

RWorksheet_Leysa#4b

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1. for loop

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

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

Amatrix
```

```
##      [,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
```

2. Print “*” using for() function

```
for (i in 1:5){
  cat(rep("*",i),"\n")
}
```

```
## *
## * *
## * * *
## * * * *
## * * * * *
```

3. Get an Input from user to print Fibonacci Sequence from 1st input to 500

```
#start_num <- as.numeric(readline(prompt = "Enter a number from 1-500: "))
# a <- start_num
# b <- start_num + 1

# while (b < 500) {
#   cat(a, " ")
```

```
#   next_numb <- a+b
#   a <- b
#   b<- next_numb
# }

#   cat("\n")

#I make this chunk to a comment because this actually runs, however,when I knit it,
#it stops from knitting to pdf because it says that the while loop needs a true or false.
```

4. Import the dataset of Sizes,Height,Gender.csv

```
data_list <- read.table("Sizes,Height,Gender.csv", sep = ",", header = TRUE)
data_list
```

```
##      Shoe.size Height Gender
## 1          6.5   66.0      F
## 2          9.0   68.0      F
## 3          8.5   64.5      F
## 4          8.5   65.0      F
## 5         10.5   70.0      M
## 6          7.0   64.0      F
## 7          9.5   70.0      F
## 8          9.0   71.0      F
## 9         13.0   72.0      M
## 10         7.5   64.0      F
## 11         10.5   74.5      M
## 12          8.5   67.0      F
## 13         12.0   71.0      M
## 14         10.5   71.0      M
## 15         13.0   77.0      M
## 16         11.5   72.0      M
## 17          8.5   59.0      F
## 18          5.0   62.0      F
## 19         10.0   72.0      M
## 20          6.5   66.0      F
## 21          7.5   64.0      F
## 22          8.5   67.0      M
## 23         10.5   73.0      M
## 24          8.5   69.0      F
## 25         10.5   72.0      M
## 26         11.0   70.0      M
## 27          9.0   69.0      M
## 28         13.0   70.0      M
```

4a. Display the first 6 rows

```
head(data_list,6)
```

```
##      Shoe.size Height Gender
## 1          6.5   66.0      F
## 2          9.0   68.0      F
```

```
## 3      8.5  64.5    F
## 4      8.5  65.0    F
## 5     10.5  70.0    M
## 6      7.0  64.0    F
```

4b. Subset for male and female

```
data_of_male <- subset(data_list, Gender == "M")
data_of_female <- subset(data_list, Gender == "F")
cat("Male:", nrow(data_of_male), "\nFemale:", nrow(data_of_female), "\n")
```

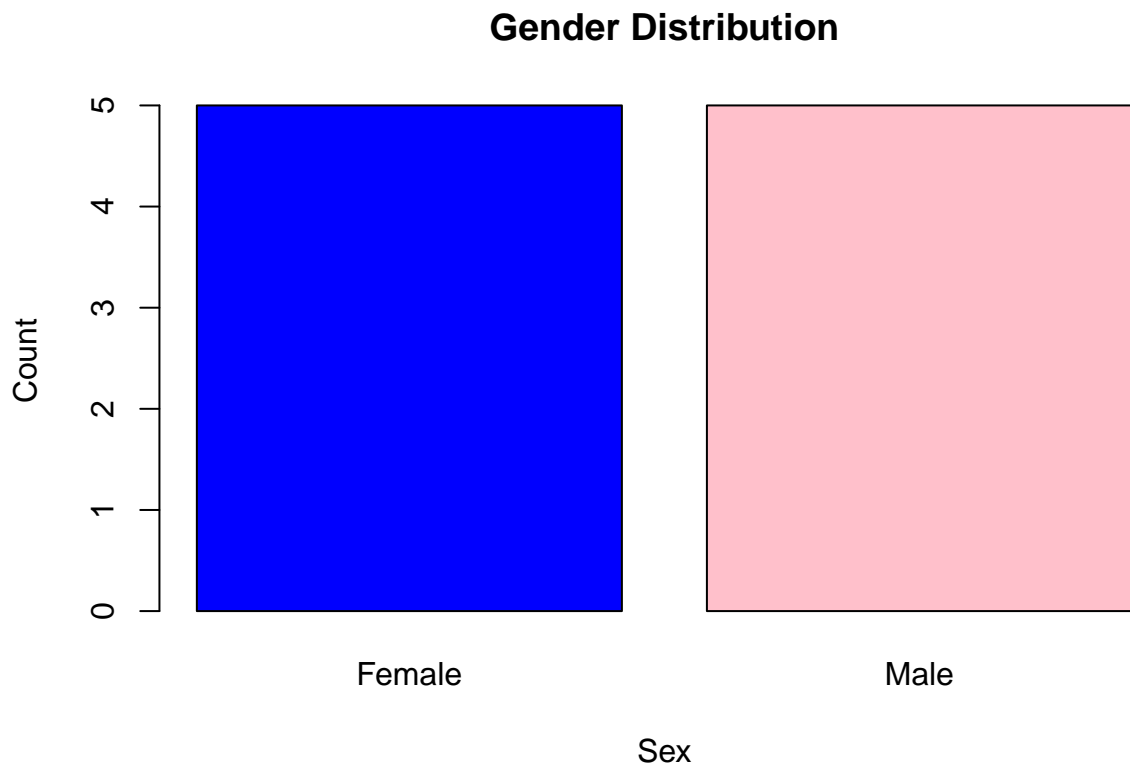
```
## Male: 14
## Female: 14
```

4c. Graph of numbers of males and females in household data

