

Confidence Intervals

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1. The sample average as a random variable
2. Sampling distributions
3. Confidence intervals

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 - ▶ We don't know which cases from the population will be sampled
 - ▶ We don't know which study participants will be randomized to the treatment/control group
- ▶ Further, *any descriptive summary* of the sample data (ie: means, proportions, correlations, etc.) is the *observed outcome* of a *random variable*

The Sample Average as a Random Variable

For a sample of n cases from a population, the sample average is calculated:

$$\bar{x} = \frac{x_1 + x_2 + x_3 + \dots + x_n}{n}$$

Notice how \bar{x} is the sum of many different randomly selected cases

Proportions are Averages

- ▶ Now, consider a *binary categorical* variable
 - ▶ Because binary variables involve only two categories, we can map their outcomes to the numeric values of 0 and 1
 - ▶ For example, consider a coin flip, we could map the outcome “Heads” to “1” and the outcome “Tails” to “0”

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 - ▶ For example, consider a coin flip, we could map the outcome “Heads” to “1” and the outcome “Tails” to “0”
- ▶ By coding the outcomes using 1s and 0s, we can see that the *sample proportion* is also an average:

$$\hat{p} = \frac{1+0+1+1+0+\dots+1}{n}$$

- ▶ If we mapped “Heads” to a value of “1”, \hat{p} would refer to the proportion of heads in our sample

The Distribution of the Sample Proportion

- ▶ According to the US Census, 27.5% of the adult population are college graduates
- ▶ Randomly sampling n adults represents a *random process*
 - ▶ The proportion of college graduates in a sample, \hat{p} , is a *random variable*

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 - ▶ The proportion of college graduates in a sample, \hat{p} , is a *random variable*
- ▶ Let's explore some different outcomes of this random variable for two different sampling protocols: random samples of size $n = 10$, and random samples of size $n = 100$
 - ▶ To begin, describe the *sample space* of \hat{p} in each scenario

Random Samples of size $n = 10$

- ▶ For a single random sample of size $n = 10$, there are exactly 11 different sample proportions that could occur
 - ▶ Thus, the sample space is: $\{0/10, 1/10, 2/10, \dots, 10/10\}$

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 - ▶ Thus, the sample space is: $\{0/10, 1/10, 2/10, \dots, 10/10\}$
- ▶ Rather than doing probability calculations, we'll simulate sampling from the population (of size $n = 10$) to judge the likelihood of each outcome
 - ▶ StatKey allows us to quickly draw many samples

Random Samples of size $n = 10$

Due to the relatively small number of discrete outcomes, it's reasonable to use a table to convey a probability model for the sample proportion:

| Sample Proportion ($n = 10$) | Probability |
|--------------------------------|-------------------|
| 0/10 | $40/1000 = 0.04$ |
| 1/10 | $150/1000 = 0.15$ |
| 2/10 | $250/1000 = 0.25$ |
| 3/10 | $270/1000 = 0.27$ |
| 4/10 | $190/1000 = 0.19$ |
| ... | ... |
| 10/10 | $0/1000 = 0$ |

Random Samples of size $n = 100$

- ▶ For $n = 100$, it's more sensible to use a Normal probability model
- ▶ StatKey can help us determine this model
 - ▶ Any Normal model depends upon two parameters, the expected value/mean (ie: μ) and the standard deviation (ie: σ)

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- ▶ StatKey can help us determine this model
 - ▶ Any Normal model depends upon two parameters, the expected value/mean (ie: μ) and the standard deviation (ie: σ)
- ▶ When the random variable our interest is a *descriptive statistic*, the term **standard error** is used to describe model's standard deviation
 - ▶ Thus, “standard deviation” is used to describe variability among individual cases, while “standard error” describe the variability of an estimate

Use StatKey to simulate drawing random samples of $n = 200$ from this population (US adults, among whom 27.5% are college graduates)

- 1) Is the *standard error* larger or smaller than it was for samples of size $n = 100$? Was the change *linear*? (ie: is the *SE* half or double what it previously had been?)
- 2) Use the mean and standard error as the basis for a Normal model. What percentage of samples would you expect to have between 26% and 29% colleges graduates?
- 3) Can you find an *interval* that you'd anticipate will contain the sample proportions found in 90% of samples?

Is simulation necessary?

The idea of reporting an *interval estimate* is very useful in describing the uncertainty involved in random processes (ie: describing sample data). However, there are two important challenges to consider:

- 1) We typically do not know anything about the population, everything we know is based upon the sample data
- 2) It's not always reasonable to do thousands of simulations

- ▶ John Kerrich, a South African mathematician, was visiting Copenhagen in 1940
- ▶ When Germany invaded Denmark he was sent to an internment camp, where he spend the next five years
- ▶ To pass time, Kerrich conducted experiments exploring sampling and probability theory
 - ▶ One of these experiments involved flipping a coin 10,000 times

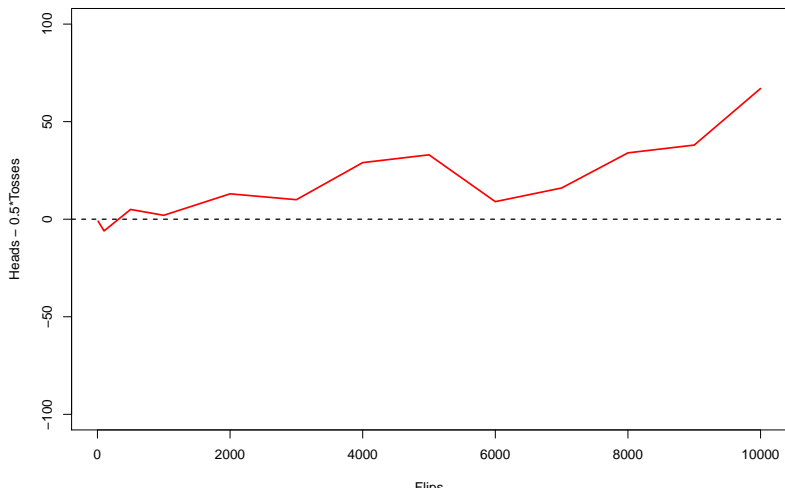
- ▶ We know that a fair coin shows “Heads” with a probability of 50%
- ▶ So, in a random sample of n coin flips, we'd expect roughly even numbers of “Heads” and “Tails”
 - ▶ We'll explore the results of Kerrich's experiment to see why the *sample average* is so special

Kerrich's Results

| Number of Tosses (n) | Number of Heads | Heads - $0.5 * \text{Tosses}$ |
|--------------------------|-----------------|-------------------------------|
| 10 | 4 | -1 |
| 100 | 44 | -6 |
| 500 | 255 | 5 |
| 1,000 | 502 | 2 |
| 2,000 | 1,013 | 13 |
| 3,000 | 1,510 | 10 |
| 4,000 | 2,029 | 29 |
| 5,000 | 2,533 | 33 |
| 6,000 | 3,009 | 9 |
| 7,000 | 3,516 | 16 |
| 8,000 | 4,034 | 34 |
| 9,000 | 4,538 | 38 |
| 10,000 | 5,067 | 67 |

