

STAT/SOC/CSSS 221
Statistical Concepts and Methods
for the Social Sciences

Making Inferences from Samples

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Motivation

How do we know what the average American thinks about an issue?

Usual approach: conduct an opinion poll, randomly sample 1000 or so people, and present the average of their opinions

But how do we know this matches the average opinion of *all* Americans?

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Without knowing anything but the sample, can we estimate the deviation between the sample mean and the population mean?

Populations & Samples

We will consider groups of observations at two distinct levels:

Population All the potential units of analysis in our chosen research design

Ideally we'd like to analyze a census, or complete set, of these observations

Example: Average support $\bar{x}^{\text{population}}$ of all Washingtonians for same-sex marriage

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Example: Average support $\bar{x}^{\text{population}}$ of all Washingtonians for same-sex marriage

Sample The units of analysis actually collected for our study
Usually a subset of the population

Example: Average support \bar{x} of 500 randomly selected Washingtonians for same-sex marriage

Sampling Frames

In an ideal situation, our sample and population will contain the same cases (a census)

Usually, we must instead make inferences about the population using a subset, or sample, of cases

Can select this sample in different ways

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Random sample Make a list of the full population and randomly select by identification number.

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If done correctly, produces something close to a random sample

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Convenience samples do *not* allow scientific inference to the population parameters

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Are convenience samples more likely to be representative as they get larger?

When sampling goes wrong

If a random sample is non-representative, will adding more random samples help make it so? Yes

If a stratified sample has the wrong weights, will adding more samples make it representative? No

Are convenience samples more likely to be representative as they get larger? NO! No matter how large a convenience sample, they are likely to be sampled with huge and unknown selection bias

Sampling Inference

Our goal is to make scientifically valid inferences from the random or representative sample we've collected

Standard scientific practice requires that we quantify the uncertainty introduced by sampling

To learn how to do this, we will eventually learn new concepts:

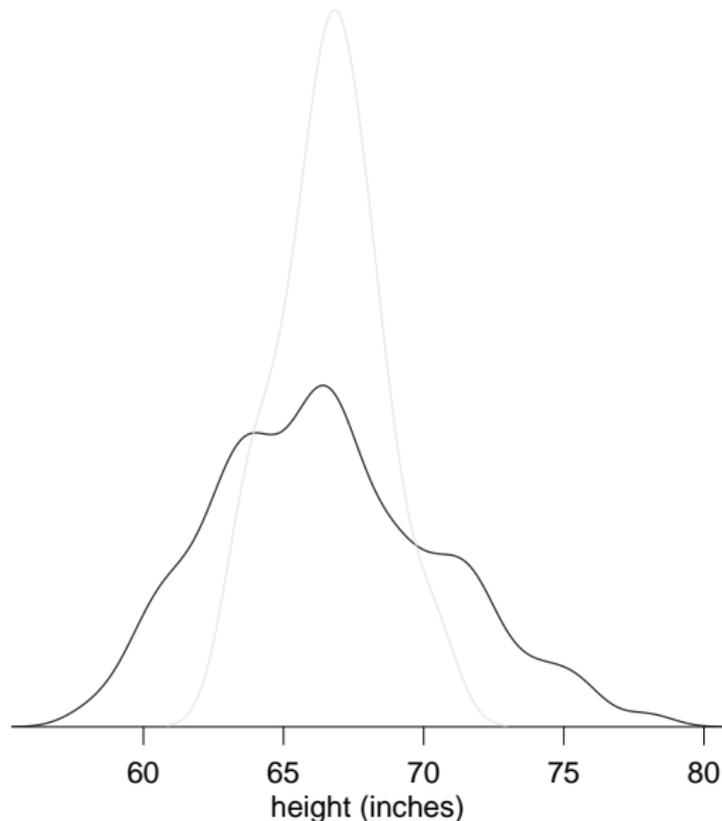
the **standard error** and **t -statistic**,

and new continuous probability distributions:

the **χ^2 distribution** and **t distribution**

Today, we focus on the *standard error*

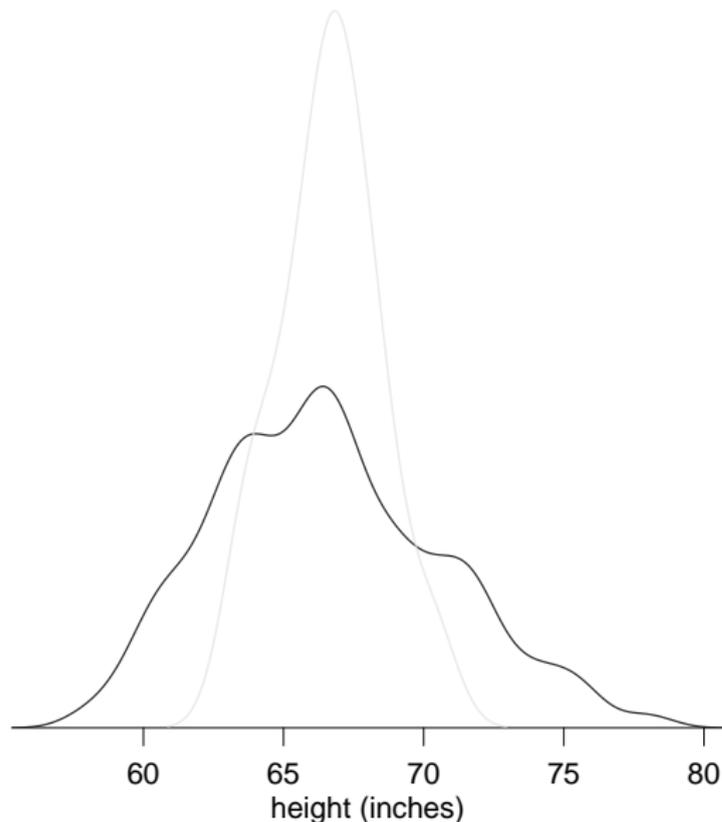
Student height example



119 students submitted their heights in inches, with a mean of 66.6 inches, and a standard deviation of 4.1 inches.

The class distribution of heights is shown

Student height example

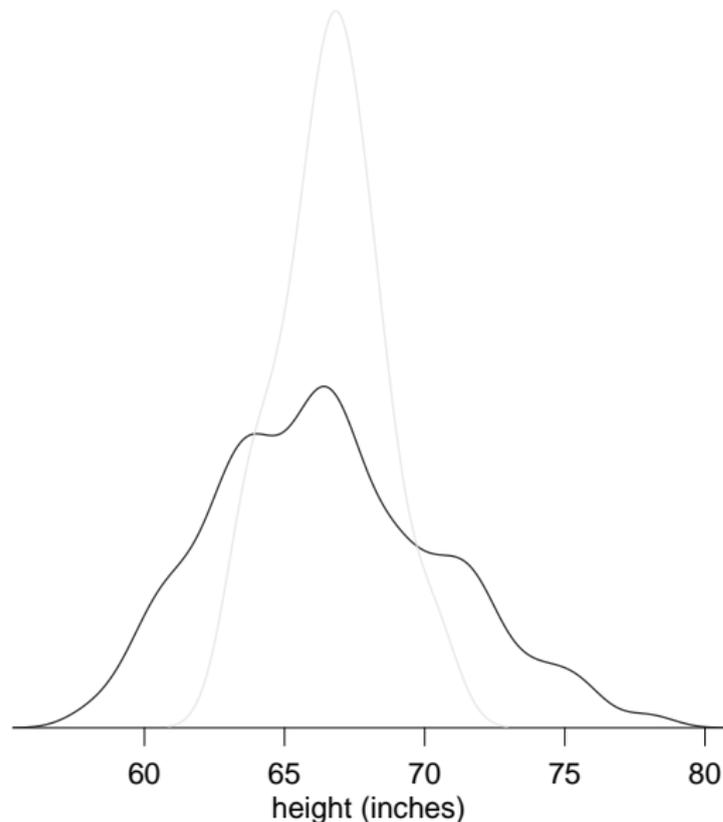


About 2/3s of the class submitted heights.

Getting that kind of response from a population is expensive and impractical in most cases

Suppose we just wanted to know the average height in the class?

Student height example



How well could we estimate the class mean from a sample of a specific size?

And how can we tell if our sample estimate of the mean is reliable?

We need a way to predict how much our sample mean will diverge from the population mean

Useful concepts for estimating uncertainty (statistical inference)

Error The difference between an estimate (based on a sample) and the true value of a quantity (in the population)

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Standard error An *estimate* of the error in our sample’s estimate of the population quantity.

Thus the standard error is the best guess from a single sample of what the RMSE would turn out to be if we could afford to take many samples

