Analysis A

Our goal is to find common hidden characteristics that may help explain or predict changes in stock prices. Thirteen variables, not including the date. I will assume that the factors are 8 because it is less than total of 13. By seeing the boxplots plotted by given data. The data needs to be standardized (fig1). And the variables are correlated highly (fig2).

Factor Analysis of both PC & ML Methods:

Employed both methods PC & MLE to comprehensively address the study goal of identifying underlying patterns in stock price behavior. Using the PC method, I observed that after considering the first 4 factors, there were no substantial effects on variables, as determined by a threshold of 0.3. Similarly, employing the MLE method, we found that after the first 3 factors, there were no significant impacts on variables, using the same threshold.

Despite the PC method typically indicating larger loadings, I opted for a 3-factor model because the consistency between the two methods and the number of significant groups identified in the correlation matrix. I finally concluded that a 3-factor model adequately captures the underlying patterns in the data.

After employing Varimax Rotation(fig3) of pc and mle for clarity, our analysis identified three key factors in stock price behavior. The first factor encompasses market leaders with diversified portfolios, indicating stability and growth potential. The second factor reflects companies influenced by consumer spending trends and online retail performance, particularly in e-commerce and consumer goods sectors. The third factor represents international giants

focusing on technology, consumer goods, and finance, showcasing their dominance across multiple industries.

By Comparing the final models obtained from both PC and MLE methods results are shown below that there is a significant similarity, particularly for Factors 1 and 2. Acknowledged some differences in the specific companies captured by Factor 3 but overall found both methods identified similar underlying patterns in Factor analysis methods.

Principal Component Analysis (PCA):

The goal of the PCA is to determine the total number of principal components necessary to explain most of the variance in the data. Standardization of data is essential due to different ranges observed in the variables, as shown by box plots (Fig1). By Plotting scree plot (Fig 4) and examined result to identify that only two principal components are enough to cover approximately 82% of the data's variability.

In our Principal Component Analysis (PCA), I identified the key variables driving PC1 are Walmart, Apple, Toyota, Dow Jones, Google, and Amazon, with Walmart emerging as the most influential. These variables exhibit strong positive relationships with PC1. And also, Walmart and Apple carry the highest weights in defining PC1. while variables like HSBC and Honda show negative weights, suggesting an inverse relationship. PC1 captures common trends among these variables.

Additionally, our analysis of variable loadings reveals that certain variables, such as Walmart, Apple, and Toyota, exhibit the highest loadings in PC1, while others like HSBC and Honda demonstrate opposing trends.

From the PCA graph (Fig5), I observe that Walmart, Coca Cola, Pepsi, S&P 500, Google, Toyota, Apple, Chase, and Dow Jones form a tight cluster near the origin. However, Honda, Amazon, eBay, and HSBC stand out from this main cluster. This suggests that these companies that set them apart to share similar characteristics on the two principal components analyzed.

Comparison between the two parts PCA & FA:

By comparing the both PCA & FA methods. I found that both methods agree with each other in the results. Because they have similar patterns and groups. And also, both coveys the same interpretations. among the variables, companies clustering near the origin in the PCA graph, corresponding to the market leaders identified in the factor analysis. The outliers in the PCA graph align with the e-commerce and international giants factors.

Analysis B1:

The study goal is to identify if there is difference in Age, sleep duration and Heart Rate between the different genders Male, Female. The chi-sq QQ Plots & Royston test imply that all variables appear to be close enough to normality with some outliers.

For univariate analysis of the Age, sleep duration and Heart Rate plotted the boxplots. And for multivariate analysis also have plotted the 3d scatterplot and graph are in fig 8 to 12.

To address the study goal, we chose to analyze the hoteling t2 test. The hoteling's T2 test gives a p-value 1.624e-07 which is less than $0.05(\alpha)$ stating that there is significant difference in atleast one of the variables between the groups at a significance of 0.05.

By checking the confidence intervals of 95% of Bonferroni. I found that there is difference in all the variables because there is no zero in between the resulted interval values.

It shows that three variable sleep duration, age, heart rate varies in gender male to female.

