## WorkSheet5

## Carl Frederick Delicana, BSIT 2-A

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#Accomplish this worksheet by answering the questions being asked and writing the code manually.

```
#1. The table shows the enrollment of BS in Computer Science, SY 2010-2011.
   Course Year 2019 - 2020
   1st
   2nd
                         75
                         70
   3rd
   4th
                         60
#a. Plot the data using a bar graph. Write the codes and copy the result.
date2019_2020 <- c(80,75,70,60)
numb1a <- 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"
course <- c("1st","2nd","3rd","4th")</pre>
numb1b <- barplot(date2019_2020,</pre>
                  main = "Enrollment of BS Computer Science",
                  xlab = "Curriculum Year", names.arg = course)
numb1b
#Output:
     [,1]
#[1,] 0.7
#[2,] 1.9
#[3,] 3.1
#[4,] 4.3
#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.
expenses <-c(60,10,5,25)
barplot(expenses,names.arg = c("Food", "Electricity", "Savings", "Misclaneous"))
#b. Plot the data using a pie chart. Add labels, colors and legend.
  Write the codes and its result.
pie(expenses)
```

```
numb2b <- pie(expenses,</pre>
               col = rainbow(length(expenses)),
               labels = c(60,10,5,25))
ex_labels <- round(expenses/sum(expenses) * 100, 1)</pre>
ex labels <- paste(ex labels, "%", sep = "")</pre>
pie(expenses, main = "Expenses",col=rainbow(length(expenses)),labels = ex_labels, cex=0)
legend(1, c("Food", "Electricity", "Savings", "Miscllaneous"),
       cex = 0.8,fill = rainbow((length(expenses))))
#3. Open the mtcars dataset.
data("mtcars")
numb3 <- mtcars$mpg</pre>
#a. Create a simple histogram specifically for mpg (miles per gallon) variable.
#Use $ to select the mpg only. Write the codes and its result.
numb3a <-hist(numb3, xlab="Miles Per Gallon",</pre>
              main="Histogram of mpg")
#b. Colored histogram with different number of bins.
numb3b <-hist(numb3, breaks=12, col="red", xlab="Miles Per Gallon",</pre>
              main="Histogram of mpg")
#Note: breaks= controls the number of bins
#c. Add a Normal Curve
numb3c <-hist(numb3, breaks=12, col="red", xlab="Miles Per Gallon",</pre>
               main="Histogram with Normal Curve")
xfit <- seq (min(numb3), max(numb3), length=40)
yfit<-dnorm(xfit,mean=mean(numb3),sd=sd(numb3))</pre>
yfit <- yfit*diff(numb3c$mids[1:2])*length(numb3)</pre>
lines(xfit, yfit, col="blue", lwd=2)
#Copy 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
#Output:
#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
#Output:
```

```
#Sepal.Length Sepal.Width Petal.Length Petal.Width
        5.936
                                   4.260
                     2.770
                                                1.326
vir <- subset(iris, Species == "virginica")</pre>
virginica <- colMeans(vir[sapply(vir,is.numeric)])</pre>
virginica
#Output:
#Sepal.Length Sepal.Width Petal.Length Petal.Width
        6.588
                     2.974
                                   5.552
                                                2.026
#Example: setosa <- colMeans(setosa[sapply(setosaDF,is.numeric)])</pre>
#c. Combine all species by using rbind()
#The table should be look like this:
trans3 <- rbind(setosa,</pre>
                versicolor,
                virginica)
trans3
#Output:
            Sepal.Length Sepal.Width Petal.Length Petal.Width
                   5.006
                                             1.462
                                                          0.246
#setosa
                               3.428
#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"))
#Figure 1: Iris Data using Barplot
```