```
# Title: Mercury Case Study ------
library(readxl)
library(janitor)
library(tidyverse)
library(lubridate)
library(dplyr)
library(zoo)
library(readxl)
library(reshape2)
library(corrplot)
library (broom)
#load Data
file_path <- "mercury_case_study dataset.xlsx"</pre>
if (!file.exists(file path)) {
 cat("File not found! Please select the correct file.\n")
  file path <- file.choose() # Opens file explorer for user selection</pre>
sheets <- excel sheets(path = file path)</pre>
# Load sheet names
excel sheets(file path)
pointguard data <- read excel(file path, sheet = 1)</pre>
# Read data
catapult <- read excel(file path, sheet = "Catapult & Wellness")
cmj <- read_excel(file path, sheet = "CMJ data")</pre>
calf raise <- read excel(file path, sheet = "SL Standing Isometric Calf Rais")
sl cmj <- read excel(file path, sheet = "Single leg CMJ")</pre>
## Data Cleaning -----
# Remove empty rows/columns
catapult <- catapult %>% remove empty("cols") %>% remove empty("rows")
cmj <- cmj %>% remove empty("cols") %>% remove empty("rows")
calf raise <- calf raise %>% remove empty("cols") %>% remove empty("rows")
sl cmj <- sl cmj %>% remove empty("cols") %>% remove empty("rows")
# Handle missing values, wellness imputed with forward fill
## unsure if she played in All-Star game
### skip and leave as NA for all star break ------
# Define and assume All-Star Break Dates: Assume (July 18 - July 26, 2024)
all star break <- seq(as.Date("2024-07-18"), as.Date("2024-07-26"), by="days")
# Define wellness variables to forward fill
wellness vars <- c("Achilles Soreness", "General Soreness", "Sleep Quantity (hr)", "Sleep
Quality (10 best)")
# Apply forward fill only for wellness columns, but NOT during All-Star Break
catapult <- catapult %>%
 arrange(Date) %>% # Ensure data is sorted
 mutate(across(all of(wellness vars),
                ~ ifelse(Date %in% all star break, .x, na.locf(.x, na.rm = FALSE)))) %>%
 mutate(across(all_of(wellness_vars),
               ~ na.locf(.x, fromLast = TRUE, na.rm = FALSE)))  # Backward fill for
leading NAs
# Ensure performance metrics remain NA for All-Star, unsure if tracked remote
# and continued to train
performance vars <- c("Load", "Distance", "intensity pl min", "Jumps",
```

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"explosive efforts")
catapult[performance vars] <- lapply(catapult[performance vars],</pre>
                                         function(x) ifelse(catapult$Date %in%
all star break, NA, x))
# Read data and clean
catapult <- catapult %>% clean names()
cmj <- cmj %>% clean names()
calf raise <- calf raise %>% clean names()
sl cmj <- sl cmj %>% clean names()
### Convert dates, decimals and on court to minutes ----
catapult$date <- as.Date(catapult$date)</pre>
cmj$date <- as.Date(cmj$date)</pre>
calf raise$date <- as.Date(calf raise$date)</pre>
sl cmj$date <- as.Date(sl cmj$date)</pre>
# convert decimals in my load per distance
catapult <- catapult %>%
 mutate(load per distance = round(load / distance, 1))
#convert catapult time on court to minutes
catapult <- catapult %>%
 mutate(on court time = ifelse(!is.na(on court time),
                                 hour (on court time) * 60 + minute (on court time),
                                 NA)) # Convert to minutes while keeping NAs
# Form an only catapult, sleep and wellness dataframe
catapult wellness sleep <- catapult %>%
  select(date, on court time, intensity pl min, load per distance, achilles soreness,
