



COMILLAS

UNIVERSIDAD PONTIFICIA

ICAI

ICADE

CIHS

# Inferential Statistics

DID THE SUN JUST EXPLODE?  
(IT'S NIGHT, SO WE'RE NOT SURE.)

THIS NEUTRINO DETECTOR MEASURES  
WHETHER THE SUN HAS GONE NOVA.

THEN, IT ROLLS TWO DICE. IF THEY  
BOTH COME UP SIX, IT LIES TO US.  
OTHERWISE, IT TELLS THE TRUTH.

LET'S TRY.

DETECTOR! HAS THE  
SUN GONE NOVA?

(ROLL)  
YES.



FREQUENTIST STATISTICIAN:

THE PROBABILITY OF THIS RESULT  
HAPPENING BY CHANCE IS  $\frac{1}{36} = 0.027$ .  
SINCE  $p < 0.05$ , I CONCLUDE  
THAT THE SUN HAS EXPLODED.



BAYESIAN STATISTICIAN:

BET YOU \$50  
IT HASN'T.

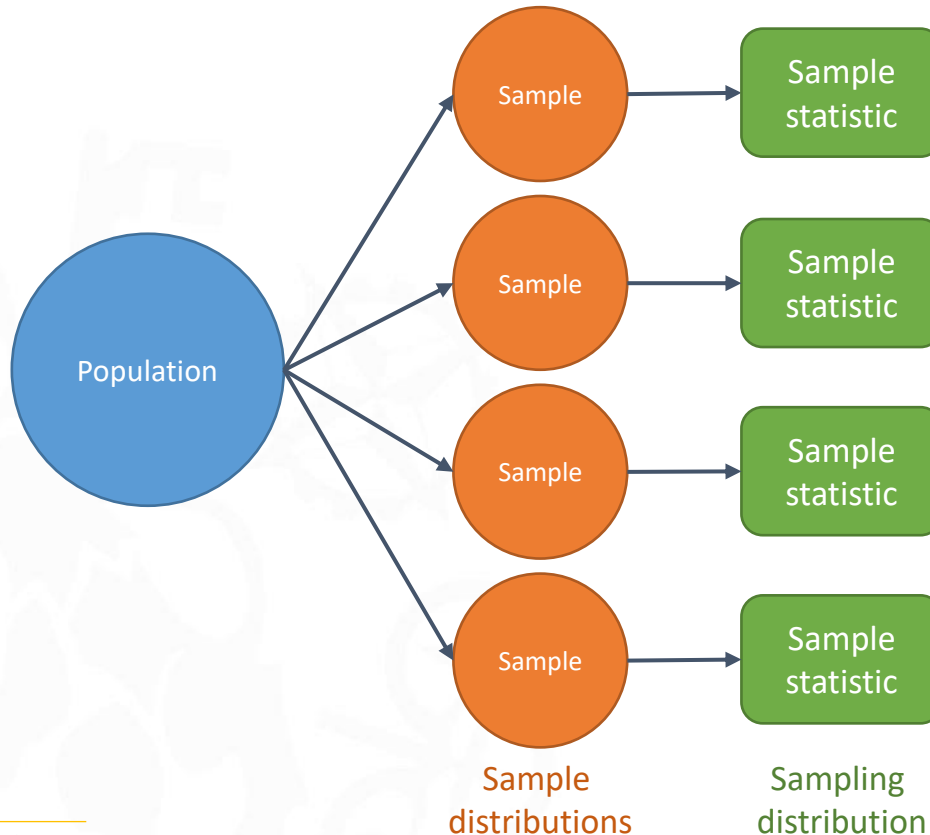


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4. Comparing two means
5. Inference for proportions

# Central Limit Theorem (CLT)

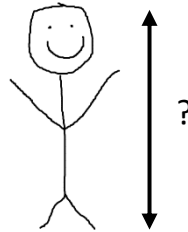
## Sample distributions



# Central Limit Theorem (CLT)

## Standard Error

We want to obtain information about the height of men in Spain.



AR:  $x_{AR,1}, x_{AR,2} + \dots + x_{AR,1000} \longrightarrow \bar{x}_{AR}$

...  
MD:  $x_{MD,1}, x_{MD,2} + \dots + x_{MD,1000} \longrightarrow \bar{x}_{MD}$

...  
ZM:  $x_{ZM,1}, x_{ZM,2} + \dots + x_{ZM,1000} \longrightarrow \bar{x}_{ZM}$

$$\mu = \frac{x_1 + x_2 + \dots + x_N}{N}$$

$$\sigma = \sqrt{\frac{\sum_{i=1}^N (x_i - \mu)^2}{N}}$$

Sampling  
distribution

$mean(\bar{x}) \approx \mu$

$\uparrow n$

$\downarrow$  Standard error

$SD(\bar{x}) < \sigma$

# Central Limit Theorem (CLT)

## Definition

**Central Limit Theorem (CLT):** The distribution of sample statistics is nearly normal, centered at the population mean, and with a standard deviation equal to the population standard deviation divided by square root of the sample size.

$$\bar{x} \sim N(\text{mean} = \mu, SE = \frac{\sigma}{\sqrt{n}})$$

Usually, standard deviation of the population is not known, so standard deviation of sample is used:

$$\bar{x} \sim N(\text{mean} = \mu, SE = \frac{s}{\sqrt{n}})$$

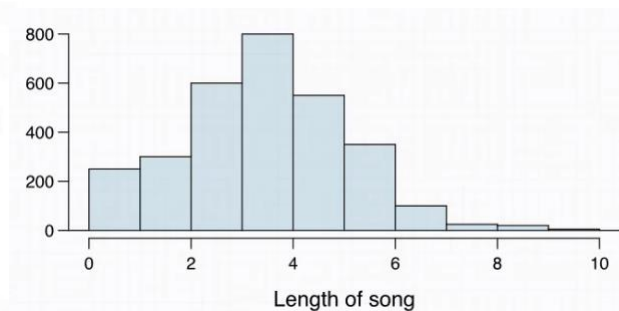
**Conditions for the CLT:**

1. **Independence:** Sampled observations must be independent.
  - Random sample/assignment
  - If sampling with replacement,  $n < 10\%$  of population
2. **Sample size/skew:** Either the population distribution is normal, or the sample size is large (rule of thumb:  $n > 30$ )

# Central Limit Theorem (CLT)

## Example

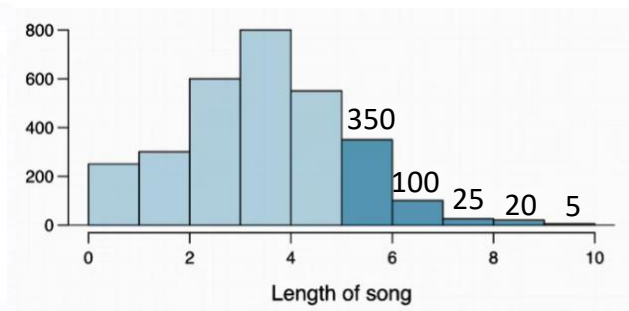
Suppose my iPod has 3,000 songs. The histogram below shows a distribution of the lengths of these songs. We also know that, for this iPod, the mean length is 3.45 minutes and the standard deviation is 1.63 minutes. Calculate the probability that a randomly selected song lasts more than five minutes.



# Central Limit Theorem (CLT)

## Example

Suppose my iPod has 3,000 songs. The histogram below shows a distribution of the lengths of these songs. We also know that, for this iPod, the mean length is 3.45 minutes and the standard deviation is 1.63 minutes. Calculate the probability that a randomly selected song lasts more than five minutes.



$x$  = length of one song

$$P(x > 5) = \frac{350 + 100 + 25 + 20 + 5}{3000} = \frac{500}{3000} \approx 0.17$$

# Central Limit Theorem (CLT)

## Example

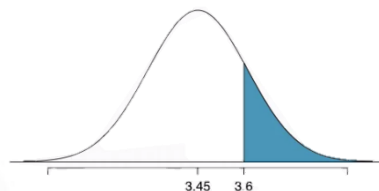
Another question. I'm about to take a trip to visit my parents and the drive is six hours. I make a random playlist of 100 songs. What is the probability that my playlist lasts the entire drive?

