

Rworksheet_Castigador

2023-11-17

1

```
vectorA <-c (1,2,3,4,5)
seroMatrix<- matrix(0,nrow=5, ncol=5)

for (i in 1:5){
  for (j in 1:5){

    yne <- abs(vectorA[i]-j)

    cat(yne," ")
  }
  cat ("\n")
}
```

```
## 0  1  2  3  4
## 1  0  1  2  3
## 2  1  0  1  2
## 3  2  1  0  1
## 4  3  2  1  0
```

2

```
for (i in 1:5){
  yne <- rep ("*",i)
  print(yne)
}
```

```
## [1] "*"
## [1] "*" "*"
## [1] "*" "*" "*"
## [1] "*" "*" "*" "*"
## [1] "*" "*" "*" "*" "*"
```

3

```
userInp <- as.numeric(readline("Enter a number to start the Fibonacci sequence: "))

## Enter a number to start the Fibonacci sequence:

a <- 0
b <- 1

cat("Fibonacci sequence starting from", userInp, ": ")

## Fibonacci sequence starting from NA :

cat(userInp, " ")
```

```
## NA
```

```

repeat {

  nextFib <- a + b

  if (nextFib > 500) {
    break
  }

  cat(nextFib, " ")

  a <- b
  b <- nextFib
}

```

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

4

```
shoesizes <- read.csv("Shoe_Sizes")
```

4b

```

Male <- shoesizes[shoesizes$Gender=="M",]
Female <- shoesizes[shoesizes$Gender=="F",]

male_Count <- nrow(Male)
female_Count <- nrow(Female)

cat("Numbers of Male: ", male_Count, "\n")

```

```
## Numbers of Male: 14
```

```
cat("Numbers of Female: ", female_Count, "\n")
```

```
## Numbers of Female: 14
```

4c

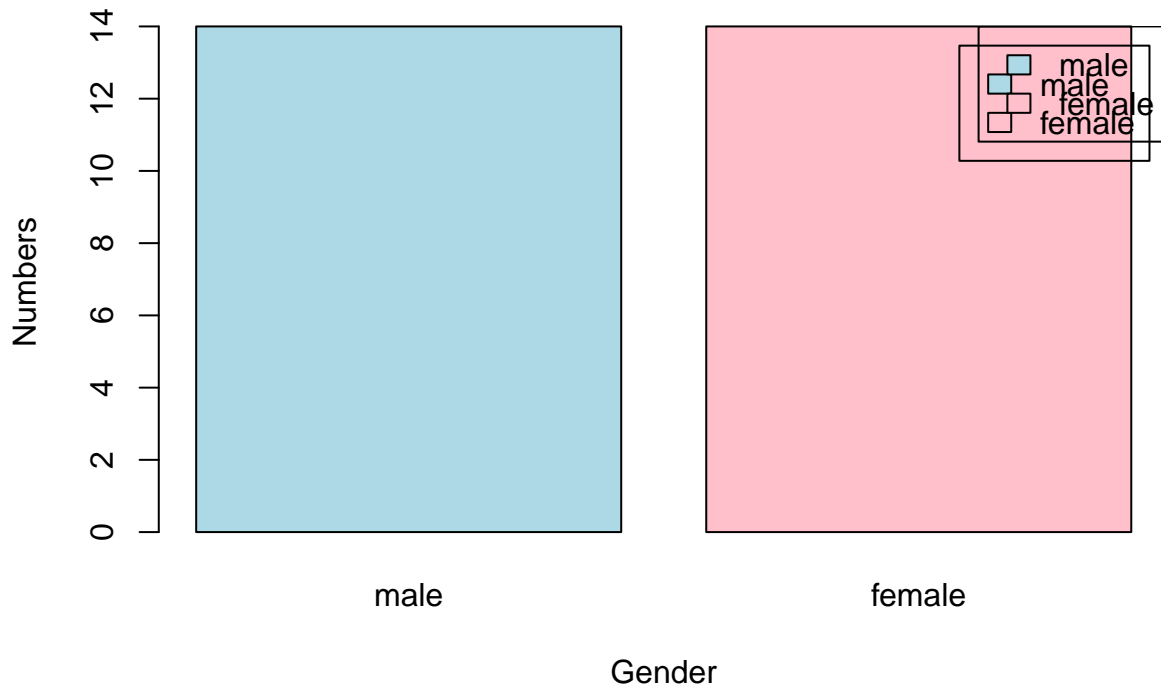
```

gender_Count <- c(male = male_Count, female = female_Count)

barplot(gender_Count, main = "Numbers of Male and Female",
        xlab = "Gender", ylab = "Numbers", col = c("lightblue", "pink"),
        legend.text = TRUE)
legend("topright", legend = names(gender_Count), fill = c("lightblue", "pink"))

```

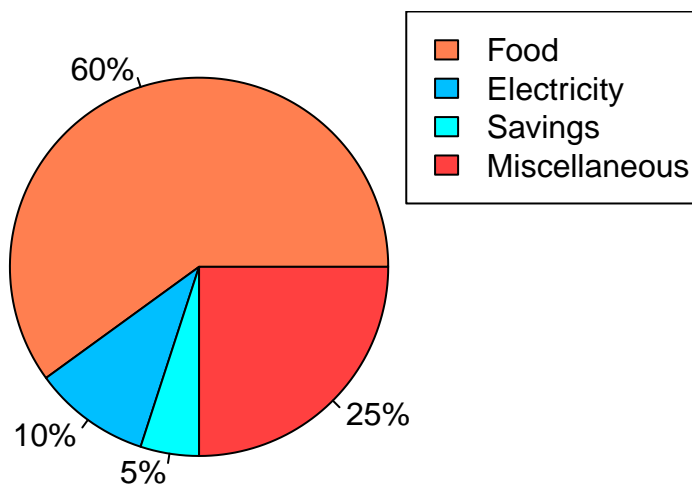
Numbers of Male and Female



5

```
# Define the data
dapie <- c(60, 10, 5, 25)
#Create the pie chart
pie(dapie,
    main = "Monthly Income of Dela Cruz family",
    col = c("coral", "deepskyblue", "cyan", "brown1"),
    labels = c("60%", "10%", "5%", "25%"),
)
legend("topright", legend = c("Food", "Electricity", "Savings", "Miscellaneous"), fill = c("coral", "deepskyblue", "cyan", "brown1"))
```

Monthly Income of Dela Cruz family



6

```
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 ...
```

#This output shows the structure of the data(iris) that contains the sepal length, sepal width, petal length, petal width, and species

6b.

```
lengthsep <- mean(iris$Sepal.Length)

widthsep <- mean(iris$Sepal.Width)

lengthpe <- mean(iris$Petal.Length)

widthpe <- mean(iris$Petal.Width)

print(lengthsep)
```

```
## [1] 5.843333
```

```
print(widthsep)
```

```
## [1] 3.057333
```

```
print(lengthpe)
```

```
## [1] 3.758
```

```
print(widthpe)
```

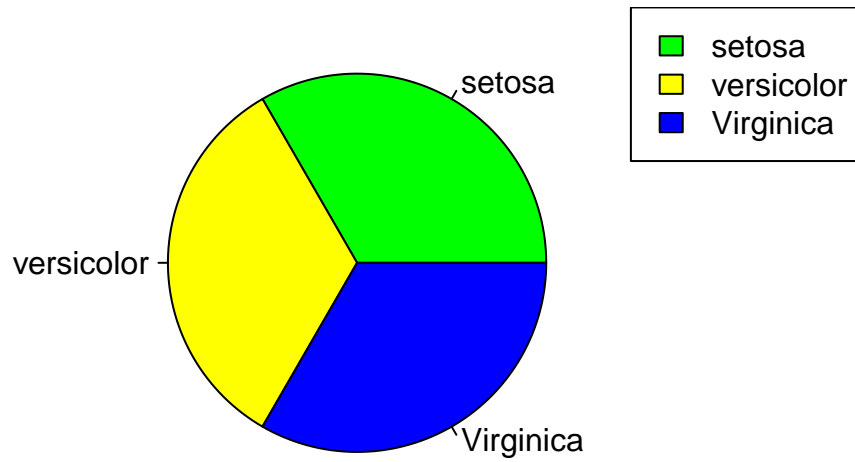
```
## [1] 1.199333
```

6c.

```
specount <- table(iris$Species)

pie(specount,
    main = "Species",
    col = c("green", "yellow", "blue"),
    labels = c("setosa", "versicolor", "Virginica")
)
legend("topright", legend = c("setosa", "versicolor", "Virginica"), fill = c("green", "yellow", "blue"))
```

Species



6d.

```
subSpec <- iris[iris$Species == "setosa" | iris$Species == "Versicolor" | iris$Species == "virginica", ]
subSpec
```

##	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
## 1	5.1	3.5	1.4	0.2	setosa
## 2	4.9	3.0	1.4	0.2	setosa
## 3	4.7	3.2	1.3	0.2	setosa
## 4	4.6	3.1	1.5	0.2	setosa
## 5	5.0	3.6	1.4	0.2	setosa
## 6	5.4	3.9	1.7	0.4	setosa
## 7	4.6	3.4	1.4	0.3	setosa
## 8	5.0	3.4	1.5	0.2	setosa
## 9	4.4	2.9	1.4	0.2	setosa
## 10	4.9	3.1	1.5	0.1	setosa
## 11	5.4	3.7	1.5	0.2	setosa
## 12	4.8	3.4	1.6	0.2	setosa
## 13	4.8	3.0	1.4	0.1	setosa
## 14	4.3	3.0	1.1	0.1	setosa
## 15	5.8	4.0	1.2	0.2	setosa
## 16	5.7	4.4	1.5	0.4	setosa
## 17	5.4	3.9	1.3	0.4	setosa
## 18	5.1	3.5	1.4	0.3	setosa
## 19	5.7	3.8	1.7	0.3	setosa
## 20	5.1	3.8	1.5	0.3	setosa
## 21	5.4	3.4	1.7	0.2	setosa
## 22	5.1	3.7	1.5	0.4	setosa
## 23	4.6	3.6	1.0	0.2	setosa
## 24	5.1	3.3	1.7	0.5	setosa
## 25	4.8	3.4	1.9	0.2	setosa
## 26	5.0	3.0	1.6	0.2	setosa
## 27	5.0	3.4	1.6	0.4	setosa
## 28	5.2	3.5	1.5	0.2	setosa
## 29	5.2	3.4	1.4	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 31	4.8	3.1	1.6	0.2	setosa
## 32	5.4	3.4	1.5	0.4	setosa

## 33	5.2	4.1	1.5	0.1	setosa
## 34	5.5	4.2	1.4	0.2	setosa
## 35	4.9	3.1	1.5	0.2	setosa
## 36	5.0	3.2	1.2	0.2	setosa
## 37	5.5	3.5	1.3	0.2	setosa
## 38	4.9	3.6	1.4	0.1	setosa
## 39	4.4	3.0	1.3	0.2	setosa
## 40	5.1	3.4	1.5	0.2	setosa
## 41	5.0	3.5	1.3	0.3	setosa
## 42	4.5	2.3	1.3	0.3	setosa
## 43	4.4	3.2	1.3	0.2	setosa
## 44	5.0	3.5	1.6	0.6	setosa
## 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
## 101	6.3	3.3	6.0	2.5	virginica
## 102	5.8	2.7	5.1	1.9	virginica
## 103	7.1	3.0	5.9	2.1	virginica
## 104	6.3	2.9	5.6	1.8	virginica
## 105	6.5	3.0	5.8	2.2	virginica
## 106	7.6	3.0	6.6	2.1	virginica
## 107	4.9	2.5	4.5	1.7	virginica
## 108	7.3	2.9	6.3	1.8	virginica
## 109	6.7	2.5	5.8	1.8	virginica
## 110	7.2	3.6	6.1	2.5	virginica
## 111	6.5	3.2	5.1	2.0	virginica
## 112	6.4	2.7	5.3	1.9	virginica
## 113	6.8	3.0	5.5	2.1	virginica
## 114	5.7	2.5	5.0	2.0	virginica
## 115	5.8	2.8	5.1	2.4	virginica
## 116	6.4	3.2	5.3	2.3	virginica
## 117	6.5	3.0	5.5	1.8	virginica
## 118	7.7	3.8	6.7	2.2	virginica
## 119	7.7	2.6	6.9	2.3	virginica
## 120	6.0	2.2	5.0	1.5	virginica
## 121	6.9	3.2	5.7	2.3	virginica
## 122	5.6	2.8	4.9	2.0	virginica
## 123	7.7	2.8	6.7	2.0	virginica
## 124	6.3	2.7	4.9	1.8	virginica
## 125	6.7	3.3	5.7	2.1	virginica
## 126	7.2	3.2	6.0	1.8	virginica
## 127	6.2	2.8	4.8	1.8	virginica
## 128	6.1	3.0	4.9	1.8	virginica
## 129	6.4	2.8	5.6	2.1	virginica
## 130	7.2	3.0	5.8	1.6	virginica
## 131	7.4	2.8	6.1	1.9	virginica
## 132	7.9	3.8	6.4	2.0	virginica
## 133	6.4	2.8	5.6	2.2	virginica
## 134	6.3	2.8	5.1	1.5	virginica
## 135	6.1	2.6	5.6	1.4	virginica
## 136	7.7	3.0	6.1	2.3	virginica

```
## 137      6.3      3.4      5.6      2.4 virginica
## 138      6.4      3.1      5.5      1.8 virginica
## 139      6.0      3.0      4.8      1.8 virginica
## 140      6.9      3.1      5.4      2.1 virginica
## 141      6.7      3.1      5.6      2.4 virginica
## 142      6.9      3.1      5.1      2.3 virginica
## 143      5.8      2.7      5.1      1.9 virginica
## 144      6.8      3.2      5.9      2.3 virginica
## 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
```

```
tail(subSpec, 6)
```

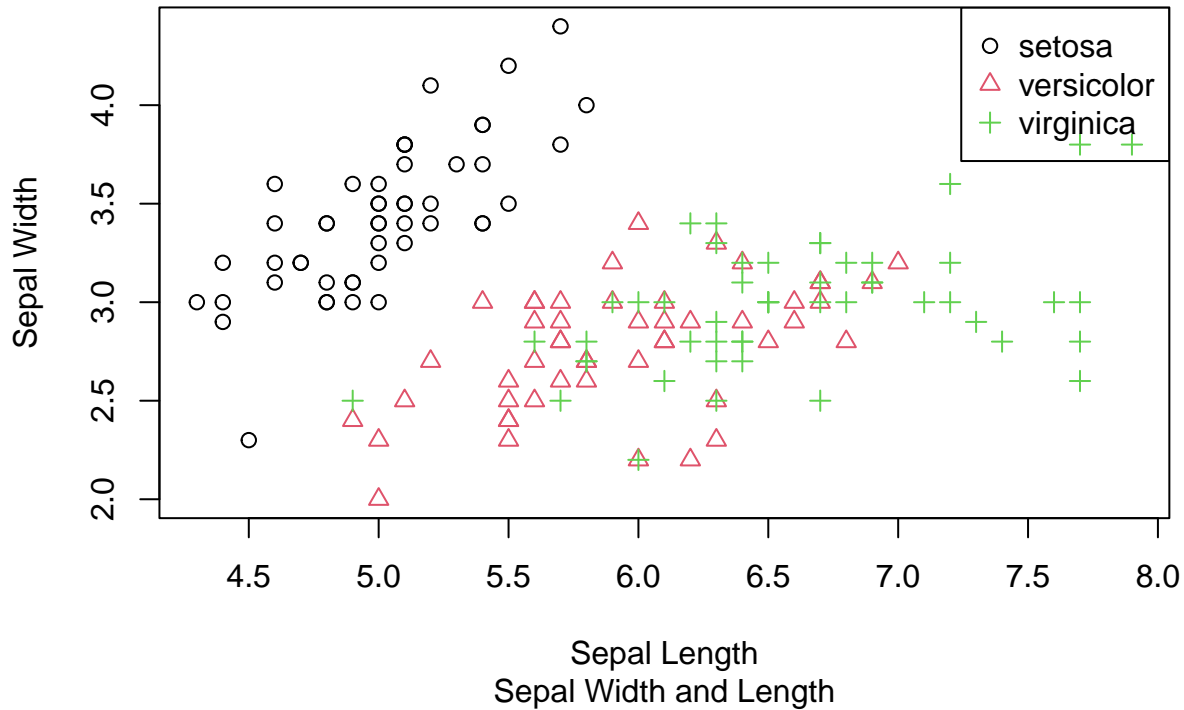
```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width  Species
## 145      6.7      3.3      5.7      2.5 virginica
## 146      6.7      3.0      5.2      2.3 virginica
## 147      6.3      2.5      5.0      1.9 virginica
## 148      6.5      3.0      5.2      2.0 virginica
## 149      6.2  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

6e.

```
plot(iris$Sepal.Length, iris$Sepal.Width,
     col = as.numeric(iris$Species), # Color based on species
     pch = as.numeric(iris$Species), # Different symbol for each species
     main = "Iris Dataset",
     sub = "Sepal Width and Length",
     xlab = "Sepal Length", ylab = "Sepal Width"
)
```

```
legend("topright", legend = levels(iris$Species), col = unique(as.numeric(iris$Species)), pch = unique(
```

Iris Dataset



#6f the scatterplot displays the relationship between the sepal length and width.

7

```
library(readr)
file_show <- read_csv("file_show.csv")

## Rows: 3150 Columns: 5
## -- Column specification -----
## Delimiter: ","
## chr (3): date, variation, verified_reviews
## dbl (2): rating, feedback
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
file_show

## # A tibble: 3,150 x 5
##   rating date      variation verified_reviews feedback
##   <dbl> <chr>    <chr>         <chr>         <dbl>
## 1     5 31-Jul-18 Charcoal Fabric Love my Echo!      1
## 2     5 31-Jul-18 Charcoal Fabric Loved it!          1
## 3     4 31-Jul-18 Walnut Finish Sometimes while playing a game~ 1
## 4     5 31-Jul-18 Charcoal Fabric I have had a lot of fun with t~ 1
## 5     5 31-Jul-18 Charcoal Fabric Music              1
## 6     5 31-Jul-18 Heather Gray Fabric I received the echo as a gift.~ 1
## 7     3 31-Jul-18 Sandstone Fabric Without having a cellphone, I ~ 1
## 8     5 31-Jul-18 Charcoal Fabric I think this is the 5th one I'~ 1
## 9     5 30-Jul-18 Heather Gray Fabric looks great 1
```



```
## 10      5 30-Jul-18 Heather Gray Fabric Love it! I've listened to song~ 1
## # i 3,140 more rows
```

7a.

```
file_show$variation <- gsub("Black Dot", "BlackDot", file_show$variation)
file_show$variation <- gsub("Black Plus", "BlackPlus", file_show$variation)
file_show$variation <- gsub("Black Show", "BlackShow", file_show$variation)
file_show$variation <- gsub("Black Spot", "BlackSpot", file_show$variation)

file_show$variation <- gsub("White Dot", "WhiteDot", file_show$variation)
file_show$variation <- gsub("White Plus", "WhitePlus", file_show$variation)
file_show$variation <- gsub("White Show", "WhiteShow", file_show$variation)
file_show$variation <- gsub("White Spot", "WhiteSpot", file_show$variation)
```

7b.

```
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
```

```
var <- file_show %>%
  count(file_show$variation)
```

var

```
## # A tibble: 16 x 2
##   `file_show$variation`      n
##   <chr>                  <int>
## 1 Black                  261
## 2 BlackDot               516
## 3 BlackPlus              270
## 4 BlackShow              265
## 5 BlackSpot              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 WhiteDot              184
## 14 WhitePlus              78
## 15 WhiteShow             85
## 16 WhiteSpot            109
```

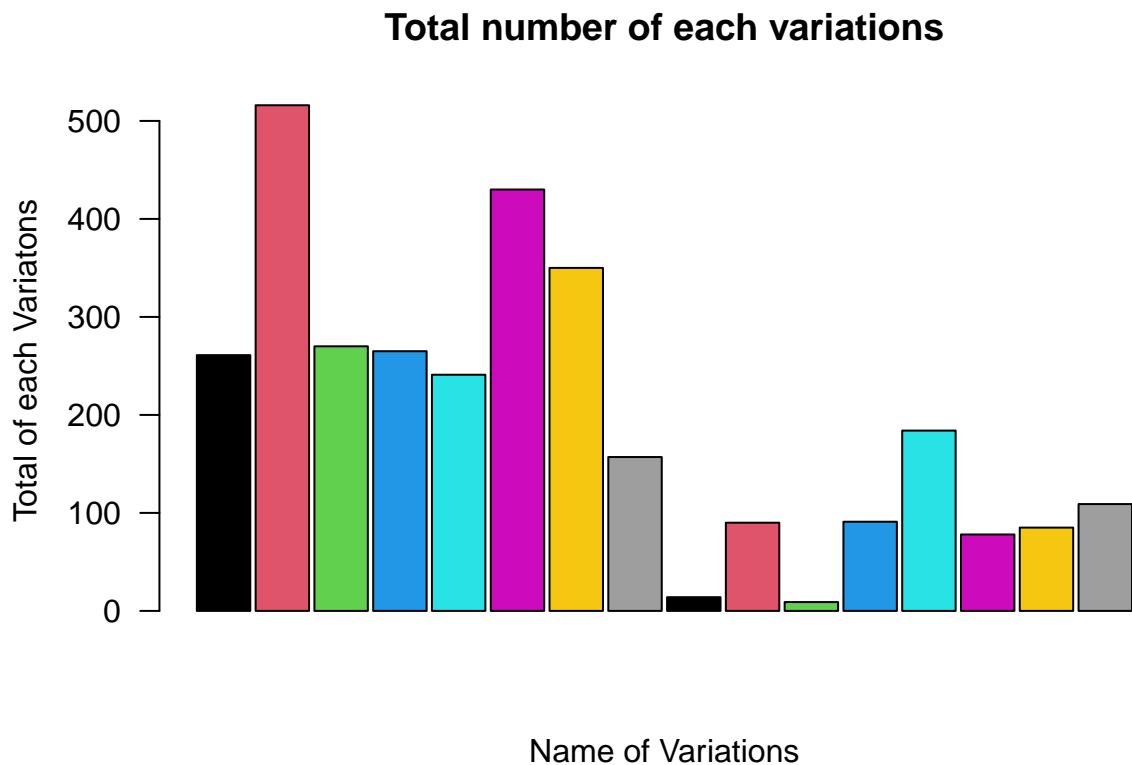
```
save(var, file = "variations.RData")
```

7c.

```
load("variations.RData")
namevar <- var$`excimp$variation`
```

```
## Warning: Unknown or uninitialised column: `excimp$variation`.
```

```
plot <- barplot(var$n,
  names.arg = namevar,
  main = "Total number of each variations",
  xlab = "Name of Variations",
  ylab = "Total of each Variatons",
  col = 1:16,
  space = 0.1,
  cex.names = 0.5,
  las = 2)
```



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

```
# Load the required libraries
```

```
library(ggplot2)
library(magrittr)
```

```
# Subset the data for Black variations
```

```
Blackplot <- var[var$`file_show$variation` %in% c("Black", "BlackDot", "BlackShow", "BlackPlus", "BlackMinus")]
```

```
# Subset the data for White variations
```

```
Whiteplot <- var[var$`file_show$variation` %in% c("White", "WhiteDot", "WhiteShow", "WhitePlus", "WhiteMinus")]
```

```
#layout one frame
```

```

par(mfrow = c(1, 2), mar = c(2, 2, 2, 2))

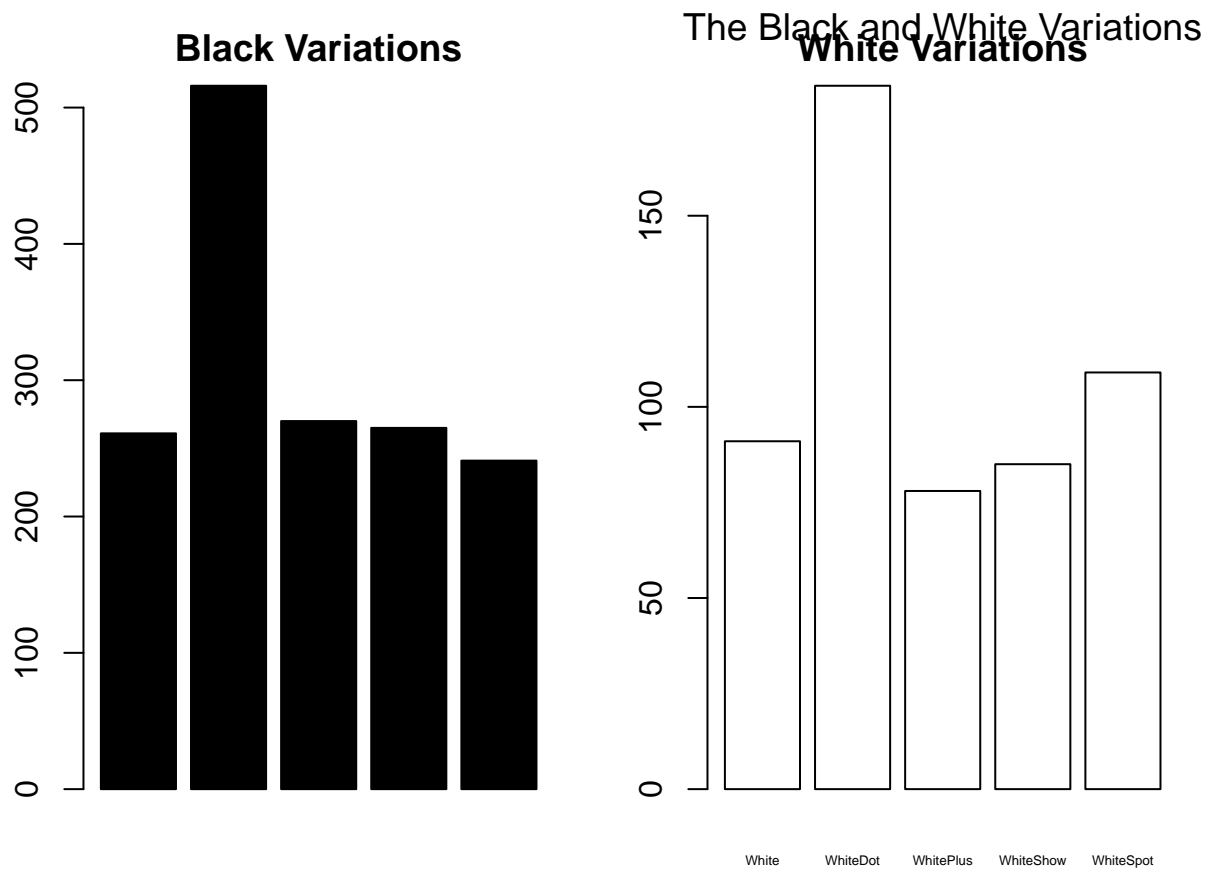
# Barplot for Black variations
Black <- barplot(height = Blackplot$n,
  names.arg = Blackplot$file_show$var`,
  col = "Black",
  main = "Black Variations",
  xlab = "Number of Variation",
  ylab = "Count",
  cex.names = 0.4,
  space = 0.2
)

## Warning: Unknown or uninitialised column: `file_show$var`.

# Barplot for White variations
White <- barplot(height = Whiteplot$n,
  names.arg = Whiteplot$file_show$variation`,
  col = "White",
  main = "White Variations",
  xlab = "Number of Variation",
  ylab = "Count",
  cex.names = 0.4,
  space = 0.2
)

mtext("The Black and White Variations", side = 3, line = 1, cex = 1.2)

```



```
knitr::include_graphics("/cloud/project/RWorksheet$4/BLACKandWHITE.png")
```