Concepts for statistical inference applied to the *sample mean*

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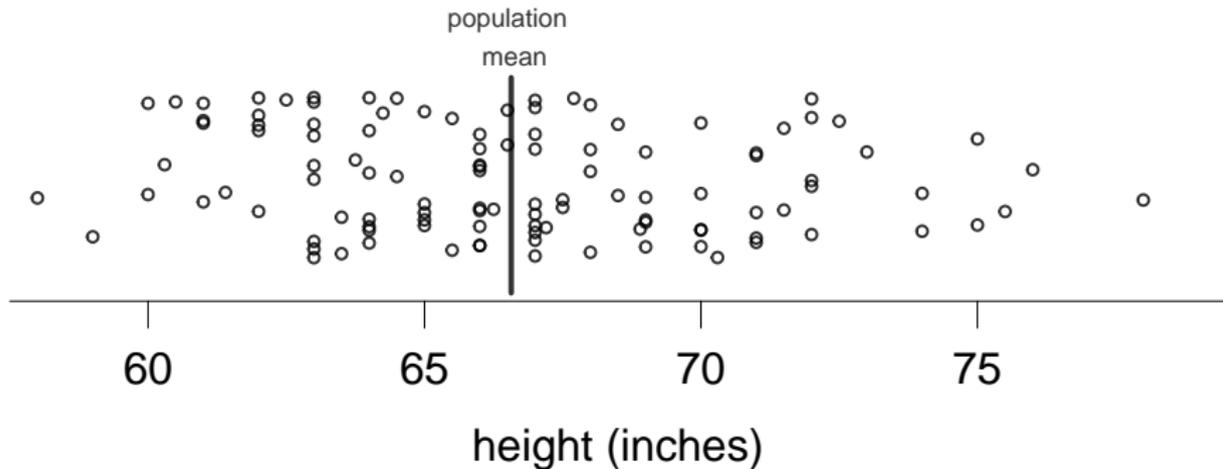
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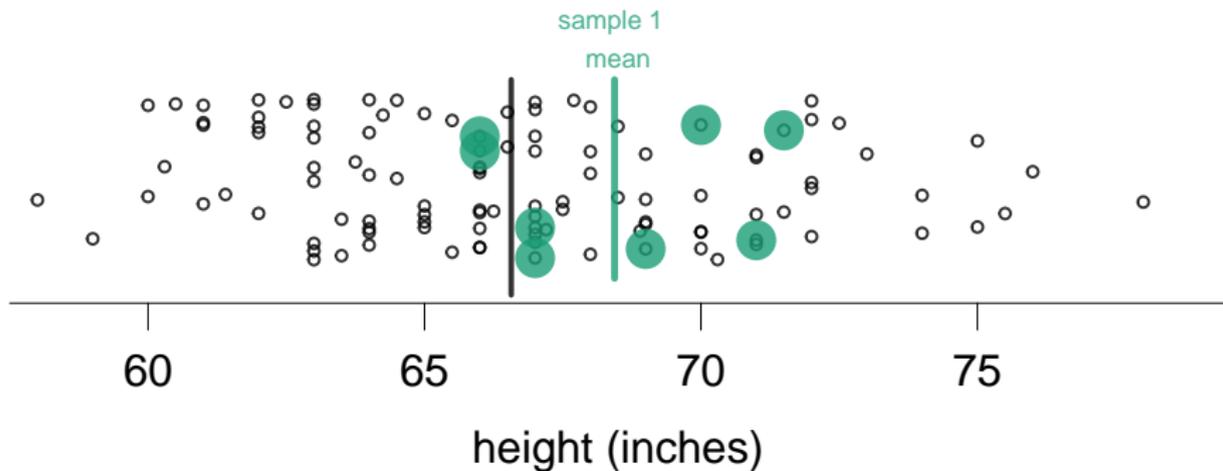
Your book also calls this the *standard deviation of the sampling mean* (Chapter 11)



Above are the submitted height data and their mean

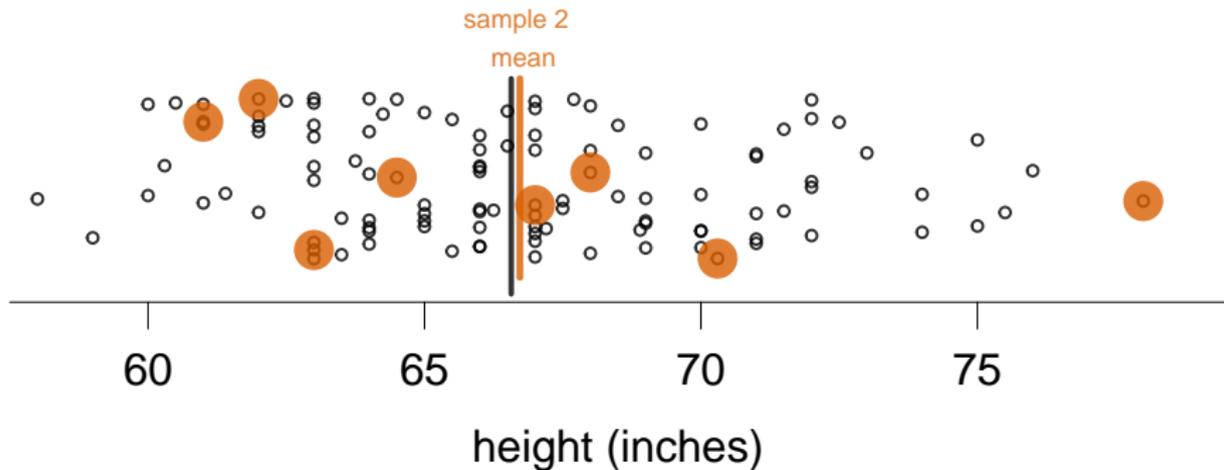
We will treat the rows of this classroom as samples

While you collect your data on your row...



I will explore a set of 14 pre-selected samples of 8 heights each

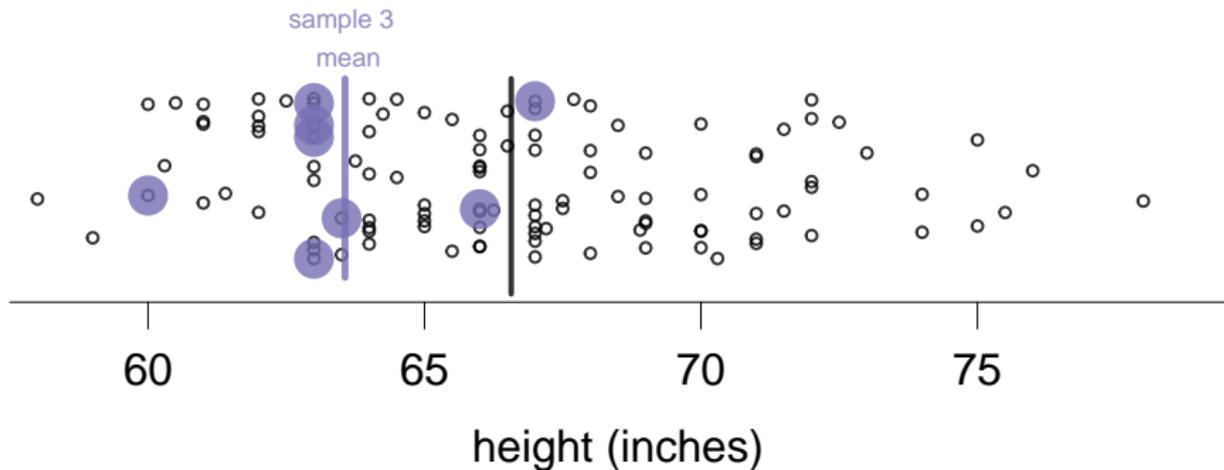
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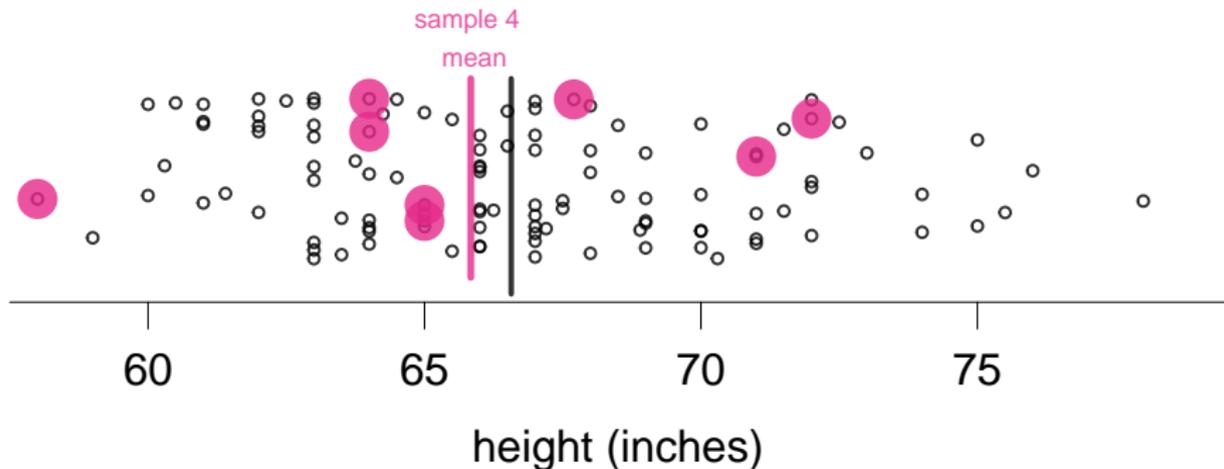
Note that the means and variances of the samples can differ from the full population...

...but can also resemble it fairly closely



Sometimes an individual sample can be so far off that it would mislead us considerably

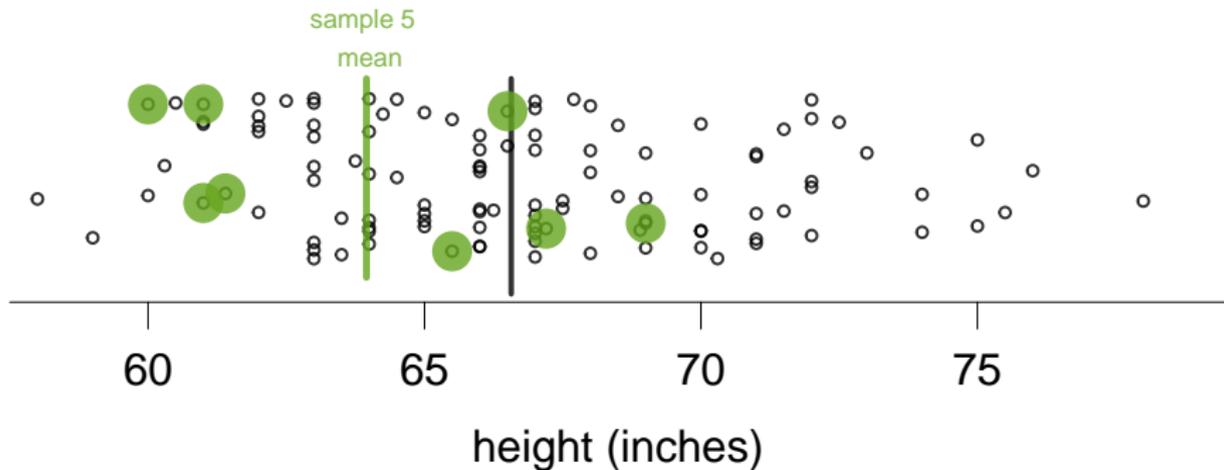
This is more likely the smaller the sample