*6 hours = 360 minutes*

$$P(x_1 + x_2 + \dots + x_{100} > 360 \text{ min})? = P(\bar{x} > 3.6 \text{ min})?$$

$$\text{By CLT: } \bar{x} \sim N(\text{mean} = \mu = 3.45, SE = \frac{\sigma}{\sqrt{n}} = \frac{1.63}{\sqrt{100}} = 0.163)$$

$$Z = \frac{3.6 - 3.45}{0.163} = 0.92$$



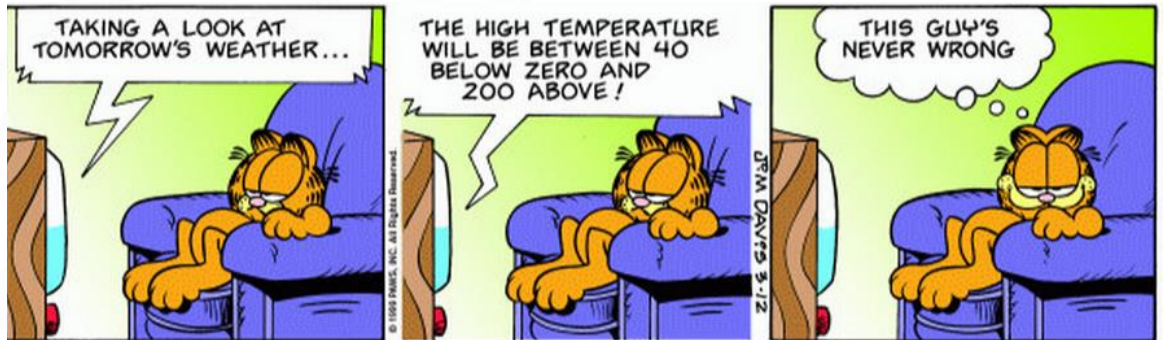
So we only need to calculate  $P(Z > 0.92)$



# Confidence Interval Definition

A plausible range of values for the population parameter is called a **confidence interval**. Why are they important?

- If we report a point estimate, hitting the exact value for the population parameter is highly unlikely.
- If we report a range of plausible values, having the exact value for the population inside the range is much more probable.



# Confidence Interval For a mean

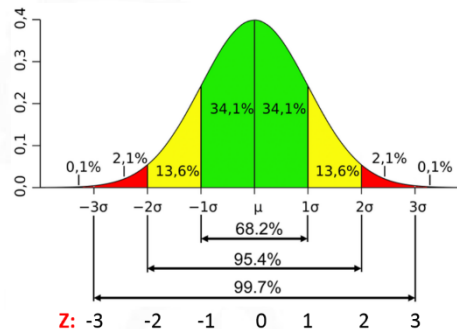
By CLT:

$$\bar{x} \sim N(\text{mean} = \mu, SE = \frac{s}{\sqrt{n}})$$

So an approximate 95% confidence interval for  $\mu$  is:

$$CI_{(95\%)} \approx \bar{x} \pm 2 \cdot SE$$

Where  $2 \cdot SE$  is called the margin of error (ME)



# Confidence Interval For a mean

For different confidence levels, the confidence interval for a population mean is defined as the sample mean plus/minus a margin of error calculated as the critical value corresponding to the middle XX% of the normal distribution times the standard error of the sampling distribution:

$$CI_{(XX\%)} = \bar{x} \pm z^* \cdot \frac{s}{\sqrt{n}}$$

A confidence value of XX% determines that for XX out of 100 random samples, the true value of the population parameter would fall inside the confidence interval of each random sample.

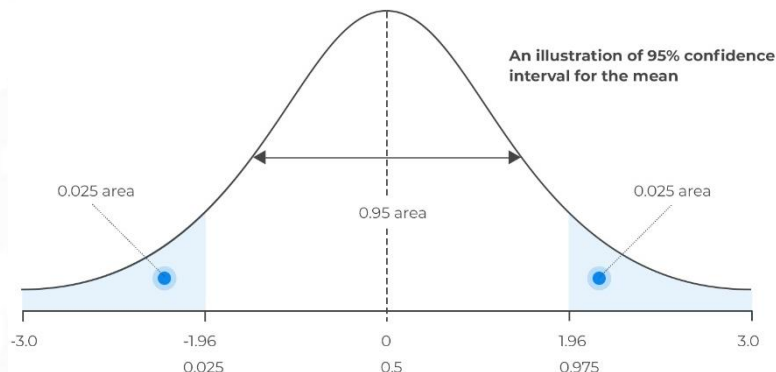
## Conditions for this confidence interval:

1. **Independence:** Sampled observations must be independent.
  - Random sample/assignment
  - If sampling with replacement,  $n < 10\%$  of population
2. **Sample size/skew:**  $n > 30$ , larger if the population is severely skewed

# Confidence Interval

## Finding the critical value

For the 95% confidence interval:



Second decimal place					Z
0.07	0.06	0.05	0.04	0.00	
0.0003	0.0003	0.0003	0.0003	0.0003	-3.4
0.0004	0.0004	0.0004	0.0004	0.0005	-3.3
0.0005	0.0006	0.0006	0.0006	0.0007	-3.2
0.0008	0.0008	0.0008	0.0008	0.0010	-3.1
0.0011	0.0011	0.0011	0.0012	0.0013	-3.0
0.0015	0.0015	0.0016	0.0016	0.0019	-2.9
0.0021	0.0021	0.0022	0.0023	0.0026	-2.8
0.0028	0.0029	0.0030	0.0031	0.0035	-2.7
0.0038	0.0039	0.0040	0.0041	0.0047	-2.6
0.0051	0.0052	0.0054	0.0055	0.0062	-2.5
0.0068	0.0069	0.0071	0.0073	0.0082	-2.4
0.0089	0.0091	0.0094	0.0096	0.0107	-2.3
0.0116	0.0119	0.0122	0.0125	0.0139	-2.2
0.0150	0.0154	0.0158	0.0162	0.0179	-2.1
0.0192	0.0197	0.0202	0.0207	0.0228	-2.0
0.0244	0.0250	0.0256	0.0262	0.0287	-1.9
0.0307	0.0314	0.0322	0.0329	0.0359	-1.8

$$z^* = 1.96$$

# Confidence Interval

## Accuracy vs precision

Commonly used critical values are 90%, 95%, 98% and 99%.

What happens to the confidence Interval when the confidence level increases?

1. Confidence Interval is wider, as it captures more values
2. Confidence Interval is narrower, as it is more confident in the calculation

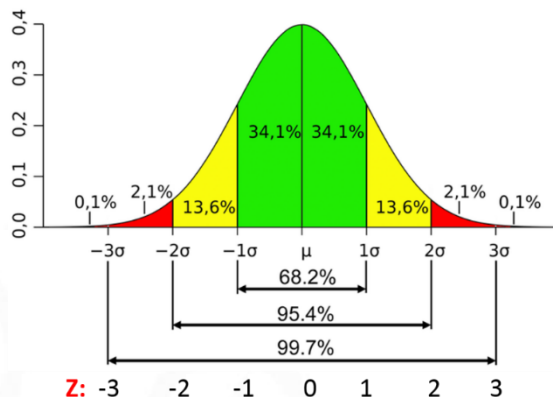
# Confidence Interval

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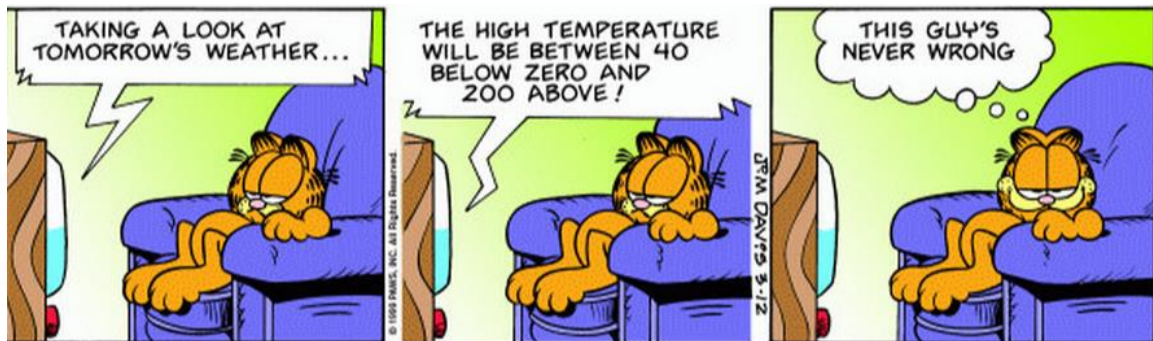


A **wider interval is more accurate**, as it is more probable to capture the output statistic true value.

# Confidence Interval

## Accuracy vs precision

However, a wider interval is less informative, i.e., it loses precision. Remember:



To get the best of both worlds we need to **increase the sample size**, as the margin error would decrease:

$$CI_{(XX\%)} = \bar{x} \pm z^* \cdot \frac{s}{\sqrt{n}}$$

# Confidence Interval

## Accuracy vs precision

The General Social Survey, the GSS, is a sociological survey used to collect data on demographic characteristics and attitudes of residents of the United States. In 2010, the survey collected responses from 1,154 U.S. residents. Based on the survey results, a 95% confidence interval for the average number of hours Americans have to relax or pursue activities that they enjoy after an average workday, was found to be 3.53 to 3.83 hours.

Determine if each of the following statements are true or false:

1. 95% of Americans spend between 3.53 to 3.83 hours relaxing after a work day.
2. 95% of random samples of 1,154 Americans will yield confidence intervals that contain the true average number of hours Americans spend relaxing after a work day.
3. 95% of the time the true average number of hours Americans spend relaxing after a work day is between 3.53 and 3.83 hours.



# Confidence Interval

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# Confidence Interval

## Backtracking to $n$ from ME

Given a target margin of error, confidence level, and information on the variability of the sample or the population, we want to determine the required sample size to achieve the desired margin of error.

Let's check an example

# Confidence Interval

## Backtracking to n from ME

Suppose a group of researchers want to test the possible effect of an epilepsy medication taken by pregnant mothers on the cognitive development of their children. As evidence, they want to estimate the IQs of three-year-old children born to mothers who were on this medication during their pregnancy.

Previous studies suggest that the standard deviation of IQ scores of three-year-old children is 18 points.

How many such children should the researches sample in order to obtain a 90% confidence interval with a margin of error less than or equal to four points?

$$ME = 4$$

$$CL = 90\%$$

$$z^* = 1.96$$

$$\sigma = 18$$

# Confidence Interval

## Backtracking to n from ME

Suppose a group of researchers want to test the possible effect of an epilepsy medication taken by pregnant mothers on the cognitive development of their children. As evidence, they want to estimate the IQs of three-year-old children born to mothers who were on this medication during their pregnancy.

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$$ME = 4$$

$$CL = 90\%$$

$$z^* = 1.96$$

$$\sigma = 18$$

$$4 = 1.65 \cdot \frac{18}{\sqrt{n}} \rightarrow n = \left( \frac{1.65 \cdot 18}{4} \right)^2 = 55.13 \rightarrow 56 \text{ children}$$

# Confidence Interval

## Backtracking to n from ME

Earlier we found that we needed at least 56 children in the sample to achieve a maximum margin of error of four points. How would the required sample size change if we want to further decrease the margin of error by half?

$$\frac{1}{2} \cdot ME = z^* \cdot \frac{s}{\sqrt{n}} \cdot \frac{1}{2}$$

# Confidence Interval

## Backtracking to n from ME

Earlier we found that we needed at least 56 children in the sample to achieve a maximum margin of error of four points. How would the required sample size change if we want to further decrease the margin of error by half?

$$\frac{1}{2} \cdot ME = z^* \cdot \frac{s}{\sqrt{n}} \cdot \frac{1}{2}$$
$$\frac{1}{2} \cdot ME = z^* \cdot \frac{s}{\sqrt{4 \cdot n}}$$

So, to reduce the ME by half, we need to duplicate the sample size!

# Hypothesis testing

## Hypothesis

Null – $H_0$	Often either a skeptical perspective or a claim to be tested	=
Alternative – $H_a$	Represents an alternative claim under consideration and is often represented by a range of possible parameter values	<, >, $\neq$

The skeptic will not abandon  $H_0$  unless the evidence in favor of  $H_a$  is so strong that  $H_0$  must be rejected in favor of  $H_a$

# Hypothesis testing

## Hypothesis via CI

A previous study found that a 95% confidence interval for the average number of exclusive relationships college students have been in to be 2.7 to 3.7. Based on this confidence interval, do these data support the hypothesis that college students on average have been in more than three exclusive relationships?



# Hypothesis testing

## Hypothesis via CI

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Being  $X$  the number of exclusive relationships college students have been in, what are the hypothesis?

# Hypothesis testing

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Being  $X$  the number of exclusive relationships college students have been in, what are the hypothesis?

$H_0: \mu = 3$  College students have been in 3 exclusive relationships, on average

$H_a: \mu > 3$  College students have been in more than 3 exclusive relationships, on average.

# Hypothesis testing

## Hypothesis via CI

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$H_0: \mu = 3$  College students have been in 3 exclusive relationships, on average

$H_a: \mu > 3$  College students have been in more than 3 exclusive relationships, on average.

Note that hypothesis are always about population parameters, never about sample statistics. Also... could you reject  $H_0$  based on your knowledge?

# Hypothesis testing

## p-value

The p-value is the probability of observed or more extreme outcome, given that the null hypothesis is true.

$$p - value = P(\text{observed or more extreme outcome} \mid H_0)$$

In the previous example, the CI interval comes from a sample of 50 students with a mean of 3.2 and a standard deviation of 1.74:

$$n = 50$$

$$\bar{x} = 3.2$$

$$s = 1.74$$

With this data and using CLT, we know that:

$$\bar{x} \sim N(\mu = 3, SE = \frac{1.74}{\sqrt{50}} = 0.246)$$

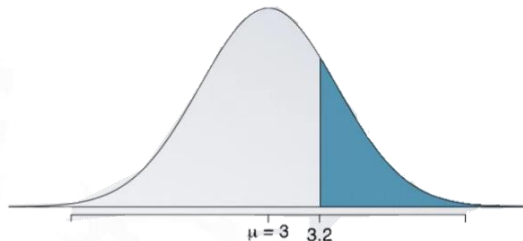
So, what is the probability of observing the sample mean or a more extreme outcome, given the null hypothesis?

$$P(\bar{x} > 3.2 \mid \mu = 3)$$

# Hypothesis testing

## p-value

So let's represent the probability we want to calculate:



The z-score of the point of interest is:

$$z = \frac{3.2 - 3}{0.246} = 0.81$$

This z-score would be the test statistic, because is the statistic used to calculate the p-score:

$$P(z > 0.81) = 0.209$$

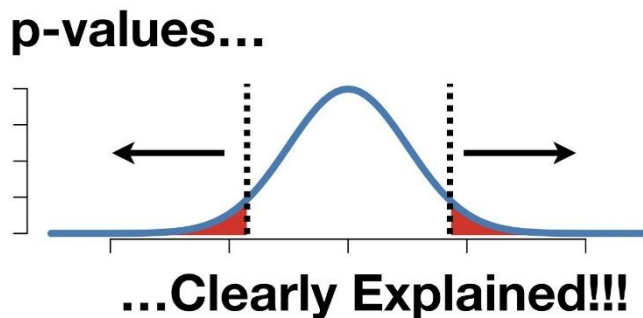
Since p-value is high,  $H_0$  can not be rejected.

# Hypothesis testing

## Interpreting the p-value

For this example, the p-value means that if in fact college students have been in 3 exclusive relationships on average, that's the equivalent of saying if in fact the null hypothesis is true, there is a 21% chance that a random sample of 50 college students would still yield a sample mean of 3.2 or higher.

