Data211-Midterm Exam

Peyton Hall

02/29/2024

Required libraries

```
library(ggplot2)
library(readxl)
```

Question 01

```
# Open R Markdown
```

```
# Create R data structure:
# a) Create and print a numeric vector
x <- c(1, 3, 5, 12, 20)
x
## [1] 1 3 5 12 20
# b) Create and print a character vector
y <- c("red", "yellow", "blue")
y

## [1] "red" "yellow" "blue"
# c) Create and print a list, z.
z <- list(a = x, b = y)
z

## $a
## [1] 1 3 5 12 20
## $b
## [1] "red" "yellow" "blue"</pre>
```

```
# d) Convert the vector y in b) to a factor
y_factor <- factor(y)
y_factor

## [1] red yellow blue
## Levels: blue red yellow</pre>
```

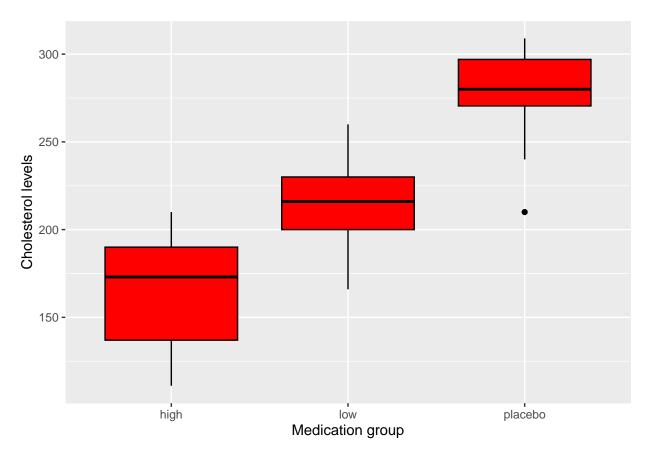
Question 03

```
# Create a function named triangle_identifier with the following requirements:
# a) Have three inputs (arguments): x1, x2, and x3. The x1, x2, and x3 are the
    lengths of the three sides of a triangle.
triangle_identifier <- function(x1, x2, x3) {</pre>
  # stub
# b) Include if-else statements in the function to show whether the x1, x2 and
   x3 can create a triangle.
    • If x1+x2 < x3, then it is not a triangle.
     • If x1+x3<x2, then it is not a triangle.
     • If x2+x3 < x1, then it is not a triangle.
    • Otherwise, it is a triangle.
   Return the result of whether it is a triangle.
triangle_identifier <- function(x1, x2, x3) {</pre>
  if (x1 + x2 < x3 || x1 + x3 < x2 || x2 + x3 < x1) {
    result <- "It is not a triangle"</pre>
  } else {
    result <- "It is a triangle"
 return(result)
# c) Call the function using x1=3, x2=5, x3=6 and print the output; and call the
    function using x1=2, x2=3, x3=7 and print the output.
output1 <- triangle_identifier(3, 5, 6)</pre>
output1
## [1] "It is a triangle"
```

```
output2 <- triangle_identifier(2, 3, 7)
output2</pre>
```

