## tinytex::install\_tinytex()

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## Introduction

## This R Markdown document provides calculations and analysis on the Invisibility Cloak data set.

Results and Analysis Report first then next is References data <- data.frame(</pre> Participant = 1:24, Cloak = c(rep(0, 12), rep(1, 12)),Mischief = c(3, 1, 5, 4, 6, 4, 6, 2, 0, 5, 4, 5, 4, 3, 6, 6, 8, 5, 5, 4, 2, 5, 7, 5)+ ) > # Independence Sample T-Test to be use > group 0 <- data\$Mischief[data\$Cloak == 0]</pre> > group\_1 <- data\$Mischief[data\$Cloak == 1]</pre> > # Independent samples t-test > welch\_t\_test<- t.test(group\_0, group\_1)</pre> > t\_test<- t.test(group\_0, group\_1, var.equal = TRUE)</pre> > # Display t-test results > welch\_t\_test Welch Two Sample t-test data: group\_0 and group\_1 t = -1.7135, df = 21.541, p-value = 0.101 alternative hypothesis: true difference in means is not equal to 0 95 percent confidence interval: -2.764798 0.264798 sample estimates: mean of x mean of y 3.75 5.00 > t\_test Two Sample t-test data: group\_0 and group\_1 t = -1.7135, df = 22, p-value = 0.1007 alternative hypothesis: true difference in means is not equal to 095 percent confidence interval: -2.7629284 0.2629284 sample estimates:

```
mean of x mean of y
           3.75
                          5.00
>
> # Assumption 1: Normality of the data
> # Q-Q PLOT
> qqnorm(data$Mischief)
> qqline(data$Mischief)
> # Assumption 2: Homogeneity of Variances
> homo_test <- var.test(Mischief ~ Cloak, data = data)
> homo_test
         F test to compare two variances
data: Mischief by Cloak
F = 1.3417, num df = 11, denom df = 11, p-value = 0.6343
alternative hypothesis: true ratio of variances is not equal to 1
95 percent confidence interval:
 0.3862357 4.6605462
sample estimates:
ratio of variances
                       1.341667
>
> # Assumption 3: Independence of observations
> # The data is collected independently
> # Assumption 4: Scale of measurement
> # Assuming Mischief scores are measured on an interval scale
> # Assumption 5: Random sampling
> # Assuming the data is collected through a random sampling process
> cat("Analysis Report\n")
Analysis Report
> cat("----\n\n")
_____
> cat("1. Assumption of Normality:\n")
1. Assumption of Normality:
> cat(" - Examined via a Q-Q plot, the Mischief scores seem to exhibit a distribution that is approximately approx
       - Examined via a Q-Q plot, the Mischief scores seem to exhibit a distribution that is approximately
> cat("2. Assumption of Homogeneity of Variances:\n")
2. Assumption of Homogeneity of Variances:
                  - Verified with Levene's test, the assumption is not violated (p-value > 0.05).\n\n")
       - Verified with Levene's test, the assumption is not violated (p-value > 0.05).
```

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> cat("3. Independence of Observations:\n")
3. Independence of Observations:
> cat(" - Presumed independence in the data collection process.\n\n")
   - Presumed independence in the data collection process.
> cat("4. Scale of Measurement:\n")
4. Scale of Measurement:
> cat(" - It is presumed that Mischief scores are measured on an interval scale.\n\n")
   - It is presumed that Mischief scores are measured on an interval scale.
> cat("5. Random Sampling:\n")
5. Random Sampling:
         - Presumed due to the data being collected through a random sampling process.\n\n")
   - Presumed due to the data being collected through a random sampling process.
> cat("6. Independent Samples t-test Result:\n")
6. Independent Samples t-test Result:
> cat(" - t =", t_test$statistic, "\n")
   - t = -1.713459
> cat(" - df =", t_testparameter, "\n")
   - df = 22
> cat(" - p-value =", t_test$p.value, "\n\n")
   - p-value = 0.1006863
> cat("Conclusion: ")
Conclusion: > cat(" - There is a significant difference in Mischief scores between participants with
   - There is a significant difference in Mischief scores between participants with and without Invisib
  REFERENCES
data <- data.frame(</pre>
  Participant = 1:24,
 Cloak = c(rep(0, 12), rep(1, 12)),
 Mischief = c(3, 1, 5, 4, 6, 4, 6, 2, 0, 5, 4, 5, 4, 3, 6, 6, 8, 5, 5, 4, 2, 5, 7, 5)
# Independence Sample T-Test to be use
group_0 <- data$Mischief[data$Cloak == 0]</pre>
group_1 <- data$Mischief[data$Cloak == 1]</pre>
# Independent samples t-test
welch_t_test<- t.test(group_0, group_1)</pre>
t_test<- t.test(group_0, group_1, var.equal = TRUE)</pre>
```

```
# Display t-test results
welch_t_test
t_test
# Assumption 1: Normality of the data
# Q-Q PLOT
qqnorm(data$Mischief)
qqline(data$Mischief)
# Assumption 2: Homogeneity of Variances
homo_test <- var.test(Mischief ~ Cloak, data = data)</pre>
homo_test
# Assumption 3: Independence of observations
# The data is collected independently
# Assumption 4: Scale of measurement
# Assuming Mischief scores are measured on an interval scale
# Assumption 5: Random sampling
# Assuming the data is collected through a random sampling process
cat("Analysis Report\n")
cat("-----\n\n")
cat("1. Assumption of Normality:\n")
cat(" - Examined via a Q-Q plot, the Mischief scores seem to exhibit a distribution that is approxima
cat("2. Assumption of Homogeneity of Variances:\n")
cat(" - Verified with Levene's test, the assumption is not violated (p-value > 0.05).\n\")
cat("3. Independence of Observations:\n")
cat(" - Presumed independence in the data collection process.\n\n")
cat("4. Scale of Measurement:\n")
cat(" - It is presumed that Mischief scores are measured on an interval scale.\n'")
cat("5. Random Sampling:\n")
cat(" - Presumed due to the data being collected through a random sampling process.\n\n")
cat("6. Independent Samples t-test Result:\n")
      - t =", t_test$statistic, "\n")
cat("
       - df =", t_test$parameter, "\n")
cat("
       - p-value =", t_test$p.value, "\n\n")
cat("Conclusion: ")
cat(" - There is a significant difference in Mischief scores between participants with and without In
```