Since this is a pretty high probability, a sample mean of 3.2 or more exclusive relationships is likely to happen simply by chance or sampling variability.



# Hypothesis testing

## Two-sided tests

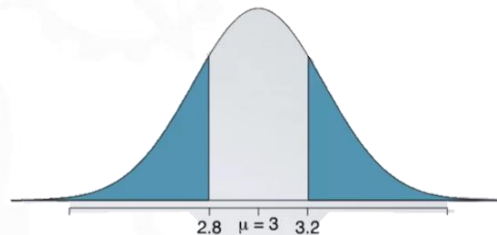
Instead of looking for a divergence from the null hypothesis in a specific direction, it might be interesting the divergence in any direction.

This type of hypothesis tests are called two-sided (or two-tailed).

The p-value “at least as extreme as the observed outcome” probability is extended in both directions.

For the example, the p-value would be:

$$P(\bar{x} > 3.2 \text{ OR } \bar{x} < 2.8 \mid H_0: \mu = 3)$$

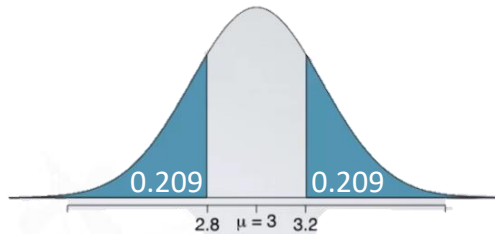


# Hypothesis testing

## Two-sided tests

The p-value would be:

$$P(\bar{x} > 3.2 \text{ OR } \bar{x} < 2.8 \mid H_0: \mu = 3)$$



$$p - value = P(z > 0.81) + P(z < -0.81) = 2 \cdot 0.209 = 0,418$$



# Nearly normal sampling distributions

The previous method, doing hypothesis tests and confidence intervals, can be easily adapted for any point estimator that has a nearly normal sampling distribution:

- Sample mean:  $\bar{x}$
- Difference between sample means:  $\bar{x}_1 - \bar{x}_2$
- Sample proportion:  $\hat{p}$
- Difference between sample proportions:  $\hat{p}_1 - \hat{p}_2$

Point estimators are unbiased, i.e., they do not generally over or underestimate the population parameter true value, only gives a good estimate.

To construct CI of unbiased estimators:

$$\text{point estimate} \pm z^* \cdot SE$$

*SE* calculation might be different for different point estimators. More about this in following sections!

## Other estimators

# Example

A 2010 Pew Research foundation poll indicates that among 1,099 college graduates, 33% watch the Daily Show. An American late-night TV Show. The standard error of this estimate is 0.014. We are asked to estimate the 95% confidence interval for the proportion of college graduates who watch The Daily Show.

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$$\hat{p} = 0.33$$
$$SE = 0.014$$

## Other estimators

### Example

A 2010 Pew Research foundation poll indicates that among 1,099 college graduates, 33% watch the Daily Show. An American late-night TV Show. The standard error of this estimate is 0.014. We are asked to estimate the 95% confidence interval for the proportion of college graduates who watch The Daily Show.

$$\hat{p} = 0.33$$

$$SE = 0.014$$

$$CI_{95\%}: 0.33 \pm 1.96 \cdot 0.014 = (0.303, 0.357)$$

## Other estimators

# Second example

The third national health and nutrition examination survey NHANES, collected body fat percentage and gender data from over 13,000 subjects in ages between 20 to 80. The average body fat percentage for the 6,580 men in the sample was 23.9%. And this value was 35% for the, for the 7,021 women. The standard error for the difference between the average male and female body fat percentages was 0.114. Do these data provide convincing evidence that men and women have different average body fat percentages? You may assume that the distribution of the point estimate is nearly normal.

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$H_0: \mu_{men} = \mu_{women}; H_A: \mu_{men} \neq \mu_{women} \rightarrow$  Two-tailed test

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Point estimate:  $\bar{x}_{men} - \bar{x}_{women} = 23.9 - 35 = -11.1$

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$H_0: \mu_{men} = \mu_{women}; H_A: \mu_{men} \neq \mu_{women} \rightarrow$  Two-tailed test

Point estimate:  $\bar{x}_{men} - \bar{x}_{women} = 23.9 - 35 = -11.1$

This point estimate is distributed:

$$\bar{x}_{men} - \bar{x}_{women} \sim N(\mu_{men} - \mu_{women}, SE)$$



## Other estimators

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$H_0: \mu_{men} = \mu_{women}; H_A: \mu_{men} \neq \mu_{women} \rightarrow$  Two-tailed test

Point estimate:  $\bar{x}_{men} - \bar{x}_{women} = 23.9 - 35 = -11.1$

This point estimate is distributed:

$$\bar{x}_{men} - \bar{x}_{women} \sim N(\mu_{men} - \mu_{women}, SE) = N(0, 0.114)$$

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$H_0: \mu_{men} = \mu_{women}; H_A: \mu_{men} \neq \mu_{women} \rightarrow$  Two-tailed test

Point estimate:  $\bar{x}_{men} - \bar{x}_{women} = 23.9 - 35 = -11.1$

This point estimate is distributed:

$$\bar{x}_{men} - \bar{x}_{women} \sim N(\mu_{men} - \mu_{women}, SE) = N(0, 0.114)$$

So, the p-value is basically 0 and  $H_0$  can be rejected.

# Decision errors

## Types of errors

		Decision	
		Fail to reject $H_0$	Reject $H_0$
Truth	$H_0$ true	✓	<b>Type I error</b>
	$H_a$ true	<b>Type II error</b>	✓

- Type I error is rejecting  $H_0$  when  $H_0$  is true
- Type II error is failing to reject  $H_0$  when  $H_A$  is true

## Decision errors

# Which is worse?

In general, reducing both types of errors is not feasible, so we must try to determine which type of error we want to minimize. So, which is worse?

- Type II: Declaring the defendant innocent when they are actually guilty.
- Type I: Declaring the defendant guilty when they are actually innocent.

The answer is... depends on the problem!



## Decision errors

# Significance rate

Usually, the significance rate is chosen at  $\alpha = 0.05$ , meaning that we reject  $H_0$  when the p-value is less than 0.05, i.e., given that  $H_0$  is true there is a probability of less than 5% of seeing an observed value or a more extreme value.

This significance level means that, for those cases where  $H_0$  is actually true, no more than 5% of the times  $H_0$  is incorrectly rejected.

In other words:

$$P(\text{Type I error} \mid H_0 \text{ true}) = \alpha$$

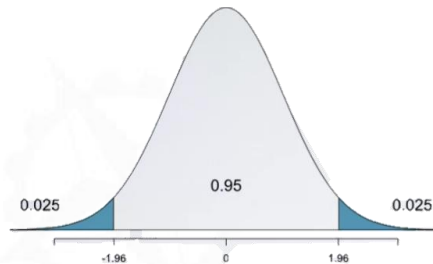
If type I errors are dangerous or especially costly, choose a smaller significance level. In the other hands, if type II errors are dangerous, choose a higher significance level.

# Significance vs confidence

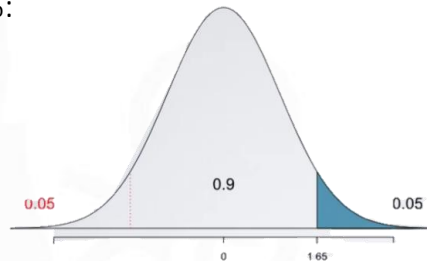
In general, significance and confidence interval are complementary:

$$1 - \alpha = \beta$$

This is related to the area under the PDF for a two-tailed test with a 95% confidence interval.



