# Confidence Intervals II

Nate Wells

Math 141, 3/17/21

# Outline

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- Implement the infer package to calculate confidence intervals
- Interpret confidence intervals

# Section 1

The infer package

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- What is the sample? What is the statistic?

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- The sample may contain many variables of interest
- We must first specify which variable(s) will be the focus of our investigation by designating a response variable
- To investigate the infection rate

```
covid %>%
   specify(response = Incubation)
```

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- The resulting data frame has a number of rows equal reps  $\times$  sample\_size

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generate( reps = 2000, type = "bootstrap") %>%
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```

• After applying calculate the resulting data frame consists of one bootstrap statistic for each replicate (saved to the variable stat)

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covid_stat<- covid %>%
   specify(response = Incubation) %>%
   calculate(stat = "mean")
covid_stat
```

• Note: we saved the value of this calculation as covid\_stat so we could use it later

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```
covid_boot<- covid %>%
  specify(response = Incubation) %>%
  generate( reps = 2000, type = "bootstrap") %>%
  calculate(stat = "mean")
head(covid_boot)
```

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```
percentile_ci<-covid_boot %>%
  get_ci(level = .95, type = "percentile")
percentile_ci
```

```
## # A tibble: 1 x 2
## lower_ci upper_ci
## <dbl> <dbl>
## 1 2.49 3.63
```

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• When using the percentile type, the first value printed is the lower and the second is the upper bound.

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#### Shade Confidence Intervals

• Once you've used get\_ci to obtain endpoints of the confidence interval, you can shade the sampling distribution with the confidence interval region.

```
covid_boot %>% visualize()+shade_ci(endpoints = percentile_ci)
```



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```
se_ci<-covid_boot %>%
  get_ci(level = .95, type = "se", point_estimate = covid_stat)
se_ci
## # A tibble: 1 x 2
## lower_ci upper_ci
## <dbl> <dbl>
```

## 1 2.46 3.60

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se_ci<-covid_boot %>%
get_ci(level = .95, type = "se", point_estimate = covid_stat)
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## # A tibble: 1 x 2
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• Note: for the se method, we also need to specify our point estimate (which is why we saved it as a variable before)

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```

• Why?

#### covid\_boot %>% visualize()



```
covid_boot %>% visualize() +
   shade_confidence_interval(endpoints = percentile_ci)+
   geom_vline(xintercept = 3.03, linetype = "dashed")
```



```
covid_boot %>% visualize() +
   shade_confidence_interval(endpoints = se_ci)+
   geom_vline(xintercept = 3.03, linetype = "dashed")
```



#### Percentile Method



Simulation-Based Bootstrap Distribution

#### SE Method



Simulation-Based Bootstrap Distribution

#### SE Method (with Percentile in blue)



#### Simulation-Based Bootstrap Distribution

# Section 2

# Interpreting Confidence Intervals

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- The problem?
  - We only have 1 sample, and we don't know if it belongs to the 95% of "good" samples, or the 5% of "bad" ones
- The consolation?
  - If I go through my life constructing 95% confidence intervals, I will be telling the truth about 95% of the time (I'll take that rate!)

# 100 Confidence Intervals



Nate Wells

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  - One sample (of 10000) had a sample mean of 4.9 and produced a confidence interval of (4.6, 5.2).

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  - At this point, the interval either does or does not contain the fixed (but unknown) parameter
  - One sample (of 10000) had a sample mean of 4.9 and produced a confidence interval of (4.6, 5.2).
  - Based on what you know about sleep patterns, do you think there is a 95% chance this interval contains the true parameter?

#### Precision

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How can we increase the precision of our confidence interval (i.e. decrease the margin of error)?

- Increase sample size.
  - The standard deviation of the sampling distribution decreases as sample size increases. More sample means are closer to the true parameter
- Decrease confidence level.
  - The margin of error is determined by the percentiles. A 95% confidence interval is formed by the 2.5th and 97.5th percentiles in the bootstrap distribution.
  - Decreasing confidence level brings the percentiles closer to the 50th percentile, decreasing the width of the interval.