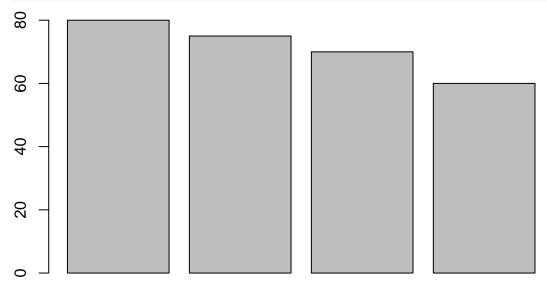
# RWorksheet\_Mirabuena#5

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#### 2022-11-22

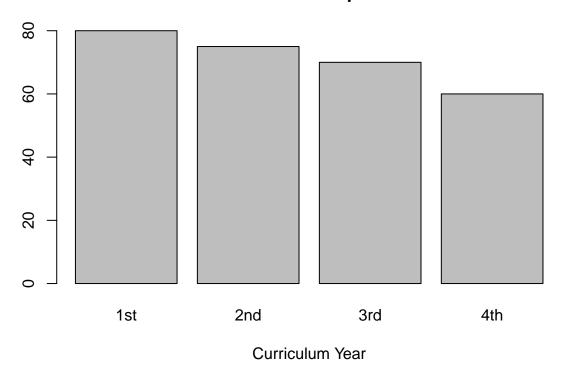
- 1. The table shows the enrollment of BS in Computer Science, SY 2010-2011. Course Year 2019 2020 1st  $80\ 2nd\ 75\ 3rd\ 70\ 4th\ 60$
- a. Plot the data using a bar graph. Write the codes and copy the result.

```
date2019_2020 <- c(80,75,70,60)
a1a <- barplot(date2019_2020)</pre>
```



#b. Using the same table, label the barchart with #Title = "Enrollment of BS Computer Science #horizontal axis = "Curriculum Year" and #vertical axis = "number of students"

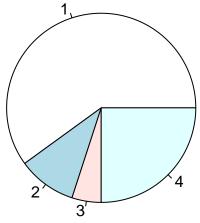
## **Enrollment of BS Computer Science**



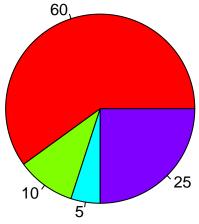
- 2. The monthly income of De Jesus family was spent on the following: 60% on Food, 10% on electricity, 5% for savings, and 25% for other miscellaneous expenses.
- a. Create a table for the above scenario. Write the codes and its result.

b. Plot the data using a pie chart. Add labels, colors and legend. Write the codes and its result.

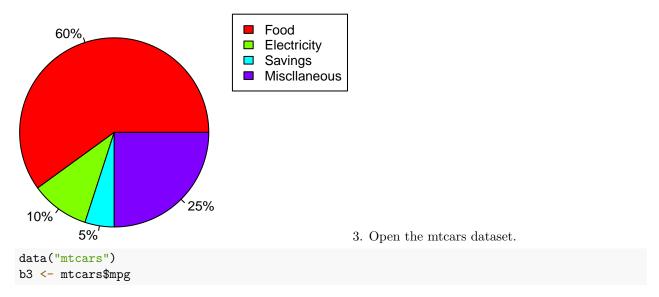
#### pie(expenses)



```
b2a <- pie(expenses,
    col = rainbow(length(expenses)),
    labels = c(60,10,5,25))</pre>
```



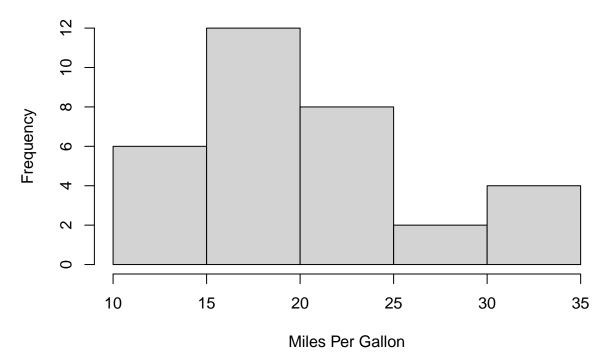
## **Expenses**



a. Create a simple histogram specifically for mpg (miles per gallon) variable. Use \$ to select the mpg only. Write the codes and its result.