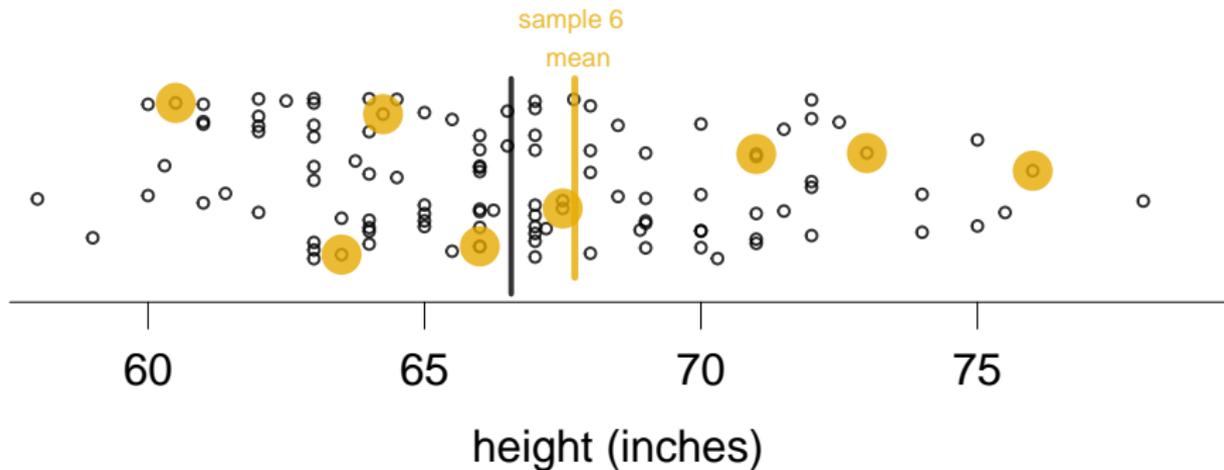
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But even with samples as tiny as 8 students, most of the time the sample mean is fairly close to the population mean

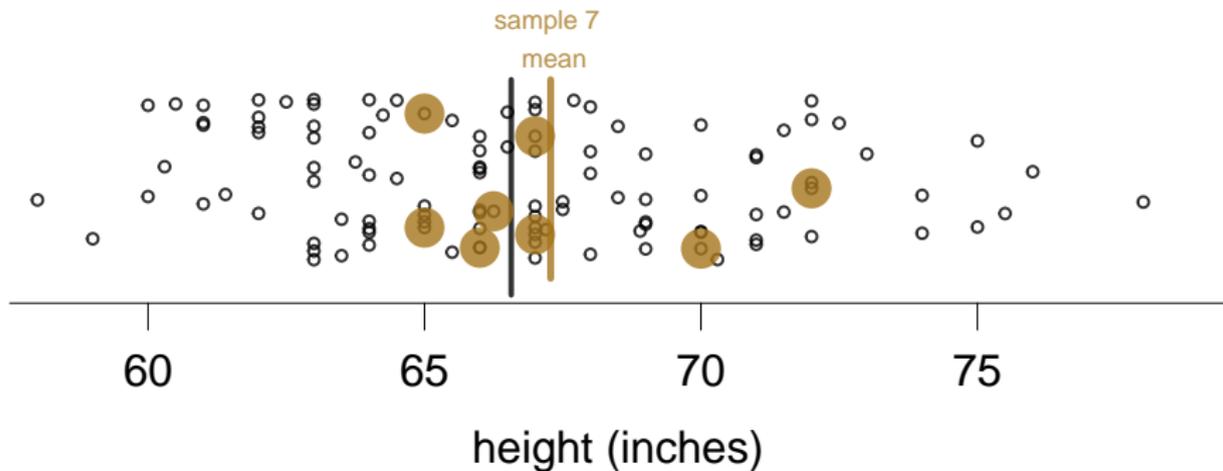


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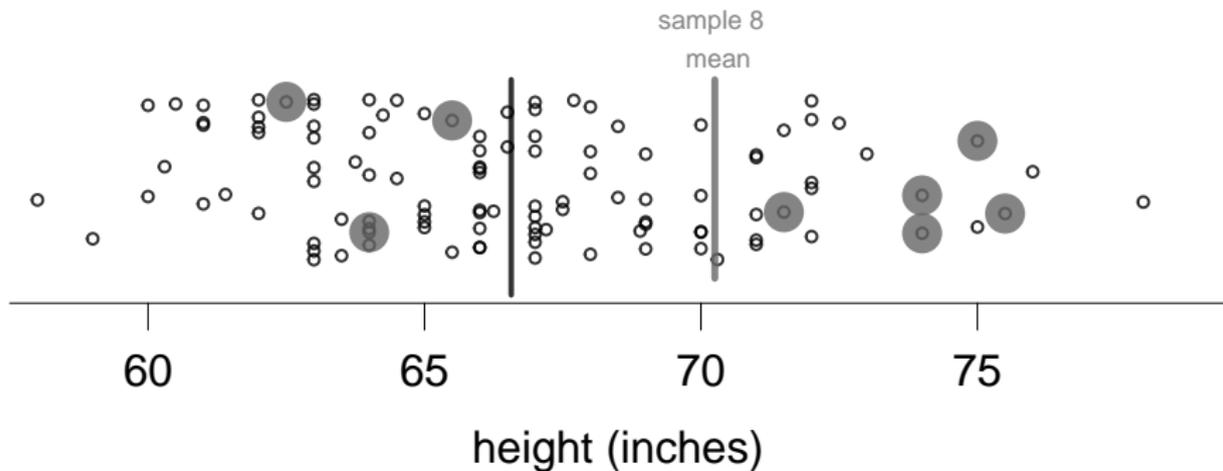
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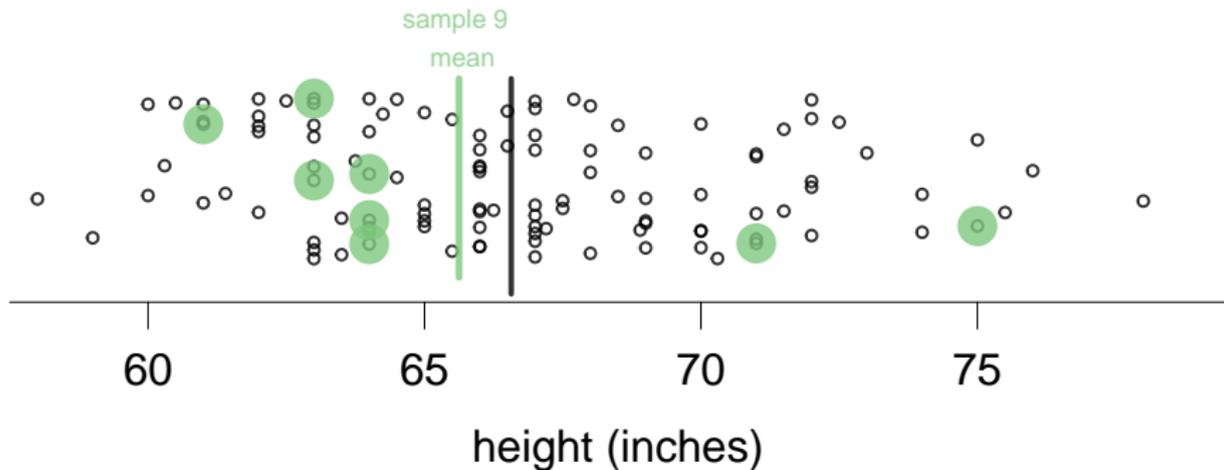
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When an estimate is neither systematically too high or too low, it is **unbiased**

