## **Bootstrap and confidence intervals**

Introduction to data analysis: Lecture 7

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#### **RE: Percentiles**

- The pth percentile is the value below which p% of the observations lie
  - The median is the 50<sup>th</sup> percentile
- Q1 = the first quartile = the 25<sup>th</sup> percentile
- Q3 = the third quartile = the 75<sup>th</sup> percentile
- IQR = Inter-quartile range = Q3-Q1

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- IQR = Inter-quartile range = Q3-Q1
- For a percentile that does not exactly correspond to an observed element, there are many interpolation methods.
- We'll take the first element that is greater.
  - 100, 92, 97, 83, 67
    - Median? Q1? Q3?
- We'll use numpy percentile (with "method='higher'")
   Or pandas quantile (with "interpolation='higher'")

#### **Estimation**

- How big is an unknown 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

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- Main question:

#### How different could the estimate have been?

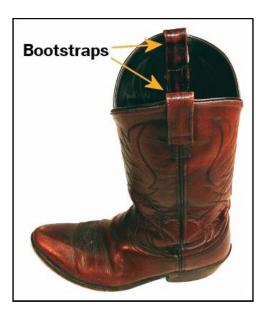
 The variability of the estimate tells us something about how accurate the estimate is: estimate = parameter + error

# Another sample?

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

- 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
    - According to?
  - That is, the empirical distribution of the sample looks like the probability distribution of the population (hopefully)

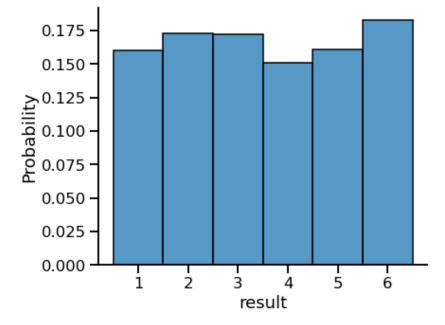
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- So we sample at random from the original sample!
  - Behave as if they are the same

• A couple of lectures ago, we simulated rolls of a die

```
# empirical distribution of roll of a die
def empirical_hist_die(num_rolls):
    die_df = pd.DataFrame({'result': np.random.choice(die, num_rolls)})
    sns.displot(die_df, x='result', bins=die_bins, stat='probability', height=5, aspect=1.3)
    return die_df

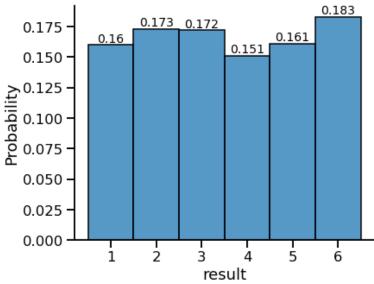
die_simulation_results = empirical_hist_die(1000)
```



• The empirical distribution of the sample of 1000 die rolls we got is not

uniform, but it is not very far from uniform

```
die_probs = []
num_rolls = die_simulation_results.shape[0]
for i in range(1,7):
    die_probs.append(np.count_nonzero(die_simulation_results == i)/num_rolls)
die_probs
[0.16, 0.173, 0.172, 0.151, 0.161, 0.183]
```

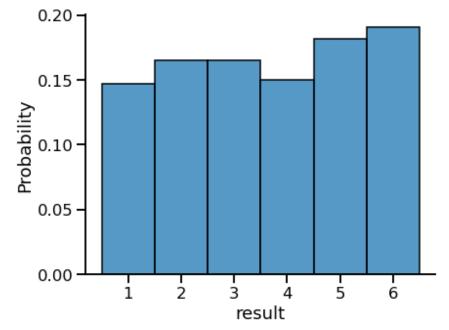


 We can use this empirical distribution (which is close to the true distribution) to generate additional simulated samples

