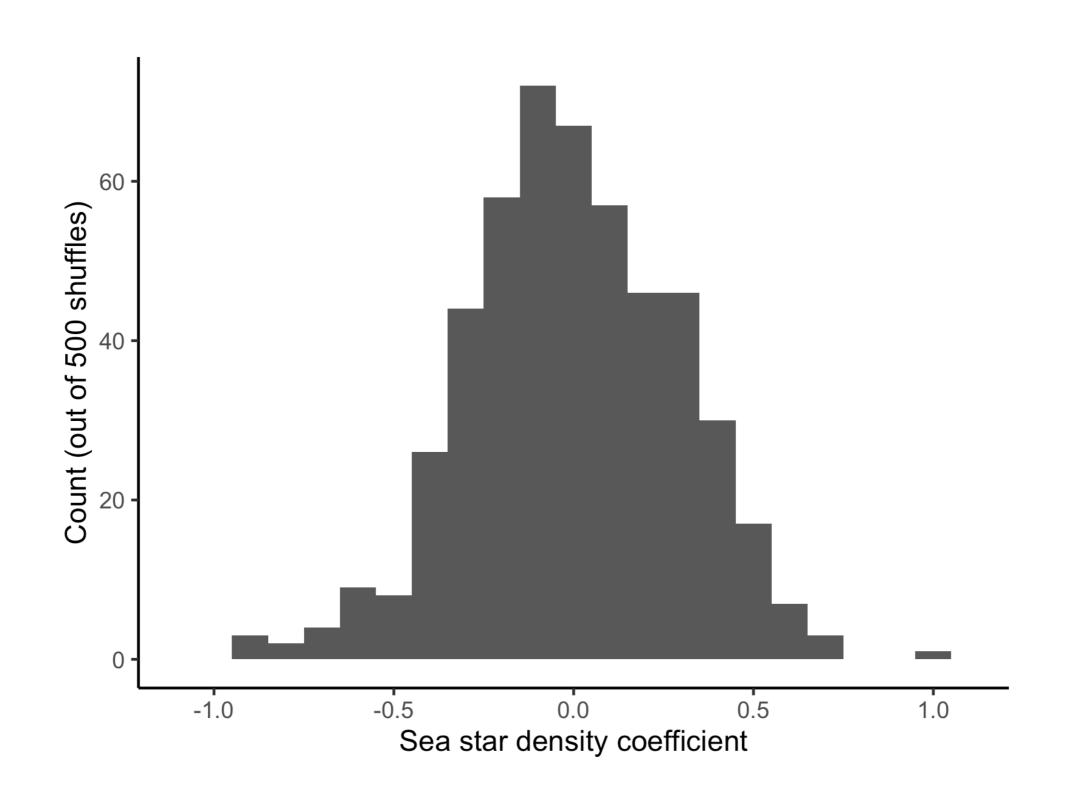


#### Recap

- 1. Formulate your hypotheses  $H_0 = no$  effect,  $H_A = some$  effect
- 2. Calculate point estimate

  Difference in means, regression coefficient, etc
- 3. Quantify uncertainty in sampling distribution Shuffle data, recalculate point estimate, repeat
- 4. Calculate p-value Probability of point estimate if null is true
- 5. Reject or fail to reject the null  $ls p \leq \alpha$ ?