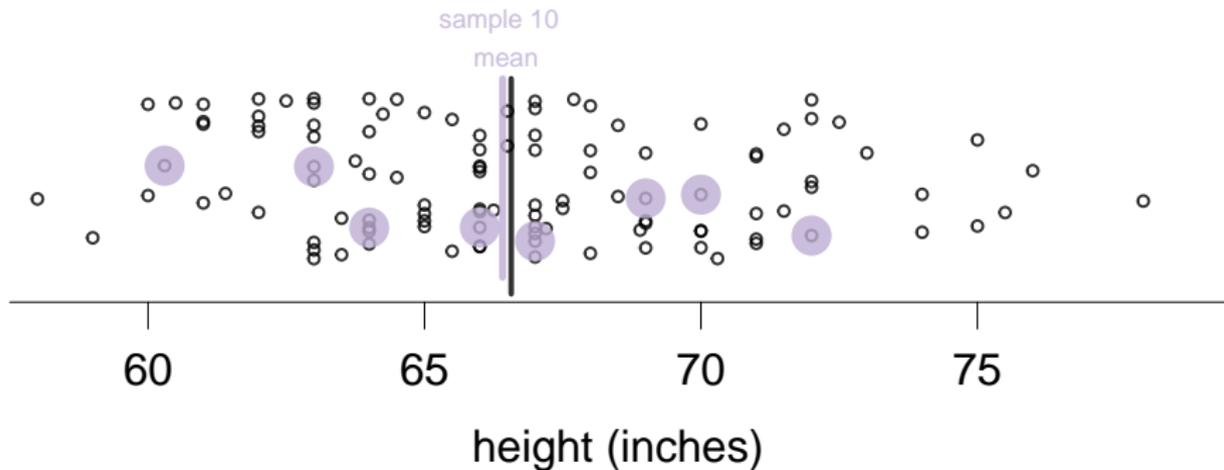


Unbiased estimators can still sometimes be far from the true value

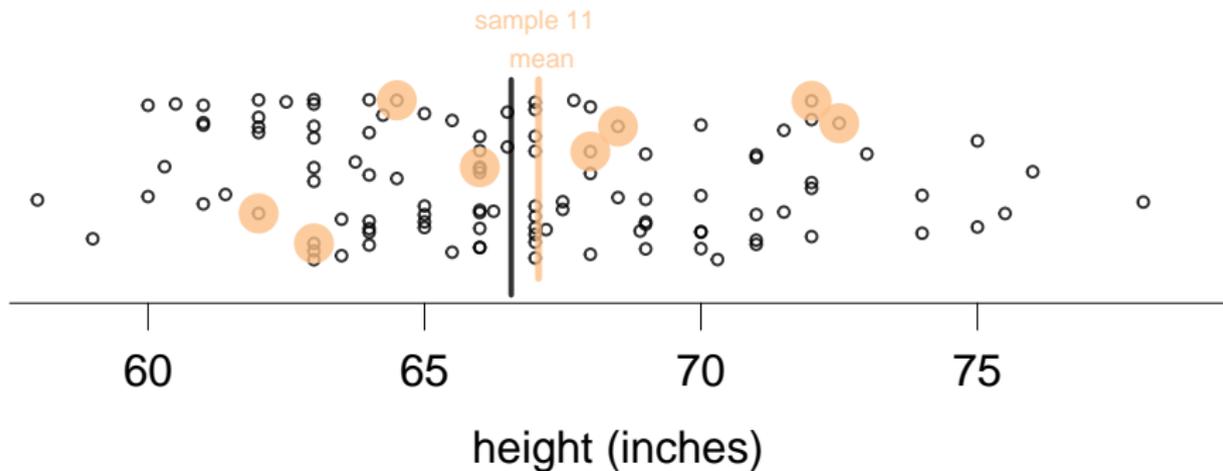


Unbiased estimators can still sometimes be far from the true value

But on average, over many samples, unbiased estimators will equal the truth, rather than show systematic *bias* up or down

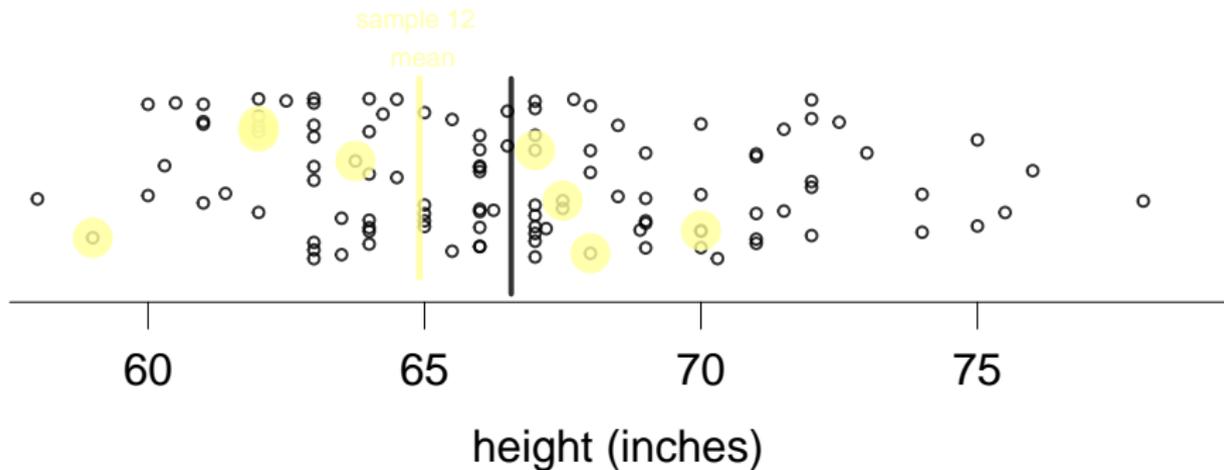


We also want our estimates to be *close* to the truth most of the time, a separate issue from bias

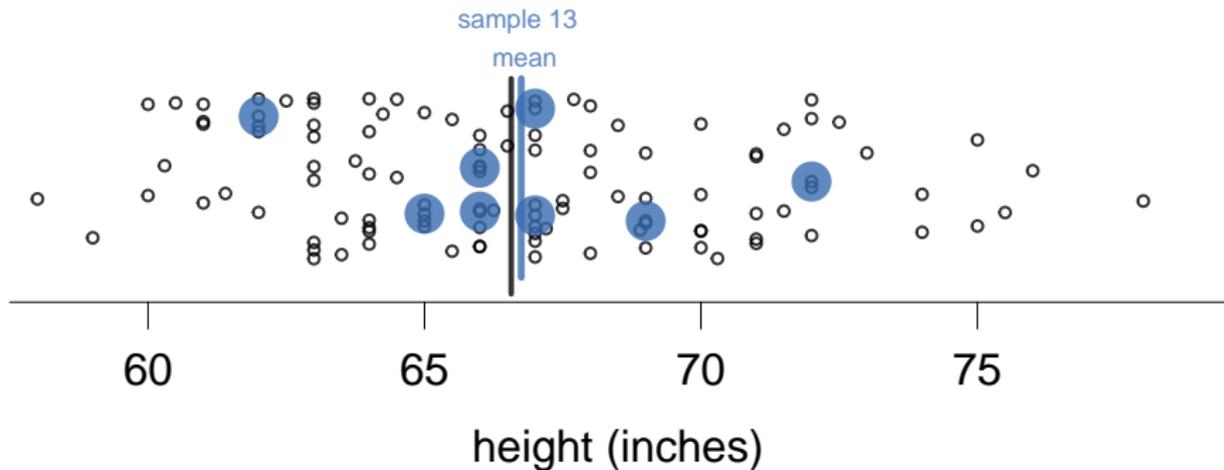


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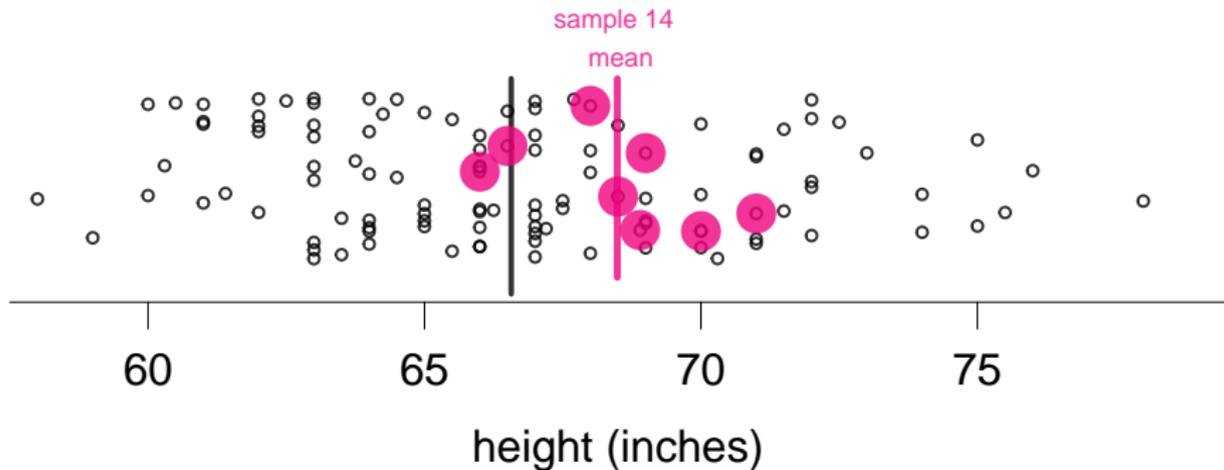
Sample estimates which are usually close to the population value are said to have low *error* or to be **efficient**



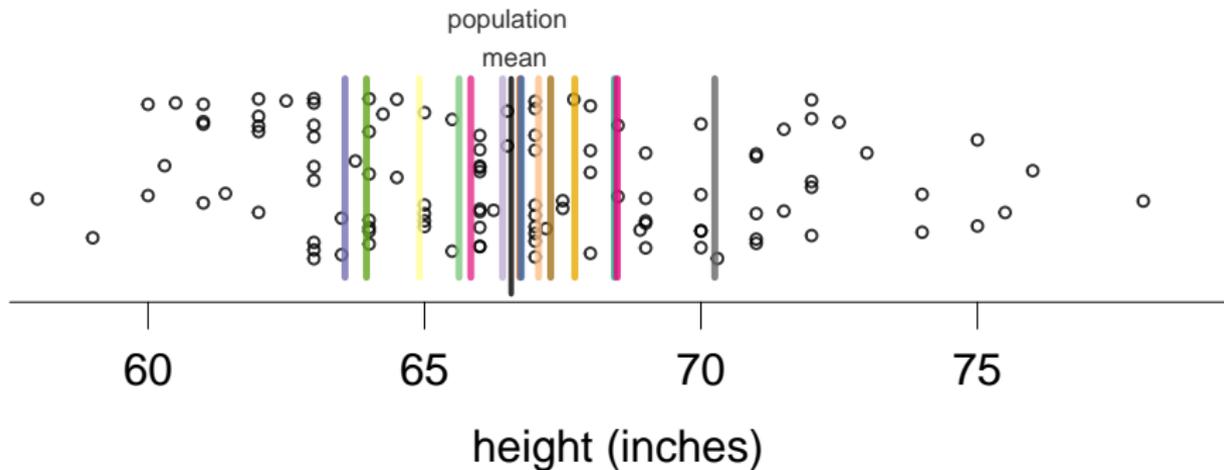
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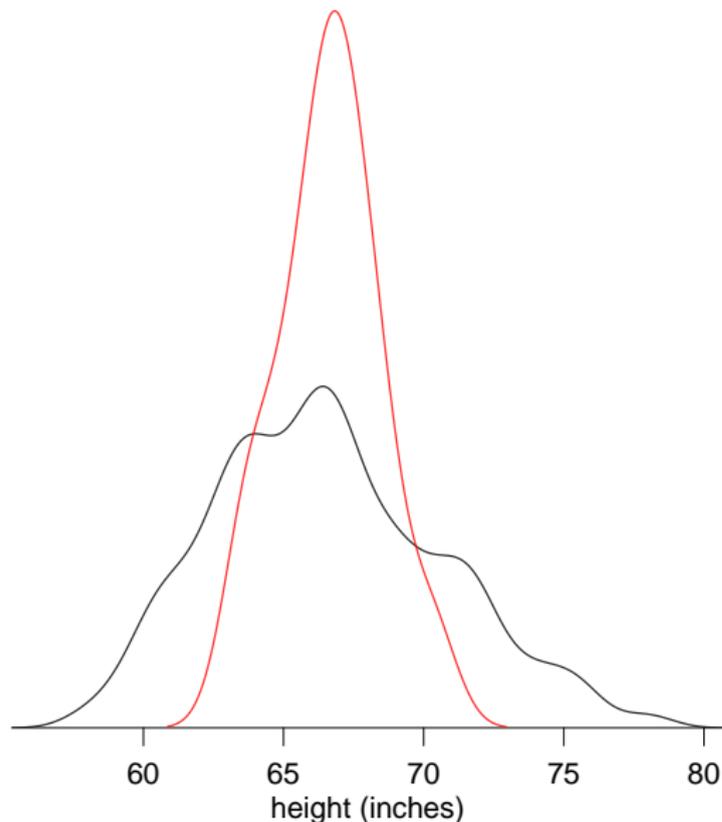
Notice that although the sample mean bounces back and forth,
it tends to stay close to the population mean
and doesn't range as far as the data itself