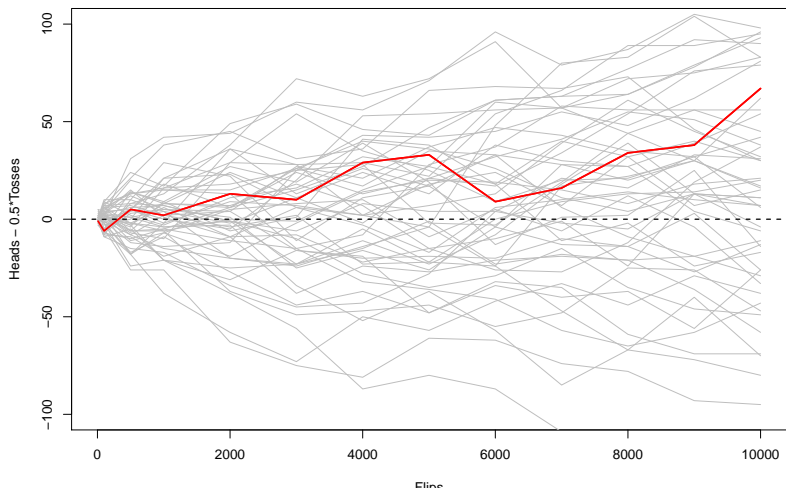
Kerrich's Results

It seems like the number of heads and tails are actually getting further apart. . . could this be a fluke?



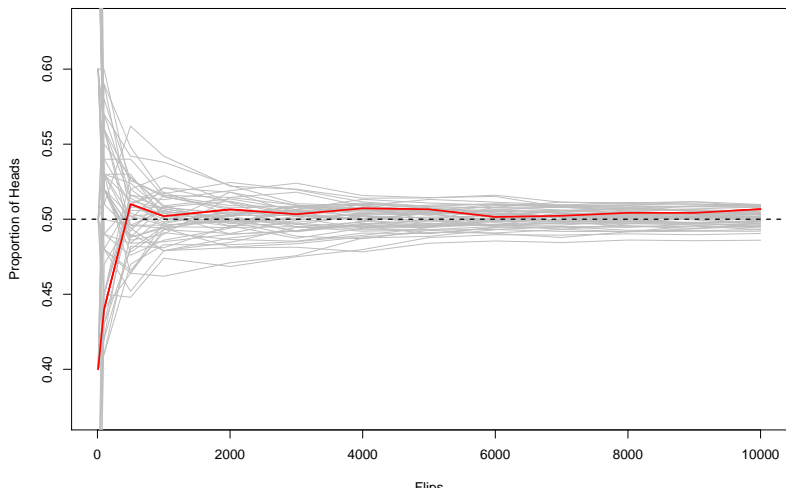
Kerrich's Experiment (repeated 50 times)

No, the phenomenon occurs systematically when repeating Kerrich's experiment



Kerrich's Experiment (sample proportions)

The *sample proportion* of heads behaves exactly as we'd expect, but why?



Central Limit Theorem

- ▶ Suppose X_1, X_2, \dots, X_n are independent random variables with a common expected value $E(X)$ and variance $Var(X)$ (see previous notes for definitions of these two terms)
- ▶ Let \bar{X} denote the average of all n random variables, **Central Limit Theorem** (CLT) states:

$$\sqrt{n} \left(\frac{\bar{X} - E(X)}{\sqrt{Var(X)}} \right) \rightarrow N(0, 1)$$

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- ▶ Often it is more useful to think of CLT in the following way (which abuses notation):

$$\bar{X} \sim N \left(E(X), \frac{SD(X)}{\sqrt{n}} \right)$$

Central Limit Theorem and Sample Proportions

- ▶ The sample proportion is comprised of n different binary variables (taking on values of 1 and 0)
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 - ▶ $Var(X) = p * (1 - p)^2 + (1 - p) * (0 - p)^2 = p * (1 - p)$

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 - ▶ $Var(X) = p * (1 - p)^2 + (1 - p) * (0 - p)^2 = p * (1 - p)$
- ▶ Thus, the *sampling distribution* of sample proportions is:

$$\hat{p} \sim N(p, \sqrt{p(1 - p)/n})$$

The Power of CLT

- ▶ Central Limit Theorem is one of the most important theoretical results in all of statistics
- ▶ In real-world applications, it is nearly impossible to know the probability distribution of something that is only observed once (remember that real researchers can only afford to collect a single sample)

The Power of CLT

- ▶ Central Limit Theorem is one of the most important theoretical results in all of statistics
- ▶ In real-world applications, it is nearly impossible to know the probability distribution of something that is only observed once (remember that real researchers can only afford to collect a single sample)
- ▶ But by focusing on the *sample average* this isn't an issue, as CLT provides us the distribution of sample averages
 - ▶ That is, we are able to use CLT to understand the *sampling variability* of our study, despite only getting to see a single sample!