#### **Motivation**

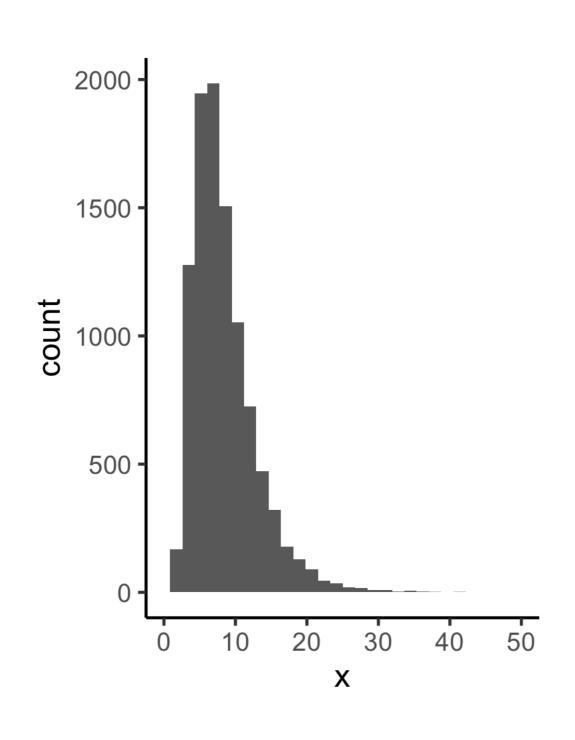


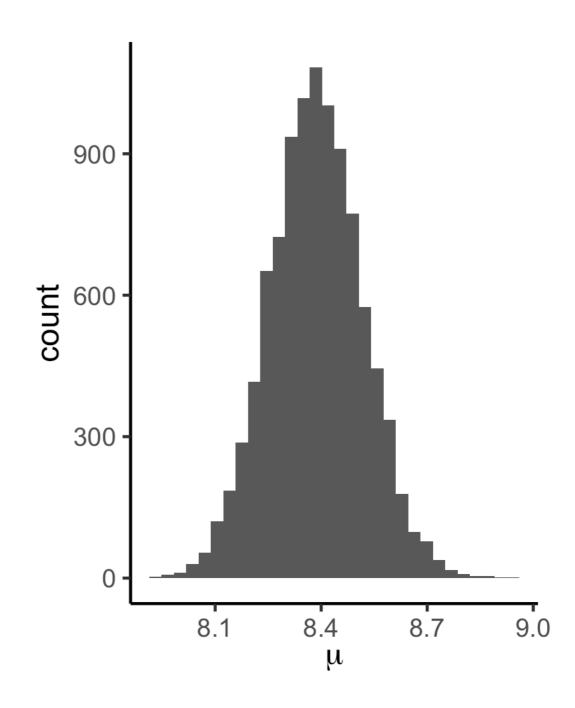
**Central limit theorem** 

The Central Limit Theorem states:

If your sample size is large enough, then the sampling distribution for many sample statistics (difference in proportions, regression coefficients, etc) are approximately normal

#### **Central limit theorem**



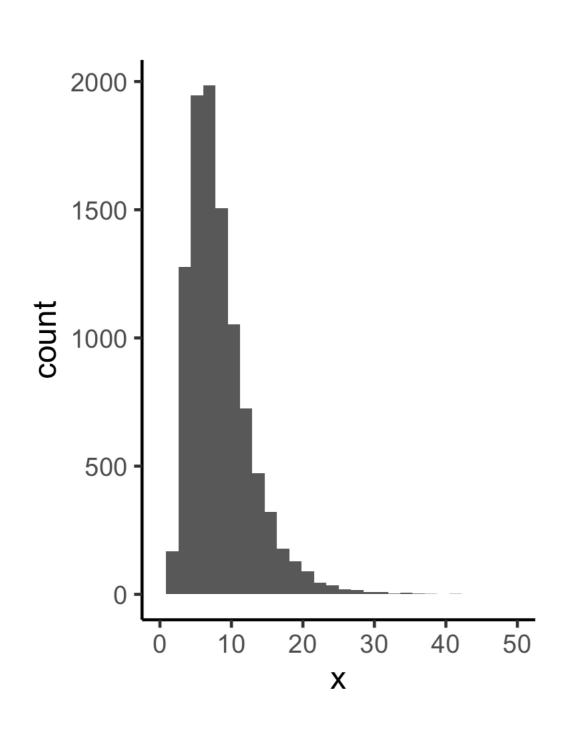


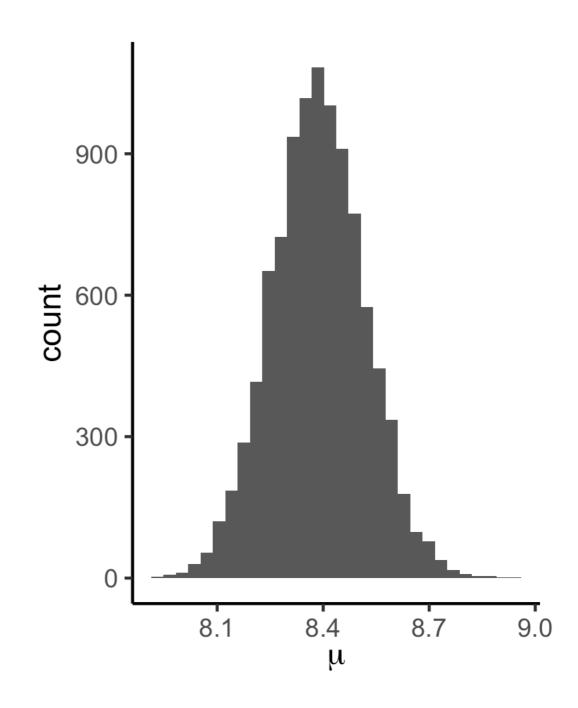
#### **Central limit theorem**

#### Try it on your own

```
# Roll a dice 10,000 times to get a non-normal population
# It's not even continuous!
x \leftarrow sample(1:6, 1e4, replace = TRUE)
qqplot(tibble(x), aes(x)) +
  geom_histogram(binwidth = 1, color = "blue", fill = NA) +
  theme_classic()
# Simulate the sampling distribution of the mean
# Do the following 1000 times
   1. Sample 50 values from your non-normal population
   2. Calculate the sample mean
mean_x <- replicate(</pre>
  1e3,
  mean(sample(x, size = 50))
ggplot(tibble(mean_x), aes(mean_x)) +
  geom_histogram(bins = 15, color = "blue", fill = NA) +
  theme_classic()
# Looks pretty normal!
```

#### **Standard errors**





#### **Standard errors**

#### Standard error

Standard deviation of the sampling statistic.

#### **Problem**

We only get one sample! Can't get the standard deviation of one data point.

#### **Solution**

Someone else solves the central limit theorem for you.

#### **Note**

Don't memorize equations! Demonstration purposes only.

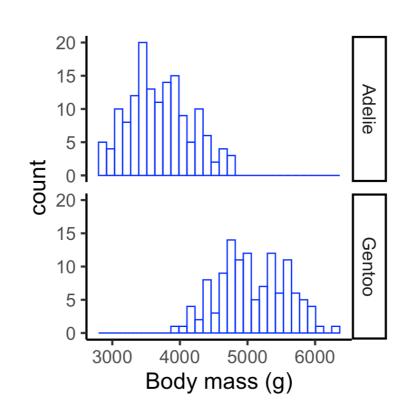
Standard error of the difference of means

**Population** 

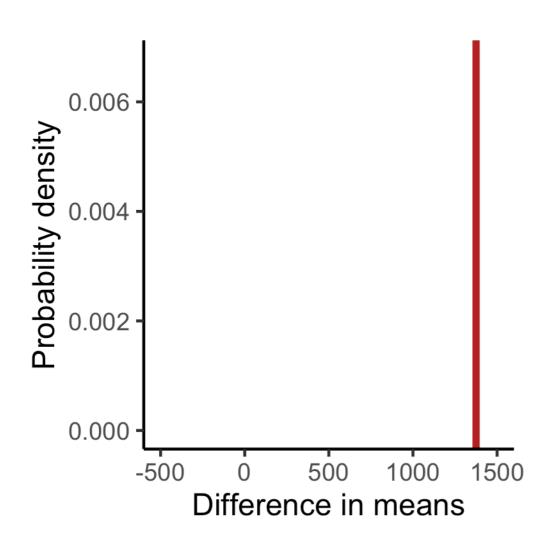
Sample

Sample statistic





Standard error of the difference of means

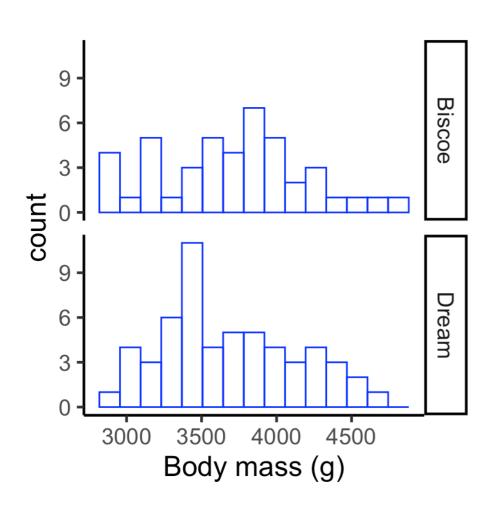


#### Standard error of the difference of means

- 1. Formulate your hypotheses  $H_0 = no$  effect,  $H_A = some$  effect
- 2. Calculate point estimate

  Difference in means, regression coefficient, etc
- 3. Quantify uncertainty in sampling distribution Shuffle data, recalculate point estimate, repeat Approximate sampling distribution using standard error
- Calculate p-value
   Probability of point estimate if null is true
   2 \* pnorm(-abs(observed), mean = 0, sd = se)
- 5. Reject or fail to reject the null  $p \le \alpha$ ?