That is, the standard deviation of the sample means is smaller than the standard deviation of the data itself:

$$\text{sd}(\bar{x}) < \text{sd}(x)$$

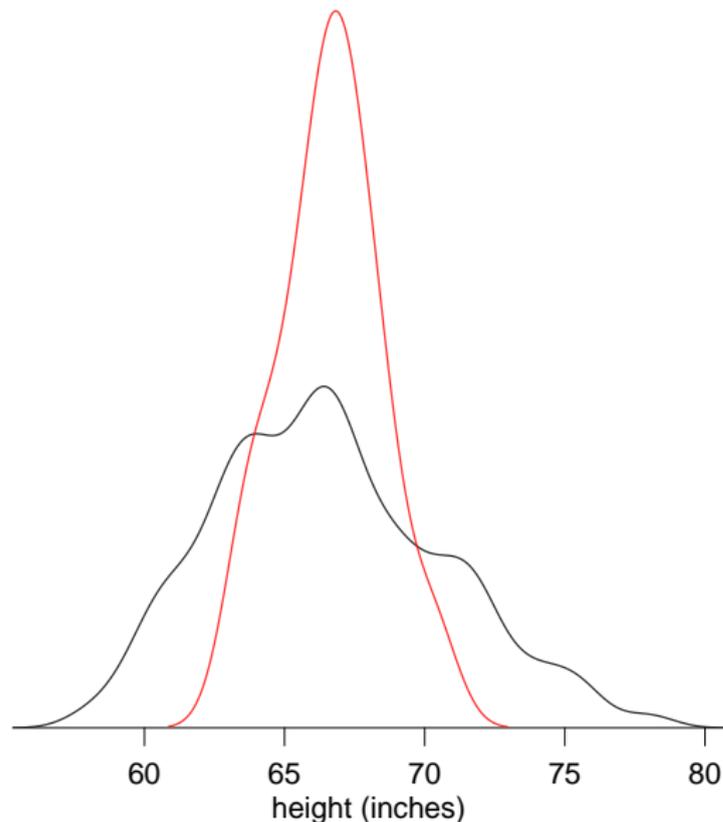
Student height example



We now overlay the distribution of sampling means in red

Note that the distribution of sampling means looks quite Normal, even though the distribution of heights is only approximately Normal.

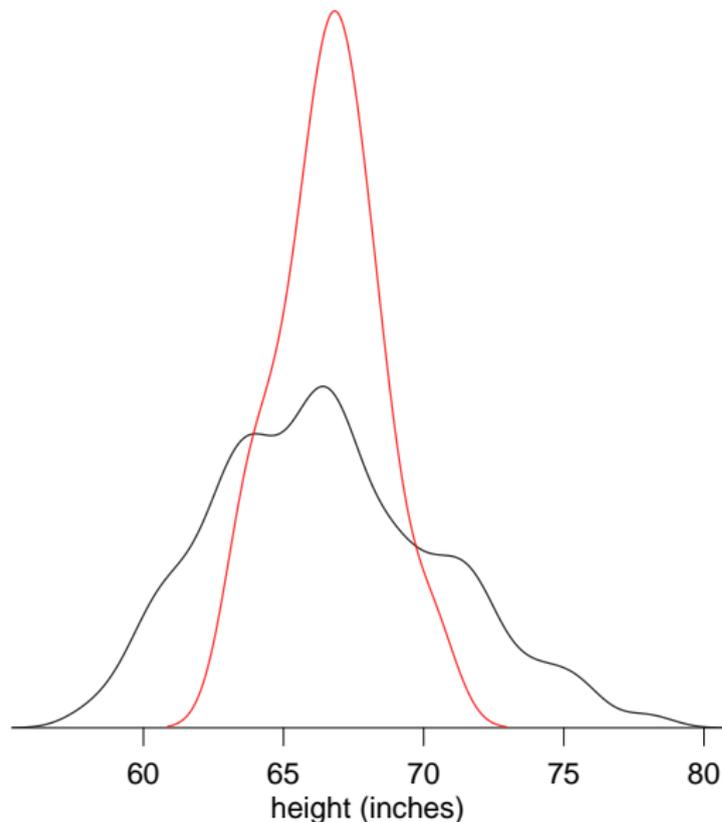
Student height example



This is the Central Limit Theorem again:

As we take more and more samples, the distribution of the sampling means of *any* random variable approaches the Normal (with few exceptions)

Student height example



To estimate the mean of population well, we want to make the distribution of sampling means (in red) as narrow as possible: ideally, a spike right over the true population mean

How can we do this?

The Law of Large Numbers

When sampling from a population, our estimates of features of that population get better the more data we sample

What do we mean by better estimates?

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Formula for the standard error of the mean:

$$se(\bar{x}) = \frac{sd(x)}{\sqrt{n}}$$

The Law of Large Numbers applies to estimating the mean of a population:

Our estimate of the mean, \bar{x} gets closer to the truth,
and its standard error, $se(\bar{x})$ gets smaller as the sampled n increases

The Square Root Law

Formula for the standard error of the mean:

$$se(\bar{x}) = \frac{sd(x)}{\sqrt{n}}$$

Remember that the smaller $se(\bar{x})$ is, the better our estimate

Making n bigger—adding more observations—will indeed shrink $se(\bar{x})$, but there are diminishing returns

Because $se(\bar{x})$ depends on \sqrt{n} , to halve the amount of error we must quadruple the amount of data

If our se is 1 inch of height with 100 observations, to reduce our expected error to 0.5 inches, we need 400 total observations

Note a surprise: the size of the population *does not appear* in this formula, and does not affect the precision of our estimates!

| | Mean | Sampling Error | Std Dev | Std Error |
|-------------|------|----------------|---------|-----------|
| Population | 66.6 | — | 4.1 | — |
| Sample 1 | 68.4 | 1.9 | 2.2 | 0.8 |
| Sample 2 | 66.7 | 0.2 | 5.5 | 2.0 |
| Sample 3 | 63.6 | -3.0 | 2.1 | 0.8 |
| Sample 4 | 65.8 | -0.7 | 4.4 | 1.6 |
| Sample 5 | 64.0 | -2.6 | 3.5 | 1.2 |
| Sample 6 | 67.7 | 1.2 | 5.2 | 1.9 |
| Sample 7 | 67.3 | 0.7 | 2.5 | 0.9 |
| Sample 8 | 70.3 | 3.7 | 5.4 | 1.9 |
| Sample 9 | 65.6 | -0.9 | 4.8 | 1.7 |
| Sample 10 | 66.4 | -0.2 | 3.9 | 1.4 |
| Sample 11 | 67.1 | 0.5 | 3.9 | 1.4 |
| Sample 12 | 64.9 | -1.7 | 3.8 | 1.3 |
| Sample 13 | 66.8 | 0.2 | 2.9 | 1.0 |
| Sample 14 | 68.5 | 1.9 | 1.7 | 0.6 |
| Sample Mean | 66.6 | | Avg SE | 1.4 |
| Pop Mean | 66.6 | | RMSE | 1.8 |

The population mean, 66.6 inches, is the average height in the class

Below it are the means of 14 samples of 8 students I drew to simulate “rows” of the classroom

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The average sample mean matches the population mean almost exactly

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Sampling Error is how much this row of students differs from the class mean:

How wrong (&in what direction) you'd be if you used your row to estimate the class average height

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RMSE is “root mean squared error”

RMSE is the average error we would make if we predicted the class from each row sample in turn

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Because errors can be + or -, we square error before averaging, then take the square root of the sum of squared errors to get RMSE

We need lots of samples to calculate RMSE

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The standard error is how much we expect this sample to miss by

$$\frac{\text{sd}(\text{height}_{\text{sample}})}{\sqrt{8}}$$

An estimate of RMSE we construct just from one sample

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| Sample 11 | 67.1 | 0.5 | 3.9 | 1.4 |
| Sample 12 | 64.9 | -1.7 | 3.8 | 1.3 |
| Sample 13 | 66.8 | 0.2 | 2.9 | 1.0 |
| Sample 14 | 68.5 | 1.9 | 1.7 | 0.6 |
| Sample Mean | 66.6 | | Avg SE | 1.4 |
| Pop Mean | 66.6 | | RMSE | 1.8 |

The average standard error is how much we expect to miss the population mean in repeated samples

Avg SE should match RMSE pretty closely

| | Mean | Sampling Error | Std Dev | Std Error |
|-------------|------|----------------|---------|-----------|
| Population | 66.6 | — | 4.1 | — |
| Sample 1 | 68.4 | 1.9 | 2.2 | 0.8 |
| Sample 2 | 66.7 | 0.2 | 5.5 | 2.0 |
| Sample 3 | 63.6 | -3.0 | 2.1 | 0.8 |
| Sample 4 | 65.8 | -0.7 | 4.4 | 1.6 |
| Sample 5 | 64.0 | -2.6 | 3.5 | 1.2 |
| Sample 6 | 67.7 | 1.2 | 5.2 | 1.9 |
| Sample 7 | 67.3 | 0.7 | 2.5 | 0.9 |
| Sample 8 | 70.3 | 3.7 | 5.4 | 1.9 |
| Sample 9 | 65.6 | -0.9 | 4.8 | 1.7 |
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Standard errors tell us how much we can trust our sample estimates

If standard errors are close to RMSE, then they are close to the true error in the estimate

Hypothesis testing

A framework for using a *sample* to test whether the mean of a population is on one side of a *reference point*

Invented by Jerzy Neyman & Egon Pearson, using concepts by R A Fisher

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For example, we might try to reject $\bar{x}^{\text{population}} = \mu_0$
in favor of $\bar{x}^{\text{population}} > \mu_0$

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Notice the italicized words. Our own theory is (somewhat) summarized by the alternative hypothesis,

But everything will hinge on an arbitrary reference point.

