Start Here

```
# Load necessary libraries
library(dplyr)
Attaching package: 'dplyr'
The following objects are masked from 'package:stats':
          filter, lag
The following objects are masked from 'package:base':
          intersect, setdiff, setequal, union
library(corrplot) # correlation matrix visualization
corrplot 0.92 loaded
library(ggcorrplot) # ggplot2 style visualization of correlation matrix
Loading required package: ggplot2
# Read datasets
data <- read.csv("../../Data/data.csv")</pre>
data2 <- read.csv("../../Data/500cities.csv")</pre>
# Prepare 'data' by padding and merging columns to create a unique identifier
data$STATEFP <- sprintf("%02d", as.numeric(data$STATEFP))</pre>
data$COUNTYFP <- sprintf("%03d", as.numeric(data$COUNTYFP))</pre>
data$TRACTCE <- sprintf("%06d", as.numeric(data$TRACTCE))</pre>
data$TractFIPS <- as.numeric(paste0(data$STATEFP, data$COUNTYFP, data$TRACTCE))</pre>
# Summarize 'data' to get mean of 'NatWalkInd' by 'TractFIPS'
averaged_data <- data %>%
     group_by(TractFIPS) %>%
     summarise(NatWalkInd = mean(NatWalkInd, na.rm = TRUE)) %>%
     ungroup()
# Select relevant columns from 'data2'
data2_relevant <- data2 %>%
     select(TractFIPS, StateAbbr, DIABETES_CrudePrev, BPHIGH_CrudePrev, OBESITY_CrudePrev, LPA_CrudePrev, CrudePrev, CrudePrev
# Merge 'averaged_data' with 'data2_relevant' on 'TractFIPS'
merged_data <- merge(averaged_data, data2_relevant, by = "TractFIPS", all.x = TRUE, all.y = TRUE)
# Filter for specific condition
merged_data <- merged_data %>% filter(StateAbbr == "CA")
```

```
# Calculate correlation matrix
cor_matrix <- cor(merged_data %>% select(-TractFIPS, -StateAbbr), use = "complete.obs") # Handling miss
# Output variable names (column names) used for correlation matrix
cat("Variables used for correlation matrix:\n")
```

Variables used for correlation matrix:

