

```

#1.
A <- c(1, 2, 3, 4, 5)
MatrixA <- matrix(0, nrow = 5, ncol = 5)
print(MatrixA)

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

for (i in 1:5) {
  for (j in 1:5) {
    MatrixA[i, j] <- abs(A[i] - A[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

num.lines <- 5

for (a in 1:num.lines) {
  for (b in 1:a) {
    cat("*")
  }
  cat("\n")
}

## *
## **
## ***
## ****
## *****

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

## Enter a number to start the Fibonacci sequence:

assume.number <- 0
x <- 0
y <- 1
repeat {
  if (x > 500) {
    break
  }
  if (x >= assume.number) {
    cat(x, " ")
  }
  temp <- x + y

```

```

    x <- y
    y <- temp
  }

## 0  1  1  2  3  5  8  13  21  34  55  89  144  233  377

cat("\n")

shoe <- read.csv("Shoe_sizes.csv")
shoe

##      Shoe.size Height Gender Shoe.size.1 Height.1 Gender.1
## 1         6.5   66.0      F         13.0        77         M
## 2         9.0   68.0      F         11.5        72         M
## 3         8.5   64.5      F          8.5        59         F
## 4         8.5   65.0      F          5.0        62         F
## 5        10.5   70.0      M         10.0        72         M
## 6         7.0   64.0      F          6.5        66         F
## 7         9.5   70.0      F          7.5        64         F
## 8         9.0   71.0      F          8.5        67         M
## 9        13.0   72.0      M         10.5        73         M
## 10        7.5   64.0      F          8.5        69         F
## 11        10.5   74.5      M         10.5        72         M
## 12         8.5   67.0      F         11.0        70         M
## 13        12.0   71.0      M          9.0        69         M
## 14        10.5   71.0      M         13.0        70         M

shoe1 <- shoe[c(1:6),]
shoe1

##      Shoe.size Height Gender Shoe.size.1 Height.1 Gender.1
## 1         6.5   66.0      F         13.0        77         M
## 2         9.0   68.0      F         11.5        72         M
## 3         8.5   64.5      F          8.5        59         F
## 4         8.5   65.0      F          5.0        62         F
## 5        10.5   70.0      M         10.0        72         M
## 6         7.0   64.0      F          6.5        66         F

femaleData <- subset(shoe, Gender == "Female")
femaleData

## [1] Shoe.size   Height      Gender      Shoe.size.1 Height.1   Gender.1
## <0 rows> (or 0-length row.names)

maleData <- subset(shoe, Gender == "Male")
maleData

## [1] Shoe.size   Height      Gender      Shoe.size.1 Height.1   Gender.1
## <0 rows> (or 0-length row.names)

nrow(femaleData)

## [1] 0

nrow(maleData)

## [1] 0

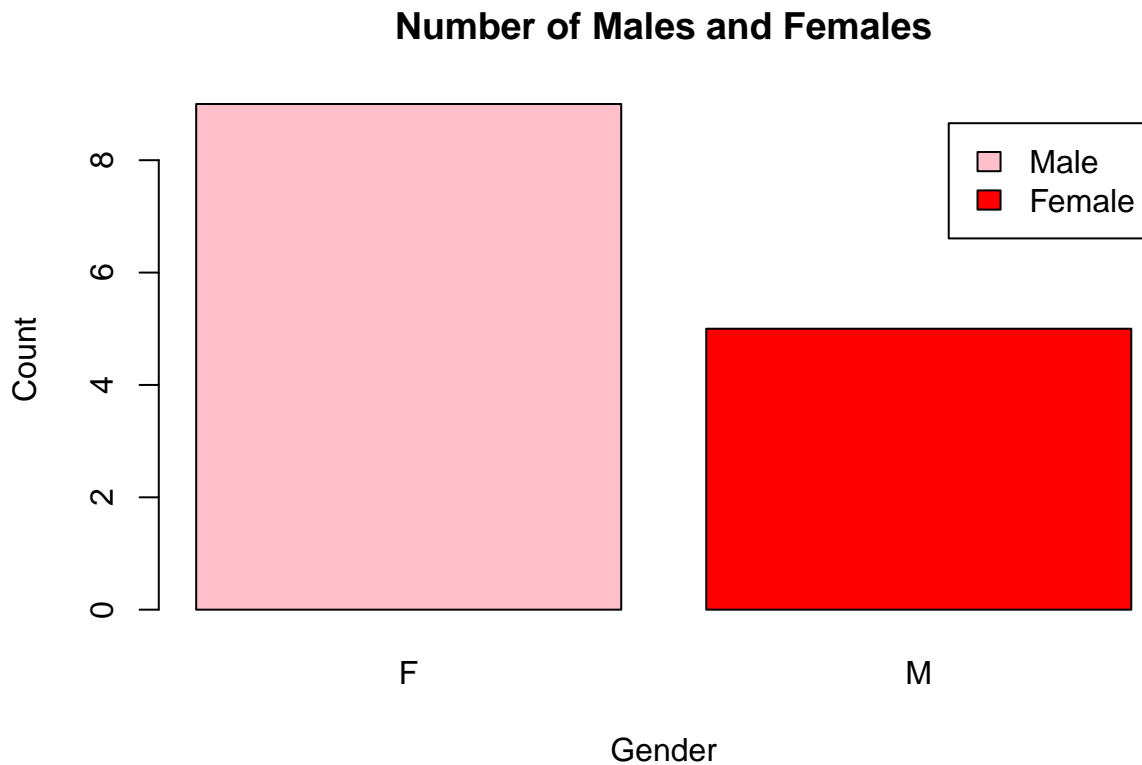
GraphMF <- table(shoe$Gender)
barplot(GraphMF,