Analysis B2:

The study goal is to identify if there is difference in Age, sleep duration and Heart Rate between the different Occupations Accountant, Doctor & Lawyer. The chi-sq QQ Plots & Royston test imply that all variables appear to be close enough to normality with some outliers.

For univariate analysis of the Age, sleep duration and Heart Rate plotted the boxplots. And for multivariate analysis also have plotted the 3d scatterplot and graph.

To address the study goal, we chose to analyze the Manova test. The Manova test gives a p-value 5.143e-14 which is less than $0.05(\alpha)$ stating that there is significant difference in atleast one of the variables between the groups at a significance of 0.05.

By checking the confidence intervals of 95% of Bonferroni. I found that there is difference in all the variables because there is no zero in between the resulted interval values.

It shows that three variable sleep duration, age, heart rate varies in different occupations (Accountant, Doctor, Lawyer).

variable	p-val
Age	9.94E-09
Sleep duration	3.55E-10
Heart rate	1.09E-05

Table: Anova test for occupation for other variables

Appendix:

Analysis A:

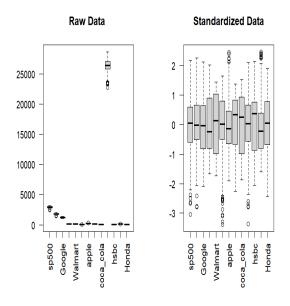


Fig1: Raw & Standardized data boxplots

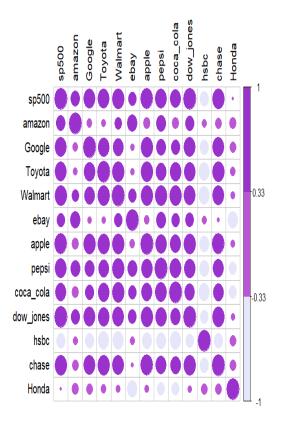


Fig2: Correlation plot



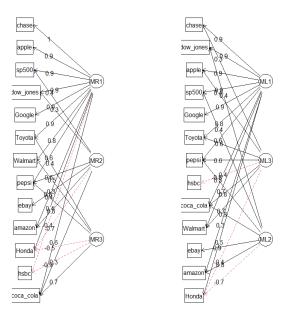


Fig3: Varimex Rotation

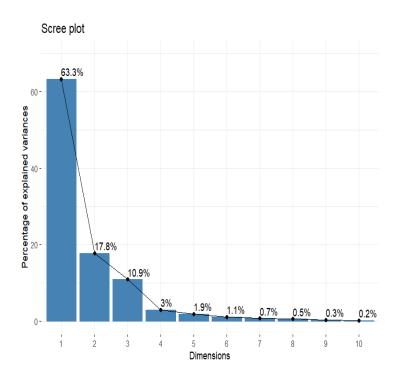


Fig4: screeplot

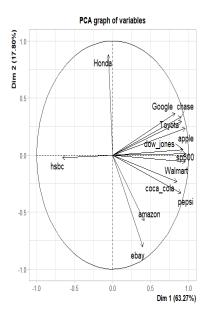


Fig5: PCA Graph

Analysis b1:

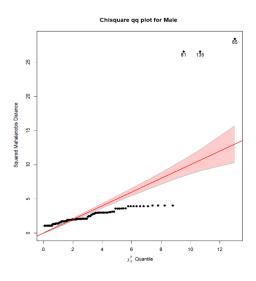


Fig 6: chisq qaplot for male

Chisquare qq plot for female

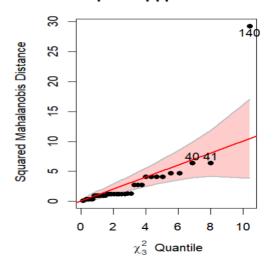


Fig7: chi-sq QQ-plot for female

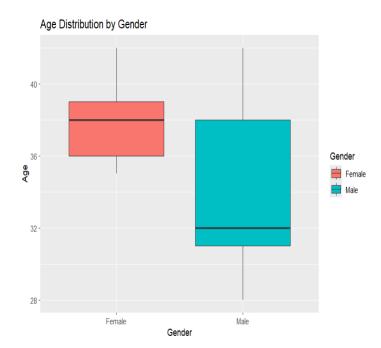


Fig 8: boxplot of age distribution by gender

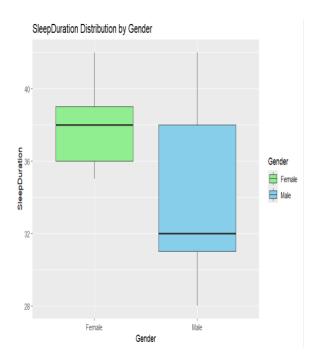


Fig 9: boxplot of sleep duration by gender

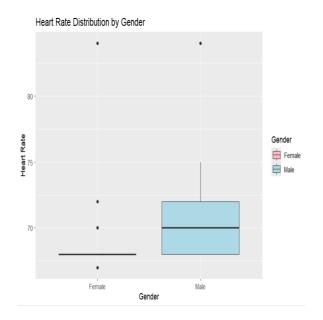


Fig 10: boxplot of heartrate distributed by gender

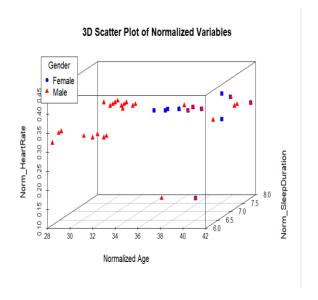


Fig 11: 3d sctterplot of numerical variables with gender

Scatterplot Matrix with Colors and Shapes for Gender

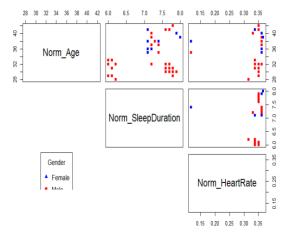


Fig 12 scatter plot matrix with gender

Analysis b2:

Chisquare qq plot of Occupation Type Accountant

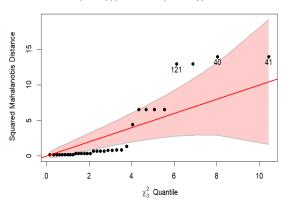


Fig 13: chi-sq of occupation type accountant

Chisquare qq plot of Occupation Type Doctor

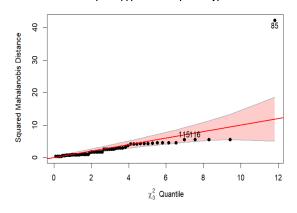


Fig 14: chisq of occupation type doctor

Chisquare qq plot of Occupation Type lawyer

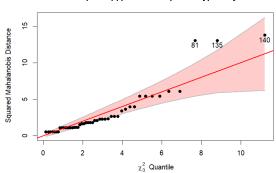


Fig 15: chi-sq of occupation type lawyer

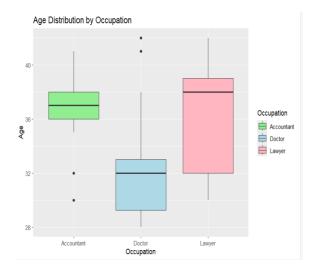


Fig 16: boxplot of age distribution by occupation

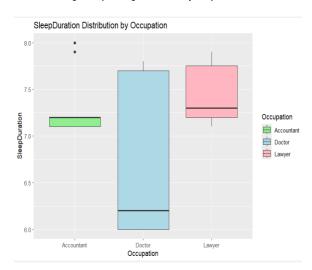


Fig 17: boxplot of sleep duration distribution by occupation

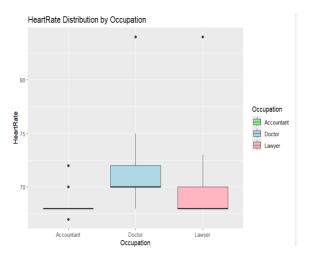


Fig 18: boxplot of heart rate distribution by occupation

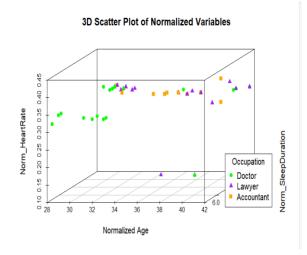


Fig 19: 3d scatterplot of occupation

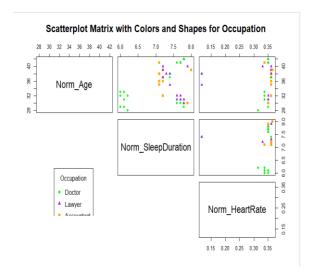


Fig 20: 3d scatterplot matrix of occupation

Index	Factor 1	Factor 2	Factor 3
1	sp500	amazon	amazon
2	amazon	pepsi	coca cola
3	Google	ebay	hsbc
4	Toyota	coca cola	
5	Walmart	Honda	
6	apple		
7	pepsi		
8	coca cola		
9	dow jones		
10	hsbc		
11	chase		

Table1: Companies under by each factor by ML method

Index	Factor 1	Factor 2	Factor 3
1	sp500	sp500	Toyota
2	Google	amazon	Walmart
3	Toyota	ebay	pepsi
4	Walmart	pepsi	coca cola
5	apple	coca cola	hsbc
6	pepsi	dow jones	Honda
7	coca cola	Honda	
8	dow jones		
9	hsbc		
10	chase		
11	Honda		

Table2: Companies under by each factor by pc method

Company	Dim.1	Dim.2
sp500	0.34	0.01
amazon	0.15	-0.38
Google	0.29	0.24
Toyota	0.32	0.19
Walmart	0.33	-0.04
eBay	0.14	-0.53
Apple	0.33	0.15
Pepsi	0.31	-0.22
Coca-Cola	0.30	-0.15
Dow Jones	0.32	0.03
HSBC	-0.23	-0.02
Chase	0.32	0.22
Honda	-0.02	0.58

Table 3: PCAWeights

MLE	P-Val
MLE1	NA
MLE2	0
	7.5 ×
MLE3	10-224
	1.6 ×
MLE4	10-157
MLE5	4.1 × 10-71
MLE6	1.0 × 10-45
MLE7	5.1 × 10-28
MLE8	3.9 × 10-23
T	1 (0()

Table 4: p-values for 8 factors

Pc method loadings for factors 9

sp500 -0.962 -0.011 -0.241 0.064 -0.012 -0.054 -0.017 0.073 -0.036 -0.417 0.575 -0.625 -0.199 -0.098 0.226 -0.061 0.026 -0.003 Amazon -0.822 -0.366 -0.130 0.305 -0.210 0.113 0.104 -0.098 0.032 Google -0.907 -0.293 0.173 -0.089 0.156 0.128 0.005 0.010 0.082 Toyota -0.956 0.058 0.160 -0.122 0.116 -0.042 -0.074 -0.057 0.105 Walmert -0.400 0.807 -0.160 0.315 0.239 -0.012 0.048 -0.038 0.003 eBay -0.955 -0.233 -0.051 0.038 -0.114 -0.055 -0.009 0.025 -0.018 Apple -0.899 0.332 0.021 -0.120 -0.105 -0.132 -0.096 -0.154 -0.048 Pepsi Coca-Cola -0.848 0.231 0.345 -0.235 0.081 0.054 0.189 -0.014 -0.086 Dow Janes -0.927 -0.046 -0.297 0.129 0.100 -0.076 -0.036 0.093 -0.027 0.659 0.025 -0.704 -0.160 0.029 -0.134 0.136 -0.050 0.058 H5BC -0.904 -0.337 -0.132 -0.124 -0.055 -0.105 0.065 0.070 0.028 Chase 0.058 -0.880 -0.387 -0.028 0.226 0.063 -0.049 -0.086 -0.076 Honda

