STAT3355(HW-7)

Tulasi Janjanam

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Problem 1

```
# Step 1: Define the data
fill_data <- c(15.997, 16.005, 15.981, 15.954, 15.986,
               16.021, 15.985, 16.001, 16.018, 16.056)
# Step 2: Specify the hypothesized mean and significance level
mu_0 <- 16.00 # Hypothesized mean
alpha <- 0.05 # Significance level
# Step 3: Perform calculations
sample_mean <- mean(fill_data) # Sample mean</pre>
sample_sd <- sd(fill_data)  # Sample standard deviation
n <- length(fill_data)
                                 # Sample size
# Test statistic
t_stat <- (sample_mean - mu_0) / (sample_sd / sqrt(n))
# Degrees of freedom
df <- n - 1
# p-value for the one-sided test
p_value <- pt(t_stat, df)</pre>
# Critical value for a one-sided t-test
critical_value <- qt(alpha, df)</pre>
# Display results
cat("Sample Mean:", sample_mean, "\n")
## Sample Mean: 16.0004
cat("Sample Standard Deviation:", sample_sd, "\n")
## Sample Standard Deviation: 0.02755278
cat("Test Statistic (t):", t_stat, "\n")
## Test Statistic (t): 0.04590866
```

```
cat("Degrees of Freedom:", df, "\n")

## Degrees of Freedom: 9

cat("Critical Value (t):", critical_value, "\n")

## Critical Value (t): -1.833113

cat("p-value:", p_value, "\n")

## p-value: 0.5178072

# Step 4: Decision
if (p_value <= alpha) {
   cat("Decision: Reject the null hypothesis. The mean fill is less than 16.00 ounces.\n")
} else {
   cat("Decision: Fail to reject the null hypothesis. Insufficient evidence that the mean fill is less than 16.00 ounces.\n")
}</pre>
```

Decision: Fail to reject the null hypothesis. Insufficient evidence that the mean fill is less than

Problem 2

```
# Step 1: Given data
x_bar <- 8.412 # Sample mean
               # Sample standard deviation
s <- 1.512
n <- 60
              # Sample size
mu_0 <- 8.2 # Hypothesized mean
alpha <- 0.05 # Significance level
# Step 2: Calculate the test statistic (t)
t_stat <- (x_bar - mu_0) / (s / sqrt(n))
# Degrees of freedom
df <- n - 1
# Step 3: Calculate the p-value for the one-sided t-test
p_value <- 1 - pt(t_stat, df)</pre>
# Step 4: Critical value for a one-sided t-test
critical_value <- qt(1 - alpha, df)</pre>
# Step 5: Display results
cat("Test Statistic (t):", t_stat, "\n")
```

Test Statistic (t): 1.086075

```
cat("Critical Value (t):", critical_value, "\n")

## Critical Value (t): 1.671093

cat("p-value:", p_value, "\n")

## p-value: 0.1409317

# Step 6: Decision
if (p_value <= alpha) {
    cat("Decision: Reject the null hypothesis. There is sufficient evidence to suggest that the mean engage eat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that the eat the period of the suggest that the mean engage eat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that the mean engage eat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that the mean engage eat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that the mean engage eat that the mean engage eat the null hypothesis. There is insufficient evidence to suggest that the mean engage eat the null hypothesis.</pre>
```

Problem 3

```
# Given data
x <- 130 # Number of stressed students
n <- 200 # Sample size
p0 <- 0.70 # Hypothesized population proportion
# Sample proportion
p_hat <- x / n
# Step 1: Calculate the z-test statistic
z_{stat} \leftarrow (p_{nat} - p0) / sqrt((p0 * (1 - p0)) / n)
# Step 2: Calculate the p-value for a two-tailed test
p_value <- 2 * (1 - pnorm(abs(z_stat))) # Two-tailed test</pre>
# Step 3: Display results
cat("Test Statistic (z):", z_stat, "\n")
## Test Statistic (z): -1.543033
cat("p-value:", p_value, "\n")
## p-value: 0.1228226
# Step 4: Decision
alpha <- 0.05
if (p_value <= alpha) {</pre>
  cat("Decision: Reject the null hypothesis. There is evidence to suggest the proportion of stressed st
  cat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest the prop
```

Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest the proporti

Problem 4

```
# Given data
x1 <- 2.00 # Mean for ages 18-50
s1 <- 0.812 # Standard deviation for ages 18-50
n1 <- 350 # Sample size for ages 18-50
x2 < -1.85 # Mean for ages > 50
s2 <- 0.837 # Standard deviation for ages > 50
n2 <- 150
           # Sample size for ages > 50
# Step 1: Calculate the t-statistic
t_stat \leftarrow (x1 - x2) / sqrt((s1^2 / n1) + (s2^2 / n2))
# Step 2: Calculate the degrees of freedom using Welch-Satterthwaite equation
df \leftarrow ((s1^2 / n1 + s2^2 / n2)^2) / ((s1^2 / n1)^2 / (n1 - 1) + (s2^2 / n2)^2 / (n2 - 1))
# Step 3: Calculate the p-value for a one-sided test
p_value <- 1 - pt(t_stat, df)</pre>
# Step 4: Display results
cat("Test Statistic (t):", t_stat, "\n")
## Test Statistic (t): 1.852798
cat("Degrees of Freedom:", df, "\n")
## Degrees of Freedom: 274.3818
cat("p-value:", p_value, "\n")
## p-value: 0.03249249
# Step 5: Decision
alpha <- 0.05
if (p_value <= alpha) {</pre>
  cat("Decision: Reject the null hypothesis. There is evidence to suggest that people over 50 years old
  cat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that peo
```