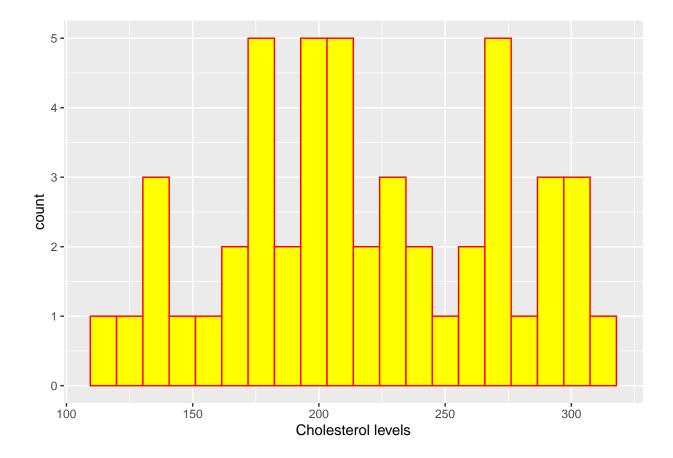
[1] "It is not a triangle"

```
# The data Cholesterol on D2L recorded 49 high cholesterol patients' cholesterol
# levels after taking different medications. Among those 49 patients, some of
# them were assigned to take the placebo, some were assigned to take the
# low-dosage medication, and the rest took high dosages.
# press "Import Dataset" -> press "From Excel..." -> press "Browse..." ->
# navigate & select .xlsx file -> copy (command + c) code from "Code Preview:"
# -> press "Import" -> paste (command + v) code to the RStudio Integrated
# Development Environment.
Cholesterol <- read_excel("/Users/peytonhall/Desktop/Data211Midterm/Cholesterol.xlsx")</pre>
# View(Cholesterol)
# a) Use ggplot() to generate an appropriate graph to show the five-number
     summary of cholesterol levels at each medication group. Label the x-axis as
     "Medication group" and y as "Cholesterol levels."
     Color the inside red and the borders black.
ggplot(Cholesterol, aes(x = Dosage, y = Chol)) +
 labs(x = "Medication group", y = "Cholesterol levels") + # labels
 geom_boxplot(fill = "red", color = "black") # red insides & black borders
```



```
# the boxplot by the dot in the placebo column on the x axis.

# c) Use ggplot() to generate an appropriate graph to show the distribution of
# all cholesterol levels regardless of medication groups, with 20 bins. Label
# the x axis as "Cholesterol levels". Color the border with red and the
# inside of the bars with yellow.
# create a histogram because bars are required
ggplot(Cholesterol, aes(x = Chol)) +
labs(x = "Cholesterol levels") + # label the x axis
geom_histogram(fill = "yellow", color = "red", bins = 20) # red border
```

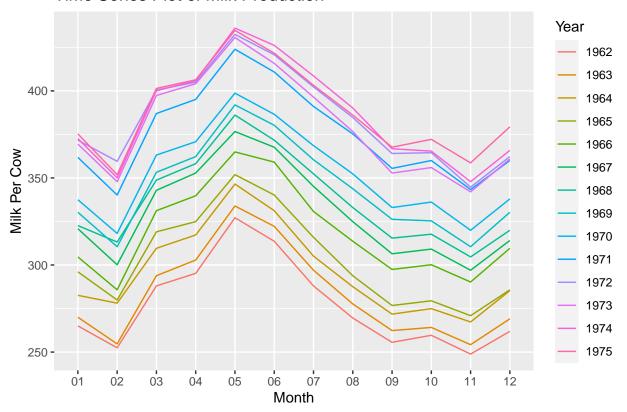


```
# Create the following table as a data frame
# and complete the following questions:
# a) Create the data frame with the name PatientInfo and print it
#
     PatientID, Diastolic Blood Pressure (DBP), Systolic Blood Pressure (SBP)
     Α,
                90.
                                                 120
#
     Β,
                98,
                                                 135
     С,
                76,
                                                  109
     D,
                112,
                                                 141
PatientInfo <- data.frame(
```

```
PatientID = c("A", "B", "C", "D"),
  Diastolic_Blood_Pressure_DBP = c(90, 98, 76, 112),
  Systolic_Blood_Pressure_SBP = c(120, 135, 109, 141)
PatientInfo
     PatientID Diastolic_Blood_Pressure_DBP Systolic_Blood_Pressure_SBP
## 1
             Α
                                          90
                                                                      120
## 2
             В
                                          98
                                                                      135
             С
## 3
                                          76
                                                                      109
## 4
                                         112
                                                                      141
# b) Use rbind() to add one row of new individuals with the information:
    PatientID: E
#
     Diastolic Blood Pressure: 118
     Systolic Blood Pressure: 129.
    Name the new data frame with the new E as BloodPressure and print it.
new_individual <- data.frame(</pre>
  PatientID = "E",
  Diastolic_Blood_Pressure_DBP = 118,
  Systolic_Blood_Pressure_SBP = 129
BloodPressure <- rbind(PatientInfo, new_individual) # row bind adds the new row
BloodPressure # print the updated result
    PatientID Diastolic_Blood_Pressure_DBP Systolic_Blood_Pressure_SBP
##
## 1
             Α
                                          90
## 2
             В
                                          98
                                                                      135
## 3
             С
                                          76
                                                                      109
## 4
                                         112
                                                                      141
             D
## 5
                                         118
                                                                      129
# c) Add a new column named Age to the above data frame (BloodPressure)
   and name the new data frame as NewPatientInfo.
     Following are the data in column Age: 28, 35,42, 40, 39.
Age \leftarrow c(28, 35, 42, 40, 39)
BloodPressure$Age <- Age # adds a new column
NewPatientInfo <- BloodPressure # assigns new data to a new data frame
NewPatientInfo # print the new data frame
     PatientID Diastolic_Blood_Pressure_DBP Systolic_Blood_Pressure_SBP Age
##
## 1
             Α
                                          90
                                                                      120 28
## 2
             В
                                          98
                                                                      135 35
## 3
             С
                                          76
                                                                      109 42
## 4
                                         112
                                                                      141 40
             D
## 5
                                         118
                                                                      129 39
```

```
# Use the milk data to generate a time series plot.
# press "Import Dataset" -> press "From Excel..." -> press "Browse..." ->
# navigate & select .xlsx file -> copy (command + c) code from "Code Preview:"
# -> press "Import" -> paste (command + v) code to the RStudio Integrated
# Development Environment.
milk <- read_excel("/Users/peytonhall/Desktop/Data211Midterm/milk.xlsx")</pre>
# View(milk)
# a) Use the following to extract the month information and year information:
milk$year<-format( milk$timep , format="%Y")</pre>
milk$month<-format( milk$timep , format="%m")</pre>
# b) Generate a time series plot with the axis being a month and the
     axis being milk per cow and use color to show different years.
ggplot(milk, aes(x = month, y = milk_per_cow_kg)) +
  geom_line(aes(group = year, color = year)) + # aes function to represent the grouping and the columns
  labs(x = "Month", y = "Milk Per Cow", color = "Year") +
  ggtitle("Time Series Plot of Milk Production")
```

Time Series Plot of Milk Production

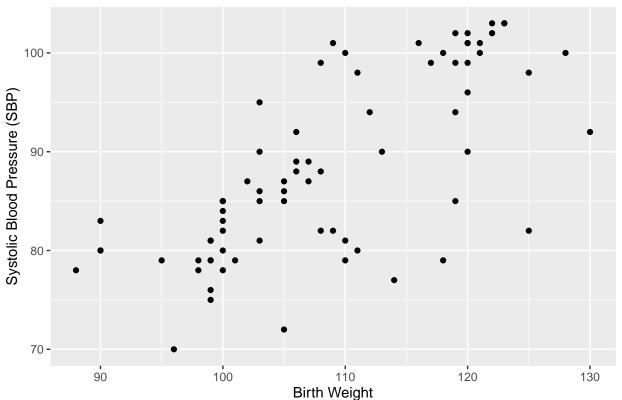


```
# The data birthweightSBP1 on D2L recorded the systolic blood pressure of 78
# children, together with their birth weight and age. Use the data
# birthweightSBP on D2L to do the following:

# press "Import Dataset" -> press "From Excel..." -> press "Browse..." ->
# navigate & select .xlsx file -> copy (command + c) code from "Code Preview:"
# -> press "Import" -> paste (command + v) code to the RStudio Integrated
# Development Environment.
birthweightSBP1 <- read_excel("/Users/peytonhall/Desktop/Data211Midterm/birthweightSBP1.xlsx")
# View(birthweightSBP1)

# a) Use ggplot() to generate an appropriate graph to show the relationship
# between birth weight(x) and SBP (y).
ggplot(birthweightSBP1, aes(x = Birthweight, y = SBP)) +
geom_point() +
labs(x = "Birth Weight", y = "Systolic Blood Pressure (SBP)") +
ggtitle("Relationship Between Birth Weight and SBP")</pre>
```

Relationship Between Birth Weight and SBP

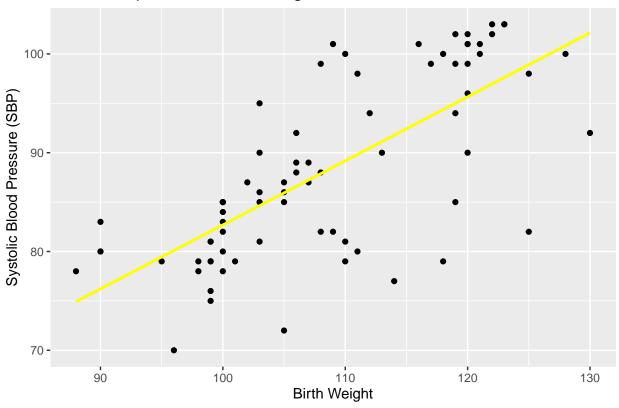


```
# b) Use appropriate functions in the ggplot2 to generate a regression line to
# the graph. Color the line yellow without
# showing the confidence interval band.
ggplot(birthweightSBP1, aes(x = Birthweight, y = SBP)) +
    geom_point() +
    geom_smooth(method = "lm", color = "yellow", se = FALSE) + # regression line
```

```
labs(x = "Birth Weight", y = "Systolic Blood Pressure (SBP)") +
ggtitle("Relationship Between Birth Weight and SBP")
```

'geom_smooth()' using formula = 'y ~ x'

Relationship Between Birth Weight and SBP



```
# Use the USMerchants data on D2L. The dataset recorded the amount of US
# merchants of Corn, Rice, and Soybeans from 1970 to 1994. However, the data
# contains some mistakes. Use the for-loop and if-else conditional statement to
# do the following fix-up:

# press "Import Dataset" -> press "From Excel..." -> press "Browse..." ->
# navigate & select .xlsx file -> copy (command + c) code from "Code Preview:"
# -> press "Import" -> paste (command + v) code to the RStudio Integrated
# Development Environment.

USMerchants <- read_excel("/Users/peytonhall/Desktop/Data211Midterm/USMerchants.XLSX")
# View(USMerchants)

# a) For the years from 1970 to 1979, the corn prices
# are actually the original amount +10
for (i in 1:nrow(USMerchants)) { # loop through the rows</pre>
```

```
# if the year is between 1970 and 1979
 if (USMerchants$Year[i] >= 1970 && USMerchants$Year[i] <= 1979) {
    # Add 10 to the corn prices for those years
   USMerchants$Corn[i] <- USMerchants$Corn[i] + 10</pre>
 } # end if
} # end for
USMerchants
## # A tibble: 300 x 4
##
      Year Corn Rice Soybeans
##
      <dbl> <dbl> <dbl>
  1 1970 137. 91.7
                           70.4
##
##
   2 1970 137. 91.9
                           70.6
## 3 1970 137. 92.1
                           70.7
## 4 1970 137. 92.2
                          71.2
## 5 1970 138. 92.0
                           71.4
## 6 1970 139. 91.9
                           71.5
## 7 1970 139. 92.0
                          71.6
## 8 1970 139. 93.3
                           71.7
## 9 1970 138. 93.9
                           71.8
## 10 1970 138. 94.1
                           71.9
## # i 290 more rows
# b) For the years from 1980 to 1989, the rice prices
    are actually the original amount*1.05
for (i in 1:nrow(USMerchants)) { # loop through the rows
 # if the year is between 1980 and 1989
 if (USMerchants$Year[i] >= 1980 && USMerchants$Year[i] <= 1989) {
    # Multiply the rice prices by 1.05 for those years
   USMerchants$Rice[i] <- USMerchants$Rice[i] * 1.05</pre>
 } # end if
} # end for
USMerchants
## # A tibble: 300 x 4
##
      Year Corn Rice Soybeans
      <dbl> <dbl> <dbl>
                          <dbl>
##
  1 1970 137. 91.7
                           70.4
## 2 1970 137. 91.9
                           70.6
## 3 1970 137. 92.1
                           70.7
## 4 1970 137. 92.2
                           71.2
## 5 1970 138. 92.0
                          71.4
## 6 1970 139. 91.9
                          71.5
## 7 1970 139. 92.0
                           71.6
## 8 1970 139. 93.3
                          71.7
## 9 1970 138. 93.9
                          71.8
## 10 1970 138. 94.1
                           71.9
## # i 290 more rows
# c) For the years from 1990 to the end, the soybean prices
    are the original amount-15
for (i in 1:nrow(USMerchants)) { # loop through the rows
```

```
# if the year is between 1990 and 1994
if (USMerchants$Year[i] >= 1990 && USMerchants$Year[i] <= 1994) {
    # Subtract 15 from the soybean prices for those years
    USMerchants$Soybeans[i] <- USMerchants$Soybeans[i] - 15
} # end if
} # end for
USMerchants</pre>
```

```
## # A tibble: 300 x 4
##
      Year Corn Rice Soybeans
     <dbl> <dbl> <dbl>
                        <dbl>
##
  1 1970 137. 91.7
##
                         70.4
   2 1970 137. 91.9
##
                         70.6
## 3 1970 137. 92.1
                         70.7
## 4 1970 137. 92.2
                         71.2
## 5 1970 138. 92.0
                         71.4
## 6 1970 139. 91.9
                         71.5
##
  7 1970 139. 92.0
                         71.6
## 8 1970 139. 93.3
                         71.7
## 9 1970 138. 93.9
                         71.8
## 10 1970 138. 94.1
                         71.9
## # i 290 more rows
```