```
hdata <- read.table("HouseholdData.csv", sep = ",", header = TRUE)
hdata
```

```
## Respondents Sex Fathers.Occupation Persons.at.Home Siblings.at.School
## 1      1 Male Farmer 5 2
## 2      2 Female Driver 7 3
## 3      3 Female Others 3 0
## 4      4 Male Others 8 5
## 5      5 Male Farmer 6 2
## 6      6 Female Driver 4 3
## 7      7 Female Driver 4 1
## 8      8 Male Others 2 2
## 9      9 Female Farmer 11 6
## 10     10 Male Others 6 2
## Types.of.Houses
## 1      Wood
## 2      Congrete
## 3      Congrete
## 4      Wood
## 5      Semi-concrete
## 6      Semi-concrete
## 7      Wood
## 8      Semi-concrete
## 9      Semi-concrete
## 10     Congrete
```

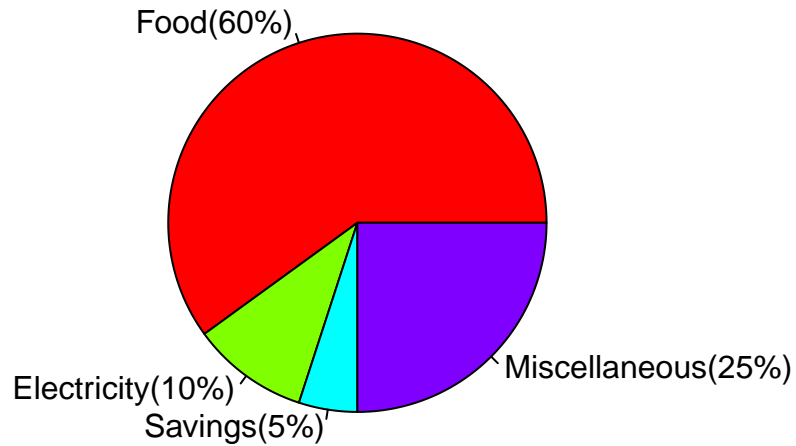
```
barplot(table(hdata$Sex), col = c("blue", "pink"), main = "Gender Distribution", xlab = "Sex", ylab = "Count")
```



5a. Pie Chart for Dela Cruz Family

```
expenses <- c(Food=60, Electricity=10,Savings=5,Miscellaneous=25)
percentage <- paste0(names(expenses),"(", round(expenses/sum(expenses)*100,1),"%)")
pie(expenses, labels= percentage, main = "Dela Cruz Family Monthly Spending", col=rainbow(length(expenses)))
```

Dela Cruz Family Monthly Spending



6a. Dataset of Iris

```
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 ...
```

6b. Mean of sepals and petals (length & width)

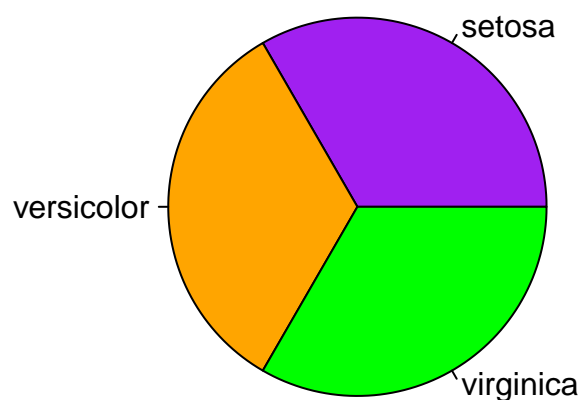
```
means <- colMeans(iris[,1:4])
means
```

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

6c. Pie Chart of Species Distribution