Decision: Reject the null hypothesis. There is evidence to suggest that people over 50 years old use

Problem 5

```
# Given data

x1 <- 22 # Number of male students who play video games

n1 <- 30 # Sample size for males
```

```
x2 <- 24 # Number of female students who play video games
n2 <- 40 # Sample size for females
# Step 1: Calculate sample proportions
p1_hat <- x1 / n1
p2_hat <- x2 / n2
# Step 2: Calculate the pooled proportion
p_hat \leftarrow (x1 + x2) / (n1 + n2)
# Step 3: Calculate the z-statistic
z_stat <- (p1_hat - p2_hat) / sqrt(p_hat * (1 - p_hat) * (1/n1 + 1/n2))
# Step 4: Calculate the p-value for a one-sided test
p_value <- 1 - pnorm(z_stat)</pre>
# Step 5: Display results
cat("Test Statistic (z):", z_stat, "\n")
## Test Statistic (z): 1.163038
cat("p-value:", p_value, "\n")
## p-value: 0.1224071
# Step 6: Decision
alpha <- 0.05
if (p_value <= alpha) {</pre>
 cat("Decision: Reject the null hypothesis. There is evidence to suggest that more male students play
  cat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that mor
```

Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that more ma

Problem 6

```
# Given data
n <- 75  # Sample size
x <- 30  # Number of college graduates without insurance
p0 <- 0.281  # Nationwide proportion of adults without insurance

# Step 1: Calculate the sample proportion
p_hat <- x / n

# Step 2: Calculate the z-statistic
z_stat <- (p_hat - p0) / sqrt(p0 * (1 - p0) / n)

# Step 3: Calculate the p-value for a two-sided test</pre>
```

```
p_value <- 2 * (1 - pnorm(abs(z_stat))) # Multiply by 2 for two-tailed test

# Step 4: Display results
cat("Test Statistic (z):", z_stat, "\n")

## Test Statistic (z): 2.292767

cat("p-value:", p_value, "\n")

## p-value: 0.0218614

# Step 5: Decision
alpha <- 0.05
if (p_value <= alpha) {
    cat("Decision: Reject the null hypothesis. There is evidence that the proportion of college graduates }
} else {
    cat("Decision: Fail to reject the null hypothesis. There is insufficient evidence to suggest that the }
}</pre>
```

Decision: Reject the null hypothesis. There is evidence that the proportion of college graduates with

Problem 7

```
# Given data
n1 <- 150 # Number of iPhones sold
x1 <- 14 # Number of iPhones returned
n2 <- 125 # Number of Samsung phones sold
x2 <- 15 # Number of Samsung phones returned
# Step 1: Calculate the sample proportions
p1_hat <- x1 / n1
p2_hat <- x2 / n2
# Step 2: Calculate the pooled proportion
p_{pool} \leftarrow (x1 + x2) / (n1 + n2)
# Step 3: Calculate the z-statistic
z_{stat} \leftarrow (p_{hat} - p_{hat}) / sqrt(p_{pool} * (1 - p_{pool}) * (1 / n1 + 1 / n2))
# Step 4: Calculate the p-value for a left-tailed test
p_value <- pnorm(z_stat)</pre>
# Step 5: Display results
cat("Test Statistic (z):", z_stat, "\n")
```

Test Statistic (z): -0.7169175

```
cat("p-value:", p_value, "\n")
## p-value: 0.2367125
# Step 6: Decision
alpha <- 0.05
if (p_value <= alpha) {</pre>
  cat("Decision: Reject the null hypothesis. There is evidence that Apple has a smaller chance of being
  cat("Decision: Fail to reject the null hypothesis. There is insufficient evidence that Apple has a sm
## Decision: Fail to reject the null hypothesis. There is insufficient evidence that Apple has a smalle
Problem 8
# Load the UsingR package and dataset
library(UsingR)
## Loading required package: MASS
## Loading required package: HistData
## Loading required package: Hmisc
##
## Attaching package: 'Hmisc'
## The following objects are masked from 'package:base':
##
##
       format.pval, units
data(babies)
# Perform a two-sample t-test to compare mother's age and father's age
t_test_result <- t.test(babies$age, babies$dage)</pre>
# Display the test result
t_test_result
##
## Welch Two Sample t-test
## data: babies$age and babies$dage
## t = -11.067, df = 2301.5, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -3.962068 -2.769323
## sample estimates:
## mean of x mean of y
## 27.37136 30.73706
```

```
# Extract p-value and make a decision
p_value <- t_test_result$p.value
alpha <- 0.05

# Print the conclusion
if (p_value <= alpha) {
   cat("Decision: Reject the null hypothesis. The mean age of mothers is significantly different from th
} else {
   cat("Decision: Fail to reject the null hypothesis. There is no significant difference between the mean
}</pre>
```

Decision: Reject the null hypothesis. The mean age of mothers is significantly different from the me