

RWork-sheet_Sabio#4b.Rmd

2023-11-08

1. Using the for loop, create an R script that will display a 5x5 matrix as shown in Figure 1. It must contain vectorA = [1,2,3,4,5] and a 5 x 5 zero matrix.

```
vectorA<- c(1,2,3,4,5)
matrixA<- matrix( nrow = 5, ncol = 5)

for (i in 1:5){
  for (j in 1:5){
    matrixA[i, j] <- abs(vectorA[i] - vectorA[j])
  }
}
print(matrixA)
```

```
##      [,1] [,2] [,3] [,4] [,5]
## [1,]    0    1    2    3    4
## [2,]    1    0    1    2    3
## [3,]    2    1    0    1    2
## [4,]    3    2    1    0    1
## [5,]    4    3    2    1    0
```

2. Print the string "*" using for() function. The output should be the same as shown in Figure 2.

```
n <- 5
for ( i in 1:n){
  for (j in 1:i){
    cat("*")
  }
  cat("\n")
}
```

```
## *
## **
## ***
## ****
## *****
```

3. Get an input from the user to print the Fibonacci sequence starting from the 1st input up to 500. Use repeat and break statements. Write the R Scripts and its output.

```
fibSequence <- as.integer(readline(prompt = "Enter the number of terms: "))
```

```
## Enter the number of terms:
```

```
a <- 0
b <- 1
cat("Fibonacci Sequence:", a, b)
```

```
## Fibonacci Sequence: 0 1
```

```
repeat {
  c <- a + b
  if (c > 500) {
    break
  }
  cat(" ", c)
  a <- b
  b <- c
}
```

```
## , 1, 2, 3, 5, 8, 13, 21, 34, 55, 89, 144, 233, 377
```

4. Import the dataset as shown in Figure 1 you have created previously. a. What is the R script for importing an excel or a csv file? Display the first 6 rows of the dataset? Show your codes and its result

```
install.packages("readxl")
```

```
## Installing package into '/cloud/lib/x86_64-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
```

```
library(readxl)
data <- read_excel("Shoe Sizes.xlsx")
```

```
## New names:
## * 'Height' -> 'Height...2'
## * 'Gender' -> 'Gender...3'
## * 'Height' -> 'Height...5'
## * 'Gender' -> 'Gender...6'
```

```
data
```

```
## # A tibble: 14 x 6
##   'Shoe Size' Height...2 Gender...3 'Shoe size' Height...5 Gender...6
##   <dbl>      <dbl> <chr>      <dbl>      <dbl> <chr>
## 1      6.5      66  F          13        77  M
## 2       9      68  F         11.5      72  M
## 3      8.5     64.5 F          8.5      59  F
## 4      8.5     65  F          5        62  F
## 5     10.5     70  M          10      72  M
## 6       7     64  F          6.5     66  F
## 7      9.5     70  F          7.5     64  F
## 8       9     71  F          8.5     67  M
## 9      13     72  M         10.5     73  M
## 10     7.5     64  F          8.5     69  F
```

```
## 11      10.5      74  M      10.5      72 M
## 12       8.5      67  F       11       70 M
## 13      12       71  M       9        69 M
## 14      10.5      71  M      13       70 M
```

```
head(data, n=6)
```

```
## # A tibble: 6 x 6
##   'Shoe Size' Height...2 Gender...3 'Shoe size' Height...5 Gender...6
##   <dbl>      <dbl> <chr>      <dbl>      <dbl> <chr>
## 1      6.5      66  F       13       77  M
## 2       9      68  F      11.5      72  M
## 3      8.5     64.5 F       8.5      59  F
## 4      8.5     65  F       5       62  F
## 5     10.5     70  M      10      72  M
## 6       7     64  F       6.5     66  F
```