general_soreness, sleep_quantity_hr, sleep_quality_10_best,
         load, distance, jumps, accelerations, decelerations) # Add relevant columns
# drop NA's
catapult_wellness_sleep <- catapult_wellness_sleep %>% drop na()
# convert to just on court minutes
catapult wellness sleep <- catapult wellness sleep %>%
 mutate(on_court_time = as.POSIXct(on_court_time, origin = "1899-12-30", tz = "UTC")) %>%
# Convert to proper date-time
 mutate(on_court_time = hour(on_court_time) * 60 + minute(on court time)) # Extract
total minutes
# merge two bodyweight kg columns, remove old and make just one
cmj <- cmj %>%
 mutate(bodyweight kg = ifelse(!is.na(athlete standing weight kg),
                                 athlete standing weight kg,
                                 bodyweight in kilograms kg)) %>%
  select(-bodyweight in kilograms kg, -athlete standing weight kg)
# Ensure date column is formatted correctly
cmj <- cmj %>% arrange(date)
cmj$date <- as.Date(cmj$date, format = "%Y-%m-%d")</pre>
# make sure bodyweight is numeric
cmj$bodyweight kg <- as.numeric(cmj$bodyweight kg)</pre>
## Bodyweight z scores detected that may be input error with losing
# 15 pounds in 3 days etc. Estimated weight based on nearby values in order to
# still use metrics
# z scores for outlier detectttion
#Detect outliers using Z-scores after data is cleaned
detect outliers with zscores <- function(data, df name, threshold = 1.5) {
```

```
# Select numeric columns only
  numeric data <- data %>% select(where(is.numeric))
  # Ensure dataset has numeric columns before proceeding
  if (ncol(numeric data) == 0) {
    stop(paste0("Error: No numeric columns found in dataset: ", df name))
  # Compute mean and standard deviation BEFORE filtering or modifying dataset
 mean values <- colMeans(numeric data, na.rm = TRUE)</pre>
  sd_values <- apply(numeric_data, 2, sd, na.rm = TRUE)</pre>
  # Remove columns with near-zero standard deviation
  valid columns <- sd values > 1e-5
  numeric data <- numeric_data[, valid_columns, drop = FALSE]</pre>
  mean values <- mean values[valid columns]</pre>
  sd values <- sd values[valid columns]</pre>
  # Compute Z-scores, keeping original row count
  z scores <- as.data.frame(scale(numeric data, center = mean values, scale = sd values))</pre>
  # Restore original row count (keeping NA values where applicable)
  z scores <- data.frame(Date = data$date, z scores)</pre>
  # Identify outliers where abs(Z-score) > threshold
  outlier mask <- abs(z scores[,-1]) > threshold
  # Convert Z-scores to data frame and keep only outliers
  outlier values <- z scores
  outlier values[,-1] <- outlier values[,-1] * outlier mask # Keep only flagged values
  # Convert to long format
  outlier long <- melt(outlier values, id.vars = "Date", variable.name = "Metric",
value.name = "Z Score")
  # Ensure no filtering removes valid outliers
  outlier long <- outlier long %>% filter(!is.na(Z Score) & Z Score != 0)
  # If no outliers found, return an empty DataFrame with correct structure
  if (nrow(outlier long) == 0) {
   print(paste0("No outliers detected for: ", df name))
    return(data.frame(Date = as.Date(character()), Metric = character(), Z Score =
numeric(), Dataset = character()))
  # Attach dataset name
 outlier long$Dataset <- df name</pre>
 return(outlier long)
# Apply the function to each dataset
catapult outliers <- detect outliers with zscores(catapult, "Catapult & Wellness")
