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| **Performing a t-test for a Difference in Means**  t\_test(x = <NAME OF DATASET>,  response = <NAME OF NUMERICAL VARIABLE>,  explanatory = <NAME OF CATEGORICAL VARIABLE>,  conf\_int = TRUE,  conf\_level = 0.90,  alternative = "two-sided")  ***Note:*** If you want a 95% confidence interval, you change conf\_level to 0.95  ***Note:*** If you are doing a one-sided hypothesis test, you change alternative to either “greater” or “less” |
| **Obtaining 1000 Bootstrap Differences in Means**  bootstrap <- <NAME OF DATASET> %>%  specify(response = <NAME OF NUMERICAL VARIABLE>,  explanatory = <NAME OF CATEGORICAL VARIABLE>) %>%  generate(reps = 1000, type = "bootstrap") %>%  calculate(stat = "diff in means",  order = c(“<NAME OF FIRST GROUP>”, “<NAME OF FIRST GROUP>”)  )  ***Note:*** The quotation marks in the c() function are important! They need to be there even after you replace the values!  ***Note:*** Spelling and capitalization are important. You need to be 100% certain what the names of each group are when you specify them in the order = step! |
| **Plotting the Bootstrap Distribution**  visualize(data = bootstrap,  method = “simulation”)  ***Note:*** *This step* ***must*** *come after you have obtained the bootstrapped differences in means!* |

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| **Obtaining the Sample Difference in Means**  obs\_diff\_in\_means <- <NAME OF DATASET> %>%  specify(response = <NAME OF NUMERICAL VARIABLE>,  explanatory = <NAME OF CATEGORICAL VARIABLE>) %>%  calculate(stat = "diff in means",  order = c(“<NAME OF FIRST GROUP>”, “<NAME OF FIRST GROUP>”)  )  ***Note:*** *This step* ***must*** *be done* ***before*** *you find your confidence interval and before finding your p-value!* |
| **Obtaining a Confidence Interval from a Bootstrap Distribution**  get\_confidence\_interval(x = bootstrap,  level = 0.95,  type = “percentile”,  point\_estimate = obs\_mean)  ***Note:*** *This step* ***must*** *come after you have obtained the bootstrapped differences in means* ***and*** *the observed difference in means!*  ***Note:*** If you want a 90% confidence interval, you change level to 0.90 |
| **Obtaining 1000 Permuted Differences in Means – Assuming the Null Hypothesis is True**  null\_dist <- <NAME OF DATASET> %>%  specify(response = <NAME OF NUMERICAL VARIABLE>,  explanatory = <NAME OF CATEGORICAL VARIABLE>) %>%  generate(reps = 1000, type = "permute") %>%  calculate(stat = "diff in means",  order = c(“<NAME OF FIRST GROUP>”, “<NAME OF FIRST GROUP>”)  ) |

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| **Plotting the Simulated Null Distribution**  visualize(data = null\_dist,  method = “simulation”)  ***Note:*** *This step* ***must*** *come after you have obtained the permuted differences in means!* |
| **Obtaining a p-value from a Null Distribution**  get\_pvalue(x = null\_dist,  obs\_stat = obs\_diff\_in\_means,  direction = “two-sided”)  ***Note:*** *This step* ***must*** *come after you have obtained the bootstrapped differences in means* ***and*** *the observed difference in means!*  ***Note:*** If you are doing a one-sided hypothesis test, you change alternative to either “greater” or “less” |
| **Faceted Histograms**  ggplot(data = <NAME OF DATASET>,  mapping = aes(x = <NAME OF NUMERICAL VARIABLE>)) +  geom\_histogram(binwidth = <WIDTH OF BINS>) +  facet\_wrap(~<NAME OF CATEGORICAL VARIABLE>) +  labs(x = "<TITLE FOR THE X-AXIS>")  ***Note:*** A histogram **must** have the variable on the x-axis! |
| **Side-by-Side Boxplots**  ggplot(data = <NAME OF DATASET>,  mapping = aes(x = <NAME OF VARIABLE>, y = <NAME OF VARIABLE>)  ) +  geom\_boxplot() +  labs(x = "<TITLE FOR THE X-AXIS>",  y = "<TITLE FOR THE Y-AXIS>")  ***Note:*** For **horizontally stacked** boxplots, the categorical variable should be on the **x-axis.** For **vertically stacked** boxplots, the categorical variable should be on the **y-axis**. |

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| **Calculating Summary Statistics for One Numerical Variable and One Categorical Variable**  favstats(<NAME OF NUMERICAL VARIABLE> ~ <NAME OF CATEGORICAL VARIABLE>,  data = <NAME OF DATASET>)  ***Note:*** The ~ **must** be included! |
| **Obtaining an ANOVA Table**  aov(<NAME OF NUMERICAL VARIABLE> ~ <NAME OF CATEGORICAL VARIABLE>,  data = <NAME OF DATASE>) |

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| **Plotting the Bootstrap Distribution**  visualize(data = bootstrap,  method = “simulation”)  ***Note:*** *This is* ***the same*** *as plotting the bootstrap for one mean!* |
| **Obtaining the Sample Slope**  obs\_mean <- <NAME OF DATASET> %>%  specify(response = <NAME OF Y-VARIABLE>,  explanatory = <NAME OF X-VARIABLE>) %>%  calculate(stat = "mean")  ***Note:*** *This step* ***must*** *be done* ***before*** *you find your confidence interval!* |
| **Obtaining a Confidence Interval from a Bootstrap Distribution**  get\_confidence\_interval(x = bootstrap,  level = 0.95,  type = “percentile”,  point\_estimate = obs\_mean)  ***Note:*** *This is* ***the same*** *as how you found a confidence interval for one mean!* |