# STA304 The Outliers Code Appendix\*

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## Introduction

This appendex contains all code used in our technical report. It is ordered how it appears in the technical report, accompanied by any relevant comments.

```
library(tidyverse)
library(treemapify)
library(gridExtra)
library(patchwork)
# Split into Single Items
majTab=table(gsub("\\s+", "", unlist(strsplit(data\major,","))))
# Convert to Data Frame
majDF=as.data.frame(majTab)
# Refactor for Correct Plot Order
majDF$Var1=fct_relevel(majDF$Var1, c("STA", "MAT", "CSC", "ECO", "LSC", "NLA"))
majLabs=c("Applied Statistics", "Mathematics", "Computer Science", "Economics",
          "Life Sciences", "Not Listed")
statusCounts=as.data.frame(table(data$studentStatus))
statusLabs=c("Domestic", "International")
distanceDF=as.data.frame(table(data$campusDistance))
distanceLabs=c("On Campus", "0-5km", "6-10km", "11-15km", "16-20km", "21+km")
status <- ggplot(data=statusCounts, aes(area=Freq, fill=Var1, label=paste(Freq))) +
  geom treemap(color="black", size=1.0) +
  labs(title="Student Status", x="n = 63", caption = "Figure 1") +
```

<sup>\*</sup>Data and Report Available at: https://github.com/vanshikav2/Extra\_Curricular\_Activities\_Research

```
theme minimal() +
  scale_fill_brewer(name="Options", labels=statusLabs, palette = "Set2") +
  geom treemap text(colour="black", place="center")
dist <- ggplot(data=distanceDF, aes(area=Freq, fill=Var1, label=paste(Freq))) +
  geom_treemap(color="black", size=1.0) +
  labs(title = "Distance From Campus", x="n = 63", caption = "Figure 2") +
  theme_minimal() +
  scale_fill_brewer(name="Distance", labels=distanceLabs, palette = "Set2") +
  geom_treemap_text(color="black", place="center") +
  theme(plot.title = element_text(hjust = 0.5))
maj <- ggplot(data=majDF, aes(x=Var1, y=Freq, fill=Var1)) +</pre>
  geom_bar(stat="identity", color="Black") +
  labs(x="Majors", y="Count", title = "Sample Majors Breakdown",
       caption = "Figure 3") +
  theme light() +
  scale_fill_brewer(name="Options:", labels=majLabs, palette="Set2") +
  geom_text(aes(label=Freq), vjust=-.3, color="black") +
  theme(plot.title = element_text(hjust = 0.5),
        axis.title.y = element_text(size=12, angle=90),
        axis.text.x = element text(size= 6))
(status + dist + maj) + plot_layout(widths = c(1,1,1))
```

### Research Question 1

```
library(tidyverse)
library(dplyr)
library(ggplot2)
library(car)
library(knitr)
library(kableExtra)
data <- read_csv("STA304_TheOutliers_CleanedData.csv")

#Deleting the studentID column and replacing NA values
data[is.na(data)] <- "NA"
data <- data %>% select(-studentID, -lectureSection)
```

```
#Split `activityType` and `major` into individual categories
data <- data %>%
  separate_rows(activityType, sep = ",") %>%
 separate_rows(major, sep = ",")
data <- data %>% mutate(activityType = trimws(data$activityType),
                        major = trimws(data$major))
# Descriptive Analysis (What is the most preferred Extra Curricular Activity)
data <- data %>% mutate(activityType = as.factor(activityType))
activity_counts <- data %>%
 count(activityType, name = "Frequency") %>%
 arrange(desc(Frequency))
# Plot distribution of activity types
activity_labels <- c(
 "AAS" = "Athletics and Sports",
 "NEC" = "No Extracurricular Activities",
 "CLU" = "Clubs",
 "LSG" = "Leadership/Student Governments, Councils & Unions",
 "ACS" = "Academic Societies"
)
#Graph Showing the Activity Counts
ggplot(data = activity_counts, aes(x = activityType, y = Frequency,
                                  fill = activityType)) +
 geom_bar(stat = "identity", color = "black") +
 labs(
   x = "Activity Type",
   y = "Frequency",
   title = "Distribution of Extracurricular Activity Types"
  ) +
 theme_light() +
 scale_fill_brewer(
   name = "Activity Options:",
   labels = activity_labels,
   palette = "Set2"
 ) +
  geom_text(aes(label = Frequency), vjust = -0.3, size = 4)
```

