

# Housekeeping

• Office hours tomorrow: 10:30am-12:00pm

#### Recap

- We have seen how to perform hypothesis tests for questions involving the following:
  - A single proportion (STAT 201 sleep)
  - Independence of two categorical variables (banker sex discrimination)
    - Think of as one population
  - Difference in two proportions (blood thinner)
    - Think of as two populations
- We are now going to see another hypothesis test, this time for *numerical* data

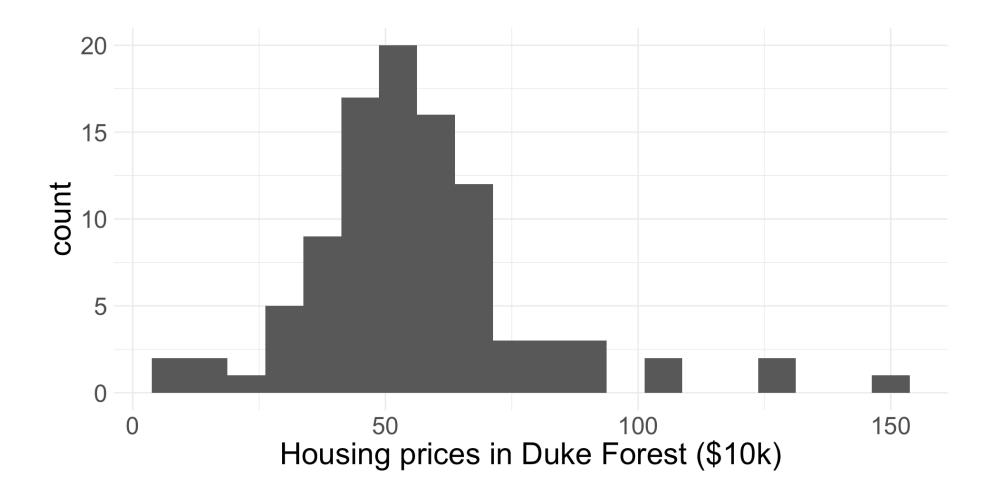
# Test for a single mean

#### Running example + form hypotheses

We will use the duke\_forest dataset provided in openintro. It provides data on some houses that were sold in the Duke Forest neighborhood of Durham, NC in November 2020.

- Before we look at the data, we should form our hypotheses. Suppose I am interested in learning if the average price of houses in Duke Forest is \$500,000 or not.
- What might our hypotheses be?
  - : versus:, where is the average house of prices in Duke Forest in \$10,000
  - Terminology: I will refer to as my "null hypothesized value". (i.e. the specific value of in )

#### Collect data



The observed/sample mean housing price is \$55.99k from a sample of 98 houses.

• Now we must determine if we have "convincing evidence"! Choose

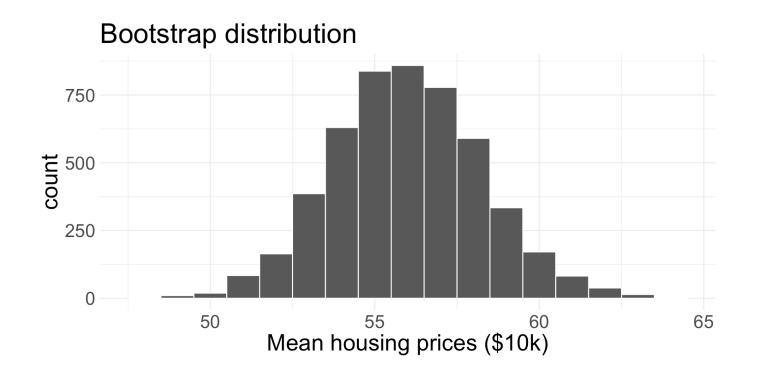
## Simulating null distribution

To simulate from the null distribution, we need to operate in a world where is true

- So, I need to repeatedly simulate data sets of size 98 where the true mean is, without change anything else about the data sets.
- If I don't want to make any assumptions about how the data behave, how might I do that?