ML method Loadings for 9 factors

sp500 0.969 0.147 0.145 0.096 -0.009 -0.069 -0.011 -0.019 0.021 Amezon 0.413 0.834 0.168 -0.299 0.063 -0.090 -0.002 -0.005 -0.008 Google 0.824 -0.185 0.219 0.201 0.106 -0.233 0.087 0.149 -0.060 Toyota 0.914 -0.31 -0.007 -0.085 0.159 0.143 0.043 0.030 -0.001 Walmart 0.952 -0.01 -0.176 -0.122 -0.079 0.152 -0.104 0.010 -0.022 eBay 0.383 0.75 -0.477 0.209 0.000 0.069 0.054 0.018 -0.021 0.959 -0.19 0.151 0.053 -0.051 -0.120 0.008 0.100 0.009 Apple 0.879 0.23 -0.189 -0.157 -0.254 -0.034 -0.058 0.034 -0.013 Pensi CocaCola 0.824 -0.32 -0.364 -0.267 -0.135 0.140 0.145 -0.026 0.180 DowJones 0.942 0.13 0.174 0.202 0.057 -0.005 -0.065 -0.051 0.032 -0.640 0445 0.591 0.063 -0.048 0.174 0.004 0.038 0.014 HSBC 0.916 -0.86 0.316 -0.004 -0.085 0.023 0.090 -0.057 -0.048 -0.013 -0424 0.741 0.206 0.332 0.162 -0.062 0.028 -0.035 Honda

B1 Analysis:

multivariateNormality

Test Η p value

Royston 124.4435 1.279754e-15

Univariate Normality

variable	statistic	p-val
Healthsleep_data.Age	0.8265	0.0294
Healthsleep_data.SleepDuration	4.6335	<0.001
Healthsleep_data.HeartRate	7.4626	<0.001

CI (bonferroni):

LCBS	UCBS
-5.857	-2.24
-0.647	-0.012
0.374	3.143

B2 Analysis:

Multivariate Normality

Test Η p value 77.58203 1.011546e-16 Royston

Univariate normality

variable	statistic	P-Val
Healthsleep_data.Age	1.6605	2.00E- 04
Healthsleep_data.SleepDuration	6.5557	< 0.001
Healthsleep_data.HeartRate	6.6117	< 0.001

CI (Bonferroni):