#### Your turn



```
adelie_biscoe <- with(penguins,</pre>
                        body_mass_g[species == "Adelie" &
                                        island == "Biscoe"])
adelie_dream <- with(penguins,</pre>
                       body_mass_g[species == "Adelie" &
                                       island == "Dream"])
obs_diff <- mean(adelie_biscoe - adelie_dream)</pre>
obs_diff <- mean(adelie_biscoe) - mean(adelie_dream)</pre>
se <- function(a, b) {</pre>
  a <- na.omit(a)</pre>
  b <- na.omit(b)</pre>
  sqrt(sd(a)^2 / length(a) + sd(b)^2 / length(b))
se_diff <- se(adelie_biscoe, adelie_dream)</pre>
pval <- 2 * pnorm(-abs(observed_difference),</pre>
                   mean = 0,
                    sd = se_difference)
pval <- 2 * pnorm(0,
                   mean = -abs(observed_difference),
                    sd = se_difference)
```

#### Your turn

1. Which obs\_diff is the difference of the means?

2. Which pval is the probability of the observed difference, if the null is true?

3. Sketch the null distribution of the sample statistic. Indicate the observed difference, the standard error, and the p-value.

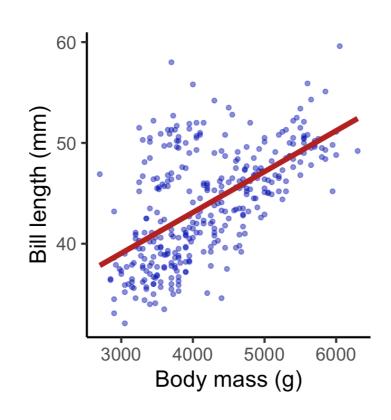
Standard error of a regression coefficient

**Population** 

Sample

Sample coefficient





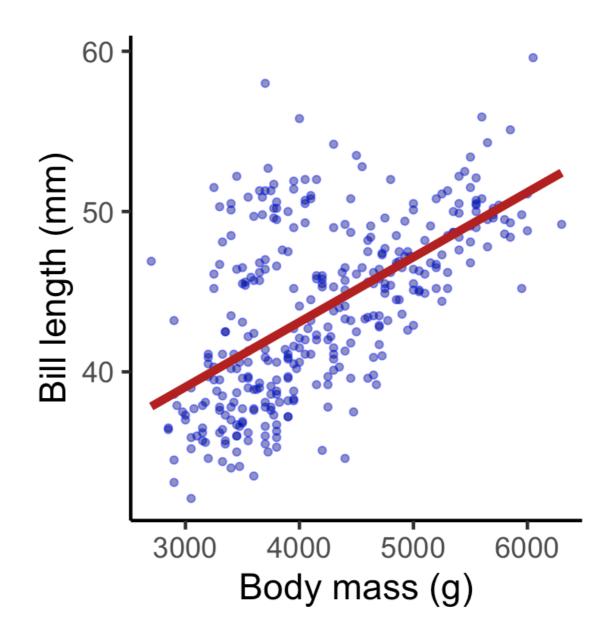
Standard error of a regression coefficient