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Suppose we wanted to know if the average UW student works more than 10 hours a week.

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To see if $\bar{H}^{\text{population}} > 10$,

we could test against the null hypothesis that $\bar{H}^{\text{population}} = 10$ exactly

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Suppose we reject the null hypothesis of 10 hours or less.
We can say that we have rejected that possibility,
or that $H^{\text{population}}$ is **statistically significantly** greater than 10 hours.

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Could the true population mean hours still be 10.01?

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Could the true population mean hours still be 10.01?

Yes. Hypothesis tests are sharp and arbitrary.

The truth could be *any* value on this side of the null hypothesis.

Take care in selecting the null and interpreting the meaning of the test.

From z -scores to t -statistics

How do we perform a hypothesis test?

We need some way to standardize the distance between our sample mean \bar{x} and the null hypothesis μ_0

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From z -scores to t -statistics

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Here, we do something similar, using the standard error, to standardize the gap:

$$t = \frac{\text{sample statistic of interest} - \mu_0}{\text{se}(\text{sample statistic of interest})}$$

The t statistic of an estimate is:
the estimate itself, minus a hypothetical level,
divided by the standard error of the estimate

The t -statistic

In the case of the sample mean,

$$t = \frac{\bar{x} - \mu_0}{\text{se}(\bar{x})}$$

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In the case of the sample mean,

$$t = \frac{\bar{x} - \mu_0}{\text{se}(\bar{x})} = \frac{\bar{x} - \mu_0}{\text{sd}(x)/\sqrt{n}}$$

We are converting the gap between the data and the null into standard error units

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We will often set our hypothetical comparison level $\mu_0 = 0$, so this frequently reduces to:

$$t = \frac{\bar{x}}{\text{sd}(x)/\sqrt{n}}$$

The t -statistic

$$t = \frac{\bar{x} - \mu_0}{\text{se}(\bar{x})}$$

As with z -scores, our goal is to say how extreme or “unusual” the observed t is with reference to the distribution of t

t is a random variable with mean 0; the further away t is from its mean, the lower the chance of seeing a sample like our if the null hypothesis is true

But t isn't Normally distributed, so can't use the method we used for z -scores (looking up the quantiles of the Normal distribution)

Instead, we need to look up quantiles of the t -distribution (Table C in Moore 6th ed.)

The t distribution

Originally discovered by William Gosset, a statistician working at Guinness Brewery in the 1908 on the problem of measuring the quality of beer

If you sample random bottles of beer, how many bad samples do you need before you decide the production line is faulty?

Guinness was a pioneer of early statistical quality control, but forbade its statisticians from publishing (trade secrets!)

Gosset published his discovery under the pseudonym “Student”.
Hence this is Student’s t -test

The t distribution

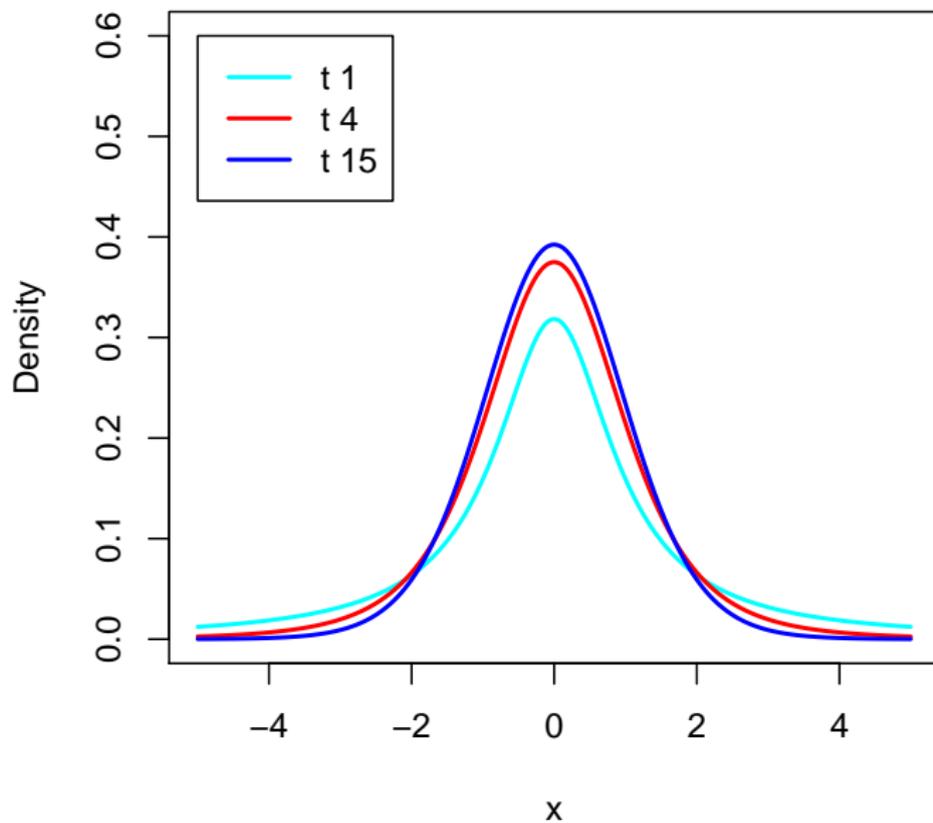
t distribution is a continuous distribution, with density from $-\infty$ to $+\infty$

Usually assume mean is 0

t distribution has a “degrees of freedom” parameter

As the degrees of freedom does up, the t distribution looks like the Normal

Example t distributions



The t distribution

Suppose we have a variable t that is t -distributed with mean 0 and 5 degrees of freedom

That is, $P(t) = t(5)$

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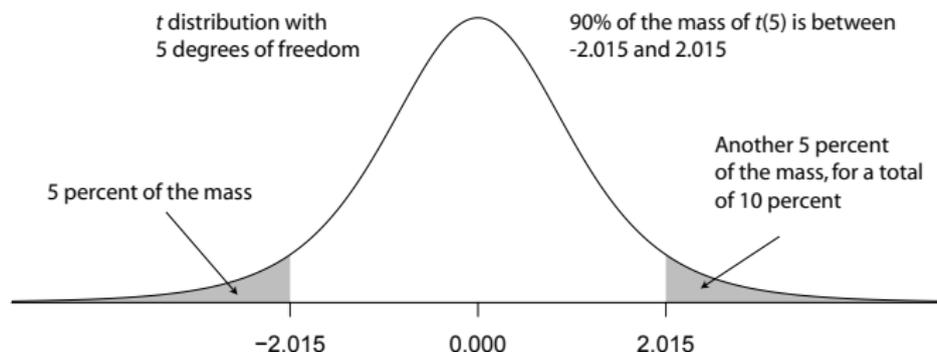
That is, $P(t) = t(5)$

What are the “critical” values of t we would see just

- once in 10 draws?
- once in 20 draws?
- once in 100 draws?

Put still another way,
which critical values will bound the 90% (or 95%, or 99%)
most ordinary t draws?

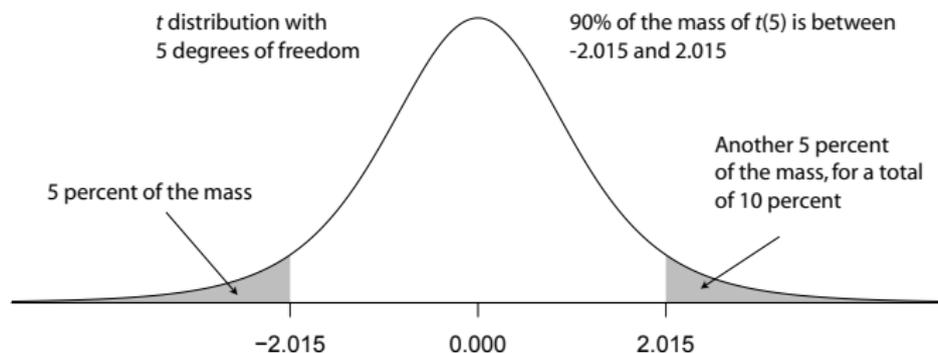
Areas under the t



Values outside the critical values are “unusual”.

