#### Social Statistics

Introducing Hypothesis Testing

October 24, 2023

#### Introducing Significance Testing

- Confidence Intervals: Based on our sample estimate and confidence level, what is the range of possible values for population parameter?
- Shift to significance: How unusual how unlikely is our sample estimate?
- More specifically: What is the probability (.1, .05, .01, etc.) of getting a value at least as extreme as our estimate?
- Can we be confident at a certain level of probability that the difference between our estimate and another value is statistically significant, meaning not just from random chance?

### Parts of the Significance Test

- Assumptions: Random, Normal, Large, Increasing Validity
- Build two hypotheses covering the entire range
- Null Hypothesis: What we are testing against
  - → Null is usually 0 but can be any value
  - $\rightarrow$  H<sub>0</sub>:  $\mu$  = 0
- Alternative Hypothesis: What we estimate from our data
  - → Estimated mean is not 0
  - $\rightarrow$  H<sub>A</sub>:  $\mu \hat{} \neq 0$

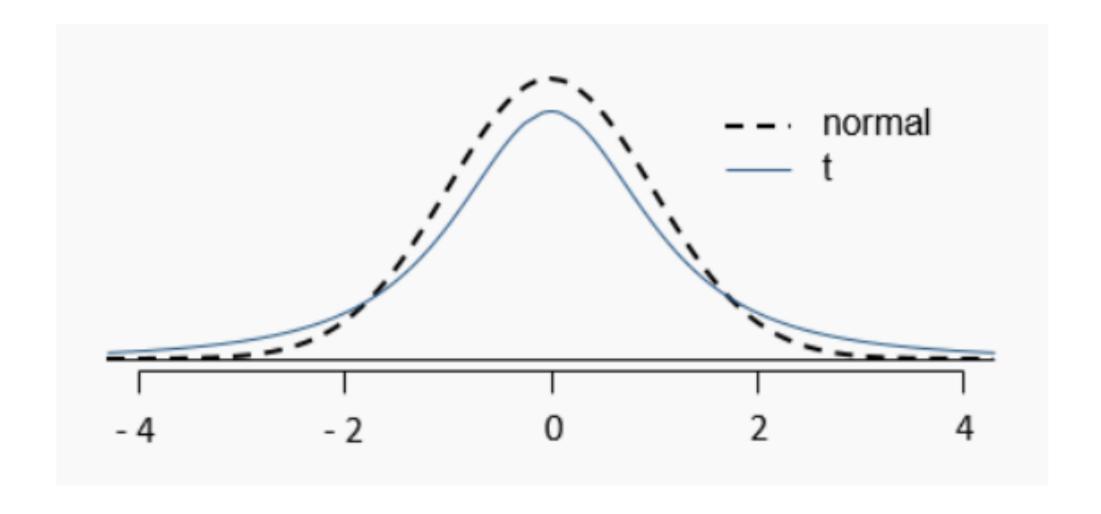
### Parts of the Significance Test

- Compute a test statistic for our estimate
- Previously we wanted distance from the mean in SDs

$$\rightarrow$$
 That formula:  $Z = \frac{x-\mu}{\sigma}$ 

- Now we want distance from the null hypothesis' value in SEs
  - $\rightarrow$  This formula:  $t = \frac{\mu_x \mu_0}{SD_x / \sqrt{n}}$
  - → t adjusts for degrees of freedom
  - → Denominator = standard error
  - $\rightarrow$  Fill in  $\mu_0$  from null hypothesis (0)

- When the population standard deviation is unknown (almost always), the uncertainty of the standard error estimate is addressed by using a new distribution: the t distribution.
- This distribution also has a bell shape, but its tails are thicker than the normal model's
- Therefore observations are more likely to fall beyond two SDs from the mean than under the normal distribution
- These extra thick tails are also helpful for resolving our problem with a less reliable estimate and a bigger standard error (if n is small)



- When the sample size is big enough, z and t are the same.
- When in doubt, use t and assert the degrees of freedom. It will always work.

```
1 qnorm(.975)
[1] 1.959964
1 qt(.975, df = 1000)
[1] 1.962339
1 qt(.975, df = 100)
[1] 1.983972
```

• Other norm() functions also work with t():

```
1 pt(1.962339, df = 1000)
[1] 0.975
1 1 - pt(1.983972, df = 100)
[1] 0.02499997
```

#### Back to the Significance Test

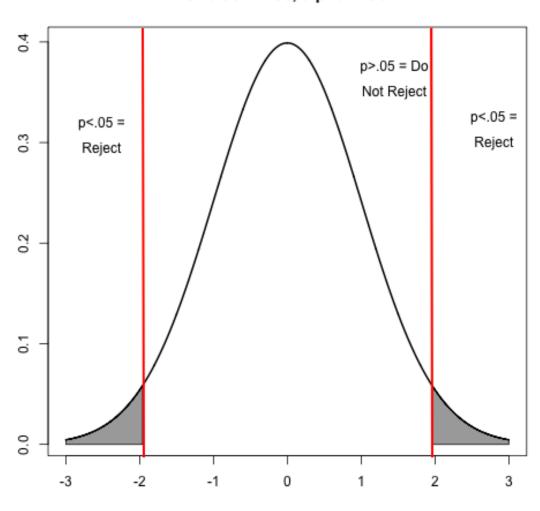
- Calculate the test statistic, and convert it to a probability (a p-value) using pt() if negative or 1 pt() if positive.
  - → Multiply this probability by two to get the probability for each tail.
- That p-value is probability of observing a value of the test statistic at least as extreme as the absolute value of our estimate
  - ightarrow A smaller p-value presents stronger evidence against  $H_0$ .
  - → Careful: not the probability that the null hypothesis is true