### Testing for the Significance of Majors on Activity Preference using the ANOVA Test

```
# 2. ANOVA for Major
#Convert ActivityType to NumericValues
data <- data %>% mutate(activityTypeNumeric = as.numeric(factor(activityType)))
anova_major <- aov(activityTypeNumeric ~ major, data = data)</pre>
#summary(anova_major)
#Q-Q plot of Residuals
residuals <- residuals(anova_major)</pre>
qqnorm(residuals, col = "purple", main = "Q-Q Plot of Residuals for ANOVA Model
       for Major")
qqline(residuals, col = "black", lwd = 1)
library(car)
library(knitr)
##Testing for Homogenity of Variances
data <- data %>%
  mutate(major = as.factor(major))
levene_test_major <- leveneTest(activityTypeNumeric ~ major, data = data)</pre>
levene_df <- as.data.frame(levene_test_major)</pre>
levene_df[is.na(levene_df)] <- ""</pre>
kable(levene_df, caption = "Levene Test on The Activity Preference by Major")
anova_summary <- summary(anova_major)[[1]]</pre>
anova_df <- data.frame(anova_summary)</pre>
anova df[is.na(anova df)] <- ""
kable(anova_df, caption = "Anova Test on The Activity Preference by Major")
activity_lbls <- c(</pre>
  "1" = "ACS",
  "2" = "AAS",
  "3" = "CLU",
  "4" = "LSG",
```

```
"5" = "NEC"
```

#### Testing for the Significance of Gender on Activity Preference using the T-Test

```
##GenderIdentity
gender_data_F <- data %% filter(genderIdentity == "F") %>% pull(activityTypeNumeric)
gender_data_M <- data %% filter(genderIdentity == "M") %>% pull(activityTypeNumeric)
data <- data %>%
  mutate(genderIdentity = as.factor(genderIdentity))
shapiro_F <- shapiro.test(gender_data_F)</pre>
shapiro_M <- shapiro.test(gender_data_M)</pre>
shapiro_M df <- as.data.frame(t(c(shapiro_M$statistic, shapiro_M$p.value)))</pre>
colnames(shapiro_M_df) <- c("W statistic", "p-value")</pre>
shapiro_F_df <- as.data.frame(t(c(shapiro_F$statistic, shapiro_F$p.value)))</pre>
colnames(shapiro_F_df) <- c("W statistic", "p-value")</pre>
shapiro_results <- data.frame(</pre>
  Gender = c("Males", "Females"),
  `W Statistic` = c(shapiro_F_df$`W statistic`, shapiro_M_df$`W statistic`),
  `p-value` = c(shapiro_F_df$`p-value`, shapiro_M_df$`p-value`)
kable(shapiro_results, col.names = c("Gender", "W Statistic", "p-value"),
      caption = "Shapiro-Wilk Test Results for Activity Preferences
      by Gender Groups")
levene_gender <- leveneTest(activityTypeNumeric ~ genderIdentity, data = data)</pre>
levene_gender_df <- as.data.frame(levene_gender)</pre>
levene_gender_df[is.na(levene_gender_df)] <- ""</pre>
kable(levene_gender_df, caption = "Levene Test for Equality of Variances in
      Activity Preferences by Gender Identity")
```