We expect to see these values rarely, and may even suspect we have the wrong distribution if we see them often

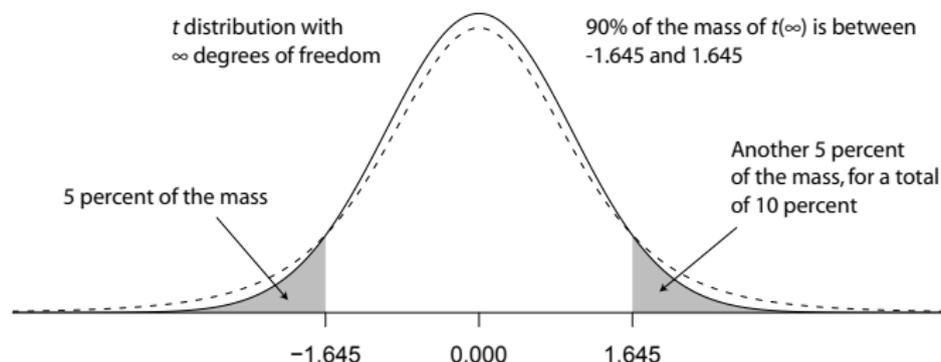
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Note that the above distribution is a t with 5 degrees of freedom

Degrees of freedom roughly here reflect how many independent pieces of information helped create the t -ratio

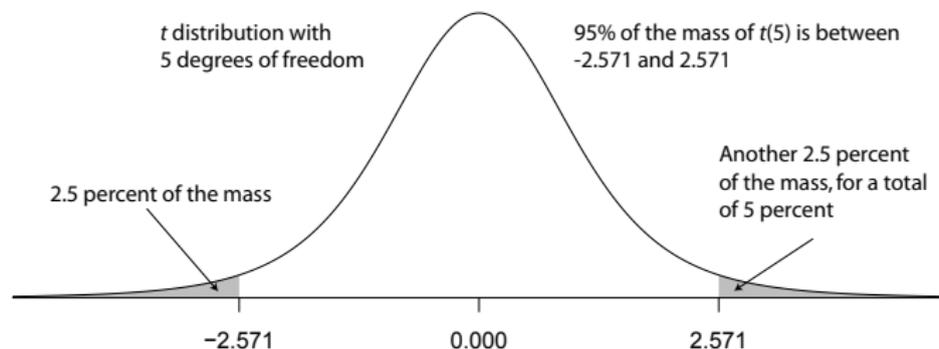
Areas under the t



More information makes t “better behaved”, so that extreme values occur less often,

More dfs thus make the tails thinner, and make critical values smaller

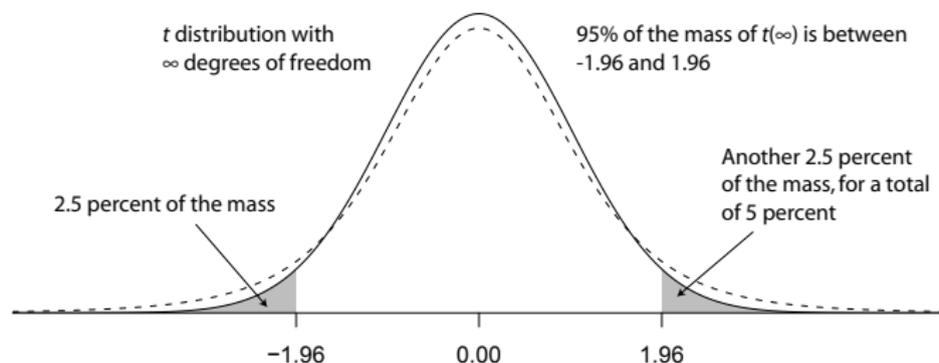
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Going back to the $df = 5$ case, notice we can choose what constitutes unusual

Here, we've raise the bar: only the 5% most extreme values are unusual, so the critical values increase

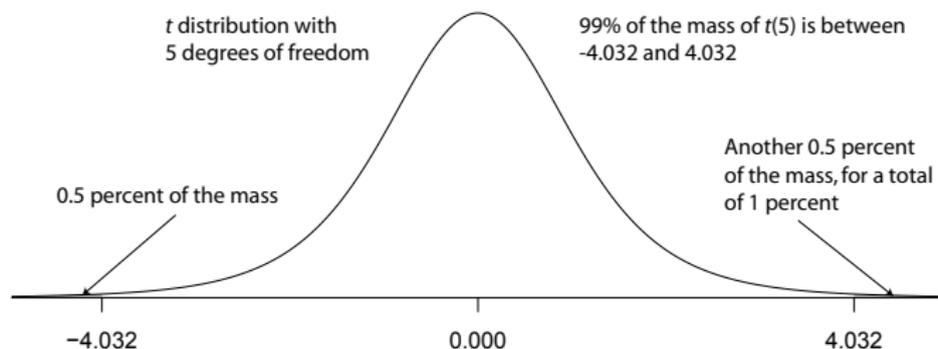
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The infinite degrees of freedom critical values for the 95% case

This is the most widely used standard for whether a t -ratio is unusual

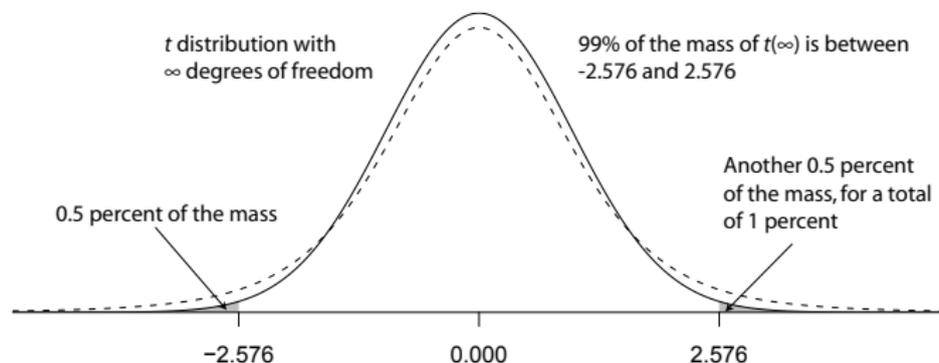
Areas under the t



The most stringent commonly used standard is 99%

In this case, a draw from the t must be in the 1% most extreme region to be considered unusual

Areas under the t



The infinite degrees of freedom case for 99%

Quick check: what do the critical values here mean?

Critical values of the t distribution

We can state how unusual a t -ratio is under the assumption that it is distributed $t(n)$

| Test level | Interval | df = 5 | df = ∞ |
|------------|----------|--------|---------------|
| 0.1 level | 90% | 2.015 | 1.645 |
| 0.05 level | 95% | 2.571 | 1.960 |
| 0.01 level | 99% | 4.032 | 2.576 |

These will be very useful for quantifying the uncertainty of estimates

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Large t could occur in one of two ways:

- 1 A unusual random sample far from the true population mean, which happens to be close to μ_0
- 2 A typical sample from a population mean that is larger than μ_0

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Roughly, dfs are how much information we have, in this case, $n - 1$, since calculating \bar{x} uses up a degree of freedom

Significance tests

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Significance tests are tests against a specific null hypothesis, and are “conservative” in the sense of being likely to favor the null over our own hypothesis

Are significance tests “really” conservative?’

Type I error Probability of falsely rejecting the null

Type II error Probability of falsely accepting the null

Significance tests minimize the chance of Type I error at the expense of allowing for more Type II error

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Is this a good idea?

The null hypothesis is usually arbitrary,
and our prior belief is usually that it is unlikely.

Significance tests may lead to excessive contrarianism,
which is not “conservative” at all

Confidence intervals

An better alternative to p -values which conveys the same information is the **confidence interval**

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Warning! We cannot say the truth lies in the confidence interval we calculate with 95% probability—we don't know in this specific case

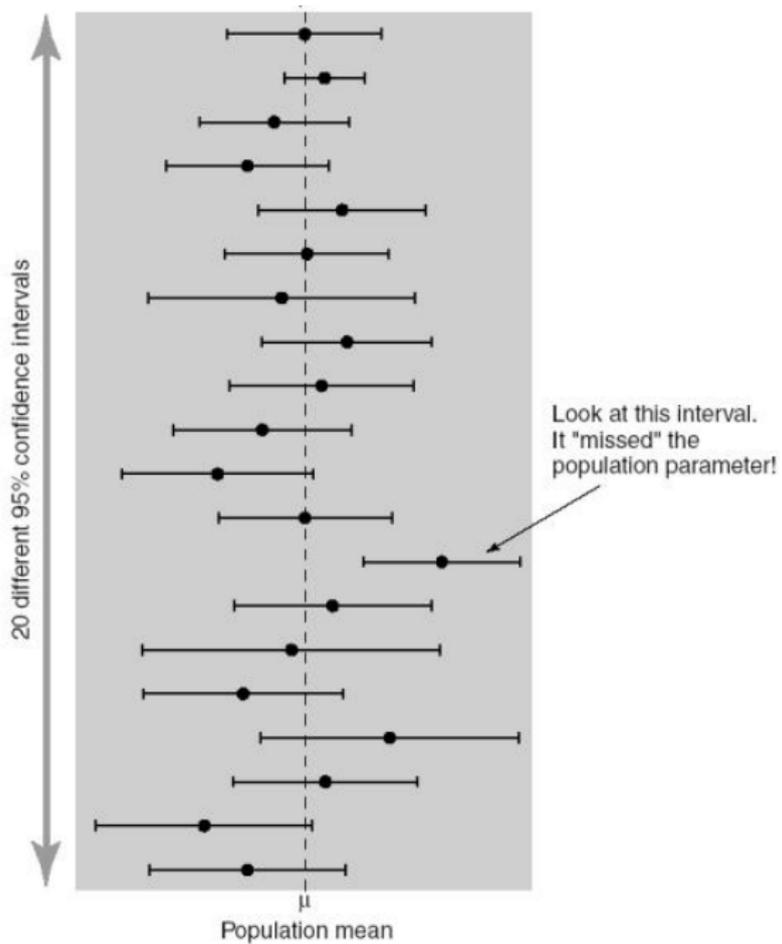
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But if we conduct 20 studies, and in each report a 95% confidence interval, we will expect to be “wrong” in only one study (1 in 20)



Calculating the confidence interval

We pick a confidence level, such as 95%

Then, we look up the critical value of t containing that 95% of the t distribution, and calculate:

$$\bar{x}^{\text{lower}} = \bar{x} - t_{n-1}^* \times \text{se}(\bar{x})$$

$$\bar{x}^{\text{upper}} = \bar{x} + t_{n-1}^* \times \text{se}(\bar{x})$$

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Note that for the 95% CI, the critical value with infinite degrees of freedom is ± 1.96 , so 95% CIs are roughly ± 2 standard errors from the estimate

Example: Washington Same-Sex Marriage Referendum

This week, Governor Gregoire signed legislation recognizing same-sex marriage in Washington State.

Opponents promised to petition to put the law on the November ballot.

Will same-sex marriage stand, or be repealed?