- b. Create a subset for gender(female and male). How many observations are there in Male? How about in Female? Write the R scripts and its output.

```
maleSubset <- subset(data, Gender...3 == "M")
femaleSubset <- subset(data, Gender...3 == "F")

Male <- nrow(maleSubset)
Female <- nrow(femaleSubset)

cat("Number of observations of Male in Gender...3 is: ", Male, "\n")
```

```
## Number of observations of Male in Gender...3 is: 5
```

```
cat("Number of observations of Female in Gender...3 is: ", Female, "\n")
```

```
## Number of observations of Female in Gender...3 is: 9
```

```
maleSubset1 <- subset(data, Gender...6 == "M")
femaleSubset1 <- subset(data, Gender...6 == "F")

Male1 <- nrow(maleSubset1)
Female1 <- nrow(femaleSubset1)

cat("Number of observations of Male in Gender...6 is: ", Male1, "\n")
```

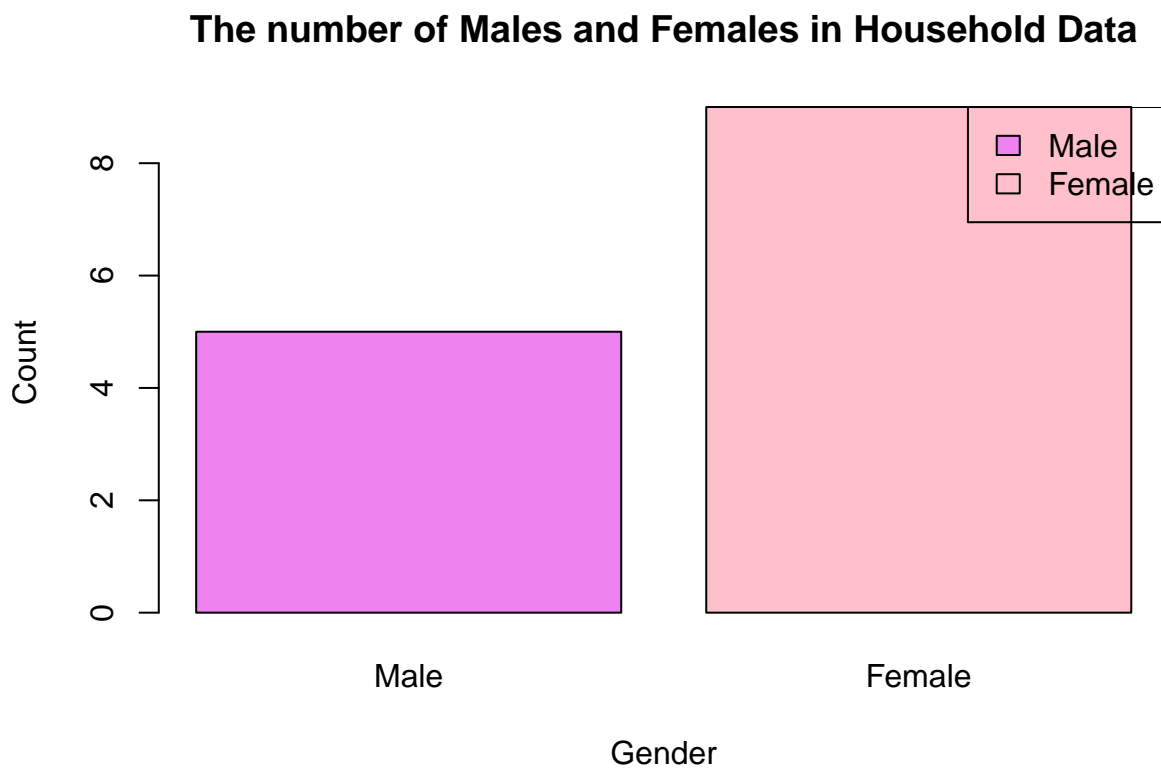
```
## Number of observations of Male in Gender...6 is: 9
```

```
cat("Number of observations of Female in Gender...6 is: ", Female1, "\n")
```

```
## Number of observations of Female in Gender...6 is: 5
```

- c. Create a graph for the number of males and females for Household Data. Use plot(), chart type = barplot. Make sure to place title, legends, and colors. Write the R scripts and its result.

```
count <- c(Male, Female)
gender <- c("Male", "Female")
barplot(count,
names.arg = gender,
main = "The number of Males and Females in Household Data",
xlab = "Gender",
ylab = "Count",
col = c("violet", "pink"))
legend("topright",
legend = gender,
fill = c("violet", "pink"))
```



5a.