However, this is not the case for a one-sided test. For the same covered area, the confidence level is 90%:



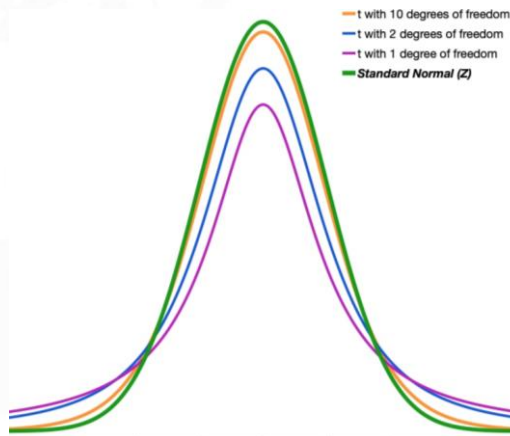
# t-distribution

## Introduction

When **population standard deviation  $\sigma$  is unknown**, t-distribution can be used to address the **uncertainty of the standard error estimate**. t-distribution is usually called student's t.

**Observations are more likely to fall beyond 2 SDs from the mean** than in an standard normal distribution, helping mitigating the effect of a less reliable estimate for the SE.

While normal distribution has two parameters ( $\mu$  and  $\sigma$ ), student's t has only one: **degrees of freedom (df)**.



# Inference for a mean

In a study, researchers evaluated the **relationship between being distracted and recall of food consumed and snacking**, with the idea that if you are distracted while you are eating, you may **not remember what you eat**. They also hypothesized that failure to recall food consumed might lead to **increased snacking later on**. The sample for this study consisted of **44 volunteer patients, half men, half women**. These 44 patients were **randomized into two groups**. **One group** was asked to play solitaire on the computer while eating and **was asked to win as many games as possible**, and the **other group** was asked to **eat lunch without any distractions**. Both groups were provided the **same amount of lunch** and then **after lunch they were offered biscuits** to snack on.

Biscuit intake	$\bar{x}$	$s$	$n$
Solitaire	52.1 g	45.1 g	22
No distraction	27.1 g	26.4 g	22



# Inference for a mean

The goal in this example is to estimate the **average snacking level for distracted eaters**:

Biscuit intake	$\bar{x}$	$s$	$n$
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Confidence interval using the t-distribution:

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Confidence interval using the t-distribution:

*point estimate  $\pm$  margin of error*

$$\bar{x} \pm t_{df}^* \cdot SE_{\bar{x}}$$

$$\bar{x} \pm t_{df}^* \cdot \frac{s}{\sqrt{n}}$$

The degrees of freedom for inference on one sample mean is the number of samples minus 1:  $df = n - 1$  ( $t_{df}^*$ : t-score)

## Inference for a mean

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Biscuit intake	$\bar{x}$	$s$	$n$
Solitaire	52.1 g	45.1 g	22
No distraction	27.1 g	26.4 g	22

In the example:

$$\bar{x} = 52.1 \text{ g}$$

$$s = 45.1 \text{ g}$$

$$n = 22$$

$$t_{21}^* = 2.08$$

$$CI: \bar{x} \pm t_{21}^* \cdot \frac{s}{\sqrt{n}} = 52.1 \pm 2.08 \cdot \frac{45.1}{\sqrt{22}} \approx 52.1 \pm 20$$

$$CI: (32.1 \text{ g}, 72.1 \text{ g})$$

## Hypothesis testing for a mean

Suppose the suggested serving size of these biscuits is 30 g. Do these data provide convincing evidence that the amount of snacks consumed by distracted eaters post-lunch is different than the suggested serving size?

$$\bar{x} = 52.1 \text{ g}$$

$$s = 45.1 \text{ g}$$

$$n = 22$$

$$df = 21$$

$$SE = \frac{45.1}{\sqrt{22}} = 9.62$$

# Hypothesis testing for a mean

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Hypothesis:

$$H_0: \mu = 30$$

$$H_A: \mu \neq 30$$



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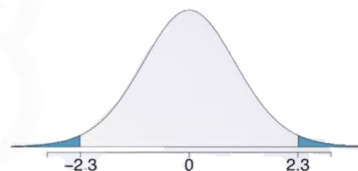
Hypothesis:

$$H_0: \mu = 30$$

$$H_A: \mu \neq 30$$

t-statistic (not the same as t-score):

$$T = \frac{\bar{x} - \mu}{SE} = \frac{52.1 - 30}{9.62} = 2.3$$



# Hypothesis testing for a mean

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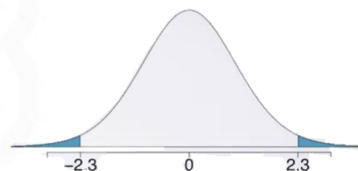
$$H_A: \mu \neq 30$$

t-statistic (not the same as t-score):

$$T = \frac{\bar{x} - \mu}{SE} = \frac{52.1 - 30}{9.62} = 2.3$$

And calculate:

$$p\text{-value}: P(T > 2.3 \text{ OR } T < -2.3 \mid t_{21}) \approx 0.0318$$



t-distribution

# Inference for difference of two independent means

Estimate the **difference of the average snacking level for distracted and non-distracted eaters**:

Biscuit intake	$\bar{x}$	$s$	$n$
Solitaire	52.1 g	45.1 g	22
No distraction	27.1 g	26.4 g	22

Confidence interval using the t-distribution for two independent means:

$$\text{point estimate} \pm \text{margin of error}$$

t-distribution

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Confidence interval using the t-distribution for two independent means:

*point estimate  $\pm$  margin of error*

$$(\bar{x}_1 - \bar{x}_2) \pm t_{df}^* \cdot SE_{\bar{x}_1 - \bar{x}_2}$$

t-distribution

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*point estimate  $\pm$  margin of error*

$$(\bar{x}_1 - \bar{x}_2) \pm t_{df}^* \cdot SE_{\bar{x}_1 - \bar{x}_2}$$

Where:

$$SE_{\bar{x}_1 - \bar{x}_2} = \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$$

$$df = \min(n_1 - 1, n_2 - 1)$$

# Inference for difference of two independent means

Estimate the **difference of the average snacking level for distracted and non-distracted eaters**:

Biscuit intake	$\bar{x}$	$s$	$n$
Solitaire	52.1 g	45.1 g	22
No distraction	27.1 g	26.4 g	22

$$\begin{aligned}
 & (\bar{x}_1 - \bar{x}_2) \pm t_{21}^* \cdot SE_{\bar{x}_1 - \bar{x}_2} \\
 &= (52.1 - 27.1) \pm 2.08 \cdot \sqrt{\frac{45.1^2}{22} + \frac{26.4^2}{22}} \\
 &= 25 \pm 2.08 \cdot 11.14 = (1.83 \text{ g}, 48.17 \text{ g})
 \end{aligned}$$

t-distribution

# Hypothesis testing for two independent means

Is there a difference between the average post-meal snack consumption between groups?

$$\bar{x}_1 = 52.1 \text{ g}$$

$$s_1 = 45.1 \text{ g}$$

$$n_1 = 22$$

$$\bar{x}_2 = 27.1 \text{ g}$$

$$s_2 = 26.4 \text{ g}$$

$$n_2 = 22$$

$$df = 21$$

$$SE = 11.14$$

t-distribution

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$$s_2 = 26.4 \text{ g}$$

$$n_2 = 22$$

$$df = 21$$

$$SE = 11.14$$

Hypothesis:

$$H_0: \mu_1 - \mu_2 = 0$$

$$H_A: \mu_1 - \mu_2 \neq 0$$



# Hypothesis testing for two independent means

Is there a difference between the average post-meal snack consumption between groups?

$$\bar{x}_1 = 52.1 \text{ g}$$

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$$s_2 = 26.4 \text{ g}$$

$$n_2 = 22$$

$$df = 21$$

$$SE = 11.14$$

Hypothesis:

$$H_0: \mu_1 - \mu_2 = 0$$

$$H_A: \mu_1 - \mu_2 \neq 0$$

t-statistic:

$$T = \frac{(\bar{x}_1 - \bar{x}_2) - (\mu_1 - \mu_2)}{SE} = \frac{25 - 0}{11.14} = 2.24$$

# Hypothesis testing for two independent means

Is there a difference between the average post-meal snack consumption between groups?

