# Rworksheet\_Gregorio#4b

#### 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 vector A = [1,2,3,4,5] and a  $5 \times 5$  zero matrix.

```
vectorA \leftarrow c(1,2,3,4,5)
matrixA <- matrix(0,nrow = 5, ncol =5)</pre>
for (i in 1:5)
  for (j in 1:5)
     matrixA[i,j] <- abs (vectorA[i] - vectorA[j])</pre>
  }
matrixA
##
          [,1] [,2] [,3] [,4] [,5]
## [1,]
             0
                    1
                                2
                                      3
## [2,]
             1
                    0
                          1
                                      2
## [3,]
             2
                   1
                          0
                                1
## [4,]
             3
                    2
                                      1
## [5,]
                          2
#2.
for (i in 1:5) {
  cat(paste0("\"", rep("*", i), "\""), "\n")
#3. n <- as.numeric(readline(prompt = "Enter a number to start the Fibonacci sequence:"))
a < \hspace{-0.07cm}\text{--} 0 b < \hspace{-0.07cm}\text{--} 1 c < \hspace{-0.07cm}\text{--} a + b
repeat { if (c > 500) { break } if (a == 0 \& b == 1) { cat(b, "") } cat(c, "") a <- b b <- c c <- a + b }
4. Import the dataset as shown in Figure 1 you have created previously. 4a. 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
ShoesData <- read.csv("Shoe sizes.csv")</pre>
head(ShoesData)
      X shoe_size height Gender
## 1 1
                       66.0
                6.5
## 2 2
                9.0
                       68.0
                                    F
                                    F
## 3 3
                8.5
                       64.5
```

```
## 4 4 8.5 65.0 F
## 5 5 10.5 70.0 M
## 6 6 7.0 64.0 F
```

4b. 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.

```
maleSub <- subset(ShoesData, Gender == "M")
femSub <- subset(ShoesData, Gender == "F")

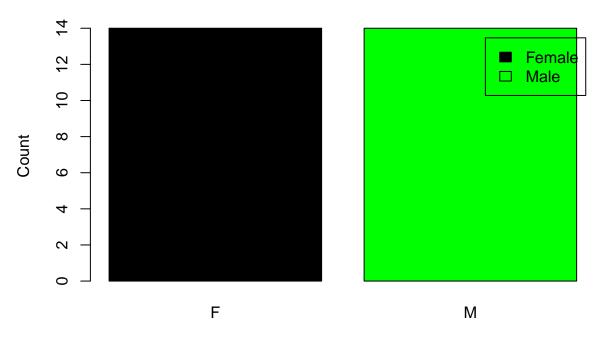
cat("The number of observation in male subset:", nrow(maleSub),"\n")</pre>
```

```
## The number of observation in male subset: 14
cat("The number of observation in female subset:", nrow(femSub),"\n")
```

## The number of observation in female subset: 14

4c. 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

### **Number of Male and Female in Household Data**



Gender

#5.

```
spending_data <- data.frame(
   Category = c("Food", "Electricity", "Savings", "Miscellaneous"),
   Value = c(60, 10, 5, 25)
)

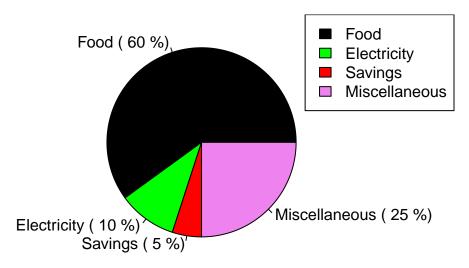
spending_data$Percentage <- spending_data$Value / sum(spending_data$Value) * 100

colors <- c("black", "green", "red", "violet")

pie(spending_data$Value,
   labels = paste(spending_data$Category,"(",spending_data$Percentage,"%)"),
   col = colors,
   main = "Monthly Income Spending of Dela Cruz Family")

legend("topright", spending_data$Category, fill = colors)</pre>
```

### **Monthly Income Spending of Dela Cruz Family**



6. Use the iris dataset

data(iris)

```
## function (..., list = character(), package = NULL, lib.loc = NULL,
##
       verbose = getOption("verbose"), envir = .GlobalEnv, overwrite = TRUE)
## {
##
       fileExt <- function(x) {</pre>
           db <- grepl("\\.[^.]+\\.(gz|bz2|xz)$", x)
##
##
           ans <- sub(".*\\.", "", x)
           ans[db] <- sub(".*\\.([^.]+\\.)(gz|bz2|xz)$", "\\1\\2",
##
                x[db])
##
##
           ans
##
##
       my_read_table <- function(...) {</pre>
##
           lcc <- Sys.getlocale("LC_COLLATE")</pre>
           on.exit(Sys.setlocale("LC_COLLATE", lcc))
##
##
           Sys.setlocale("LC_COLLATE", "C")
           read.table(...)
##
```

```
}
##
##
       stopifnot(is.character(list))
##
       names <- c(as.character(substitute(list(...))[-1L]), list)</pre>
##
       if (!is.null(package)) {
##
            if (!is.character(package))
##
                stop("'package' must be a character vector or NULL")
##
##
       paths <- find.package(package, lib.loc, verbose = verbose)</pre>
##
       if (is.null(lib.loc))
##
            paths <- c(path.package(package, TRUE), if (!length(package)) getwd(),</pre>
##
                paths)
##
       paths <- unique(normalizePath(paths[file.exists(paths)]))</pre>
##
       paths <- paths[dir.exists(file.path(paths, "data"))]</pre>
##
       dataExts <- tools:::.make_file_exts("data")</pre>
##
       if (length(names) == OL) {
##
            db <- matrix(character(), nrow = OL, ncol = 4L)</pre>
##
            for (path in paths) {
##
                entries <- NULL
##
                packageName <- if (file_test("-f", file.path(path,</pre>
##
                     "DESCRIPTION")))
##
                     basename(path)
                else "."
##
                if (file_test("-f", INDEX <- file.path(path, "Meta",</pre>
##
##
                     "data.rds"))) {
##
                     entries <- readRDS(INDEX)</pre>
##
                }
##
                else {
##
                     dataDir <- file.path(path, "data")</pre>
##
                     entries <- tools::list_files_with_type(dataDir,</pre>
##
                       "data")
##
                     if (length(entries)) {
##
                       entries <- unique(tools::file_path_sans_ext(basename(entries)))</pre>
##
                       entries <- cbind(entries, "")</pre>
                    }
##
                }
##
                if (NROW(entries)) {
##
##
                     if (is.matrix(entries) && ncol(entries) == 2L)
##
                       db <- rbind(db, cbind(packageName, dirname(path),</pre>
                         entries))
##
                     else warning(gettextf("data index for package %s is invalid and will be ignored",
##
                       sQuote(packageName)), domain = NA, call. = FALSE)
##
                }
##
            }
##
            colnames(db) <- c("Package", "LibPath", "Item", "Title")</pre>
##
##
            footer <- if (missing(package))</pre>
                paste0("Use ", sQuote(paste("data(package =", ".packages(all.available = TRUE))")),
##
##
                     "\n", "to list the data sets in all *available* packages.")
##
##
            y <- list(title = "Data sets", header = NULL, results = db,
##
                footer = footer)
##
            class(y) <- "packageIQR"</pre>
##
            return(y)
##
##
       paths <- file.path(paths, "data")</pre>
```

```
##
       for (name in names) {
##
           found <- FALSE
           for (p in paths) {
##
                tmp_env <- if (overwrite)</pre>
##
##
                    envir
##
                else new.env()
                if (file_test("-f", file.path(p, "Rdata.rds"))) {
##
                    rds <- readRDS(file.path(p, "Rdata.rds"))</pre>
##
##
                    if (name %in% names(rds)) {
                      found <- TRUE
##
##
                      if (verbose)
                        message(sprintf("name=%s:\t found in Rdata.rds",
##
##
                           name), domain = NA)
                      thispkg <- sub(".*/([^/]*)/data$", "\\1", p)
##
##
                      thispkg <- sub("_.*$", "", thispkg)</pre>
##
                      thispkg <- paste0("package:", thispkg)</pre>
##
                      objs <- rds[[name]]
##
                      lazyLoad(file.path(p, "Rdata"), envir = tmp_env,
##
                        filter = function(x) x %in% objs)
##
                      break
##
                    }
                    else if (verbose)
##
##
                      message(sprintf("name=%s:\t NOT found in names() of Rdata.rds, i.e.,\n\t%s\n",
                        name, paste(names(rds), collapse = ",")),
##
                        domain = NA)
##
##
##
                if (file_test("-f", file.path(p, "Rdata.zip"))) {
                    warning("zipped data found for package ", sQuote(basename(dirname(p))),
##
                      ".\nThat is defunct, so please re-install the package.",
##
##
                      domain = NA)
##
                    if (file_test("-f", fp <- file.path(p, "filelist")))</pre>
##
                      files <- file.path(p, scan(fp, what = "", quiet = TRUE))</pre>
##
##
                      warning(gettextf("file 'filelist' is missing for directory %s",
##
                        sQuote(p)), domain = NA)
##
                      next
##
                    }
##
               }
                else {
##
                    files <- list.files(p, full.names = TRUE)
##
##
##
                files <- files[grep(name, files, fixed = TRUE)]</pre>
##
                if (length(files) > 1L) {
##
                    o <- match(fileExt(files), dataExts, nomatch = 100L)</pre>
##
                    paths0 <- dirname(files)</pre>
                    paths0 <- factor(paths0, levels = unique(paths0))</pre>
##
##
                    files <- files[order(paths0, o)]
##
                }
##
                if (length(files)) {
##
                    for (file in files) {
##
                      if (verbose)
                        message("name=", name, ":\t file= ...", .Platform$file.sep,
##
##
                           basename(file), "::\t", appendLF = FALSE,
##
                           domain = NA)
```

```
##
                      ext <- fileExt(file)</pre>
##
                      if (basename(file) != paste0(name, ".", ext))
                        found <- FALSE
##
                      else {
##
##
                        found <- TRUE
##
                        zfile <- file
                        zipname <- file.path(dirname(file), "Rdata.zip")</pre>
##
##
                        if (file.exists(zipname)) {
##
                          Rdatadir <- tempfile("Rdata")</pre>
##
                          dir.create(Rdatadir, showWarnings = FALSE)
##
                          topic <- basename(file)</pre>
                          rc <- .External(C_unzip, zipname, topic,</pre>
##
##
                             Rdatadir, FALSE, TRUE, FALSE, FALSE)
##
                           if (rc == 0L)
##
                             zfile <- file.path(Rdatadir, topic)</pre>
##
##
                        if (zfile != file)
##
                          on.exit(unlink(zfile))
                        switch(ext, R = , r = {
##
##
                          library("utils")
##
                          sys.source(zfile, chdir = TRUE, envir = tmp_env)
                        }, RData = , rdata = , rda = load(zfile,
##
                          envir = tmp_env), TXT = , txt = , tab = ,
##
##
                          tab.gz = , tab.bz2 = , tab.xz = , txt.gz = ,
##
                          txt.bz2 = , txt.xz = assign(name, my_read_table(zfile,
##
                            header = TRUE, as.is = FALSE), envir = tmp_env),
##
                          CSV = , csv = , csv.gz = , csv.bz2 = ,
##
                          csv.xz = assign(name, my_read_table(zfile,
                            header = TRUE, sep = ";", as.is = FALSE),
##
##
                             envir = tmp_env), found <- FALSE)</pre>
                      }
##
##
                      if (found)
##
                        break
                    }
##
##
                    if (verbose)
##
                      message(if (!found)
##
                        "*NOT* ", "found", domain = NA)
##
                if (found)
##
                    break
##
           }
##
##
           if (!found) {
##
                warning(gettextf("data set %s not found", sQuote(name)),
##
                    domain = NA)
##
           else if (!overwrite) {
##
##
                for (o in ls(envir = tmp_env, all.names = TRUE)) {
##
                    if (exists(o, envir = envir, inherits = FALSE))
##
                      warning(gettextf("an object named %s already exists and will not be overwritten",
##
                        sQuote(o)))
##
                    else assign(o, get(o, envir = tmp_env, inherits = FALSE),
##
                      envir = envir)
##
                }
##
                rm(tmp_env)
```

```
##  }
##  }
## invisible(names)
## }
## <bytecode: 0x55c8177de848>
## <environment: namespace:utils>
```

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

```
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 1 ...
```

# The dataset contains information on iris blossoms. It contains information on the length and width of

6b.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?

```
meanOfFlowers <- colMeans(iris[,1:4])
meanOfFlowers

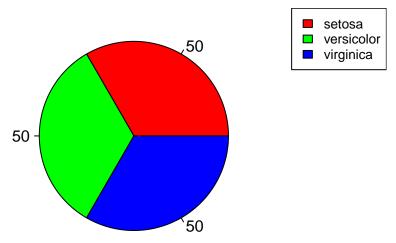
## Sepal.Length Sepal.Width Petal.Length Petal.Width</pre>
```

## 5.843333 3.057333 3.758000 1.199333

6c. Create a pie chart for the Species distribution. Add title, legends, and colors. Write the R script and its result.

```
species_count <- table(iris$Species)
pie(species_count, labels = species_count, col = rainbow(length(species_count)), main = "Species Distrilegend("topright", names(species_count), cex = 0.8, fill = rainbow(length(species_count)))</pre>
```

## **Species Distribution**



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

```
# Subset the iris data set into the three species.
setosa_subset <- subset(iris, Species == "setosa")
versicolor_subset <- subset(iris, Species == "versicolor")
virginica_subset <- subset(iris, Species == "virginica")
setosa_subset</pre>
```

##		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
##		5.5	3.5	1.3	0.2	setosa
	38	4.9	3.6	1.4	0.1	setosa
##		4.4	3.0	1.3	0.2	setosa
##		5.1	3.4	1.5	0.2	setosa
##		5.0	3.5	1.3	0.3	setosa
##		4.5	2.3	1.3	0.3	setosa
##		4.4	3.2	1.3	0.2	setosa
##		5.0	3.5	1.6	0.6	setosa
##		5.1	3.8	1.9	0.4	setosa
##	40	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

 ${\tt versicolor\_subset}$ 

##		Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
##	51	7.0	3.2	4.7	1.4	versicolor
##	52	6.4	3.2	4.5	1.5	versicolor
##	53	6.9	3.1	4.9	1.5	versicolor
##	54	5.5	2.3	4.0	1.3	versicolor
##	55	6.5	2.8	4.6	1.5	versicolor
##	56	5.7	2.8	4.5	1.3	versicolor
##	57	6.3	3.3	4.7	1.6	versicolor
##	58	4.9	2.4	3.3	1.0	versicolor
##	59	6.6	2.9	4.6	1.3	versicolor
##	60	5.2	2.7	3.9	1.4	versicolor
##	61	5.0	2.0	3.5	1.0	versicolor
##	62	5.9	3.0	4.2	1.5	versicolor
##	63	6.0	2.2	4.0	1.0	versicolor
##	64	6.1	2.9	4.7		versicolor
##	65	5.6	2.9	3.6		versicolor
##	66	6.7	3.1	4.4		versicolor
##	67	5.6	3.0	4.5		versicolor
##	68	5.8	2.7	4.1		versicolor
##	69	6.2	2.2	4.5		versicolor
##	70	5.6	2.5	3.9		versicolor
##	71	5.9	3.2	4.8		versicolor
##	72	6.1	2.8	4.0		versicolor
##	73	6.3	2.5	4.9		versicolor
##	74	6.1	2.8	4.7	1.2	versicolor
##	75	6.4	2.9	4.3		versicolor
##	76	6.6	3.0	4.4		versicolor
##	77	6.8	2.8	4.8		versicolor
##	78	6.7	3.0	5.0		versicolor
##	79	6.0	2.9	4.5		versicolor
##	80	5.7	2.6	3.5		versicolor
##	81	5.5	2.4	3.8		versicolor
##	82	5.5	2.4	3.7		versicolor
##	83	5.8	2.7	3.9		versicolor
##	84	6.0	2.7	5.1		versicolor
##	85	5.4	3.0	4.5		versicolor
##		6.0	3.4	4.5		versicolor
##		6.7	3.1	4.7		versicolor
##		6.3	2.3	4.4		versicolor
##		5.6	3.0	4.1		versicolor
##		5.5	2.5	4.0		versicolor
##		5.5	2.6	4.4		versicolor
##		6.1	3.0	4.6		versicolor
##		5.8	2.6	4.0		versicolor
##		5.0	2.3	3.3		versicolor
##		5.6	2.7	4.2		versicolor
##		5.7	3.0	4.2		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

virginica\_subset

##		Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
##	101	6.3	3.3	6.0		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		virginica
	120	6.0	2.2	5.0		virginica
	121	6.9	3.2	5.7		virginica
	122	5.6	2.8	4.9		virginica
	123	7.7	2.8	6.7		virginica
	124	6.3	2.7	4.9		virginica
	125	6.7	3.3	5.7		virginica
	126	7.2	3.2	6.0		virginica
	127	6.2	2.8	4.8		virginica
	128	6.1	3.0	4.9		virginica
	129	6.4	2.8	5.6		virginica
## ##	130 131	7.2	3.0	5.8 6.1		virginica
##	132	7.4 7.9	2.8	6.4		virginica
	133	6.4	3.8 2.8	5.6		virginica
	134	6.3	2.8	5.1		virginica virginica
##	135	6.1	2.6	5.6		virginica
	136	7.7	3.0	6.1		virginica
	137	6.3	3.4	5.6		virginica
##	138	6.4	3.1	5.5		virginica
##	139	6.0	3.0	4.8		virginica
##	140	6.9	3.1	5.4		virginica
##	141	6.7	3.1	5.6		virginica
##	142	6.9	3.1	5.1		virginica
##	143	5.8	2.7	5.1		virginica
##	144	6.8	3.2	5.9		virginica
	145	6.7	3.3	5.7		virginica
##	146	6.7	3.0	5.2		virginica
##	147	6.3	2.5	5.0		virginica
##	148	6.5	3.0	5.2	2.0	virginica

```
## 149
                 6.2
                              3.4
                                            5.4
                                                         2.3 virginica
## 150
                 5.9
                                                         1.8 virginica
                              3.0
                                            5.1
# Display the last six rows of each species.
tail(setosa_subset, 6)
      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
                                                        0.2 setosa
                5.0
                                           1.4
## 50
                             3.3
tail(versicolor_subset, 6)
##
       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
                              2.9
                                            4.2
                                                         1.3 versicolor
                 5.7
## 98
                                                         1.3 versicolor
                 6.2
                              2.9
                                            4.3
## 99
                 5.1
                              2.5
                                            3.0
                                                         1.1 versicolor
                                                         1.3 versicolor
## 100
                 5.7
                              2.8
                                            4.1
tail(virginica_subset, 6)
##
       Sepal.Length Sepal.Width Petal.Length Petal.Width
                                                               Species
## 145
                              3.3
                                            5.7
                 6.7
                                                         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. 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.
# Convert the "Species" column to a factor
iris$Species <- as.factor(iris$Species)</pre>
# Create a scatterplot
plot(
  Sepal.Length ~ Sepal.Width,
  data = iris,
  pch = as.integer(iris$Species), # Use different pch symbols for each species
  col = as.integer(iris$Species), # Use different colors for each species
  xlab = "Sepal Length",
  ylab = "Sepal Width",
```

legend("topright", legend = levels(iris\$Species), col = 1:3, pch = 1:3, cex = 0.8, title = "Species")

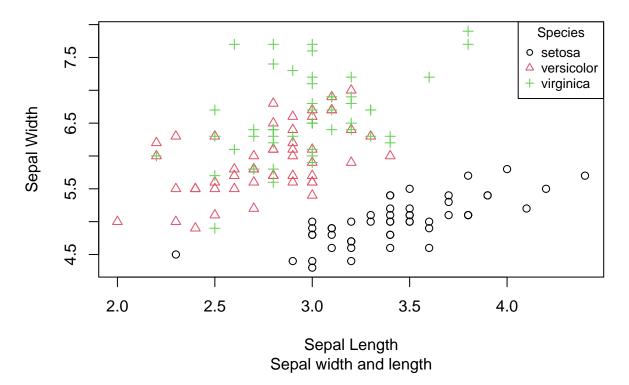
main = "Iris Dataset",

# Add a legend

)

sub = "Sepal width and length"

#### **Iris Dataset**



6f.Interpret the result.

```
# The dataset consists of five variables (columns) and 150 observations (rows) in a data frame format.
# Petal.Length, Petal.Width, Sepal.Length, and Sepal. Width are the names of the four numerical variabl
# The factor variable Species, which represents the species of iris flowers, is the sixth variable. The
```

7.Import the alexa-file.xlsx. Check on the variations. Notice that there are extra whitespaces among black variants (Black Dot, Black Plus, Black Show, Black Spot). Also on the white variants (White Dot, White Plus, White Show, White Spot).

```
library(readxl)
alexa_file <- read_excel("alexa_file.xlsx")
alexa_file</pre>
```

```
# A tibble: 3,150 x 5
##
      rating date
                                   variation
                                                        verified_reviews
                                                                               feedback
##
       <dbl> <dttm>
                                   <chr>
                                                        <chr>
                                                                                  <dbl>
##
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                        Love my Echo!
    1
                                                                                       1
##
    2
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                        Loved it!
                                                                                       1
           4 2018-07-31 00:00:00 Walnut Finish
##
    3
                                                        Sometimes while play~
                                                                                       1
##
    4
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                        I have had a lot of ~
                                                                                       1
##
    5
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                        Music
                                                                                       1
           5 2018-07-31 00:00:00 Heather Gray Fabric I received the echo \sim
                                                                                       1
##
    6
##
           3 2018-07-31 00:00:00 Sandstone Fabric
                                                        Without having a cel~
                                                                                       1
##
    8
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                        I think this is the ~
                                                                                       1
##
           5 2018-07-30 00:00:00 Heather Gray Fabric looks great
                                                                                       1
##
           5 2018-07-30 00:00:00 Heather Gray Fabric Love it! I've listen~
                                                                                       1
  10
  # i 3,140 more rows
```

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

```
alexa_file$variation <- gsub("Black Dot", "BlackDot", alexa_file$variation)
alexa_file$variation <- gsub("Black Plus", "BlackPlus", alexa_file$variation)
alexa_file$variation <- gsub("Black Show", "BlackShow", alexa_file$variation)
alexa_file$variation <- gsub("Black Spot", "BlackSpot", alexa_file$variation)
alexa_file$variation <- gsub("White Dot", "WhiteDot", alexa_file$variation)
alexa_file$variation <- gsub("White Plus", "WhitePlus", alexa_file$variation)
alexa_file$variation <- gsub("White Show", "WhiteShow", alexa_file$variation)
alexa_file$variation <- gsub("White Spot", "WhiteSpot", alexa_file$variation)
alexa_file$
## # A tibble: 3,150 x 5</pre>
```

```
feedback
##
     rating date
                                 variation
                                                     verified_reviews
##
       <dbl> <dttm>
                                 <chr>
                                                     <chr>>
                                                                               <dbl>
##
  1
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                     Love my Echo!
                                                                                   1
## 2
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                     Loved it!
                                                                                   1
##
           4 2018-07-31 00:00:00 Walnut Finish
                                                     Sometimes while play~
                                                                                   1
## 4
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                     I have had a lot of ~
                                                                                   1
## 5
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                     Music
                                                                                   1
## 6
           5 2018-07-31 00:00:00 Heather Gray Fabric I received the echo ~
                                                                                   1
## 7
           3 2018-07-31 00:00:00 Sandstone Fabric
                                                     Without having a cel~
                                                                                   1
## 8
           5 2018-07-31 00:00:00 Charcoal Fabric
                                                     I think this is the ~
                                                                                   1
           5 2018-07-30 00:00:00 Heather Gray Fabric looks great
## 9
                                                                                   1
           5 2018-07-30 00:00:00 Heather Gray Fabric Love it! I've listen~
## 10
                                                                                   1
## # i 3,140 more rows
```

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

var_total <- alexa_file %>%

    count(alexa_file$variation)
```

```
## # A tibble: 16 x 2
##
      `alexa_file$variation`
                                        n
##
      <chr>
                                    <int>
##
   1 Black
                                      261
## 2 BlackDot
                                      516
## 3 BlackPlus
                                      270
## 4 BlackShow
                                      265
## 5 BlackSpot
                                      241
## 6 Charcoal Fabric
                                      430
```

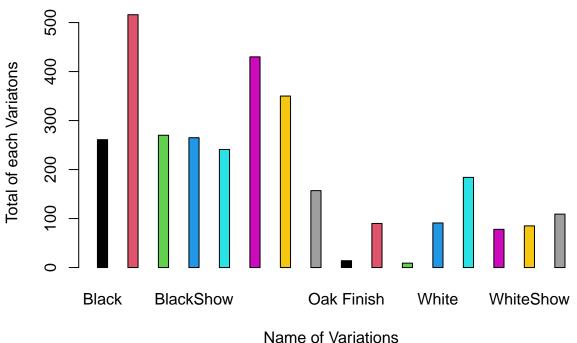
```
## 7 Configuration: Fire TV Stick
## 8 Heather Gray Fabric
                                     157
## 9 Oak Finish
                                      14
## 10 Sandstone Fabric
                                      90
## 11 Walnut Finish
                                       9
## 12 White
                                      91
## 13 WhiteDot
                                     184
## 14 WhitePlus
                                      78
## 15 WhiteShow
                                      85
                                     109
## 16 WhiteSpot
save(var_total, file = "var_total.RData")
```

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

```
load("var_total.RData")
var_total
```

```
## # A tibble: 16 x 2
##
      `alexa_file$variation`
                                        n
##
      <chr>
                                    <int>
## 1 Black
                                      261
## 2 BlackDot
                                      516
## 3 BlackPlus
                                      270
## 4 BlackShow
                                      265
## 5 BlackSpot
                                      241
## 6 Charcoal Fabric
                                      430
## 7 Configuration: Fire TV Stick
                                      350
## 8 Heather Gray Fabric
                                      157
## 9 Oak Finish
                                       14
## 10 Sandstone Fabric
                                       90
## 11 Walnut Finish
                                        9
## 12 White
                                       91
## 13 WhiteDot
                                      184
## 14 WhitePlus
                                       78
## 15 WhiteShow
                                       85
## 16 WhiteSpot
                                      109
varNames <- var_total$`alexa_file$variation`</pre>
```

#### Total number of each variations



7d. Create a barplot() for the black and white variations. Plot it in 1 frame, side by side. Complete t blackVars <- var\_total[var\_total\$`alexa\_file\$variation` %in% c("Black", "BlackPlus" , "BlackShow" , "BlackShow , "BlackSho whiteVars <- var\_total[var\_total\$`alexa\_file\$variation` %in% c("White", "WhiteDot", "WhitePlus", "White par(mfrow = c(1,2))barplot(height = blackVars\$n, names.arg = blackVars\$`alexa\_file\$variation`, col = c("black"), main = "Black Variations", xlab = "Variation", ylab = "Count", border = "black") barplot(height = whiteVars\$n,

names.arg = whiteVars\$`alexa\_file\$variation`,

col = c("black"),

xlab = "Variation", ylab = "Count", border = "black")

main = "White Variations",