cmj outliers <- detect outliers with zscores(cmj, "CMJ")</pre>
calf outliers <- detect outliers with zscores(calf raise, "Calf Raise")
# Combine outlier reports
all outliers <- bind rows(catapult_outliers, cmj_outliers, calf_outliers) %>%
 mutate(Date = as.Date(Date)) # Ensure Date remains in Date format
# Save the Z-score outlier report
write.csv(all outliers, file.path("C:/Users/prist/OneDrive/Desktop/mercury",
"zscore outlier report.csv"), row.names = FALSE)
```

```
# Ensure date is in Date format
cmj <- cmj %>%
  mutate(date = as.Date(date))
# Check basic statistics for bodyweight
mean bw <- mean(cmj$bodyweight kg, na.rm = TRUE)</pre>
sd bw <- sd(cmj$bodyweight kg, na.rm = TRUE)</pre>
max change <- max(abs(diff(cmj$bodyweight kg)), na.rm = TRUE)</pre>
# use 86.31 as mean imputation for bodybweight kg outliers
cmj <- cmj %>%
  mutate(bodyweight kg = if else(date == as.Date("2024-05-31", "2024-5-13"), 86.31,
bodyweight kg))
# convert kg to pounds
# Ensure bodyweight kg is numeric
cmj$bodyweight kg <- as.numeric(cmj$bodyweight kg)</pre>
# Convert kg to pounds and round to one decimal place
cmj <- cmj %>%
  mutate(bodyweight in pounds lbs = round(bodyweight kg * 2.20462, 1))
# Recalculate CMJ metrics that depend on bodyweight now that imputed correctly
cmj <- cmj %>%
  mutate(
    # Recalculate Concentric Impulse (overwrite existing values)
    concentric impulse n s = if else(!is.na(concentric impulse n s) &
!is.na(bodyweight kg),
                                      concentric impulse n s / bodyweight kg *
bodyweight_kg, # Correcting for bodyweight errors
                                      concentric impulse n s),
    # Recalculate RSI (Reactive Strength Index)
    rsi_modified_m_s = if_else(!is.na(jump_height_flight_time_in_inches_in) &
!is.na(eccentric duration ms),
                                jump height flight time in inches in /
(eccentric duration ms / 1000), # RSI = Jump Height / Eccentric Time
                               rsi modified m s),
    # Recalculate Eccentric Braking Impulse (overwrite existing values)
    eccentric braking impulse n s = if else(!is.na(eccentric braking impulse n s) &
!is.na(bodyweight kg),
                                             eccentric braking impulse n s / bodyweight kg
* bodyweight kg, # Correcting for bodyweight changes
                                             eccentric braking impulse n s)
  )
# Delete cmj date for 8-4 due to missing bodyweight originally, peak power/bm at 3.26 and
#multiple metrics at >1.5 z scores errors.
# Remove CMJ data for 8/4/2024
cmj <- cmj %>%
  filter(date != as.Date("2024-08-04"))
#### Data Visualizations and Stats -----
# Achilles soreness over time
ggplot(catapult wellness sleep, aes(x = date, y = achilles soreness)) +
  geom line(color = "red", size = 1) +
```

```
geom point(color = "black", size = 2) +
  labs(title = "Achilles Soreness Over Time", x = "Date", y = "Soreness Level") +
  theme minimal()
# Sleep Quality and Quantity over time
ggplot(catapult wellness_sleep, aes(x = date)) +
  geom line (aes (y = sleep quality_10_best, color = "Sleep Quality"), size = 1) +
  geom line(aes(y = sleep quantity hr, color = "Sleep Quantity"), size = 1) +
  labs(title = "Sleep Trends Over Time", x = "Date", y = "Sleep Score") +
  scale color manual(values = c("Sleep Quality" = "blue", "Sleep Quantity" = "green")) +
  theme minimal()