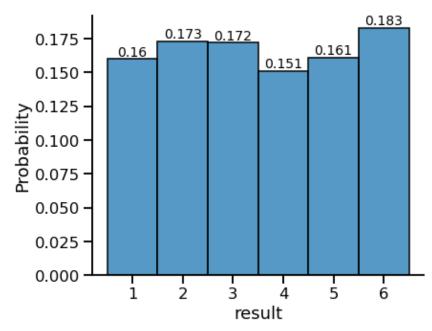
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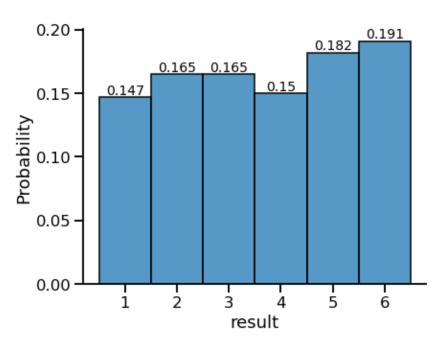
bootstrap_die_simulation_results = empirical_hist_die(1000)
```



The distribution of the bootstrapped sample we got is similar to the distribution of the sample we got when using the true distribution of the die



1000 simulations of a die roll using its true probability distribution



1000 simulations of a die roll using the empirical distribution of our initial sample

# Key to resampling

From the original sample

- Draw at random
- With replacement
- As many value as the original sample contained

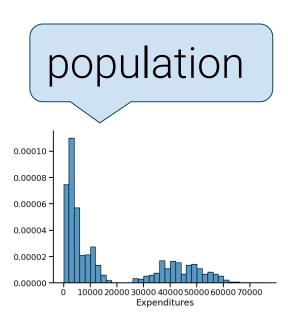
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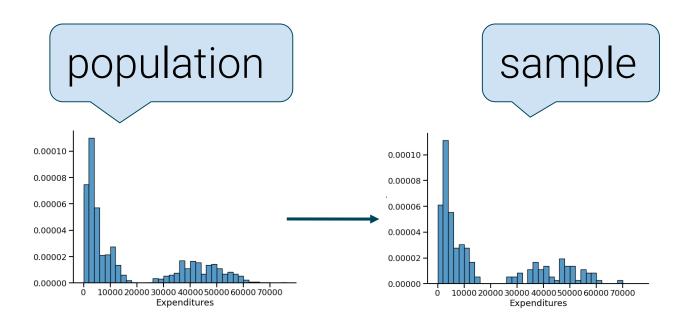
- Draw at random
- With replacement
- As many value as the original sample contained
- The size of the new sample has to be the same as the original so that the two estimates we get are comparable
  - The variability of the estimate is linked to the sample size

[notebook]

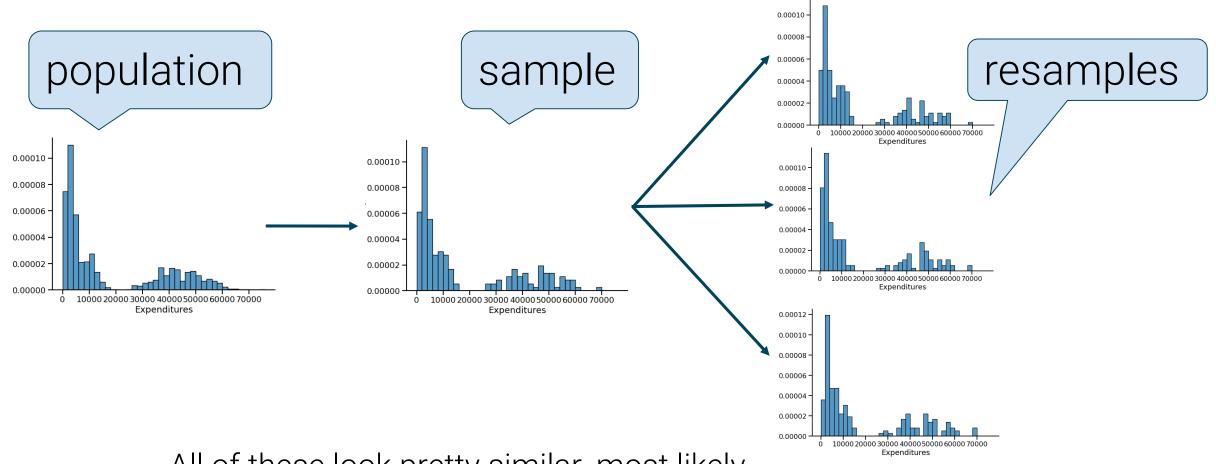
## Why the bootstrap works



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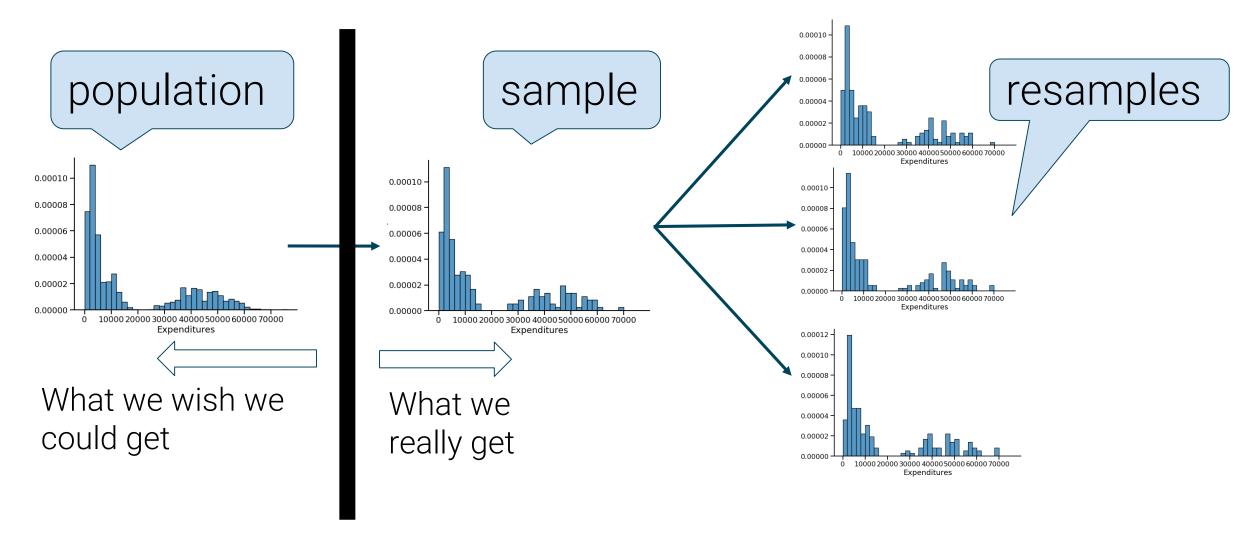


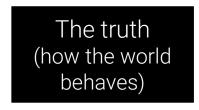