```

```

    main = "Number of Males and Females",
    xlab = "Gender",
    ylab = "Count",
    col = c("pink", "red"),
    legend.text = c("Male", "Female"),
    beside = TRUE
)

```



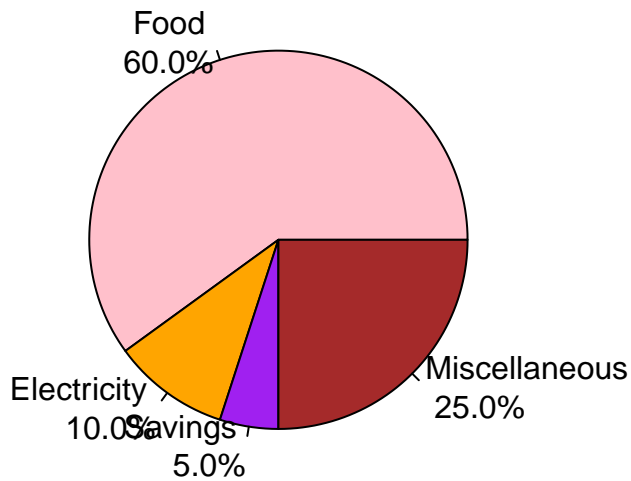
```

Expenses <- c(Food = 60, Electricity = 10, Savings = 5, Miscellaneous = 25)

pie(Expenses,
    labels = paste(names(Expenses), "\n", sprintf("%.1f%%", prop.table(Expenses) * 100)),
    col = c("pink", "orange", "purple", "brown"),
    main = "Monthly Expenses of the Dela Cruz Family"
)

```

Monthly Expenses of the Dela Cruz Family



```
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 has 150 observations and 5 variables

meanIris<- colMeans(iris[, c("Sepal.Length", "Sepal.Width", "Petal.Length", "Petal.Width")])
print(meanIris)

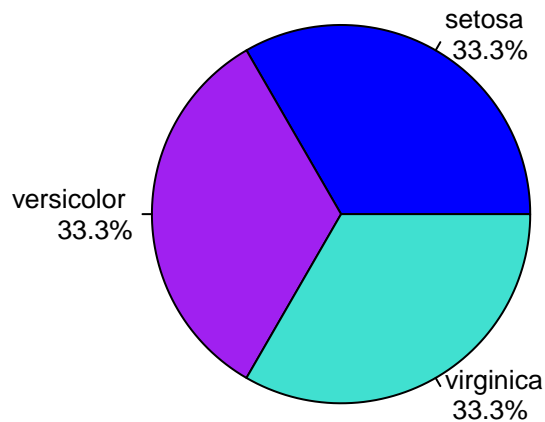
## Sepal.Length Sepal.Width Petal.Length Petal.Width
## 5.843333 3.057333 3.758000 1.199333

specs <- table(iris$Species)

clors <- c("blue", "purple", "turquoise")

pie(specs,
  labels = paste(names(specs), "\n", sprintf("%.1f%%", prop.table(specs) * 100)),
  col = clors,
  main = "Species Distribution",
  cex.main = 1.5,
  cex = 0.8
)
```