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$$s_1 = 45.1 \text{ g}$$

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$$n_2 = 22$$

$$df = 21$$

$$SE = 11.14$$

Hypothesis:

$$H_0: \mu_1 - \mu_2 = 0$$

$$H_A: \mu_1 - \mu_2 \neq 0$$

t-statistic:

$$T = \frac{(\bar{x}_1 - \bar{x}_2) - (\mu_1 - \mu_2)}{SE} = \frac{25 - 0}{11.14} = 2.24$$

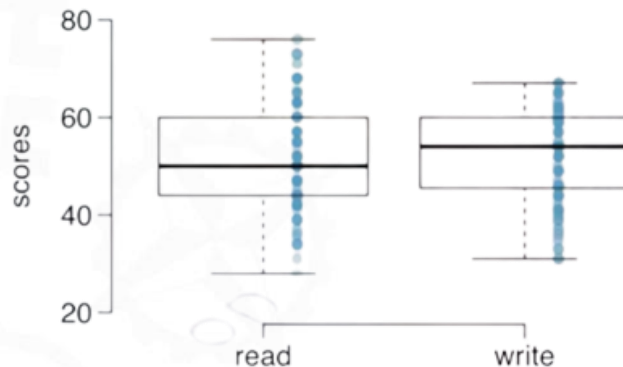
And calculate:

$$p\text{-value: } P(T > 2.24 \text{ OR } T < -2.24 \mid t_{21}) \approx 0.036$$

t-distribution

## Hypothesis testing for two paired means

200 observations were randomly sampled from the High School and Beyond survey. The same students took a reading and a writing test.



Can the reading and writing scores for a giving student be independent from each other?

# Hypothesis testing for two paired means

- When two sets of observations have this special correspondence, or in other words they are not independent, they are said **to be paired**.
- To analyze paired data, it is often **useful to look at the difference in outcomes** of each pair of observations.
  - So, in this example, for each student, we subtract their writing score from their reading score

ID	Read	Write	Diff
0	57	52	5
1	44	33	11
...	...	...	...

t-distribution

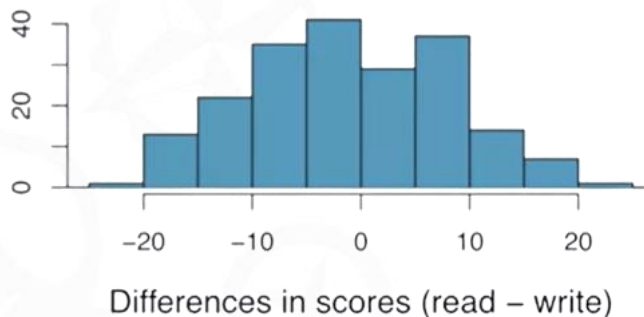
## Hypothesis testing for two paired means

If in fact there was no difference between the reading and writing scores, what would we expect the differences to be?

$$\bar{x}_{diff} = -0.545$$

$$s_{diff} = 8.887$$

$$n_{diff} = 200$$



## Hypothesis testing for two paired means

If in fact there was no difference between the reading and writing scores, what would we expect the differences to be?

$$\bar{x}_{diff} = -0.545$$

$$s_{diff} = 8.887$$

$$n_{diff} = 200$$

$$df = 199$$

Hypothesis:

$$H_0: \mu_{diff} = 0$$

$$H_A: \mu_{diff} \neq 0$$

t-statistic:

$$T = \frac{\bar{x} - \mu}{SE} = \frac{-0.545 - 0}{8.887/\sqrt{200}} = -0.87$$

And calculate:

$$p\text{-value: } P(T > 0.87 \text{ OR } T < -0.87 \mid t_{199}) \approx 0.385$$

## Interpretation of p-value

Which of the following is the correct interpretation of the p-value?

1. p-value is the probability that the average scores on the reading and writing exams are equal.
2. p-value is the probability that the average scores on the reading and writing exams are different.
3. p-value is the probability of obtaining a random sample of 200 students where the average difference between the reading and writing scores is at least 0.87 in either direction, if in fact the true average difference between the scores is zero.
4. p-value is the probability of incorrectly rejecting the null hypothesis if in fact the null hypothesis is true.

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4. p-value is the probability of incorrectly rejecting the null hypothesis if in fact the null hypothesis is true.  $\rightarrow P(\text{Type I error})$

## ANOVA

### Compare n means

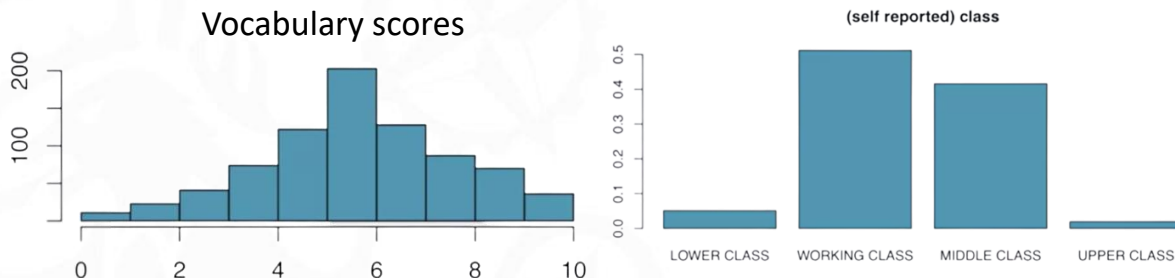
The example for this section comes from the general social survey. The variable of interest are vocabulary scores and self-identified social class. Vocabulary score is calculated based on a set of 10 question vocabulary test, where a higher score means better vocabulary, and self identified social class has four levels, lower, working, middle, and upper class.

<b>Voc. scores</b>	<b>Class</b>
6	Middle class
9	Working class
...	...
7	Upper class

# ANOVA

## Compare n means

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## ANOVA

### Compare n means

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	<b>n</b>	<b>mean</b>	<b>sd</b>
Lower class	41	5.07	2.24
Working class	407	5.75	1.87
Middle class	331	6.76	1.89
Upper class	16	6.19	2.34
Overall	795	6.14	1.98

# ANOVA

## Hypothesis

In an ANOVA test, hypothesis are:

*$H_0$ : mean outcome of a variable is the same across all  $k$  groups (categories)*

$$\mu_1 = \mu_2 = \dots = \mu_k$$

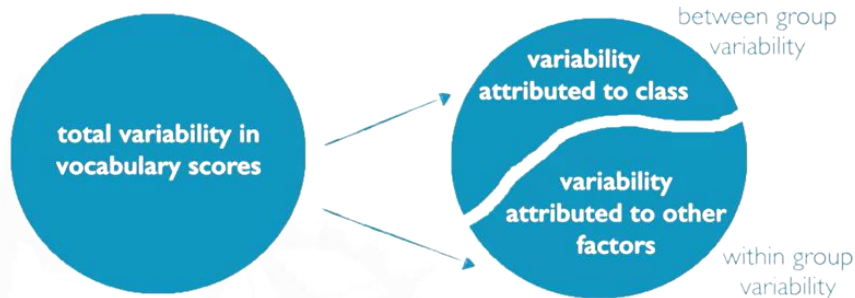
*$H_A$ : at least one pair of means are different*

t-test	ANOVA
Compare means from two groups to see whether they're so far apart that the observed difference cannot reasonably be attributed to sampling variability.	Compare means from more than two groups to see whether they're so far apart that the observed differences cannot all reasonably be attributed to sampling variability.

# ANOVA

## Variability partitioning

For the example:



		Df	Sum Sq	Mean Sq	F value	P-value
<b>Group</b>	<b>Class</b>	3	236.56	78.855	21.735	<0.0001
<b>Error</b>	<b>Residuals</b>	791	2869.80	3.628		
	<b>Total</b>	794	3106.36			



# Variability partitioning

For the example:

			Sum Sq			
<b>Group</b>	<b>Class</b>		236.56			
<b>Error</b>	<b>Residuals</b>		2869.80			
	<b>Total</b>		3106.36			

## Sum of Squares Total or SST

- Measures the total variability of the output response
- Calculated similar to variance (not scaled by sample size)

$$SST = \sum_{i=1}^n (y_i - \bar{y})^2$$

# Variability partitioning

For the example:

			Sum Sq			
<b>Group</b>	<b>Class</b>		236.56			
<b>Error</b>	<b>Residuals</b>		2869.80			
	<b>Total</b>		3106.36			

## Sum of Squares Group or SSG

- Measures the variability between groups
- Explained variability: squared deviation of group means from overall means, weighted by sample size

$$SSG = \sum_{j=1}^n n_j (\bar{y}_j - \bar{y})^2$$

# ANOVA

## Variability partitioning

For the example:

			Sum Sq			
<b>Group</b>	<b>Class</b>		236.56			
<b>Error</b>	<b>Residuals</b>		2869.80			
	<b>Total</b>		3106.36			

### Sum of Squares Errors or SSE

- Measures the variability within groups
- Unexplained variability by the group variable

$$SSE = SST - SSG$$

# ANOVA

## Variability partitioning

For the example:

		df	Sum Sq			
<b>Group</b>	<b>Class</b>	3	236.56			
<b>Error</b>	<b>Residuals</b>	791	2869.80			
	<b>Total</b>	794	3106.36			

Degrees of freedom associated with ANOVA

- Total:  $df_T = n - 1$
- Group:  $df_G = k - 1$
- Error:  $df_E = df_T - df_G$

# ANOVA

## Variability partitioning

For the example:

		df	Sum Sq	Mean Sq		
Group	Class	3	236.56	78.855		
Error	Residuals	791	2869.80	3.628		
	Total	794	3106.36			

### Mean Squares

- Average variability between and within groups.
  - Group:  $MSG = SSG/df_G$
  - Error:  $MSE = SSE/df_E$

## ANOVA

# Variability partitioning

For the example:

		df	Sum Sq	Mean Sq	F value	
<b>Group</b>	<b>Class</b>	3	236.56	78.855	21.735	
<b>Error</b>	<b>Residuals</b>	791	2869.80	3.628		
	<b>Total</b>	794	3106.36			

F statistic

- Ratio of the average between group and within group variabilities:

$$F = \frac{MSG}{MSE}$$

# ANOVA

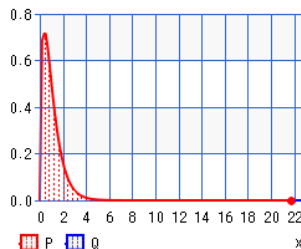
## Variability partitioning

For the example:

		df	Sum Sq	Mean Sq	F value	P-value
<b>Group</b>	<b>Class</b>	3	236.56	78.855	21.735	<0.0001
<b>Error</b>	<b>Residuals</b>	791	2869.80	3.628		
	<b>Total</b>	794	3106.36			

### P-value

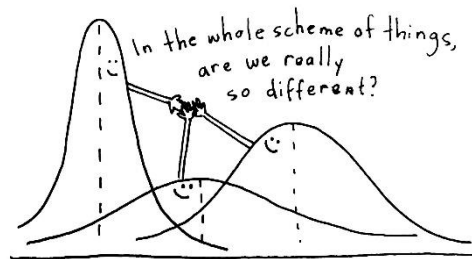
- Probability of at least as large a ratio between the “between” and “within” group variabilities if in fact the means of all groups are equal.
- Area under the F curve with degrees of freedom  $df_G$  and  $df_E$



# ANOVA

## Conditions

1. Independence
  - Within groups: sampled observations must be independent
  - Between groups: groups must be independent from each other
2. Approximate normality: distributions should be nearly normal within each group
3. Equal variance: groups should have roughly equal variability: **homoscedastic groups**.





## Multiple comparisons

# Which means differ

ANOVA is great to check if any mean differs from the rest, but does not tell which pair of means differ.

Solution, use multiple t-tests:

- Two sample t tests for differences in each possible pair of groups.
- Multiple tests → inflated Type I error rate
- Need to use a modified significance level → Bonferroni correction.
  - Adjust  $\alpha$  by the number of comparisons being considered

$$\alpha^* = \frac{\alpha}{K}; K: \text{number of comparisons} = \frac{k \cdot (k - 1)}{2}$$

## Multiple comparisons

# Which means differ

Apart from using a corrected significance level, other conditions must be considered:

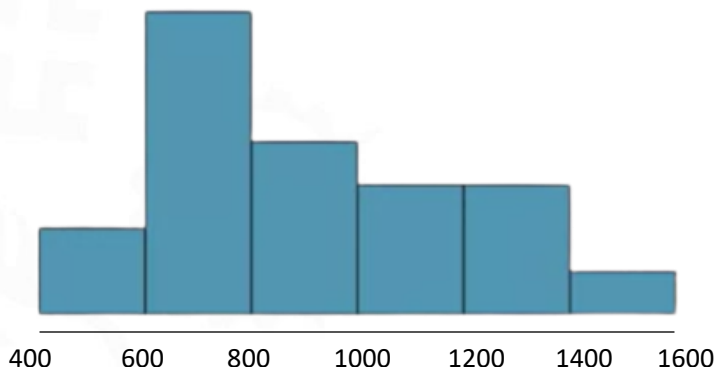
- Constant variance → corrected standard error and degrees of freedom for all tests

$$SE = \sqrt{\frac{MSE}{n_1} + \frac{MSE}{n_2}}$$

$$df = df_E$$

# Bootstrapping Example

We have a dataset of 20 apartments randomly selected from Craigslist housing ads. These are apartments with at least one bedroom in Durham, North Carolina.



Which is the best central measure?

# Bootstrapping Example

The sample median in the original sample is \$887.



In bootstrapping, we assume that, for each observation in the sample, there may be others like it in the population. The bootstrap population can be thought as a population where each observation from the sample appears many times.

# Bootstrapping

## Bootstrapping scheme

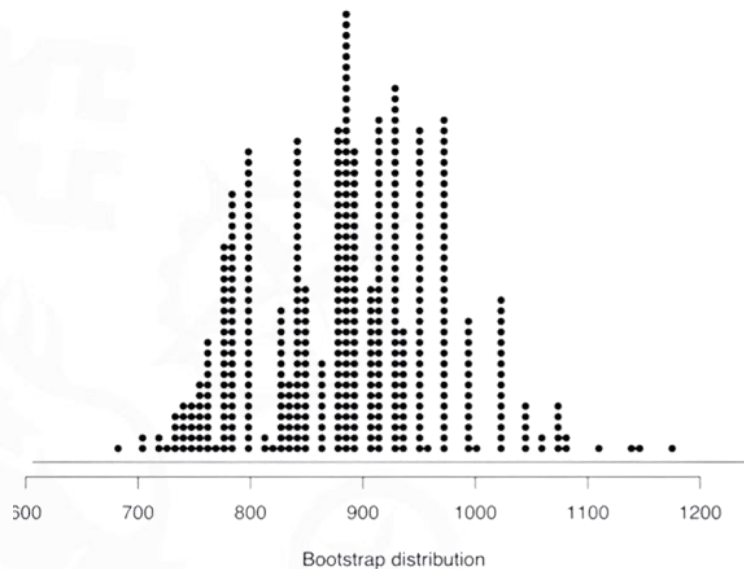
So, how is bootstrap performed:

1. Take a bootstrap sample: random sample with replacement from the original sample, of the same size as the original sample.
2. Calculate the bootstrap statistic: mean, median, proportion, ... from the bootstrap samples.
3. Repeat steps 1 and 2 many times to create a bootstrap statistics distribution.