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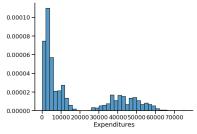


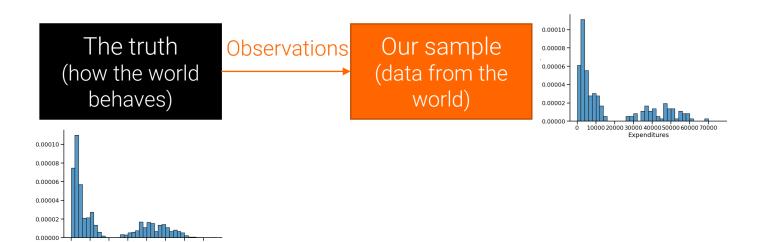
All of these look pretty similar, most likely.

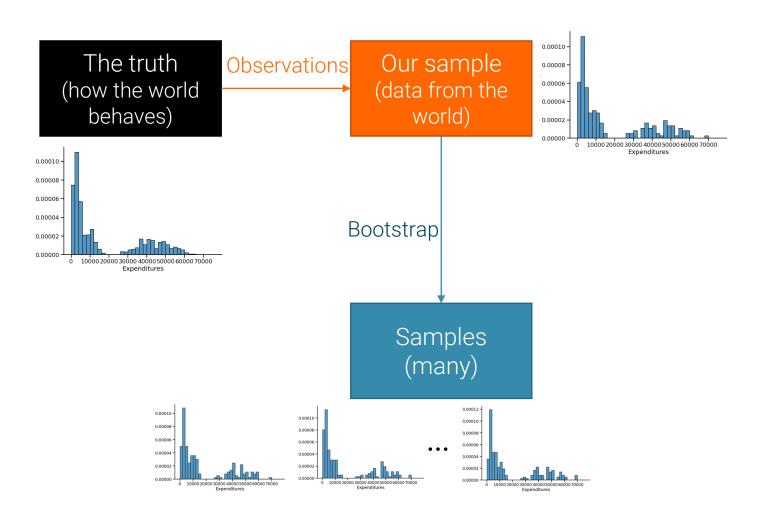
# Why we need the bootstrap

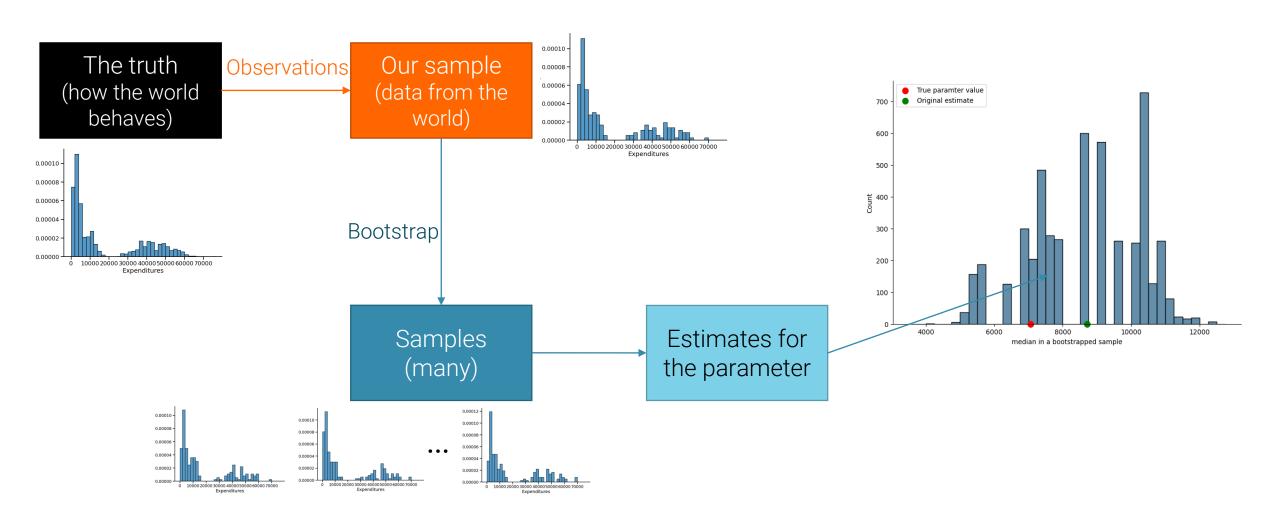


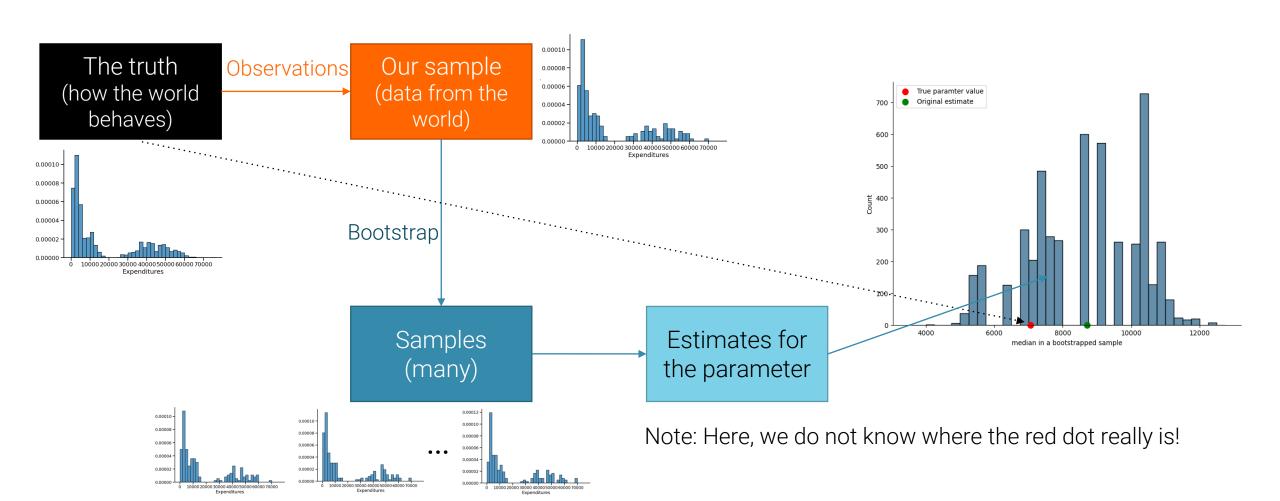












### Confidence Interval (רווח סמך)

- So, we got many possible estimates for the parameter
  - One for each bootstrapped sample
  - The distribution we get tells us how our estimate may change due to variability in data
- Some of these are more likely than others
  - Values that are in the tail of the distribution of estimates are less likely
- It is very unlikely that the one estimate we got from the original sample equals the true parameter exactly, but it is likely that true parameter lies within a range where we see most of the estimates we got

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- It is very unlikely that the one estimate we got from the original sample equals the true parameter exactly, but it is likely that true parameter lies within a range where we see most of the estimates we got
- Confidence interval = a range of values within which we expect the true parameter lies