Example

- ▶ Let's consider a random sample of $n = 100$ coin flips
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 - ▶ What proportion of heads might we expect? It'll likely be close to 50%, but we know there's sampling variability, the question is how much. . .
- ▶ Each coin flip is a random variable an expected value of 0.5, so Central Limit Theorem tells us that proportion of heads in random samples of $n = 100$ coin flips follows a Normal distribution:

$$\hat{p} \sim N(0.5, \sqrt{0.5(1 - 0.5)/100})$$

Using the Central Limit theorem to determine the distribution of sample averages is only appropriate when the following conditions are met:

- 1) *Independence* - the cases in the sample (ie: the individual contributions to the sample average) are not related to each other
- 2) *Large population* - less that 10% of the population is being sampled (otherwise removing the already sampled individuals has too much of an impact on the probability of selection)
- 3) *Large sample* - $n\hat{p} \geq 10$ and $n(1 - \hat{p}) \geq 10$

Most of the time, it's only the third condition that is problematic

Let's revisit the selection of a random sample of size $n = 100$ from the population of US adults (of which 27.5% are college graduates)

- 1) Use Central Limit theorem to come up with a Normal model for the proportion of college graduates in the sample.
- 2) How does the *standard error* in this model compare to the standard error we found by simulation? (recall it was approximately 0.045)

Statistical Inference

Recall that the *fundamental goal* of statisticians is to use information from a sample to make *reliable* statements about a population, a process known as **statistical inference**:

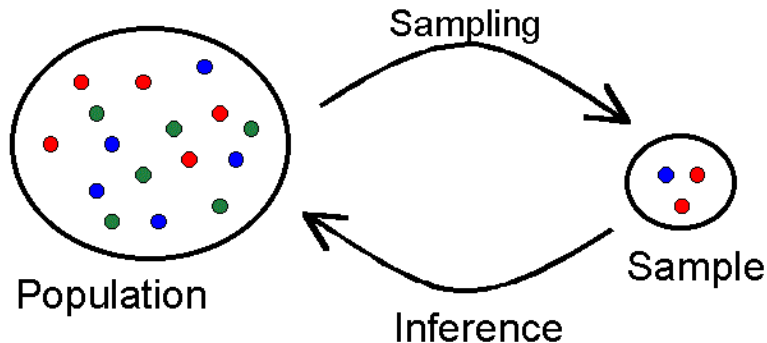


Image credit: <http://testofhypothesis.blogspot.com/2014/09/the-sample.html>

- ▶ If a sampling protocol is *unbiased*, the sample average is a sensible estimate of the population mean
 - ▶ This is called a **point estimate**, referring to the fact that it is a single value

- ▶ If a sampling protocol is *unbiased*, the sample average is a sensible estimate of the population mean
 - ▶ This is called a **point estimate**, referring to the fact that it is a single value
- ▶ From our study of *sampling distributions*, we know that the existence of sampling variability means a point estimate is almost certainly wrong (at least to some degree)
 - ▶ This suggests that we can more appropriately describe what we think is true of the population by reporting an **interval estimate** that accounts for *sampling variability*

Point vs. Interval Estimation

To summarize:

- ▶ **Point estimation** uses sample data to produce a *single “most likely” estimate* of a population characteristic, which will almost always miss the target (at least by some degree)
- ▶ **Interval estimation** uses sample data to produce a *range of plausible estimates* of a population characteristic, an approach that has a much better chance at capturing the truth

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To summarize:

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An analogy:

Using only a point estimate is like fishing in a murky lake with a spear. We can throw a spear where we saw a fish, but we will probably miss. On the other hand, if we toss a net in that area, we have a good chance of catching the fish.