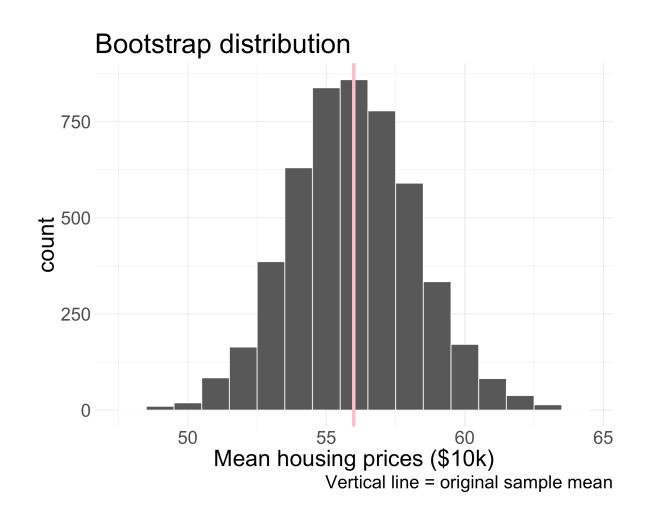
#### **Bootstrap to the rescue**

- Recall the bootstrap: we repeatedly took resamples with replacement from our original data, of same size as original data
  - Assuming the original data was representative, each one of these bootstrapped data sets gives us a plausible "new" sample of data, from which we can calculate statistics of interest



Reminder ourselves: Where is the bootstrap distribution centered?

#### Bootstrap to null distribution



- This is **not** the null distribution! The null distribution should be centered at.
- However, the null distribution should have the same variability in as the bootstrap distribution.

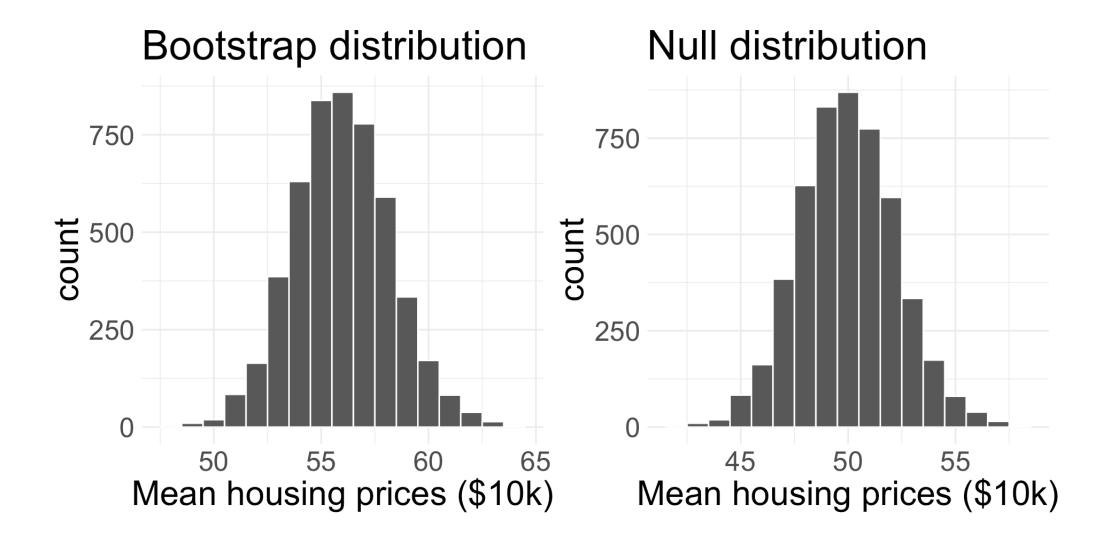
• So to get the null distribution, why not just **shift** the bootstrap distribution to be centered where we want it to be?

#### Shifting to the bootstrap distribution

- In this example, bootstrap distribution is centered at
- In order to center this distribution at , just subtract from every single bootstrapped mean
  - This will give us a simulated distribution for centered at, which is exactly the null distribution!
  - We call this "shifting the bootstrap distribution", because we simply shift where the bootstrap distribution is centered

```
1 mu0 <- 50
2 # how much to shift by, where xbar is sample mean housing prices
3 shift <- xbar - mu0
4 # shift my vector of bootstrapped sample means
5 null_dist <- boot_means - shift</pre>
```

#### **Null distribution**



Notice where the distributions are centered

# Obtain the p-value

#### : versus:

Our observed sample mean housing price is 55.99.



What does it mean to be "as or more extreme" now?

#### Two-sided alternative hypothesis

- This is the first time we've seen a two-sided hypothesis
- Since the alternative is "double sided", we can be extreme in **both** the positive and negative direction!



## Obtain the p-value (cont.)

Let represent the amount we shifted the distribution by:

Simulated means as or more extreme than or will contribute:



```
1 sum( (null_dist >= mu0 + shift) | (null_dist <= mu0 - shift))/B
[1] 0.0098</pre>
```

#### Make decision and conclusion

Make a decision and conclusion in the context of the research question.

• Since our p-value of 0.0098 is less than the significance level of 0.05, we reject. We have convincing evidence to suggest that the true average housing price of homes in Duke Forest in 2020 was not \$50k.

#### **Comprehension questions**

- Why did we shift the bootstrap distribution?
- How do we estimate the p-value in the case of a two-sided alternative hypothesis?