## Testing for the Significance of Student Status on Activity Preference using the T-Test

```
##StudentStatus
status_data_D <- data %>% filter(studentStatus == "D") %>% pull(activityTypeNumeric)
status data I <- data %>% filter(studentStatus == "I") %>% pull(activityTypeNumeric)
shapiro_D <- shapiro.test(status_data_D)</pre>
shapiro_I <- shapiro.test(status_data_I)</pre>
shapiro_results <- data.frame(</pre>
  StudentStatus = c("D", "I"),
  W_Statistic = c(shapiro_D$statistic, shapiro_I$statistic),
  P_Value = c(shapiro_D$p.value, shapiro_I$p.value)
kable(shapiro_results, caption = "Shapiro Test on studentStatus")
data <- data %>%
  mutate(studentStatus = as.factor(studentStatus))
levene_status <- leveneTest(activityTypeNumeric ~ studentStatus, data = data)</pre>
levene_status_df <- as.data.frame(levene_status)</pre>
levene_status_df[is.na(levene_status_df)] <- ""</pre>
kable(levene_status_df, caption = "Levene Test on studentStatus")
wilcox_test <- wilcox.test(activityTypeNumeric ~ studentStatus, data = data)</pre>
wilcox_test_df <- data.frame(</pre>
  Statistic = wilcox_test$statistic,
  P_Value = wilcox_test$p.value)
```

```
kable(
  wilcox_test_df,
  col.names = c("W Statistic", "P-Value"),
  caption = "Wilcoxon Rank-Sum Test Results for Activity Preferences by
  Student Status"
)
# Boxplot for Major influence
plot1 <- ggplot(data, aes(x = major, y = activityTypeNumeric, fill = major)) +
  geom boxplot() +
  labs(title = "Boxplot of Activity Preferences by Major",
       x = "Major",
       y = "Activity Type") +
  scale_y_continuous(
    breaks = 1:5,
    labels = activity_lbls
  )+
  theme_minimal() +
  theme(axis.text.y = element_text(size = 10),
    axis.title.y = element_text(size = 12),
    plot.margin = margin(10, 10, 10, 20)
  )
# Boxplot for Gender Influence
plot2 <- ggplot(data, aes(x = genderIdentity, y = activityTypeNumeric,</pre>
                          fill = genderIdentity)) +
  geom boxplot() +
  labs(title = "Boxplot of Activity Preferences by Gender",
       x = "Gender Identity",
       y = "Activity Type (Numeric)") +
  scale_y_continuous(
    breaks = 1:5,
    labels = activity_lbls)+
  theme_minimal() +
  theme(axis.text.y = element_text(size = 10),
    axis.title.y = element_text(size = 12),
   plot.margin = margin(10, 10, 10, 20)
  )
# Boxplot for StudentStatus
plot3 <- ggplot(data, aes(x = studentStatus, y = activityTypeNumeric,</pre>
                          fill = studentStatus)) +
  geom_boxplot() +
```

```
labs(title = "Boxplot of Activity Preferences by Student Status",
    x = "Student Status",
    y = "Activity Type (Numeric)") +
scale_y_continuous(
    breaks = 1:5,
    labels = activity_lbls)+
theme_minimal() +
theme(axis.text.y = element_text(size = 10),
    axis.title.y = element_text(size = 12),
    plot.margin = margin(10, 10, 10, 20)
)
grid.arrange(plot1, plot2, plot3, nrow = 2)
```

#### Research Question 2

### **Campus Distance and Activity Count**

```
# Create summaries for campusDistance and activityCount
campus <- summary(data$campusDistance)</pre>
activity <- summary(data$activityCount)</pre>
# Combine the summaries into a data frame
summary_t <- data.frame(</pre>
  Statistic = c("Min", "1st Quartile", "Median", "Mean", "3rd Quartile", "Max"),
  CampusDistance = as.numeric(campus),
 ActivityCount = as.numeric(activity)
library(knitr)
kable(summary_t, format = "pipe", col.names = c("Statistic", "CampusDistance",
                                                  "ActivityCount"))
# Load necessary libraries
library(ggplot2)
# Scatter plot with smoothing line to assess linearity
ggplot(data, aes(x = campusDistance, y = activityCount)) +
  geom_point(color ="darkgreen") +
```

