

STA304 The Outliers Code Appendix*

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Introduction

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

```
majTab=table(gsub("\\s+", "", unlist(strsplit(data$major,"")))) # Split into Single Items
# Convert to Data Frame
majDF=as.data.frame(majTab)
# Refactor for Correct Plot Order
majDF$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=0.8) +
  labs(title="Student Status", x="n = 63", caption = "Figure 1") + theme_minimal() +
  scale_fill_brewer(name="Options", labels=statusLabs, palette = "Set2") +
  geom_treemap_text(colour="black", place="center", size=10)

dist <- ggplot(data=distanceDF, aes(area=Freq, fill=Var1, label=paste(Freq))) +
  geom_treemap(color="black", size=0.8) +
```

*Data and Report Available at: https://github.com/vanshikav2/Extra_Curricular_Activities_Research

```

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", size=10) +
theme(plot.title = element_text(hjust = 0.5))

maj <- ggplot(data=majDF, aes(x=Var1, y=Freq, fill=Var1)) +
  geom_bar(stat="identity", color="black", size=0.15) +
  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", size=3) +
  theme(plot.title = element_text(hjust = 0.5),
        axis.title.y = element_text(size=12, angle=90),
        axis.text.x = element_text(size= 6)) +
  scale_y_continuous(expand = c(0.15, 0))

(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", size=0.3) +
  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) +
  scale_y_continuous(expand = c(0.1, 0))

```

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)
#summary(anova_major)

#Q-Q plot of Residuals
residuals <- residuals(anova_major)

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 Homogeneity of Variances
data <- data %>%
  mutate(major = as.factor(major))
levene_test_major <- leveneTest(activityTypeNumeric ~ major, data = data)
levene_df <- as.data.frame(levene_test_major)

levene_df[is.na(levene_df)] <- ""

kable(levene_df, caption = "Levene Test on The Activity Preference by Major")

anova_summary <- summary(anova_major)[[1]]
anova_df <- data.frame(anova_summary)
anova_df[is.na(anova_df)] <- ""
kable(anova_df, caption = "Anova Test on The Activity Preference by Major")

activity_lbls <- c(
  "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)
shapiro_M <- shapiro.test(gender_data_M)

shapiro_M_df <- as.data.frame(t(c(shapiro_M$statistic, shapiro_M$p.value)))
colnames(shapiro_M_df) <- c("W statistic", "p-value")
shapiro_F_df <- as.data.frame(t(c(shapiro_F$statistic, shapiro_F$p.value)))
colnames(shapiro_F_df) <- c("W statistic", "p-value")

shapiro_results <- data.frame(
  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)

levene_gender_df <- as.data.frame(levene_gender)
levene_gender_df[is.na(levene_gender_df)] <- ""

kable(levene_gender_df, caption = "Levene Test for Equality of Variances in
Activity Preferences by Gender Identity")

t_test_gender <- t.test(activityTypeNumeric ~ genderIdentity, data = data,
                        var.equal = FALSE)

t_test_result_df <- data.frame(T_Statistic = t_test_gender$statistic,
                              DF = t_test_gender$parameter,
```

```

      p_value = t_test_gender$p.value,
      MEAN_FEMALE = t_test_gender$estimate[1],
      MEAN_MALE = t_test_gender$estimate[2])

kable(t_test_result_df, caption = "Welch t-test on the Activity Proportion
  by Gender")

```

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)
shapiro_I <- shapiro.test(status_data_I)

shapiro_results <- data.frame(
  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)

levene_status_df <- as.data.frame(levene_status)
levene_status_df[is.na(levene_status_df)] <- ""

kable(levene_status_df, caption = "Levene Test on studentStatus")

wilcox_test <- wilcox.test(activityTypeNumeric ~ studentStatus, data = data)

wilcox_test_df <- data.frame(
  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,
                        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,
                        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$campusDistance)
activity <- summary(data$activityCount)

# Combine the summaries into a data frame
summary_t <- data.frame(
  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)) +

