

DSFA

Spring 2020

# Lecture 17

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Percentiles and the Bootstrap

# Conclusions From a Test

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**Hypothesis test**

```
graph LR; A[Hypothesis test] --> B["Fail to reject the null hypothesis  
(data are not inconsistent with the null hypothesis - inconclusive)"]; A --> C["Reject the null hypothesis  
(data are inconsistent with the null hypothesis - accept the alternative)"];
```

**Fail to reject** the null hypothesis  
(data are not inconsistent with the null hypothesis - inconclusive)

**Reject** the null hypothesis  
(data are inconsistent with the null hypothesis - accept the alternative)

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# Definition of $P$ -value

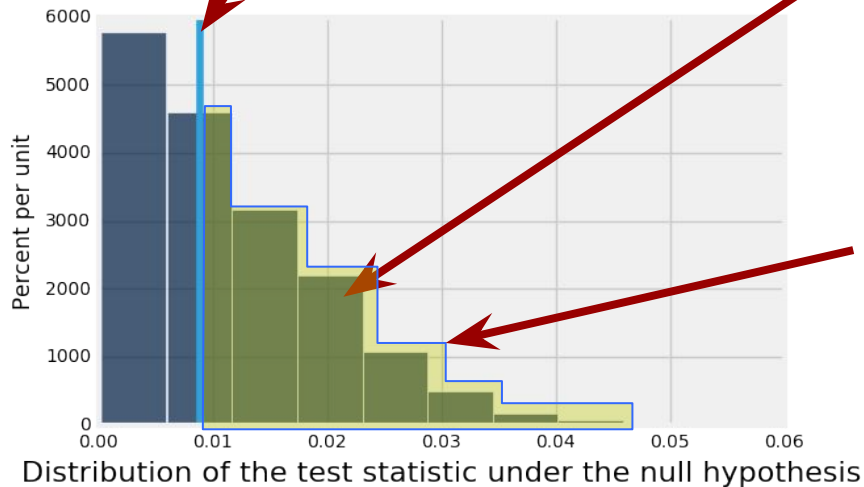
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The  $P$ -value is the chance,

- under the null hypothesis,
  - that the test statistic
  - is equal to the value that was observed in the data or is even further in the direction of the alternative.
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# Quantifying Conclusions

P(the **test statistic** would be **equal to or more extreme** than the **observed test statistic** under the null hypothesis)



Evaluating Mendel's  
pea flower hypothesis

This area is the P-value  
(approximately)

# Conventions of Consistency





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- **“Inconsistent”**: The test statistic is in the tail of the null distribution.
  - **“In the tail,” first convention**:
    - The area in the tail is less than 5%.
    - The result is “statistically significant.”
  - **“In the tail,” second convention**:
    - The area in the tail is less than 1%.
    - The result is “highly statistically significant.”
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# Can the Conclusion be Wrong?

**Yes.**

Type I error

	Null is true	Alternative is true
Test rejects the null		
Test doesn't reject the null		

(Demo)

Type II error

# An Error Probability

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- The cutoff for the P-value is an error probability.
  - If:
    - your **cutoff is 5%** (your significance level)
    - and the **null hypothesis happens to be true**
    - (but you don't know that)
  - then there is about a **5% chance** that **your test will reject the null hypothesis anyway**.
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# Type I and Type II errors

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- The significance level (or p-value cutoff) is the probability of a Type I error

Type I error = Reject null when it is true

- What if the alternative is true?

Type II error = Fail to reject null when it is false

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# More on P-Hacking

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Suppose you conduct 10 independent hypothesis test, each at a 5% significance level; i.e. the null hypothesis is rejected if  $p < 0.05$ .

The probability that at least one null hypothesis is rejected is

- A. 0.05 or less
  - B. Between 0.05 and 0.4
  - C. Between 0.4 and 0.5
  - D. Between 0.5 and 0.95
  - E. 0.95 or more
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# Percentiles

# Computing Percentiles

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The 80th percentile of a set of numbers is the smallest value in the sample that is at least as large as 80% of the sample

For  $s = [1, 7, 3, 9, 5]$ , `percentile(80, s)` is 7

The 80th percentile is ordered element 4:  $(80/100) * 5$

Percentile

Size of set

For a percentile that does not exactly correspond to an element, take the next greater element instead

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# The percentile Function

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- The  $p$ th percentile is the smallest value in the sample at least as large as  $p\%$  of the values in the sample
  - Function in the `datascience` module:  
`percentile(p, values)`
  - `p` is between 0 and 100
  - Returns the  $p$ th percentile of the array
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# Discussion Question

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Which are `True`, when `s = [1, 7, 3, 9, 5]`?

`percentile(10, s) == 0`

`percentile(39, s) == percentile(40, s)`

`percentile(40, s) == percentile(41, s)`

`percentile(50, s) == 5`

(Demo)

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# Estimation (Review)

# Inference: Estimation

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- What is the value of a population parameter?
- If you have a census (that is, the whole population):
  - Just calculate the parameter and you're done
- If you don't have a census:
  - Take a random sample from the population
  - Use a statistic as an **estimate** of the parameter

(Demo)

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# Variability of the Estimate

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- One sample → One estimate
- But the random sample could have come out differently
- And so the estimate could have been different
- Main question:
  - **How different could the estimate have been?**
- The variability of the estimate tells us something about how accurate the estimate is:

$$\text{estimate} = \text{parameter} + \text{error}$$

(Demo)

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# Where to Get Another Sample?

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- One sample → One estimate
  - To get many values of the estimate, we needed many random samples
  - Can't go back and sample again from the population:
    - No time, no money
  - Stuck?
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# The Bootstrap

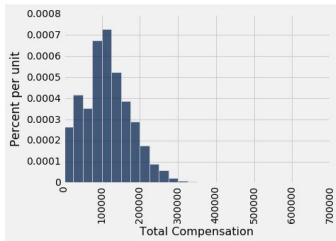
# The Bootstrap

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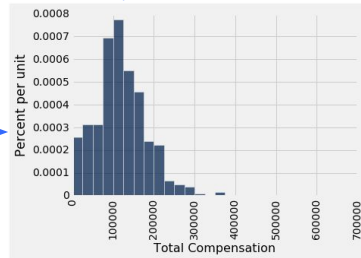
- A technique for simulating repeated random sampling
  - All that we have is the original sample
    - ... which is large and random
    - Therefore, it probably resembles the population
  - So we sample at random from the original sample!
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# Why the Bootstrap Works

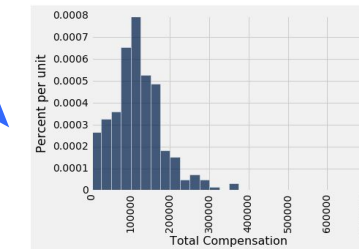
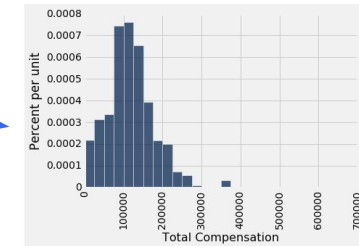
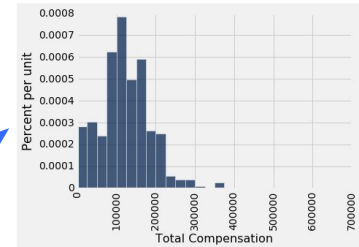
population



sample



resamples



All of these look pretty similar, most likely.

# Key to Resampling

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- From the original sample,
  - draw at random
  - with replacement
  - as many values as the original sample contained
- The size of the new sample has to be the same as the original one, so that the two estimates are comparable

(Demo)

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