Species Distribution



```
SetSub <- subset(iris, Species == "setosa")
VersiSub <- subset(iris, Species == "versicolor")
VirgiSub <- subset(iris, Species == "virginica")
```

```
# Display the last six rows of each species
cat("Last six rows of Setosa:")
```

```
## Last six rows of Setosa:
```

```
print(tail(SetSub))
```

```
##      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
```

```
cat("Last six rows of Versicolor:")
```

```
## Last six rows of Versicolor:
```

```
print(tail(VersiSub))
```

```
##      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
```

```
cat("Last six rows of Virginica:")
```

```
## Last six rows of Virginica:
```

```
print(tail(VirgiSub))
```

```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width  Species
```

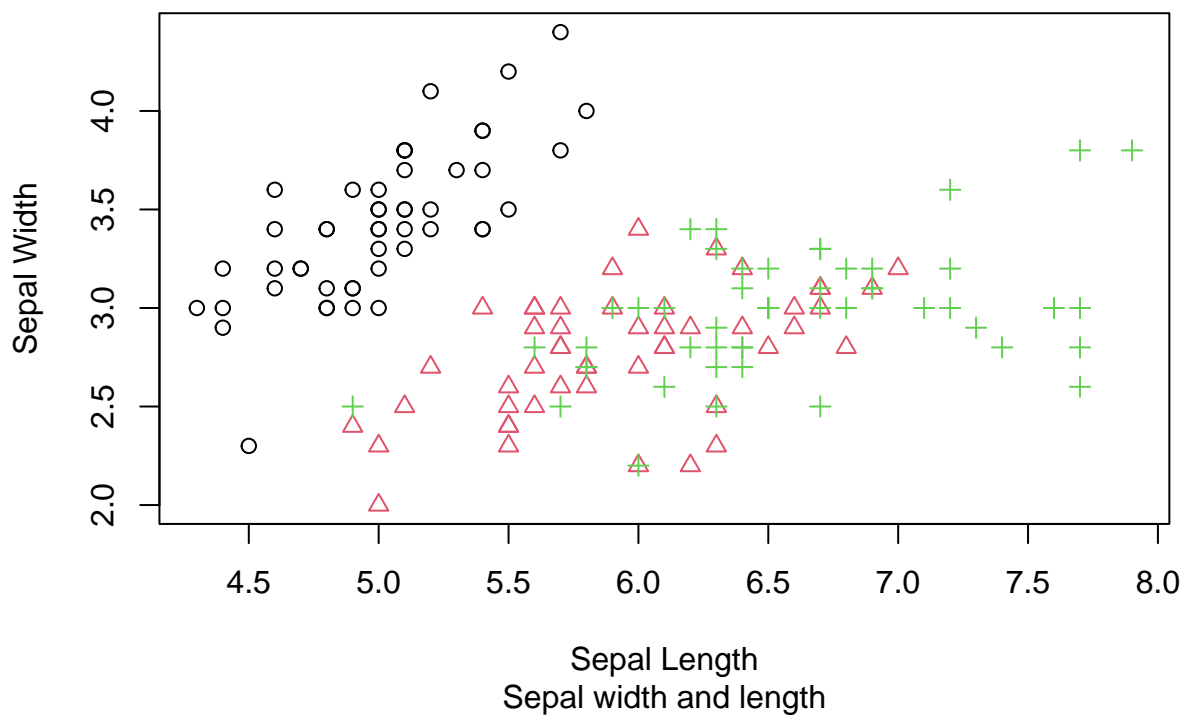
```
## 145      6.7      3.3      5.7      2.5 virginica
## 146      6.7      3.0      5.2      2.3 virginica
## 147      6.3      2.5      5.0      1.9 virginica
## 148      6.5      3.0      5.2      2.0 virginica
## 149      6.2      3.4      5.4      2.3 virginica
## 150      5.9      3.0      5.1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

```
data(iris)

iris$Species <- as.factor(iris$Species)

plot(iris$Sepal.Length, iris$Sepal.Width,
     pch = as.integer(iris$Species),
     col = iris$Species,
     main = "Iris Dataset",
     sub = "Sepal width and length",
     xlab = "Sepal Length",
     ylab = "Sepal Width"
)
```

Iris Dataset



#The scatterplot shows similarities between the sepal width and length ranging from 5.5 to 7.0

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

```
## # A tibble: 3,150 x 5
##   rating date          variation    verified_reviews    feedback
##   <dbl> <dtm>          <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
##	3	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
##	9	5	2018-07-30 00:00:00	Heather Gray Fabric	looks great	1
##	10	5	2018-07-30 00:00:00	Heather Gray Fabric	Love it! I've listen~	1
##	# i 3,140 more rows					