# Correlations: relationships between soreness, sleep, and performance.
# 10+ variables, at least 100 observations are recommended.
correlation data <- catapult wellness sleep %>%
  select (achilles soreness, general soreness, sleep quantity hr, sleep quality 10 best,
         load, distance, jumps, accelerations, decelerations, intensity pl min,
load per distance)
# Compute correlation matrix (pairwise complete)
cor matrix <- cor(correlation data, use = "pairwise.complete.obs")</pre>
# Plot heatmap
corrplot(cor matrix, method = "color", type = "lower", tl.col = "black", tl.srt = 45,
         addCoef.col = "black", number.cex = 0.7)
####Compare Pre- and Post-All-Star Break Performance
#Though this looks significant, Does Achilles Soreness Change After the All-Star
#Break? All tests Wilcoxon Rank-Sum Test, T Test, Regression) were insignificant)
ggplot(catapult wellness sleep, aes(x = date, y = load per distance, color = date >=
as.Date("2024-07-26"))) +
  geom point() +
  geom smooth(method = "lm", se = FALSE) +
  labs(title = "Load Per Distance Before and After All-Star Break",
       x = "Date", y = "Load Per Distance", color = "Post-All-Star Break") +
  theme minimal()
#### Time-Series Trends: Achilles Soreness vs. Load, Distance, Jumps, and
Intensity##############
catapult wellness sleep$date <- as.Date(catapult wellness sleep$date)</pre>
# Select relevant columns
plot data <- catapult wellness sleep %>%
  select(date, achilles soreness, load, distance, jumps, intensity pl min)
# Convert date to Date format
plot data$date <- as.Date(plot data$date)</pre>
# Plot with two y-axes
ggplot(plot_data, aes(x = date)) +
  geom_line(aes(y = load, color = "Load"), size = 1) +
  geom line(aes(y = distance, color = "Distance"), size = 1) +
  geom line(aes(y = jumps, color = "Jumps"), size = 1) +
  geom_line(aes(y = intensity_pl_min, color = "Intensity"), size = 1) +
  geom_line(aes(y = achilles_soreness * 10, color = "Achilles_Soreness"), size = 1,
linetype = "dashed") + # Adjust scale
  scale_y_continuous(sec.axis = sec axis(~ . / 10, name = "Achilles Soreness")) +
  labs(title = "Time-Series: Soreness vs Performance Metrics",
       x = "Date",
       y = "Primary Metrics",
       color = "Metric") +
  theme minimal()
```

## Pre- vs. Post-All-Star Break Comparison, skewed soreness so wilcoxson test

```
# Define pre- and post-All-Star Break groups
pre break <- catapult wellness sleep %>% filter(date < as.Date("2024-07-26"))</pre>
post break <- catapult wellness sleep %>% filter(date >= as.Date("2024-07-26"))
# Perform Wilcoxon Rank-Sum Test (Non-Parametric)
wilcox test <- wilcox.test(pre break$achilles soreness, post break$achilles soreness,</pre>
exact = FALSE)
# Print results
print(wilcox test)
ggplot(catapult wellness sleep, aes(x = date, y = achilles soreness, color = date >=
as.Date("2024-07-26"))) +
  geom point(size = 3, alpha = 0.6) +
  geom smooth(method = "lm", se = FALSE) +
  labs(title = "Achilles Soreness Before & After All-Star Break",
       x = "Date", y = "Soreness Level", color = "Post-All-Star Break") +
  theme minimal()
### Anomaly Detection: Spotting Sudden Workload Spikes
# Compute rolling average and flag spikes
catapult wellness sleep <- catapult wellness sleep %>%
  mutate(load_rolling_avg = rollmean(load, k = 7, fill = NA, align = "right"), # 7-day
rolling avg
         load spike = case when(
           load > (load rolling avg * 1.5) ~ "Severe Spike",
           load > (load rolling avg * 1.3) ~ "Moderate Spike",
           TRUE ~ "Normal"
         ))
# Plot workload with anomaly detection
ggplot(catapult wellness sleep, aes(x = date, y = load, color = load spike)) +
  geom line(size = 1) +
  geom point(size = 3, alpha = 0.7) +