# Bootstrapping

## Bootstrap distribution

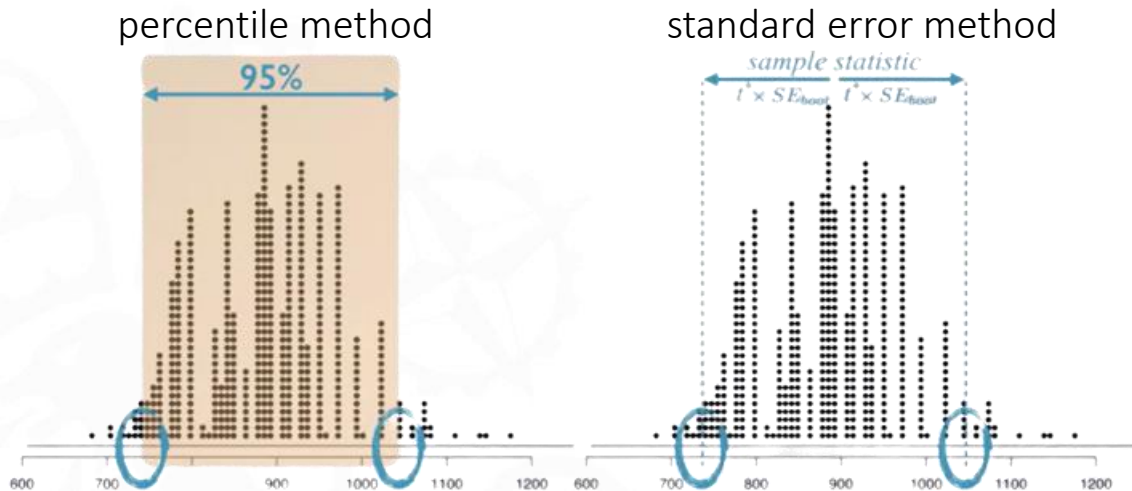
For the example, the median bootstrap distribution for 500 simulations:



# Bootstrapping

## Confidence interval

Confidence interval can be calculated using two methods:



The DF in the standard error methods would be  $n-1$  being  $n$  the original sample size.

# Bootstrapping Limitations

Bootstrap does not present as rigid conditions as CLT based methods. However:

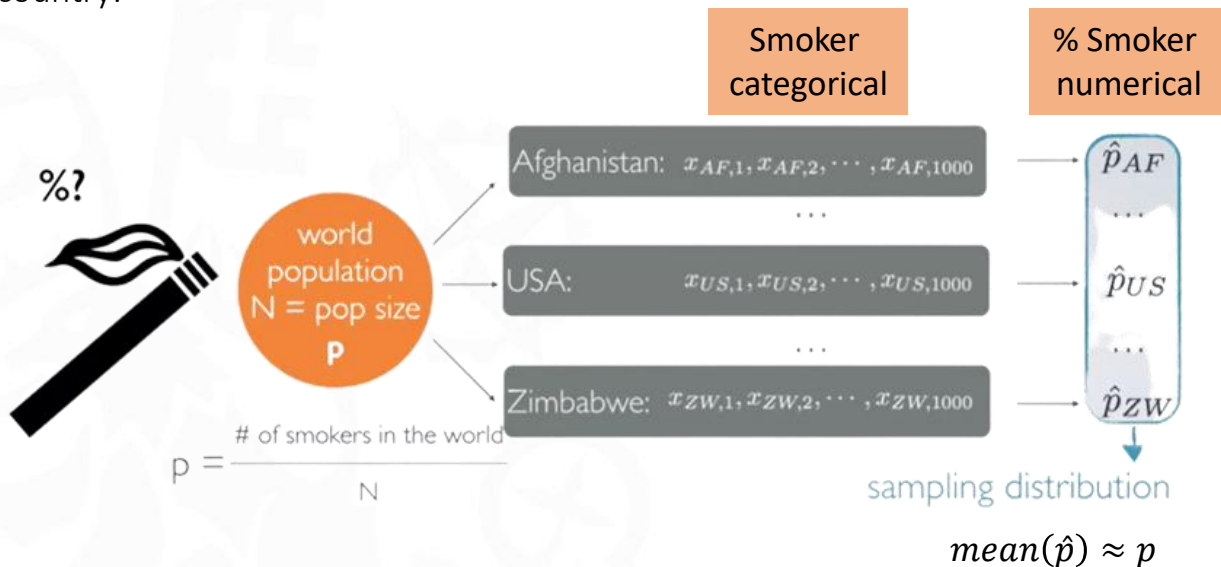
- If the bootstrap distribution is extremely skewed or sparse, the bootstrap interval might be unreliable.
- A representative sample is still required. If the sample is biased, the estimates would also be biased.



# Categorical variables

## Sampling distribution

In order to estimate the world smoker population proportion, we can ask individual people in each country and calculate the proportion of smokers for each country:



## Categorical variables

# CLT for proportions

The distribution of sample proportions is nearly normal, centered at the population proportion, and with a standard error inversely proportional to the sample size:

$$\hat{p} \sim N \left( \text{mean} = p, SE = \sqrt{\frac{p \cdot (1 - p)}{n}} \right)$$

Conditions for CLT:

1. **Independence:** Sampled observations must be independent
2. **Sample size/skew:** There should be at least 10 successes and 10 failures in the sample:  $n \cdot p \geq 10$  and  $n \cdot (1 - p) \geq 10$ 
  1. If  $p$  is unknown, use  $\hat{p}$

# Confidence interval for a proportion

Two scientists want to know if a certain drug is effective against high blood pressure. The first scientist wants to give the drug to a thousand people with high blood pressure, and see how many of them experience lower blood pressure levels. The second scientist wants to give the drug to 500 people with high blood pressure and not give the drug to another 500 people with high blood pressure, and see how many in both groups experience lower blood pressure levels. Which is the better way to test this drug?

1. All 1000 get the drug
2. 500 get the drug, 500 don't

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This same question was asked to 670 americans:

Option 1 (bad int.)	99
Option 2 (good int.)	571
Total	670

## Categorical variables

# Confidence interval for a proportion

Estimate the proportion of all americans who have good intuition about experimental design:

parameter of interest

Percentage of **all** Americans who have good intuition about experimental design.

$p$

point estimate

Percentage of **sampled** Americans who have good intuition about experimental design.

$571 / 670 \approx 0.85 \quad \hat{p}$

As always:

*point estimate  $\pm$  margin of error*

$$\hat{p} \pm z^* \cdot SE_{\hat{p}}$$

Where:

$$SE_{\hat{p}} = \sqrt{\frac{\hat{p} \cdot (1 - \hat{p})}{n}}$$

## Confidence interval for a proportion

$$\begin{aligned} CI_{95\%}: \hat{p} \pm z^* \cdot SE_{\hat{p}} \\ = 0.85 \pm 1.96 \cdot \sqrt{\frac{0.85 \cdot 0.15}{670}} \\ = 0.85 \pm 1.96 \cdot 0.0138 \\ = (0.823, 0.877) \end{aligned}$$

We are 95% confident that 82.3% to 87.7% of all americans have good intuition about experimental design.

# Required sample size for a given ME

$$ME = z^* \cdot \sqrt{\frac{\hat{p} \cdot (1 - \hat{p})}{n}}$$

Using this formula, we can obtain  $n$  for a given ME. If we do not have a sample yet:

- If there is a previous study, rely on the value calculated in that study for  $\hat{p}$ .
- If not, use  $\hat{p} = 0.5$ 
  - If you do not know any better, 50-50 is a good guess
  - Gives the most conservative estimate for  $n$

Remember to always **round up**  $n$ .

## Categorical variables

# Hypothesis testing for a proportion

1. Set the hypothesis
  - $H_0: p = \text{null value}$
  - $H_A: p < \text{or } > \text{or } \neq \text{null value}$
2. Calculate the point estimate
3. Check conditions
  1. Independence
  2. Sample size/skew
4. Calculate test statistic

$$Z = \frac{\hat{p} - p}{SE}, SE = \sqrt{\frac{p \cdot (1 - p)}{n}}$$

5. Make a decision based on p-value and  $\alpha$



## Categorical variables

# Confidence interval for a proportion

A 2013 Pew Research poll found that 60% of 1,983 randomly sampled American adults believe in evolution. Does this provide convincing evidence that majority of Americans believe in evolution?

$$H_0: p = 0.5$$

$$H_A: p > 0.5$$

$$\hat{p} = 0.6$$

$$n = 1983$$

$$\hat{p} \sim N\left(0.5, \sqrt{\frac{0.5 \cdot (1 - 0.5)}{1983}} \approx 0.0112\right)$$

$$Z = \frac{0.6 - 0.5}{0.0112} \approx 8.92 \gg 3 \rightarrow p\text{-value} \approx 0$$

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