```


Table 1: ?(caption)

```
geom_point(color = "darkgreen") +
geom_smooth(method = "lm", col = "orange", se = FALSE) +
labs(title = "Campus Distance vs. Activity Count with Linear Trend Line",
      x = "Campus Distance",
      y = "Activity Count")

# Fit preliminary model
model <- lm(activityCount ~ campusDistance, data = data)

# Q-Q plot for normality of residuals
qqnorm(residuals(model))
qqline(residuals(model), col = "red")

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

# Create a data frame to summarize the correlation test results
correlation_results <- data.frame(
  Test = c("Pearson's Correlation", "Spearman's Correlation"),
  Correlation_Coefficient = c(-0.018, 0.025),
  p_value = c(0.891, 0.847),
  Significance = c("Not Significant", "Not Significant")
)

kable(correlation_results, caption = "Summary of Correlation Test Results")
```

Research Question 3

Time Commitment vs. Student Involvement and Activity Count

```
library(rstanarm)

# Set up the plotting area for 4 plots (2 rows, 2 columns)
par(mfrow = c(1, 3))
```

```

# QQ plot for timeCommitment (from the first chunk)
qqnorm(data$timeCommitment, main = "QQ Plot (TC)",
       xlab = "Theoretical Quantiles", ylab = "Sample Quantiles")
qqline(data$timeCommitment, col = "red")

# QQ plot for studentInvolvement (from the first chunk)
qqnorm(data$studentInvolvement, main = "QQ Plot (SI)",
       xlab = "Theoretical Quantiles", ylab = "Sample Quantiles")
qqline(data$studentInvolvement, col = "blue")

# QQ plot for activityCount (from the second chunk)
qqnorm(data$activityCount, main = "QQ Plot (AC)",
       xlab = "Theoretical Quantiles", ylab = "Sample Quantiles")
qqline(data$activityCount, col = "blue")

# Reset the plotting layout to default
par(mfrow = c(1, 1))

# Load necessary libraries
library(ggplot2)
library(patchwork)

# Create the first scatter plot with smaller title
plot1 <- 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 Commit. vs. Student Involvement") +
  xlab("Time per week (hrs)") + ylab("Involvement (1-10)") +
  theme(plot.title = element_text(size = 10)) # Reduce title size

# Create the second scatter plot with smaller title
plot2 <- 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 Commit. vs. Activity Count") +
  xlab("Time per week (hrs)") + ylab("Number of Activities") +
  theme(plot.title = element_text(size = 10)) # Reduce title size

# Combine the plots side by side

```

Table 2: ?(caption)

```
plot1 + plot2
```

```
# 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$timeCommitment, data$studentInvolvement,
                    method = "spearman", exact = FALSE)

# Extract relevant values
p_value <- sprintf("%.3e", result$p.value) # Format p-value in scientific notation
rho <- sprintf("%.3f", result$estimate)      # Format rho to 3 decimal places

# Create a data frame for the table
table_results <- data.frame(
  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")
```

Time Commitment and Activity Count

```
# 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,
                    method = "spearman", exact = FALSE)
```

```

# Extract relevant values
p_value <- sprintf("%.3e", result$p.value) # Format p-value in scientific notation
rho <- sprintf("%.3f", result$estimate)      # Format rho to 3 decimal places

# Create a data frame for the table
table_results <- data.frame(
  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")

```

Generative AI

Generative AI was used for assistance in troubleshooting code, formatting, and editing. No generative AI was used to create data or make the contents of the report other than that listed above. Links to the generative AI conversations are linked below.

<https://g.co/gemini/share/67545eab5501>

<https://g.co/gemini/share/781ab6a1b9aa>

<https://g.co/gemini/share/301105d9bd4b>

<https://chatgpt.com/share/6732d55a-4ab4-8010-bbf7-f1c1a85bdc5b>

<https://chatgpt.com/share/674e6690-6060-8008-b3aa-5b3eb07ce61d>

Additional conversation outputs are available in GitHub.