  scale color manual(values = c("Severe Spike" = "red", "Moderate Spike" = "orange",
"Normal" = "\overline{black}") +
  labs(title = "Workload Spike Detection Over Time",
       x = "Date", y = "Load", color = "Spike Type") +
  theme minimal()
catapult wellness sleep %>%
  filter(load spike != "Normal") %>%
  select(date, load, load rolling avg, load spike)
#Compare Load Per Distance Before & After Spikes
ggplot(catapult_wellness_sleep, aes(x = date, y = load_per_distance, color = load_spike))
 geom line(size = 1) +
  geom point(size = 3, alpha = 0.7) +
 scale color manual (values = c("Severe Spike" = "red", "Moderate Spike" = "orange",
"Normal" = "black")) +
  labs(title = "Load Per Distance Before & After Spikes",
       x = "Date", y = "Load Per Distance", color = "Spike Type") +
 theme minimal()
## Visualization load per distance vs decels
catapult wellness sleep$date <- as.Date(catapult wellness sleep$date)</pre>
# Ensure numeric columns
catapult wellness sleep <- catapult wellness sleep %>%
```

```
mutate(load per distance = as.numeric(load per distance),
         decelerations = as.numeric(decelerations))
# Plot Load Per Distance vs Decelerations Over Time
qqplot(catapult wellness sleep, aes(x = date)) +
  geom line(aes(y = load per distance, color = "Load Per Distance"), size = 1.2) +
  geom line(aes(y = decelerations * 10, color = "Decelerations"), linetype = "dashed",
size = 1.2) + # Adjust scaling if needed
  scale color manual(values = c("Load Per Distance" = "blue", "Decelerations" = "red")) +
  labs(title = "Load Per Distance vs. Decelerations Over Time",
       x = "Date",
       y = "Metrics",
       color = "Legend") +
  theme_minimal()
# force plate asymmetry trends visual
# Read the Excel file (No sheet name required, as it's the first sheet)
force plate data <- read excel("C:/Users/prist/OneDrive/Desktop/mercury/cmj.xlsx")</pre>
# Ensure the date column is correctly formatted
force plate data <- force plate data %>%
 mutate(Date = as.Date(date))
# Function to extract numerical values from asymmetry columns (removing "R" or "L" text)
clean numeric <- function(column) {</pre>
  as.numeric(gsub("[^0-9.]", "", as.character(column)))
# Extract relevant asymmetry metrics and clean data
force plate data <- force plate data %>%
 mutate(
    `Jump Height Asymmetry (%)` =
clean numeric (eccentric braking impulse percent asym percent),
    `Peak Power Asymmetry (%)` =
clean_numeric(eccentric_deceleration_impulse_percent_asym_percent),
    `RSI Asymmetry (%)` = clean numeric(landing impulse percent asym percent),
    `Eccentric Braking Asymmetry (%)` =
clean_numeric(eccentric_peak_force_percent_asym_percent)
# Create the Force Plate Asymmetry Trend Plot
ggplot(force plate data, aes(x = Date)) +
  geom line(aes(y = `Jump Height Asymmetry (%)`, color = "Jump Height Asymmetry"), size=1)
  geom line(aes(y = `Peak Power Asymmetry (%)`, color = "Peak Power Asymmetry"), size=1) +
  geom line(aes(y = `RSI Asymmetry (%)`, color = "RSI Asymmetry"), size=1) +
  geom line (aes (y = `Eccentric Braking Asymmetry (%)`, color = "Eccentric Braking
Asymmetry"), size=1) +
  geom point(aes(y = `Jump Height Asymmetry (%)`, color = "Jump Height Asymmetry"),
size=3) +
  geom point(aes(y = `Peak Power Asymmetry (%)`, color = "Peak Power Asymmetry"), size=3)
  geom point(aes(y = `RSI Asymmetry (%)`, color = "RSI Asymmetry"), size=3) +
  geom point(aes(y = `Eccentric Braking Asymmetry (%)`, color = "Eccentric Braking
Asymmetry"), size=3) +
  scale color manual(values = c("Jump Height Asymmetry" = "blue",
                                "Peak Power Asymmetry" = "red",
                                "RSI Asymmetry" = "green",
                                "Eccentric Braking Asymmetry" = "purple")) +