	sleep duration	
ige (LL,UL)	(LL,UL)	Heart rate (LL,UL)
1.9665577	(0.2515194	
•	•	
5.3755733)	0.9533682)	(-4.6022878, -1.2677024)
-2.0069251, 2.64006498)	(-0.5194463,- 0.22028443)	(-3.4297534,-0.08481464)
1.9665577,6	(0.2515194,	
3755733)	0.9533682)	(-4.6022878, -1.2677024)
	.64006498)	Age (LL,UL) (LL,UL) 1.9665577, (0.2515194, 3.3755733) 0.9533682) -2.0069251, (-0.5194463,- 3.64006498) 0.22028443) 1.9665577,6 (0.2515194,

```
R codes:
                                                              boxplot (stocks data, las = 2, main = "Raw Data")
                                                              R = cor(Standardized Data) # correlation matrix of
# Load required libraries
                                                              standardized data
library (openxlsx)
                                                              R2 = cor( stocks data ) #correlation matrix of raw
                                                              data
library(readxl)
                                                              boxplot (Standardized_Data, las = 2, main= "
library (psych)
                                                              Standardized Data ")
library (tableone)
                                                              # Create a correlation plot
library (heplots)
                                                              par ( mfrow = c(1,1))
library (MVN)
                                                              library ("corrplot")
library (biotools)
                                                              corrplot (R, is.corr =TRUE, col =
                                                              c("lavender", "mediumorchid", "darkorchid"), tl.col =
library (car)
                                                              " black ")
library (GPArotation)
                                                              # Perform factor analysis and calculate p-values for
                                                              different numbers of factors
library (corrr)
                                                              pvals=c()
library (ggplot2)
                                                              for (k in 2:8) {
library (FactoMineR)
                                                               pvals[k]=factanal(covmat = R, n.obs = 252, factors =
library (factoextra)
                                                              k)$PVAL
# Read data from Excel file
                                                              }
stocks data = read excel("StockData.xlsx")
                                                              pvals
stocks data = stocks data [2:14] #exclude the first
                                                              # Set the number of factors
column as it 's probably an index
                                                              factors = 3
head (stocks data, 5) # display the first 5 rows
                                                              # Perform principal component analysis
# Create a table of numerical summaries
                                                              Eig.Res=eigen(R)
numerical_summaries = CreateTableOne ( data =
stocks_data, includeNA = FALSE)
                                                              loadings=Eig.Res$vectors[, 1:factors]
print ( numerical summaries )
                                                              eigenvalues = Eig.Res$values[1:factors]
# Standardize the data and create boxplots of raw
                                                              L=round(loadings %*% diag(sqrt(eigenvalues)), 3)
and standardized data
                                                              PSI=diag(diag(R - L %*% t(L)))
Standardized Data = as.data.frame ( scale
(stocks_data))
                                                              EPS = R - L \%*\% t(L) - PSI
par ( mfrow = c(1,2))
```

```
communalities=round(rowSums(L^2), 3)
                                                             for (col in names(selected companies list)) {
diag(PSI)
                                                              cat("Companies for column", col, ":",
                                                             toString(selected_companies_list[[col]]), "\n")
# Perform factor analysis using maximum likelihood
estimation (MLE)
                                                             }
FA.Res=fa(stocks data, nfactors = factors, rotate =
                                                             # Perform PCA using FactoMineR
"none", fm = "ml")
                                                             PCA.Res=PCA(Standardized_Data, graph = TRUE, ncp
L1=round(FA.Res$loadings[, 1:factors], 3)
                                                             = 2, scale.unit = TRUE)
PSI1=diag(FA.Res$uniquenesses)
                                                             # Print PCA results
EPS1=R - L1 %*% t(L1) - PSI1
                                                             PCA.Res
H1=FA.Res$communality
                                                             str(PCA.Res)
# Perform factor analysis with different rotation
                                                             # Summary of PCA analysis
methods (PC method)
                                                             summary(PCA.Res)
FA.Res.None=fa(stocks data, nfactors = factors,
rotate = "none", fm = "pc")
                                                             # Print coordinates of variables
FA.Res.Varimax=fa(stocks data, nfactors = factors,
                                                             print(PCA.Res$var$coord)
rotate = "varimax", fm = "pc")
                                                             # Create a scree plot
FA.Res.Quartimax=fa(stocks_data, nfactors = factors,
rotate = "quartimax", fm = "pc")
                                                             fviz_eig(PCA.Res, addlabels = TRUE, ylim = c(0, 70))
# Print the results
                                                             # Calculate factor loadings using PCA.Res