Most interval estimates have the form:

$$\text{Point Estimate} \pm \text{Margin of Error}$$

We often report these intervals using only their endpoints:

$$(\text{Est} - \text{MOE}, \text{Est} + \text{MOE})$$

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- ▶ We'd like the *margin of error* to be constructed in way that carries a *quantifiable* claim of precision
 - ▶ ie: 80% of the time an interval with this type of margin of error will contain the population characteristic
 - ▶ Without an accompanying claim regarding precision, reporting a margin of error is not particularly useful

So, what can we say about a population proportion, p , based upon an observed sample proportion, \hat{p} ? Consider a representative sample of 100 infants used to estimate the proportion of all babies who are born prematurely

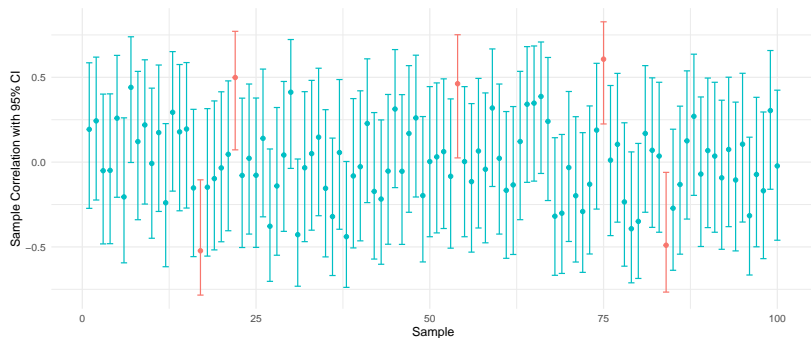
- ▶ True or false? “We observed $\hat{p} = 0.14$, so we know that 14% of all babies are born prematurely”
- ▶ True or false? “We observed $\hat{p} = 0.14$, it’s probably true 14% of all babies are born prematurely”
- ▶ True or false? “Although we don’t know p , if we attach a large margin error to our point estimate, the interval estimate $14\% \pm 10\% = (4\%, 24\%)$ probably contains p ”

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- ▶ A **confidence interval** is an interval estimate whose margin of error is based upon a procedure with a long-run “success rate” known as a *confidence level*
 - ▶ A 95% confidence interval was created using a procedure that will succeed in containing the true population parameter in 95% of different random samples (or study replications)
 - ▶ The confidence level does not describe the likelihood that particular interval succeeds, instead it describes the estimation procedure’s long-run success rate

Confidence intervals

Shown below are 95% CI estimates from 100 different random samples ($n = 20$) drawn from a population with correlation of $\rho = 0$



Notice that 5 of 100 samples resulted in a 95% CI that failed to contain the true population-level correlation!

Confidence Intervals and the Normal model

Under a *Normal probability model*, confidence intervals can be found via:

$$\text{Point Estimate} \pm c * SE$$

- ▶ “c” is a percentile of the *Standard Normal distribution* that is used to calibrate the interval (sometimes called z^*)
 - ▶ For example, $c = 1.96$ is used for a 95% CI
- ▶ SE is found using the results of the Central Limit theorem
 - ▶ For example, for the sample proportion $\hat{p} = 12/30 = 0.4$,

$$SE = \sqrt{\frac{0.4*(1-0.4)}{30}} = 0.09$$

A 2021 study looked at the true-positive rate (sensitivity) of an Abbott Diagnostics rapid test for Covid-19 (one of the earliest such tests). Of the 84 cases with symptomatic Covid-19 who took the test, 38 had a “positive” result. Our goal is to estimate p , the overall sensitivity of this test in the target population (ie: all symptomatic Covid cases).