## Table 1: Correlation Test Results

```
# Residuals vs Fitted plot to assess homoscedasticity
plot(model, which = 1)
```

#### Research Question 3

gqnorm(residuals(model))

#### **Time Commitment and Student Involvement**

# Q-Q plot for normality of residuals

qqline(residuals(model), col = "red")

```
library(rstanarm)
# Set up the plotting area for two plots side by side
par(mfrow = c(1, 2))
```

```
# QQ plot for timeCommitment
qqnorm(data$timeCommitment, main = "QQ Plot (TC)",
       xlab = "Theoretical Quantiles", ylab = "Sample Quantiles")
ggline(data$timeCommitment, col = "red")
# QQ plot for studentInvolvement
qqnorm(data$studentInvolvement, main = "QQ Plot (SI)",
       xlab = "Theoretical Quantiles", ylab = "Sample Quantiles")
qqline(data$studentInvolvement, col = "blue")
# Reset the plotting layout to default (if needed for subsequent plots)
par(mfrow = c(1, 1))
# Load necessary libraries
library(ggplot2)
# Create a scatter plot
ggplot(data, aes(x = timeCommitment, y = studentInvolvement)) +
  geom point(color = "darkgreen") + # Set points to green
  geom_smooth(method = "lm", color = "orange", se = FALSE) + # Set line to orange
  ggtitle("Time Commitment and Student Involvement") +
  xlab("Time per week (hrs)") +
  ylab("Involvement (1-10)")
```

#### **Time Commitment and Activity Count**

Table 2: Correlation Test Results

```
# Load the necessary packages
library(knitr)
# Ensure data is defined and contains the necessary columns
# Example: data <- data.frame(timeCommitment = ..., studentInvolvement = ...)</pre>
# Perform Spearman's correlation test
result <- cor.test(data$timeCommitment, data$studentInvolvement,</pre>
                   method = "spearman", exact = FALSE)
# Extract relevant values
p_value <- sprintf("%.3e", result$p.value) # Format p-value in scientific notation</pre>
rho <- sprintf("%.3f", result$estimate) # Format rho to 3 decimal places</pre>
# Create a data frame for the table
table_results <- data.frame(</pre>
  Statistic = c("Spearman Coefficient (rho)", "p-value"),
  Value = c(rho, p_value)
# Display the table with basic kable (without additional styling)
kable(table_results, caption = "Spearman's Rank Correlation Results")
```

```
qqline(data$studentInvolvement, col = "blue")
# Reset the plotting layout to default (if needed for subsequent plots)
par(mfrow = c(1, 1))
# Load necessary libraries
library(ggplot2)
# Create a scatter plot
ggplot(data, aes(x = timeCommitment, y = activityCount)) +
  geom_point(color = "darkgreen") +
                                                           # Set points to green
  geom_smooth(method = "lm", color = "orange", se = FALSE) + # Set line to orange
  ggtitle("Time Commitment and Activity Count") +
  xlab("Time per week (hrs)") +
  ylab("Number of Activities")
# Load the necessary packages
library(knitr)
# Ensure data is defined and contains the necessary columns
# Example: data <- data.frame(timeCommitment = ..., studentInvolvement = ...)
# Perform Spearman's correlation test
result <- cor.test(data$timeCommitment, data$activityCount,</pre>
                   method = "spearman", exact = FALSE)
# Extract relevant values
p_value \leftarrow sprintf("\%.3e", result$p.value) # Format p-value in scientific notation
rho <- sprintf("%.3f", result$estimate) # Format rho to 3 decimal places</pre>
# Create a data frame for the table
table results <- data.frame(</pre>
  Statistic = c("Spearman Coefficient (rho)", "p-value"),
  Value = c(rho, p value)
)
# Display the table with basic kable (without additional styling)
kable(table_results, caption = "Spearman's Rank Correlation Results")
```