  - In a certain percentage of the cases
  - Based on random sampling (and the distribution of bootstrapped estimates)

#### Confidence level

- Often, we look for "95% confidence intervals"
- 95% is called the **confidence level** 
  - Could be any percent between 0 and 100
  - Higher level means wider intervals

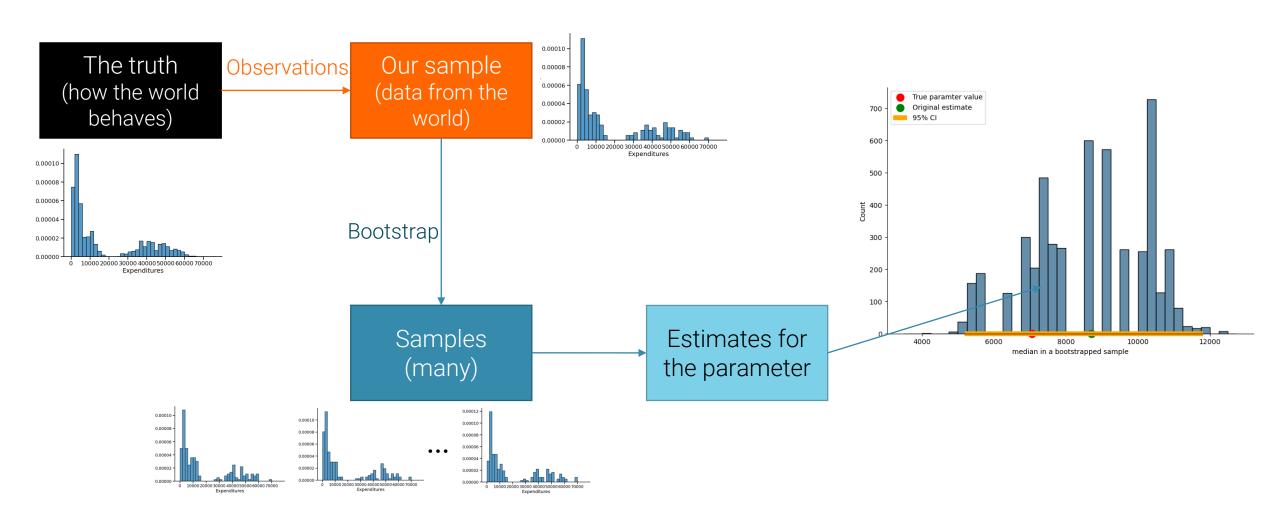
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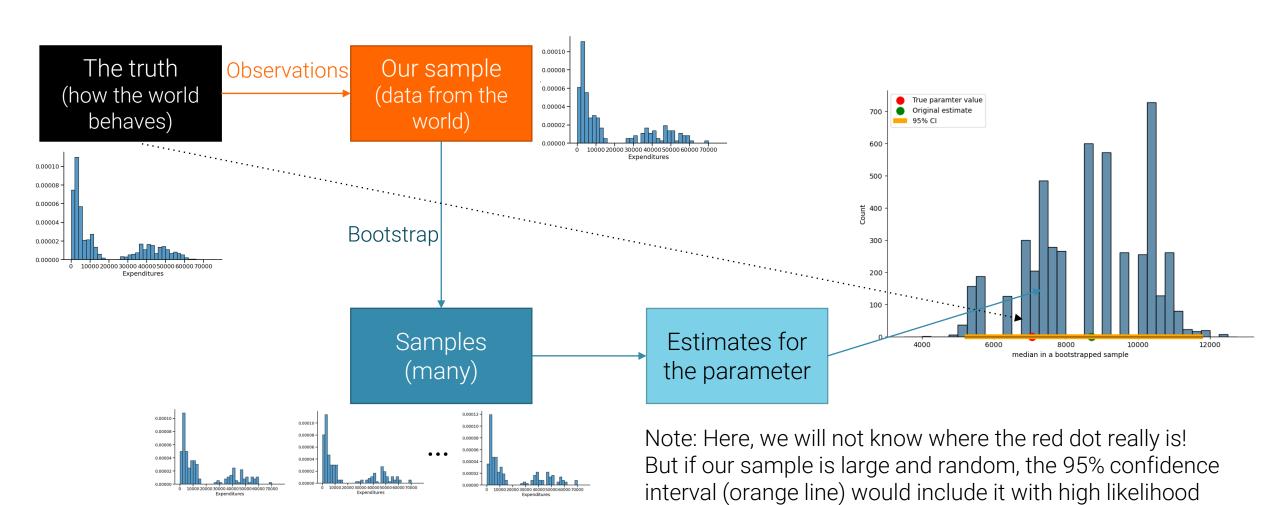
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- Default seaborn error bars (e.g. in bar plots) show 95% confidence interval based on bootstrap with 1000 replications

(notebook)





#### True or False?

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- 2. There is a 0.95 probability that the average gross income of movies in the population is in the range (48,966,237, 53,282,655).

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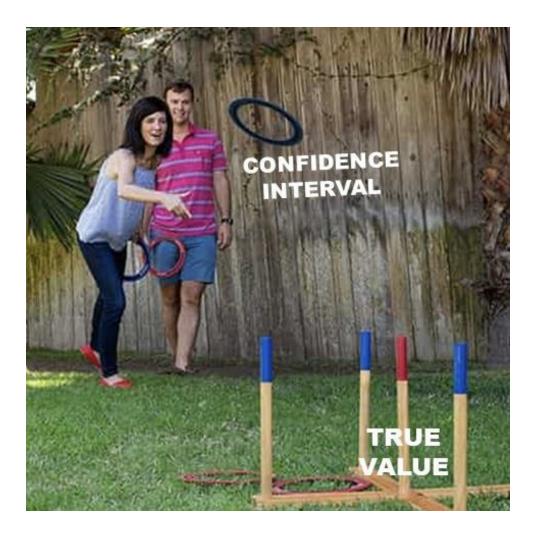
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#### True or False:

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- 2. There is a 0.95 probability that the average gross income of movies in the population is in the range (48,966,237, 53,282,655).