- 1) Verify that the conditions are met to use the CLT Normal approximation to construct a confidence interval estimate
- 2) Find the values of \hat{p} , its SE , and the value of c needed to construct a 99% CI estimate of p
- 3) Calculate and interpret the 99% CI

Practice (solution)

- 1) First, $\hat{p} = 38/84 = 0.452$. Then, $n\hat{p} = 84 * 0.452 = 38$ and $n(1 - \hat{p}) = 84 * (1 - 0.452) = 46$. Because both are larger than 10, the CLT Normal approximation is reasonable.
- 2) $\hat{p} = 38/84 = 0.452$, $SE = \sqrt{\frac{0.452(1-0.452)}{84}} = 0.054$, and $c = 2.576$ (this defines the middle 99% of a $N(0,1)$ curve)
- 3) The 99% CI is $0.452 \pm 2.576 * 0.054 = (0.313, 0.591)$. Our sample suggests, with 99% confidence, that the true sensitivity of the Abbott rapid test is somewhere between 31.3% and 59.1%

CLT Standard Errors

Without getting into the details, the CLT standard error formulas for commonly used descriptive statistics are shown below:

- ▶ Single proportion, \hat{p} , $SE = \sqrt{\frac{p(1-p)}{n}}$
 - ▶ works well when $n * p \geq 10$ and $n * (1 - p) \geq 10$
- ▶ Difference in proportions, $\hat{p}_1 - \hat{p}_2$, $SE = \sqrt{\frac{p_1(1-p_1)}{n_1} + \frac{p_2(1-p_2)}{n_2}}$
 - ▶ works well when $n_i * p_i \geq 10$ and $n_i * (1 - p_i) \geq 10$ for both groups
- ▶ Single mean, \bar{x} , $SE = \frac{\sigma}{\sqrt{n}}$
 - ▶ works well when $n \geq 30$
- ▶ Difference in means, $\bar{x}_1 - \bar{x}_2$, $SE = \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}$
 - ▶ works well when $n_i \geq 30$ for both groups
- ▶ Correlation coefficient, r , $SE = \sqrt{\frac{1-r^2}{n-2}}$
 - ▶ works well when $n \geq 30$

Note: using any of these formulas requires the replacement of unknown population parameters.

Practice #1

The previously mentioned study also examined a test produced by Siemens. Of 72 cases with symptomatic Covid-19 that took the Siemens test, 39 had a “positive” result. Recall that 38 of 84 symptomatic cases tested positive on the Abbott test. Our goal is estimate $p_1 - p_2$, the difference in sensitivity of these two tests (at the population level)

- 1) Let $\hat{p}_1 = 38/84 = 0.45$ be the sample proportion for the Abbott test, and $\hat{p}_2 = 39/72 = 0.54$ be the sample proportion for the Siemens test. Find the SE for the difference in proportions, $\hat{p}_1 - \hat{p}_2$.
- 2) Find and interpret the 95% CI estimate of $p_1 - p_2$

Practice #1 (solution)

$$1) SE = \sqrt{\frac{p_1(1-p_1)}{n_1} + \frac{p_2(1-p_2)}{n_2}} = \sqrt{\frac{0.45(1-0.45)}{84} + \frac{0.54(1-0.54)}{72}} = 0.08$$

- 2) Since $c = 1.96$, $\hat{p}_1 - \hat{p}_2 = 0.54 - 0.45 = -0.09$, and $SE = 0.08$, we calculate: $-0.09 \pm 1.96 * 0.08 = (-0.247, 0.067)$. This represents the plausible range of differences in sensitivity of these tests at the population level (estimated with 95% confidence). Because zero is included in this interval, it's plausible that the tests are actually no different.

Practice #2

A random sample of $n = 33$ adult women found a correlation coefficient of $r = 0.68$ between bodyweight and resting metabolic rate.

- 1) Can you be statistically confident that bodyweight is resting metabolic rate using only the *point estimate*?
- 2) Find and interpret the 99% CI estimate of ρ (the population-level correlation)

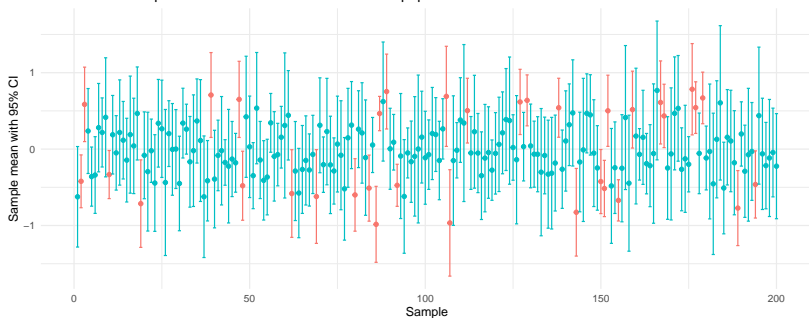
Practice #2 (solution)