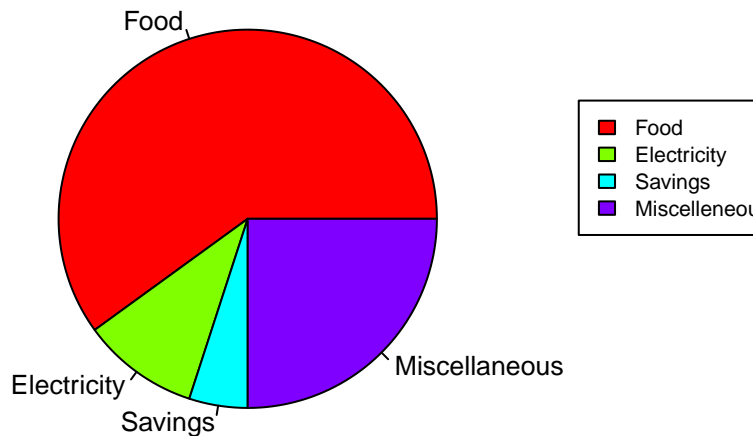
Create a piechart that will include labels in percentage. Add some colors and title of the chart. Write the R scripts and show its output.

```
expenses <- c(60, 10, 5, 25)
categories <- c("Food", "Electricity", "Savings", "Miscellaneous")

percentages <- round((expenses / sum(expenses)) * 100, 1)

pie(expenses, main="Dela Cruz Family Monthly Expenses", col = rainbow(length(expenses)),
labels = categories, cex = 0.8)
legend(1.4, 0.5,
c("Food", "Electricity", "Savings", "Miscellaneous"), cex = 0.7, fill = rainbow(length(expenses)))
```

Dela Cruz Family Monthly Expenses



6a. Check for the structure of the dataset using the `str()` function. Describe what you have seen in the output.

```
data("iris")
str(iris)
```

```
## 'data.frame': 150 obs. of 5 variables:
## $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
## $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
## $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
## $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
## $ Species : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1 ...
```

#It is a data frame that has 150 observation of 5 variables which is sepal.length, sepal.width, petal.length, petal.width, and species.

b. Create an R object that will contain the mean of the sepal.length, sepal.width, petal.length, and petal.width. What is the R script and its result?

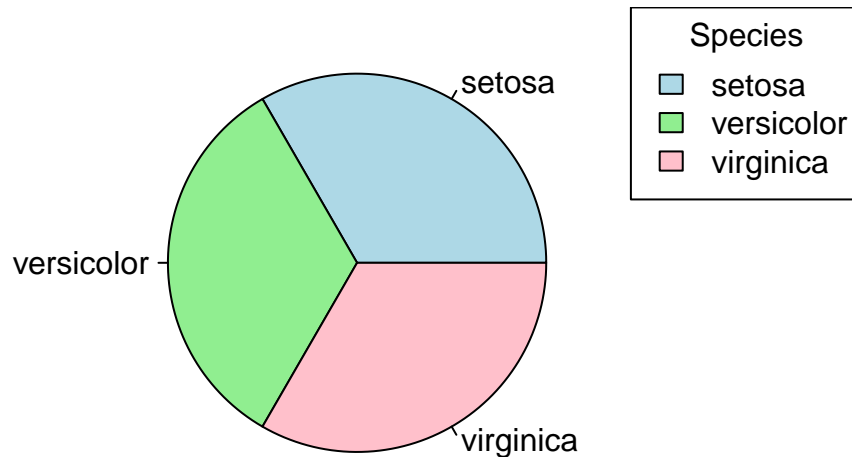
```
meanVal <- c(mean(iris$Sepal.Length), mean(iris$Sepal.Width), mean(iris$Petal.Length), mean(iris$Petal.Width))
meanVal
```

```
## [1] 5.843333 3.057333 3.758000 1.199333
```

c. Create a pie chart for the Species distribution. Add title, legends, and colors. Write the R script and its result.

```
pie(table(iris$Species),
main = "Species distribution",
labels = levels(iris$Species),
col = c("lightblue", "lightgreen", "pink"))
legend("topright", legend = levels(iris$Species), fill = c("lightblue", "lightgreen", "pink"), title = "Species distribution")
```