                                                             factor_loadings=round(sweep(PCA.Res$var$coord,
print(FA.Res.None)
                                                             2, sqrt(PCA.Res$eig[1:ncol(PCA.Res$var$coord), 1]),
                                                             FUN = "/"), 2)
print(FA.Res.Varimax)
print(FA.Res.Quartimax)
                                                             factor loadings
# Create factor diagrams for different rotation
                                                             # Perform PCA using prcomp function
methods
                                                             PCA.Res1=prcomp(Standardized_Data, scale. =
par(mfrow = c(1, 1))
                                                             TRUE)
fa.diagram(FA.Res.None, cut = 0.3, simple = FALSE,
                                                             round(PCA.Res1$rotation, 3)
main = "No Rotation")
                                                             # Analysis B1
fa.diagram(FA.Res.Varimax, cut = 0.3, simple =
FALSE, main = "Varimax Rotation")
                                                             # Load required libraries for Analysis B1
                                                             source ("projectfile.R")
fa.diagram(FA.Res.Quartimax, cut = 0.3, simple =
FALSE, main = "Quartimax Rotation")
                                                             library (heplots)
# Print selected companies for each factor
```

```
library (MVN)
library ( DescTools )
                                                           # Plot gg plots and perform multivariate normality
                                                           tests for population with heart disease
#read excel
                                                           cqplot(H2, id.n = 3, main = "Chisquare qq plot for
Healthsleep data =
                                                           female")
read excel("Healthsleepdata.xlsx")
                                                           mvn(H2, mvnTest = "royston")
head (Healthsleep data, 5) # display the first 5 rows
                                                           mvn(H2, mvnTest = "hz")
# Subset the data gender
                                                           # Reset plotting layout
H1=subset(data.frame(Healthsleep_data$Age,
Healthsleep_data$SleepDuration,
                                                           par(mfrow = c(1, 1))
Healthsleep data$HeartRate),
Healthsleep data$Gender == "Male")
                                                           # Perform Hotellings T2 Test
                                                           HotellingsT2Test (H1, H2, test = "chi")
n1=nrow(H1)
H1Bar=c(mean(H1[,1]), mean(H1[,2]), mean(H1[,3]))
                                                           library(mvtnorm)
SCOV1=cov(H1)
                                                           # Perform simultaneous confidence intervals
# Subset the data for individuals with heart disease
                                                           MVN2Sample.HT.Cls.NumSum.f(n1, H1Bar, SCOV1,
                                                           n2, H2Bar, SCOV2, conf.level=0.95, alpha = .05,
                                                           mu0 = rep (0, length( H1Bar )), ContrastMAT = NULL,
                                                           SigDig =3, var.eq = TRUE)
H2=subset(data.frame(Healthsleep_data$Age,
                                                           #b2 Analysis
Healthsleep data$SleepDuration,
Healthsleep data$HeartRate),
                                                           # Subset the data for different occupation types and
Healthsleep_data$Gender == "Female")
                                                           visualize multivariate normality
n2=nrow(H2)
                                                           new data=data.frame(Healthsleep data$Occupatio
                                                           n, Healthsleep data$Age,
H2Bar=c(mean(H2[,1]), mean(H2[,2]), mean(H2[,3]))
                                                           Healthsleep data$SleepDuration,
                                                           Healthsleep data$HeartRate)
SCOV2=cov(H2)
                                                           # Occupation type Accountant
# Set up plotting layout
                                                           O1=subset(data.frame(Healthsleep data$Age,
par(mfrow = c(1, 2))
                                                           Healthsleep_data$SleepDuration,
                                                           Healthsleep data$HeartRate),
# Plot gg plots and perform multivariate normality
                                                           Healthsleep data$Occupation == "Accountant")
tests for population without heart disease
                                                           cqplot(O1, id.n = 3, main = "Chisquare qq plot of
cqplot(H1, id.n = 3, main = "Chisquare qq plot for
                                                           Occupation Type Accountant")
Male ")
                                                           # Occupation type Doctor
mvn(H1, mvnTest = "royston")
                                                           O2= subset(data.frame(Healthsleep_data$Age,
mvn(H1, mvnTest = "hz")
                                                           Healthsleep_data$SleepDuration,
```

```
Healthsleep data$HeartRate),
                                                           VMat=statList(new data[, -1], new data[, 1], FUN =
Healthsleep data$Occupation == "Doctor")
                                                           var)
cqplot(O2, id.n = 3, main = "Chisquare qq plot of
                                                           Ns=table(new_data[, 1]) # Sample sizes
Occupation Type Doctor")
                                                           p=3 # Number of variables