- 1) No, even though a strong correlation was seen in the sample, the data consist of only 33 cases, so that correlation might be explained by sampling variability
- 2) $c = 2.576$ for 99% confidence and $SE = \sqrt{\frac{1-0.68^2}{30-2}} = 0.139$; so the 99% CI is $0.68 \pm 2.576 * 0.139$ or $(0.322, 1.00)$ - note that we'd want to cap the upper endpoint at 1.00 (as that's the maximum a correlation coefficient can be)

William Gosset and the t-distribution

- ▶ Using the CLT to calculate confidence intervals for quantitative outcomes (means, differences in means) requires an estimate of a second unknown population parameter (σ)
 - ▶ If we use the corresponding sample estimate, s , this is what happens:

200 different samples of $n = 8$ from a Standard Normal population



William Gosset and the t-distribution

- ▶ Clearly the prior 95% CI procedure is *invalid* - too many of the intervals didn't contain μ
- ▶ William Gosset, a chemist working for Guinness Brewing, became aware of this issue in the 1890s
 - ▶ His work evaluating the yields of different barley strains frequently involved small sample sizes

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 - ▶ His work evaluating the yields of different barley strains frequently involved small sample sizes
- ▶ In 1906, Gosset took a leave of absence from Guinness to study under Karl Pearson (developer of the correlation coefficient)
 - ▶ Gosset discovered the issue was due to using s interchangeably with σ

William Gosset and the t-distribution

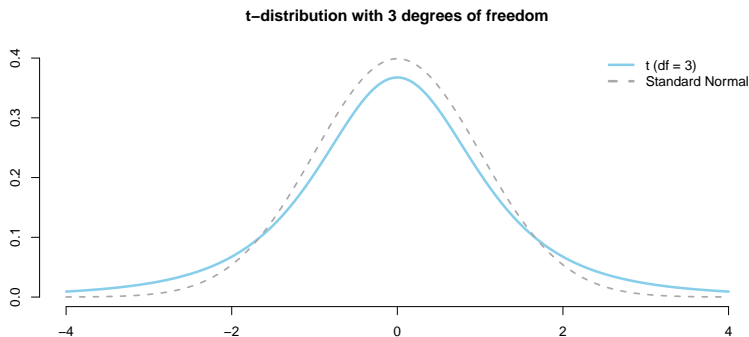
- ▶ Treating s as if it were a perfect estimate of σ results in a systematic underestimation of the total amount of variability involved in making the CI
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William Gosset and the t-distribution

- ▶ Treating s as if it were a perfect estimate of σ results in a systematic underestimation of the total amount of variability involved in making the CI
 - ▶ To account for the additional variability introduced by estimating σ using s , a modified distribution that's slightly more spread out than the Standard Normal curve must be used
- ▶ Typically the inventor of a new method gets to name it after themselves
 - ▶ However, Gosset was forced to publish his new distribution under the pseudonym “student” because Guinness didn't want it's competitors knowing they employed statisticians!
 - ▶ Student's t -distribution is now among the most widely used statistical results of all time

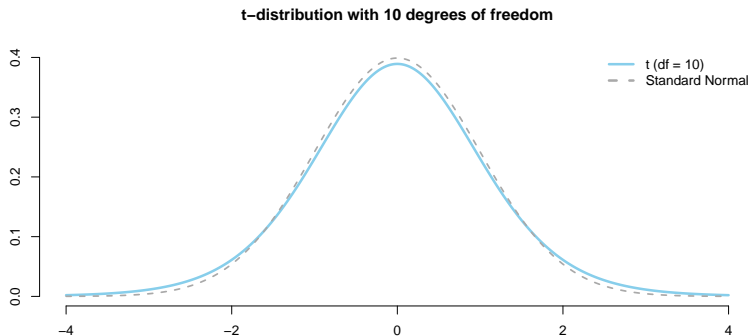
The t -distribution

The t -distribution accounts the additional uncertainty in small samples using a parameter known as *degrees of freedom*, or df :