#### Recap

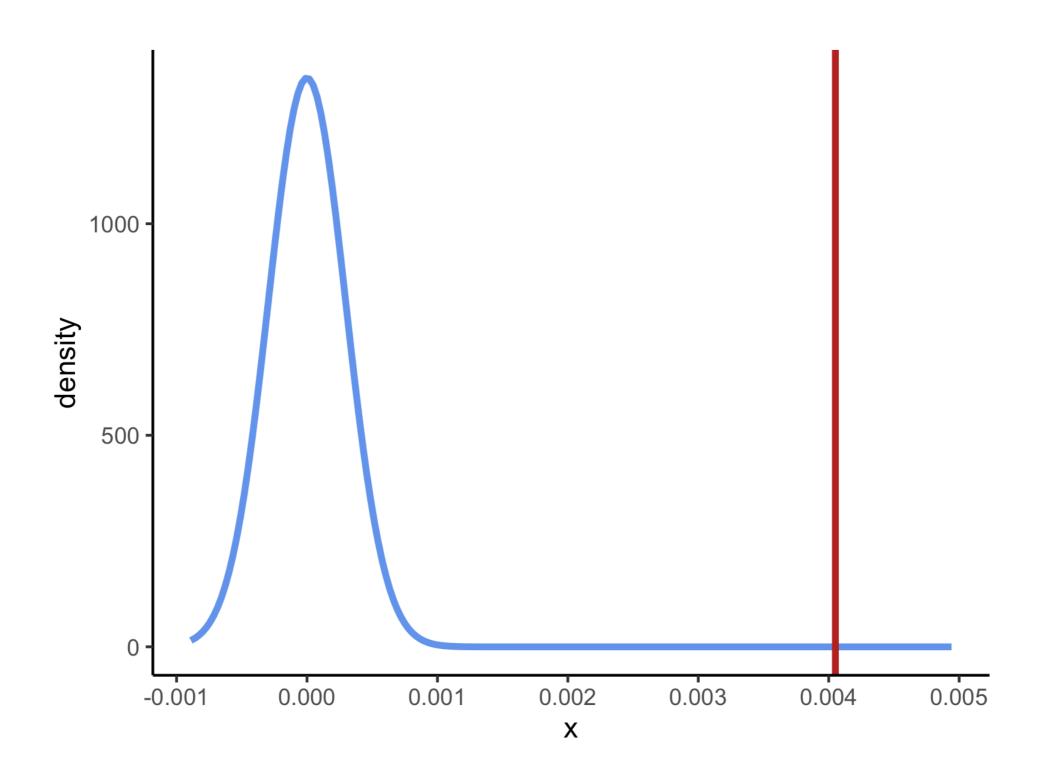
- Sampling statistics are approximately normally distributed
- 2. From the central limit theorem, we can get the standard error of the sampling distribution from just one sample
- 3. R will tell you the point estimate and the standard error when you fit a model
- 4. The p-value is the probability of getting a point estimate that many standard errors away from 0

#### **Motivation**

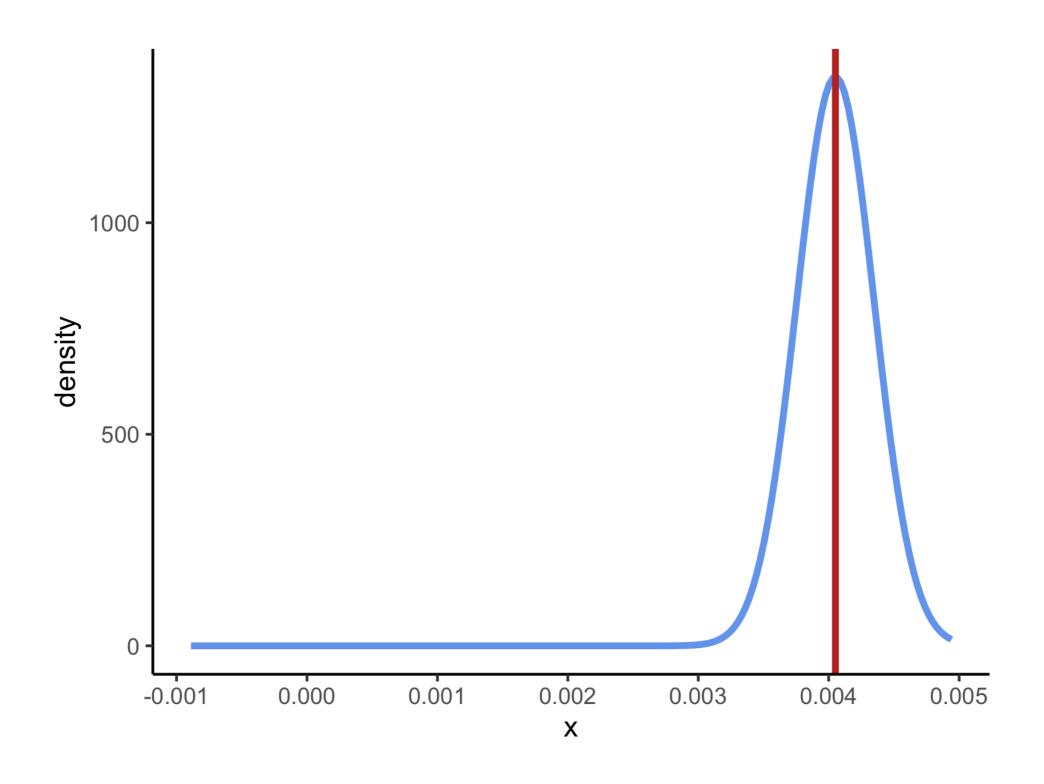
$$\hat{\beta}_1 = 0.0041$$
 $\beta_1 \in (???,???)$ 



#### Recycling standard errors



#### Recycling standard errors



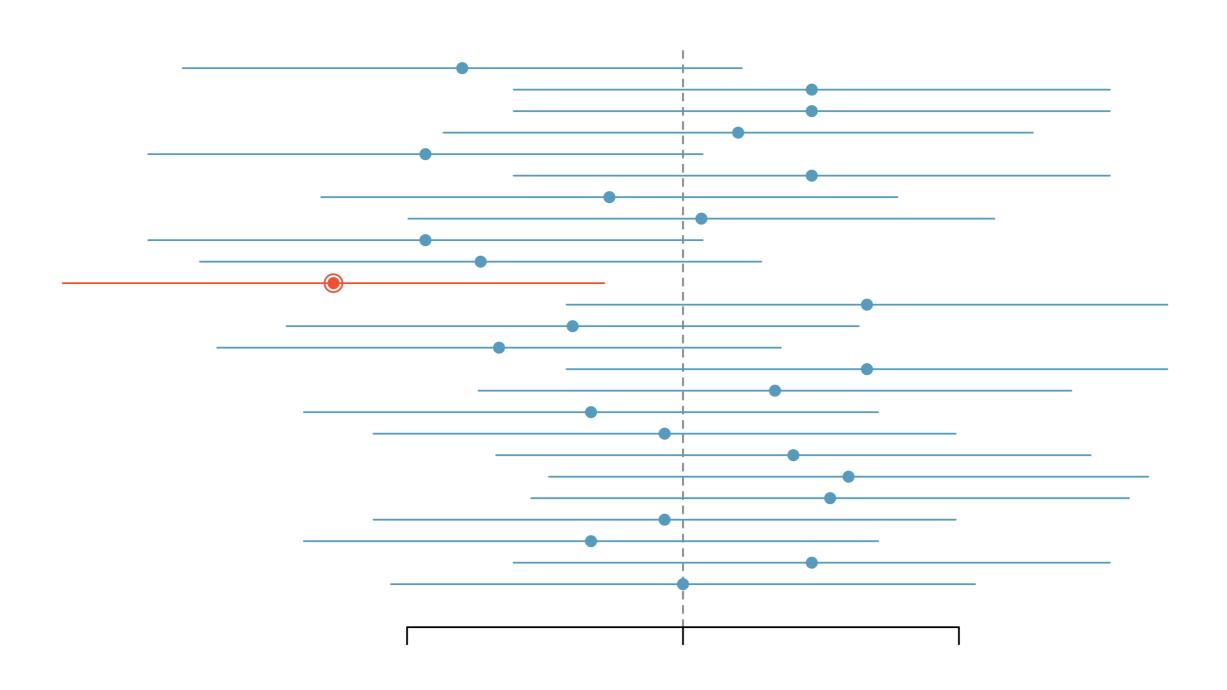
#### Interpretation

Choose the correct interpretation of the confidence interval:

"We are 95% confident the true coefficient is between 0.0035 and 0.0046."

- A. We are 95% confident the true coefficient falls in this range.
- B. The true coefficient will fall in this range 95% of the time.
- C. This range has a 95% probability of containing the true coefficient.

#### Interpretation



#### Recap

- 1. We know point estimates aren't perfect confidence intervals provide a useful bounds.
- 2. Use the standard error again, but center the distribution on the point estimate.
- 3. Be careful with interpretation! "Confidence" refers to the procedure, not to the probability the Cl contains the population parameter.

### Summary

Could our sample statistic point estimate be explained just by randomness?

#### Hypotheses

- $H_0$  no effect.  $H_A$  some effect.
- If the point estimate is improbable under the null hypothesis, reject the null. Otherwise, fail to reject.

#### Two methods for estimating null distribution

- Randomization.
- Normal approximation.

#### Confidence intervals

 An interval that we are confident contains the population parameter.