# Occupation type lawyer
                                                           g=3 # Number of groups
O3=subset(data.frame(Healthsleep_data$Age,
Healthsleep data$SleepDuration,
                                                           n=nrow(new data)
Healthsleep data$HeartRate),
                                                           # Manually computing the W matrix
Healthsleep data$Occupation == "Lawyer")
cqplot(O3, id.n = 3, main = "Chisquare qq plot of
                                                           W=(Ns[1] - 1) * VMat[[1]] + (Ns[2] - 1) * VMat[[2]] +
Occupation Type lawyer")
                                                           (Ns[3] - 1) * VMat[[3]]
# Plot qq plots and perform multivariate normality
                                                           # Load necessary packages
tests for population without heart disease
                                                           library(heplots)
cqplot(O1, id.n = 3, main = "chi")
                                                           library(MVN)
mvn(O1, mvnTest = "royston")
                                                           # Split data by group
mvn(O1, mvnTest = "hz")
                                                           new data gr=split(new data, new data[, 1])
#ANOVA test for Age
                                                           # Checking Multivariate normality for each group
anova age=aov(Healthsleep data$Age~
Healthsleep_data$Occupation)
                                                           par(mfrow = c(2, 2))
                                                           for(i in 1:g){
summary(anova_age)
# ANOVA test for sleepduration
                                                            X = new_data_gr[[i]][, -1]
anova SleepDuration=aov(Healthsleep data$SleepD
                                                            group name = names(new data gr)[i] # Get the
                                                           group name
uration ~ Healthsleep_data$Occupation)
                                                            cqplot(X, id.n=3, main = paste("Chi-square QQ plot
summary(anova SleepDuration)
                                                           for", group name))
# ANOVA test for hr
                                                            print(mvn(X, mvnTest = "royston"))
anova HeartRate=aov(Healthsleep data$HeartRate
~ Healthsleep_data$Occupation)
                                                            print(mvn(X, mvnTest = "hz"))
summary(anova HeartRate)
                                                           # MANOVA with built in function
# Manova with car package
# Computing Means and Var/Cov Mats per groups
                                                           Y = as.matrix(new_data[, -1])
                                                           Gr = as.factor(new data[, 1])
VMeans=statList(new_data[, -1], new_data[, 1], FUN
= colMeans)
                                                           # First fit a linear regression
```

```
LM.res = Im(Y \sim Gr)
                                                                v = 1 # Variable
# Call Manova, This command gives all Tests
                                                                LL = VMeans [[i]] - VMeans [[j]] - ta *sqrt ((1 /Ns[i] +
                                                                 1/Ns[j]) * diag (W) / (n - g))
# Wilk's lambda, Pillai, Hotelling-Lawley and Roy
                                                                UL = VMeans [[i]] - VMeans [[j]] + ta *sqrt ((1 /Ns[i] +
SUM = summary(Manova(LM.res), "Wilks")
                                                                 1/Ns[j]) * diag (W) / (n - g))
SUM
                                                                cbind (LL, UL)
# The W matrix using the build it functions
                                                                i = 1 # Group 2
W = SUM$SSPE
                                                                j = 2 # Group 3
# Check which groups differ from each other
                                                                v = 1 # Variable
library(biotools)
                                                                LL = VMeans [[i]] - VMeans [[j]] - ta *
mvpaircomp(LM.res, factor1 = "Gr", test = "Wilks",
                                                                 sqrt ((1/Ns[i] + 1/Ns[j]) * diag (W) / (
adjust = "bonferroni")
                                                                   n - g))
                                                                 UL = VMeans [[i]] - VMeans [[j]] + ta *
# Check each variable for differences across groups
                                                                  sqrt ((1/Ns[i] + 1/Ns[j]) * diag (W) / (
summary.aov(LM.res)
                                                                   n - g))
# Calculate simultaneous confidence intervals
                                                                 cbind (LL, UL)
k=p * g * (g - 1) / 2
ta=qt(0.05 / (2 * k), df = n - g, lower.tail = FALSE)
# Simultaneous CIs
i = 1 # Group 1
j = 2 # Group 2
v = 1 # Variable
LL = VMeans [[i]] - VMeans [[j]] - ta *sqrt ((1 /Ns[i] +
1/Ns[j]) * diag (W) / (n - g))
UL = VMeans [[i]] - VMeans [[j]] + ta *sqrt ((1 /Ns[i] +
1/Ns[j]) * diag (W) / (n - g))
cbind (LL, UL)
i = 1 # Group 1
j = 3 # Group 3
```

Sign the statement below and attach it with your project. You project will not be graded without it.

This project is entirely my work. I have not discussed this project with anybody in or out of class. I did not use any unauthorized sources such as help from tutoring services (in person or online), nor have I used generative AI/Online assignment assisting websites (such as Chat GPT, DALL-E, Chegg, course Hero, etc.). I understand and have complied with the Rowan University's academic integrity policies outlined in (https://confluence.rowan.edu/display/POLICY/Academic+Integrity+Policy).

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NAME: Akhila Vitta

DATE: 04/13/2024

SIGNATURE: V.Akhila