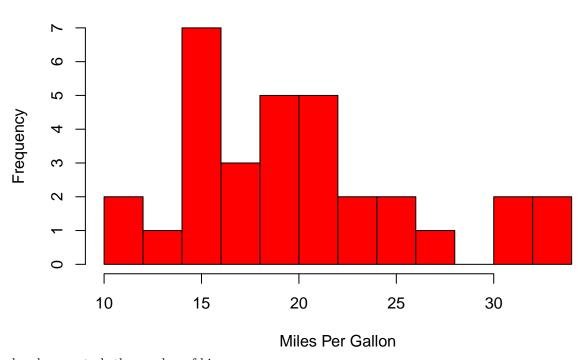
```
b3b <-hist(b3, xlab="Miles Per Gallon",
main="Histogram of mpg")
```

# Histogram of mpg



#b. Colored histogram with different number of bins.

## **Histogram of mpg**

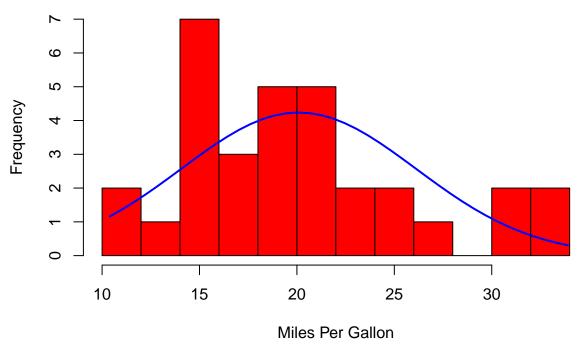


 $\label{eq:breaks} \text{breaks= controls the number of bins}$ 

#c. Add a Normal Curve

#Note:

### **Histogram with Normal Curve**



the result.

#4. Open the iris dataset. Create a subset for each species.

#a. Write the codes and its result.

```
data("iris")
set <- subset(iris, Species == "setosa")</pre>
ver <- subset(iris, Species == "versicolor")</pre>
vir <- subset(iris, Species == "virginica")</pre>
#b. Get the mean for every characteristics of each species using colMeans(). #Write the codes and its result.
set <- subset(iris, Species == "setosa")</pre>
setosa <- colMeans(set[sapply(set,is.numeric)])</pre>
setosa
## Sepal.Length Sepal.Width Petal.Length Petal.Width
           5.006
                          3.428
                                        1.462
                                                       0.246
ver <- subset(iris, Species == "versicolor")</pre>
versicolor <- colMeans(ver[sapply(ver,is.numeric)])</pre>
versicolor
## Sepal.Length
                  Sepal.Width Petal.Length
                                                Petal.Width
           5.936
                          2.770
                                                       1.326
vir <- subset(iris, Species == "virginica")</pre>
virginica <- colMeans(vir[sapply(vir,is.numeric)])</pre>
virginica
```

#Copy

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width ## 6.588 2.974 5.552 2.026
```

#Example: setosa <- colMeans(setosa[sapply(setosaDF,is.numeric)]) #c. Combine all species by using rbind()

#The table should be look like this:

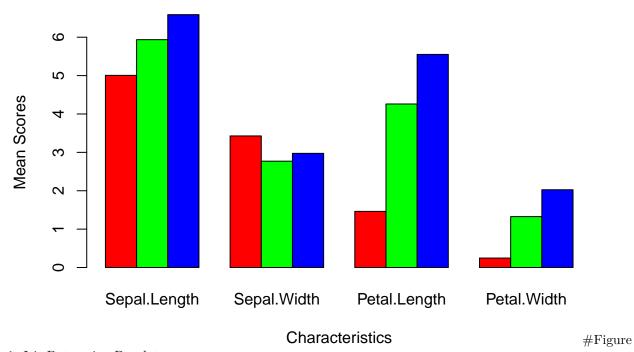
```
##
              Sepal.Length Sepal.Width Petal.Length Petal.Width
## setosa
                      5.006
                                   3.428
                                                1.462
## versicolor
                      5.936
                                   2.770
                                                4.260
                                                             1.326
## virginica
                      6.588
                                   2.974
                                                5.552
                                                             2.026
```

#Sepal.Length Sepal.Width Petal.Length Petal.Width #setosa #versicolor #virginica

#d. From the data in 4-c: Create the barplot(). #Write the codes and its result. #The barplot should be like this.

```
barplot(trans3, beside = TRUE,
    main = "Iris Mean",
    xlab = "Characteristics",
    ylab = "Mean Scores",
    col = c("red", "green", "blue"))
```

### Iris Mean



1: Iris Data using Barplot