  labs(title = "Force Plate Asymmetry Trends Over Time",
       x = "Date",
       y = "Asymmetry Percentage",
       color = "Metrics") +
  theme minimal() +
```

```
theme(axis.text.x = element text(angle = 45, hjust = 1))
# Statistics: eploratory model
# Shift soreness forward by 1-3 days to see post-spike impact
catapult wellness sleep <- catapult wellness sleep %>%
  arrange(date) %>%
 mutate(achilles soreness tomorrow = lead(achilles soreness, 1),
         achilles soreness 2days = lead(achilles soreness, 2),
         achilles soreness 3days = lead(achilles soreness, 3))
# Fit a regression model: Does a spike predict higher soreness? No
soreness model <- lm(achilles soreness tomorrow ~ load spike, data =
catapult wellness sleep)
summary(soreness model) # Check model coefficients
#Regression Load Per Distance) RSI
# Convert date columns to Date format for proper merging
catapult$date <- as.Date(catapult$date)</pre>
cmj$date <- as.Date(cmj$date)</pre>
# Merge datasets on date to align Load Per Distance and RSI
merged data <- inner join(catapult, cmj, by = "date")</pre>
# Ensure numeric columns are in the correct format
merged data <- merged data %>%
 mutate(load per distance = as.numeric(load per distance),
         rsi = as.numeric(rsi modified m s)) %>% # Rename RSI for simplicity
  drop na()
# Perform linear regression
model <- lm(rsi ~ load_per_distance, data = merged_data)</pre>
# Summary of the regression model
summary(model)
# Extract model statistics
model stats <- glance(model)</pre>
print(model stats)
# Visualization: Load Per Distance vs RSI with Regression Line
ggplot(merged data, aes(x = load_per_distance, y = rsi)) +
  geom point(alpha = 0.7, color = "blue") + # Scatter plot
  geom smooth(method = "lm", se = TRUE, color = "red") + # Regression line
  labs(title = "Load Per Distance vs. RSI Regression",
       x = "Load Per Distance",
       y = "Reactive Strength Index (RSI)") +
  theme minimal()
# regression load vs soreness
# Ensure date
catapult_wellness_sleep$date <- as.Date(catapult wellness sleep$date)</pre>
# Create a "workload spike" variable (change in load from previous day)
catapult wellness sleep <- catapult wellness sleep %>%
 arrange(date) %>%
 mutate(workload spike = load - lag(load))
# Perform linear regression
model <- lm(achilles soreness ~ workload spike, data = catapult wellness sleep)
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# Show regression summary
summary(model)
# Extract model statistics
model stats <- glance(model)</pre>
print(model stats)
# Regression model for soreness 2-3 days after spikes
soreness model 2day <- lm(achilles soreness 2days ~ load spike, data =
catapult wellness sleep)
soreness model 3day <- lm(achilles soreness 3days ~ load spike, data =
catapult wellness sleep)
summary(soreness model 2day) # Check 2-day delay impact
summary(soreness model 3day) # Check 3-day delay impact
# Full model including sleep & recovery metrics
soreness model full <- lm(achilles_soreness_tomorrow ~ load_spike + sleep_quality_10_best
+ sleep quantity hr, data = catapult wellness sleep)
summary(soreness model full) # Check if sleep amplifies soreness post-spike
# Compute rolling 7-day workload
catapult wellness sleep <- catapult wellness sleep %>%
 mutate (rolling load 7day = rollmean(load, \bar{k} = 7, fill = NA, align = "right"))
# Regression: Does cumulative load predict soreness better than spikes?
soreness model cumulative <- lm(achilles soreness tomorrow ~ rolling load 7day, data =
catapult wellness sleep)
summary(soreness_model_cumulative)
# Regression model: Does soreness predict lower intensity?
performance model <- lm(intensity pl min ~ achilles soreness, data =
catapult wellness sleep)
summary (performance model)
```