Species distribution



d. Subset the species into setosa, versicolor, and virginica. Write the R scripts and show the last six (6) rows of each species.

```
subsetSetosa <- tail(subset(iris, Species == "setosa"), 6)
subsetSetosa
```

| ## | Sepal.Length | Sepal.Width | Petal.Length | Petal.Width | Species |
|-------|--------------|-------------|--------------|-------------|---------|
| ## 45 | 5.1 | 3.8 | 1.9 | 0.4 | setosa |
| ## 46 | 4.8 | 3.0 | 1.4 | 0.3 | setosa |
| ## 47 | 5.1 | 3.8 | 1.6 | 0.2 | setosa |
| ## 48 | 4.6 | 3.2 | 1.4 | 0.2 | setosa |
| ## 49 | 5.3 | 3.7 | 1.5 | 0.2 | setosa |
| ## 50 | 5.0 | 3.3 | 1.4 | 0.2 | setosa |

```
subsetVersicolor <- tail(subset(iris, Species == "versicolor"), 6)
subsetVersicolor
```

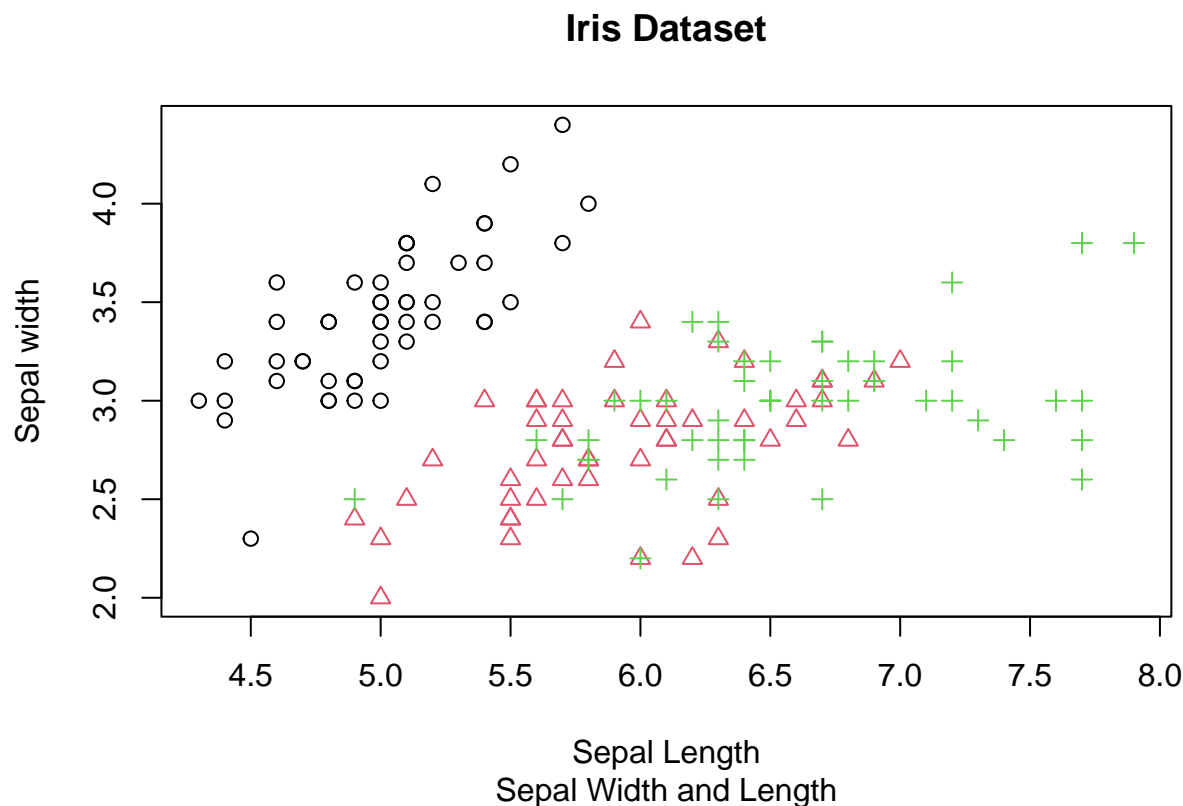
| ## | Sepal.Length | Sepal.Width | Petal.Length | Petal.Width | Species |
|--------|--------------|-------------|--------------|-------------|------------|
| ## 95 | 5.6 | 2.7 | 4.2 | 1.3 | versicolor |
| ## 96 | 5.7 | 3.0 | 4.2 | 1.2 | versicolor |
| ## 97 | 5.7 | 2.9 | 4.2 | 1.3 | versicolor |
| ## 98 | 6.2 | 2.9 | 4.3 | 1.3 | versicolor |
| ## 99 | 5.1 | 2.5 | 3.0 | 1.1 | versicolor |
| ## 100 | 5.7 | 2.8 | 4.1 | 1.3 | versicolor |

```
subsetVirginica <- tail(subset(iris, Species == "virginica"), 6)
subsetVirginica
```

| ## | Sepal.Length | Sepal.Width | Petal.Length | Petal.Width | Species |
|--------|--------------|-------------|--------------|-------------|-----------|
| ## 145 | 6.7 | 3.3 | 5.7 | 2.5 | virginica |
| ## 146 | 6.7 | 3.0 | 5.2 | 2.3 | virginica |
| ## 147 | 6.3 | 2.5 | 5.0 | 1.9 | virginica |
| ## 148 | 6.5 | 3.0 | 5.2 | 2.0 | virginica |
| ## 149 | 6.2 | 3.4 | 5.4 | 2.3 | virginica |
| ## 150 | 5.9 | 3.0 | 5.1 | 1.8 | virginica |

- e. Create a scatterplot of the sepal.length and sepal.width using the different species(setosa,versicolor,virginica). Add a title = “Iris Dataset”, subtitle = “Sepal width and length, labels for the x and y axis, the pch symbol and colors should be based on the species.

```
iris$Species<- as.factor(iris$Species)
plot(iris$Sepal.Length,iris$Sepal.Width,pch = as.integer(iris$Species),col = as.integer(iris$Species), m
```



```
as.factor(iris$Species)
```

```
## [1] setosa setosa setosa setosa setosa setosa
## [7] setosa setosa setosa setosa setosa setosa
## [13] setosa setosa setosa setosa setosa setosa
## [19] setosa setosa setosa setosa setosa setosa
## [25] setosa setosa setosa setosa setosa setosa
## [31] setosa setosa setosa setosa setosa setosa
## [37] setosa setosa setosa setosa setosa setosa
## [43] setosa setosa setosa setosa setosa setosa
## [49] setosa setosa versicolor versicolor versicolor versicolor
## [55] versicolor versicolor versicolor versicolor versicolor versicolor
## [61] versicolor versicolor versicolor versicolor versicolor versicolor
## [67] versicolor versicolor versicolor versicolor versicolor versicolor
## [73] versicolor versicolor versicolor versicolor versicolor versicolor
## [79] versicolor versicolor versicolor versicolor versicolor versicolor
## [85] versicolor versicolor versicolor versicolor versicolor versicolor
## [91] versicolor versicolor versicolor versicolor versicolor versicolor
## [97] versicolor versicolor versicolor versicolor virginica virginica
## [103] virginica virginica virginica virginica virginica virginica
```

```
## [109] virginica virginica virginica virginica virginica virginica
## [115] virginica virginica virginica virginica virginica virginica
## [121] virginica virginica virginica virginica virginica virginica
## [127] virginica virginica virginica virginica virginica virginica
## [133] virginica virginica virginica virginica virginica virginica
## [139] virginica virginica virginica virginica virginica virginica
## [145] virginica virginica virginica virginica virginica virginica
## Levels: setosa versicolor virginica
```

```
## [1] setosa setosa setosa setosa setosa setosa
## [7] setosa setosa setosa setosa setosa setosa
## [13] setosa setosa setosa setosa setosa setosa
## [19] setosa setosa setosa setosa setosa setosa
## [25] setosa setosa setosa setosa setosa setosa
## [31] setosa setosa setosa setosa setosa setosa
## [37] setosa setosa setosa setosa setosa setosa
## [43] setosa setosa setosa setosa setosa setosa
## [49] setosa setosa versicolor versicolor versicolor versicolor
## [55] versicolor versicolor versicolor versicolor versicolor versicolor
## [61] versicolor versicolor versicolor versicolor versicolor versicolor
## [67] versicolor versicolor versicolor versicolor versicolor versicolor
## [73] versicolor versicolor versicolor versicolor versicolor versicolor
## [79] versicolor versicolor versicolor versicolor versicolor versicolor
## [85] versicolor versicolor versicolor versicolor versicolor versicolor
## [91] versicolor versicolor versicolor versicolor versicolor versicolor
## [97] versicolor versicolor versicolor versicolor virginica virginica
## [103] virginica virginica virginica virginica virginica virginica
## [109] virginica virginica virginica virginica virginica virginica
## [115] virginica virginica virginica virginica virginica virginica
## [121] virginica virginica virginica virginica virginica virginica
## [127] virginica virginica virginica virginica virginica virginica
## [133] virginica virginica virginica virginica virginica virginica
## [139] virginica virginica virginica virginica virginica virginica
## [145] virginica virginica virginica virginica virginica virginica
## Levels: setosa versicolor virginica
```

f. Interpret the result.

#the scatterplot result represents the relationship between sepal length and width for different species

7a. Rename the white and black variants by using gsub() function.

```
install.packages("readxl")
```

```
## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.3'
## (as 'lib' is unspecified)
```

```
library(readxl)
alexFile <- read_excel("alexFile.xlsx")

alexFile$variation <- gsub("Black Plus", "Black Show", alexFile$variation)
alexFile$variation <- gsub("Black Show", "Black Show", alexFile$variation)
```



```

alexaFile$variation <- gsub("Black Spot", "Black Spot", alexaFile$variation)
alexaFile$variation <- gsub("Black Dot", "Black Dot", alexaFile$variation)
alexaFile$variation <- gsub("White Dot", "White Dot", alexaFile$variation)
alexaFile$variation <- gsub("White Plus", "White Plus", alexaFile$variation)
alexaFile$variation <- gsub("White Show", "White Show", alexaFile$variation)
alexaFile$variation <- gsub("White Spot", "White Spot", alexaFile$variation)

```

7b. Get the total number of each variations and save it into another object. Save the object as variations.RData. Write the R scripts. What is its result?

```
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

```

```

save(alexaFile, file = "variations.RData")
load("variations.RData")
alexaVar <- alexaFile%>%count(alexaFile$variation)
alexaVar

```

```

## # A tibble: 16 x 2
##   'alexaFile$variation'      n
##   <chr>                  <int>
## 1 Black                  261
## 2 Black Dot              516
## 3 Black Plus             270
## 4 Black Show             265
## 5 Black Spot             241
## 6 Charcoal Fabric        430
## 7 Configuration: Fire TV Stick 350
## 8 Heather Gray Fabric    157
## 9 Oak Finish              14
## 10 Sandstone Fabric       90
## 11 Walnut Finish          9
## 12 White                  91
## 13 White Dot              184
## 14 White Plus             78
## 15 White Show            85
## 16 White Spot            109

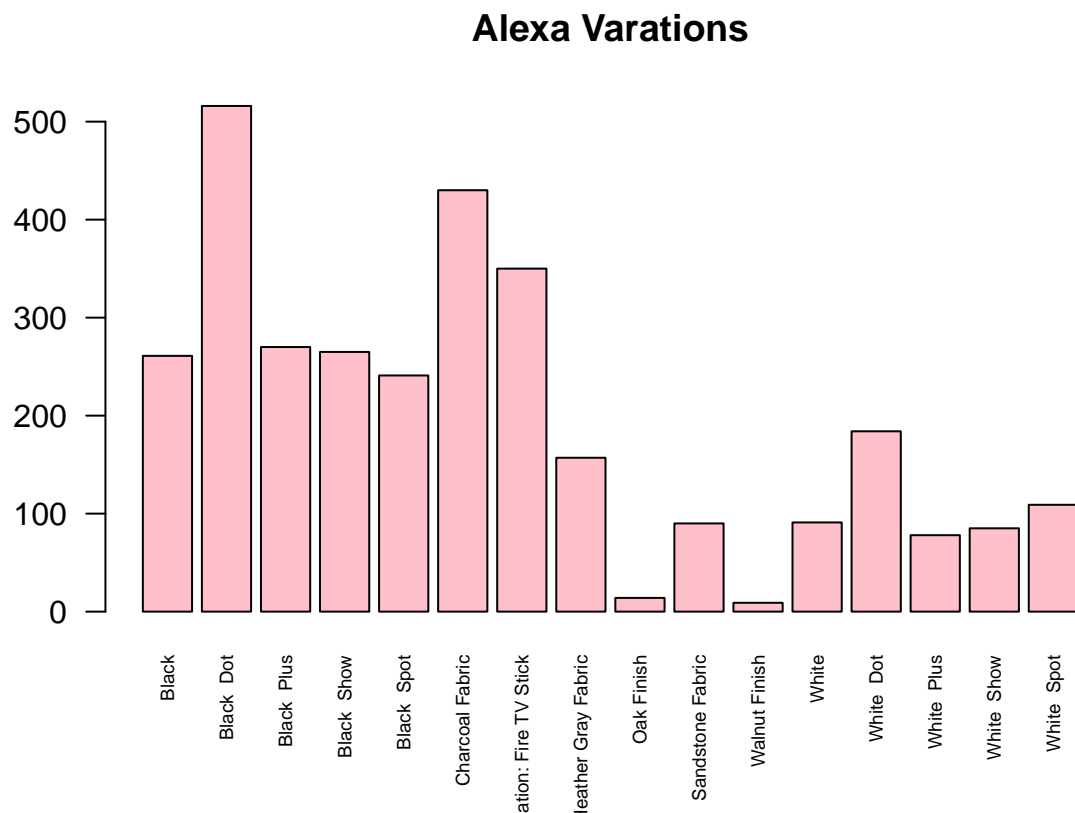
```

7c. From the variations.RData, create a barplot(). Complete the details of the chart which include the title, color, labels of each bar.

```

barplot(
  height = alexaVar$n,
  names.arg = alexaVar$`alexaFile$variation`,
  col = "pink",
  main = "Alexa Variations",
  las = 2,
  cex.names = 0.58
)

```



7d. Create a barplot() for the black and white variations. Plot it in 1 frame, side by side. Complete the details of the chart.

```

par(mfrow = c(1, 2))
blackVariants <- alexaVar[1:5,]
whiteVariants <- alexaVar[12:16,]
barplot(
  height = blackVariants$n,
  names.arg = blackVariants$`alexaFile$variation`,
  main = "Black Variants",
  col = rainbow(5),
  xlab = 'Total Numbers',
  ylab = 'Frequency',
  cex.names = 0.35,
)
barplot(
  height = whiteVariants$n,
  names.arg = whiteVariants$`alexaFile$variation`,
  main = "White Variants",

```

```
col = rainbow(5),
xlab = 'Total Numbers',
ylab = 'Frequency',
cex.names = 0.35,
)
```