The Washington Poll, October 2011, asked a prescient question of 983 Washington registered voters

Next year the legislature could pass a law allowing gay and lesbian couples to get married. If that happens, there could be a referendum in which voters would be asked to approve or reject the law.

If such a referendum were held today: Would you vote YES – that is, to keep a law in place allowing gay and lesbian couples to marry OR would you vote NO, against the law – to make it so that gay and lesbian couples could not marry?

The Washington Poll found that 55 percent of registered voters would keep same-sex marriage, and 38 percent would not.

(The Washington Poll is conducted by my colleagues, Matt Barreto and Christopher Parker, and Betsy Cooper, of UW Political Science. See www.washingtonpoll.org.)

Example: Washington Same-Sex Marriage Referendum

The Washington Poll found that 55 percent of registered voters would keep same-sex marriage, and 38 percent would not.

How certain is this result?

We will use the raw data from this survey to investigate

Some caveats:

- 1 Original survey was stratified, and weighted some groups more heavily; we will ignore weights
- 2 To simplify, we will ignore non-response and “I don't knows”.

Because of the above (especially 2) our proportions in this lecture differ from the official results of the poll.

Example: Washington Same-Sex Marriage Referendum

Our initial N of people responding YES or NO on the same-sex referendum is 979.

Of these, 61.6% say YES, they would vote to keep SSM.

Assuming the caveats above pose no problems, how certain are we the referendum will pass based on this sample?

Example: Washington Same-Sex Marriage Referendum

How likely is it that a survey of 979 random individuals from a population would find 61.6% support for a measure when really only 50% or less support the measure?

Let's use a t -test:

$$t = \frac{\bar{x} - \mu_0}{\text{se}(\bar{x})}$$

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Why is this so unlikely? Suppose that in October, a bare majority of Washington registered voters really did oppose same-sex marriage.

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Then to get 61.6% approval, instead of the correct 50% approval, the Washington Poll would have needed to sample $979 \times (0.616 - 0.500) = 114$ more supporters than we would expect on average in 979 random draws.

Example: Washington Same-Sex Marriage Referendum

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Then to get 61.6% approval, instead of the correct 50% approval, the Washington Poll would have needed to sample $979 \times (0.616 - 0.500) = 114$ more supporters than we would expect on average in 979 random draws.

That's as unlikely as flipping a coin 979 times and getting 603 heads and 376 tails.

Example: Washington Same-Sex Marriage Referendum

Another way to summarize the uncertainty in our polling results is to calculate a confidence interval

We can state with 95% confidence that the actual level of support for same-sex marriage among all Washington RVs in April was between 58.5% and 64.6%

Notice these numbers are $61.6\% \pm 3.1\%$, which also happens to be the “margin of error” for the poll (journalists’ name for a confidence interval).

Unfortunately, “margin of error” is a misleading name: errors can be bigger than this margin, & are guaranteed to be 5% of the time!

Example: Washington Same-Sex Marriage Referendum

The Washington Poll's sample of Washington voters includes 317 voters over the age of 65, 53.9% percent of whom said they would support SSM

Do a majority of older Washingtonians actually support SSM?
Or is this a sampling error?

If we made the *mistake* of judging by the “margin of error” for the whole survey, we might think a majority of older voters did support SSM:

$$53.9\% - 3.1\% = 50.8\%$$

Example: Washington Same-Sex Marriage Referendum

Let's do our own t -test to be sure:

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This is a pretty small t -statistic, one we would see by chance in 1 out of 6 random samples. The p -value is 0.161.

We find that the 95% confidence interval ranges from 48.4% to 59.5%, which is equal to our estimate of 53.9% by $\pm 5.5\%$.

We are not certain that Washington 65+'s supported SSM in October.

Example: Washington Same-Sex Marriage Referendum

- 1 Uncertainty depends on the size of the sample (which has changed)
- 2 Uncertainty depends on the variance of the sample (which has changed)

Change in Size of Sample

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The smaller the n , the bigger the confidence interval, the less certain the finding

On confidence versus significance

There are two ways we could report our finding on older voters support for same-sex marriage:

Significance test Based on a survey of Washington registered voters, we estimate 54% of voters over 65 years supported same-sex marriage in October. However, this estimate is not statistically significantly different from 50% at the 0.05 level.

On confidence versus significance

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Significance test Based on a survey of Washington registered voters, we estimate 54% of voters over 65 years supported same-sex marriage in October. However, this estimate is not statistically significantly different from 50% at the 0.05 level.

Confidence interval Based on a survey of Washington registered voters, we estimate 53.9% of voters over 65 years supported same-sex marriage in October. The 95% confidence interval for this estimate ranges from 48.4% to 59.5%, suggesting anywhere from a slight majority against same-sex marriage to a large majority in favor.

On confidence versus significance

These write-ups present the same results. They rely on the same math and the same statistical theory.

The significance test presentation obscures the substantive impact of the result in jargon, and makes it appear ignorable.

The confidence interval focuses on the substantive impact of the result, and clarifies what we can and cannot reject:

Although we aren't sure how many older voters supported same-sex marriage in October,

it is very likely that half or more do,

and very unlikely that a large percentage of older were opposed before 2012 started

On confidence versus significance

The significance test forces you to accept the author's arbitrary null hypothesis

The confidence interval allows you to choose your own null

And shows how robust your findings are to slight changes in the null

The irrelevance of population size

$$t = \frac{\bar{x} - \mu_0}{\text{sd}(x)/\sqrt{n}}$$

Notice one number that doesn't appear in this formula: the size of the population

The precision of an estimate doesn't depend on the size of the population, only the size of the sample.

That's why you tend to see polls using samples of 500 to 2000 respondents regardless of whether they are sampling from a small town population or the whole country

Comparing two means

So far, we have asked how far the mean of our sample might differ from a specific value

e.g., how much does the average support for same-sex marriage differ from 0.5?

But what if we want to compare two groups in our sample?

That is, what if we want to compare two means to each other?

e.g., how much does the average support for same-sex marriage among those with a close gay friend or family member differ from support among those without (knowledge of) close contact with someone gay?

A simple cross-tab

| | Have contact? | | Total |
|-------------|---------------|-----|-------|
| | Yes | No | |
| Support SSM | 399 | 204 | 603 |
| Oppose SSM | 183 | 193 | 376 |
| Total | 582 | 397 | 979 |

Here are the two variables, support for SSM and contact, in a cross-tabulation

Let's convert to column percentages

A simple cross-tab

| | Have contact? | | Mean |
|-------------|---------------|--------|--------|
| | Yes | No | |
| Support SSM | 68.6% | 51.4% | 61.6% |
| Oppose SSM | 31.4% | 48.6 | 38.4% |
| Total | 100.0% | 100.0% | 100.0% |

This table shows much more support for SSM among those with contact than those without (68.6% vs. 51.4%, or 17.2% more support)

Our question:

How certain are we that this difference in mean support across groups in our sample really exists in the Washington voter population?

t-test for comparison of means

To make inferences about the *difference* in means of two samples, we can do a *difference in means t*-test

Remember the form of a *t*-statistic:

$$t = \frac{\text{sample statistic of interest} - \mu_0}{\text{se}(\text{sample statistic of interest})}$$

Before, the sample statistic of interest was \bar{x} , but now it is $\bar{x} - \bar{y}$. We want to know if this difference is itself different from zero, so:

$$t = \frac{(\bar{x} - \bar{y}) - 0}{\text{se}(\bar{x} - \bar{y})}$$

t -test for comparison of means

Our difference-of-means t -statistic is:

$$t = \frac{(\bar{x} - \bar{y}) - 0}{\text{se}(\bar{x} - \bar{y})}$$

Calculating this t by hand is hard, so we'll let the computer do it.

Then we'll check if this t -statistic exceeds the chosen critical value or simply calculate the probability of seeing so large a t

Example: Washington Same-Sex Marriage Referendum

Voters in the Washington Poll sample with contact were 17.2% more likely to support same-sex marriage

How certain are we that this difference holds in the population?

That is, how certain are we that $\Pr(\text{support}|\text{contact}) - \Pr(\text{support}|\text{no contact}) > 0$?

We can do a comparison of means t -test.

We find $t = 5.425$, which implies a p -value of 0.00000007661.

The 95% confidence interval is ranges from +11.0% to +23.4%. (What does this mean?)

Example: Household Wealth and Race

In a sample of 10,000 households, we found households headed by a self-identified white earned more, on average, than households headed by a self-identified black or Hispanic.

How certain are we that these sample results hold in the full American population?

Example: Household Wealth and Race

Let's do a comparison-of-means t -test for black and white households

Average gap between black and white household wealth, in \$k: -496.7

t -stat: -19.8

p -value: 0.000000000000000022

(that's just 1 in 4,540,000,000,000,000, or 4.5 thousand trillion)

95% CI: -545.9 to -447.5

Summing up

We've added several new tools to our analytic toolkit:

- 1 Standard errors of estimates
- 2 t -tests and confidence intervals for a sample mean
- 3 t -tests and CIs for a comparison of means

Caveats

Comparison of means tests seem especially helpful for our inference about hypotheses

We can now state whether apparent differences in conditional means are likely to be mere happenstance, or real features of the population

But are there reasons to doubt findings from a comparison of means test?

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We can now state whether apparent differences in conditional means are likely to be mere happenstance, or real features of the population

But are there reasons to doubt findings from a comparison of means test?

These tests still don't control for *confounders*. So results might be spurious.

Wait! What are degrees of freedom (df)?

Degrees of freedom:

The number of separate pieces of information used to calculate a statistic

“separate” = “freely movable”

Not the same thing as the number of observations (may be the same as N or less)

Relevance: how many quantities could we estimate from a set of data?

Can't be more quantities than are left to vary!

How many things can we estimate using?

two numbers, x_1 and x_2

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fifty numbers, x_1, \dots, x_{50} , and $\bar{x} = 2$

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49 things

fifty numbers, x_1, \dots, x_{50} , $\bar{x} = 2$, and $\sigma^2 = 0.5$

Degrees of freedom (df)

Degrees of freedom (df): the remaining allowed ways you could move the data

If we make as many assumptions as there are observations, nothing left to estimate