#### Parts of the Significance Test

- Choose significance level
  - $\rightarrow$  Alpha  $\alpha$  is significance level, or level of uncertainty, against which we want to evaluate the test statistic
  - → Also known as rejection region or critical region
- We will reject the null hypothesis if test statistic falls within this rejection/critical region
- But we will not accept the null hypothesis if test statistic falls outside this region
- Described in terms of probability of being *beyond* the range of values that are more likely, so we say .05 rather than .95

# Critical Region

t-value = 1.96, alpha = .05



- Imagine we want to test if the average number of children a GSS respondent has in 2018 is different from the average from 1984. Why might we expect people have different numbers of children in these two periods?
- Using gss\_week7.csv file on Canvas, let's test if the mean value of childs in 2018 differed from the 1984 mean of 2.2 (which we know from previous research).
- What was the mean for childs in 2018?

```
1 mean(gss_week7$childs[gss_week7$year==2018])
```

- Null Hypothesis = Actual mean is 2.2 (filled in from 1984)
- Alternative Hypothesis = Actual mean is not 2.2
  - → Note, if alternative were that the mean is a specific number (like 2), hypotheses would not cover entire range!
- Significance level = .05

• 
$$t = \frac{y-y_{H0}}{SD/\sqrt{n}}$$
  
•  $t = \frac{1.998-2.2}{0.05951692}$   
•  $t = -3.389708$ 

• In words: sample mean of 1.998 is 3.39 *standard errors* away from the null hypothesis value.

- What is the t-value for the critical region for our sample size and an alpha level of .05?
- There are 573 observations in the 2018 sample, so we use 572 for the degrees of freedom

```
1 qt(.975, df = 572)
```

[1] 1.96412

- For mean to be significant at .05 level, the test statistic needs to be at least +/- 1.96412 standard errors away. Our test statistic of -3.39 is more extreme than -1.96412, so we can reject the null hypothesis that the mean number of children in 2018 does not differ from 2.2.
- In other words, the average number of children in 2018 is significantly lower than the average number of children in 1984.
   There has been a statistically significant decline in number of children during that period.

 We would reach the same conclusion using p-values. The test statistic of 3.39 is negative, so we can use pt() instead of 1 pt():

```
1 pt(-3.39, df = 572)
[1] 0.0003735504
```

Multiply this value by two (or both sides of distribution)

```
1 2 * 0.0003735504
```

[1] 0.0007471008

 The p-value for our hypothesis test is less than .05, so we can reject the null hypothesis.

 We would also reach the same conclusion using confidence intervals.

```
1 idealchilds_ll <- 1.998255 - 1.96412*0.05951692
2 idealchilds_ul <- 1.998255 + 1.96412*0.0595169
3
4 idealchilds_ci <- c(idealchilds_ll, 1.998255, idealchilds_ul)
5 idealchilds_ci</pre>
```

[1] 1.881357 1.998255 2.115153

 The null hypothesis value of 2.2 falls outside our confidence interval, so we can reject the null hypothesis.

## Big Shortcut: t.test()

95 percent confidence interval:

1.881356 2.115153

sample estimates:

mean of x 1.998255

```
One Sample t-test
      gss_week7$childs[gss_week7$year == 2018]
data:
t = -3.3897 df = 572, p-value = 0.0007479
alternative hypothesis: true mean is not equal to 2.2
95 percent confidence interval:
1.881356 2.115153
sample estimates:
mean of x
 1.998255
```

Red box gives the test statistic. Blue box gives the probability of getting a value more extreme than the test statistic. Green box is the 95% confidence interval. Orange box gives the sample mean.

#### Exercise

- Let's look at 2018's mean ideal number of children (variable = chldidel)
- Using t.test(), can we reject the null hypothesis that the 2018 mean is the same as the 1990 mean of 2.55 at the .05 alpha level?

#### Significance Test for Mean - Exercise

- No, do not reject the null hypothesis:
  - → The test statistic is not more extreme than -1.96
  - → The p-value is greater than .05
  - → The confidence interval does include null hypothesis value of 2.55

## Other t.test() Options

• Can we reject the null hypothesis that the 2016 mean is the same as 2.5 at the .01 alpha level? We can change the default level of .05 to .01 using the conf.level option (which requires the confidence level, so .99 for the .01 alpha level).

```
1 # To Change Confidence Level:
 3 t.test(gss week7$chldidel[gss week7$year==2016],
          mu = 2.5, conf.level = .99)
    One Sample t-test
data: gss week7$chldidel[gss week7$year == 2016]
t = 1.5082, df = 720, p-value = 0.1319
alternative hypothesis: true mean is not equal to 2.5
99 percent confidence interval:
 2.466898 2.626029
sample estimates:
mean of x
 2.546463
```