- 3. If we create 1000 confidence intervals using the bootstrap, we expect around 950 of them to include the true value of the average gross income of movies in the population

# Illustrating CI







## When Not to use the Bootstrap

- If you're trying to estimate very high or very low percentiles, like min or max
- If you're trying to estimate any parameter that's greatly affected by rare elements of the population
- If observations are dependent on one another (time series, spatial data)
- If the original sample is too small to reliably represent the probability distribution in the population
- In general, if your sample is biased in some way (not random), then bootstrap will not help you

# Using CI for hypothesis testing

- Our approximate 95% confidence interval for the average gross income of movies in the population is (48,966,237, 53,282,655) dollars
- This estimation process is good in 95% of the time (assuming our sample is large and random)

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- Suppose someone wants to test the following hypotheses:
  - $H_0$  (null hypothesis): The mean gross income of movies is \$55M
  - H<sub>1</sub> (alternative hypothesis): The mean gross income of movies is **not** \$55M

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- If our significance level is 0.05, we will reject the null hypothesis!
  - We are 95% confident that (48,966,237, 53,282,655) contains the true mean gross income of movies.
  - If the true mean income were 55M, it is highly unlikely we would get a sample that leads us to a CI of (48,966,237, 53,282,655)
  - In particular, we are (at least) 95% confident that \$55M is not the true mean gross income of movies → We reject H<sub>0</sub> at the 5% level

# CI for testing population mean

- Null hypothesis: Population average = x
- Alternative hypothesis: Population average ≠ x
- Cutoff for P-value: p%
- Method:
  - Construct a (100-p)% confidence interval for the population average
  - If x is not in the interval, reject the null
  - If x is in the interval, can't reject the null

#### CI for testing difference in population means

- Null hypothesis: Difference between population averages = 0
- Alternative hypothesis: Difference between population averages ≠ 0
- Cutoff for P-value: p%
- Method:
  - Construct a (100-p)% confidence interval for the difference between population averages
  - If 0 is not in the interval, reject the null
  - If 0 is in the interval, can't reject the null