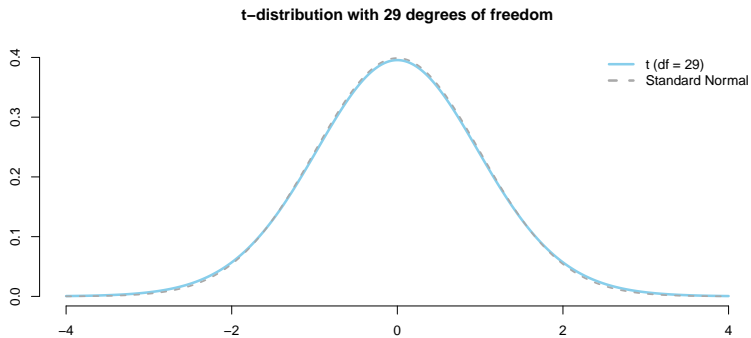


When estimating a single mean, $df = n - 1$

The t-distribution



The t-distribution



While waiting at an airport, a traveler notices 6 flights to similar a similar part of the country were delayed 6, 10, 13, 23, 45, 55 minutes. The mean delay in this sample was 25.33, with a sample standard deviation of $s = 20.2$. Assuming these data are a representative sample, answer the following:

- 1) How many degrees of freedom are involved when using the t -distribution to form a CI estimate? What is the value of c that should be used for 95% confidence?
- 2) What is the 95% CI estimate for the average delay of flights to the part of the country this traveler is heading?

Practice (solution)

- 1) Because $n = 6$, we'd use $df = n - 1 = 5$. For $df = 5$, $c = 2.571$ defines the middle 95% of the distribution.
- 2) Point Estimate \pm *MOE*, Point estimate = $\bar{x} = 25.33$, Margin of error = $c * SE = 2.571 * \frac{20.2}{\sqrt{6}}$
 - ▶ All together, 95% CI: $25.33 \pm 2.571 * \frac{20.2}{\sqrt{6}} = (4.1, 46.5)$
 - ▶ We are 95% confident the *average* delay is somewhere between 4.1 minutes and 46.5 minutes

Note: if we'd erroneously used a Normal model (instead of the t -distribution), we'd get an interval that is much narrower (9.2, 41.5), but this interval wouldn't have the confidence level we are advertising (ie: it wouldn't really be a 95% CI because it would miss too often)

When to use the t -distribution

- ▶ The t -distribution was designed for small, Normally distributed samples
 - ▶ However, it can also be reliably used on large samples, regardless of their shape

| | Sample data are approximately Normal | Sample data are non-Normal or skewed |
|--------------------------------------|--------------------------------------|--------------------------------------|
| Sample size is large ($n \geq 30$) | Use t -distribution | Use t -distribution |
| Sample size is small ($n < 30$) | Use t -distribution | <i>do not</i> use t -distribution |

- ▶ For small, non-Normal samples, more robust methods (such as bootstrapping) should be used instead

Conclusion

- ▶ Confidence intervals are preferred over point estimates because they address the question of statistical uncertainty that is inherent to random processes (such as data collection, random assignment, etc.)
- ▶ A confidence interval provides a plausible range of values for an unknown population parameter
 - ▶ The confidence level describes the success rate of the method used to calculate the interval if it were applied to many random samples
- ▶ To find a confidence interval you need three components:
 - ▶ A point estimate (calculated from the sample data)
 - ▶ The *SE* of that estimate (found using a CLT formula)
 - ▶ A calibration constant, c (found using a Normal distribution or a t -distribution)
- ▶ The t -distribution should be used for a single mean or a difference in means