```
pie(table(iris$Species), main = "Species Distribution", col=c("purple","orange","green"))
```

Species Distribution



6d. Subsets of Species

```
Setosa <- tail(subset(iris, Species == "setosa"),6)
VersiColor <- tail(subset(iris, Species == "versicolor"),6)
Virginica <- tail(subset(iris, Species == "virginica"),6)
```

Setosa

##	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

VersiColor

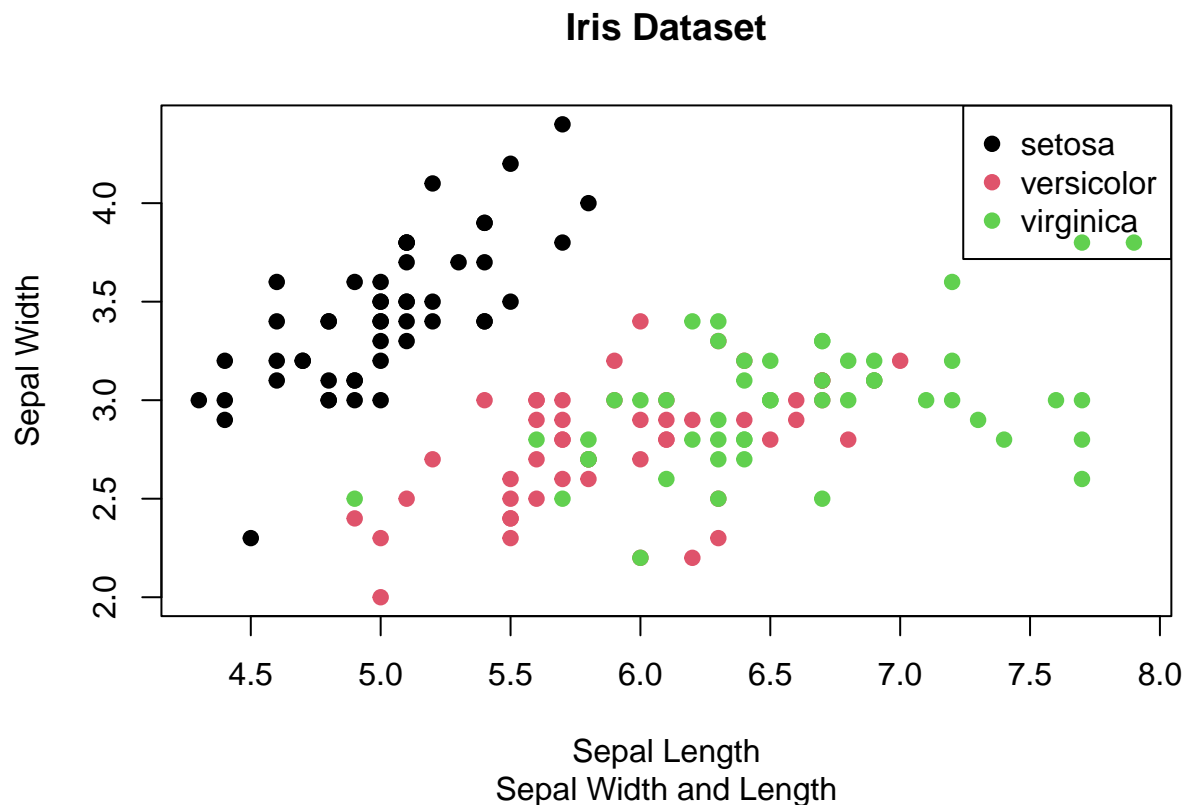
##	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
## 95	5.6	2.7	4.2	1.3	versicolor
## 96	5.7	3.0	4.2	1.2	versicolor
## 97	5.7	2.9	4.2	1.3	versicolor
## 98	6.2	2.9	4.3	1.3	versicolor
## 99	5.1	2.5	3.0	1.1	versicolor
## 100	5.7	2.8	4.1	1.3	versicolor

Virginica

```
##      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. Scatterplot

```
plot(iris$Sepal.Length, iris$Sepal.Width, col=iris$Species, pch=19, main = "Iris Dataset", xlab="Sepal Length", ylab="Sepal Width", legend="topright", legend=levels(iris$Species), col = 1:3, pch=19)
```



6f.

```
#Based on the scatter plot show, the Setosa is the most distinct species, and it can also
#be identified by its shorter sepal length and greater sepal width. On the other hand, the
#Versicolor and Virginica have more similar sepal dimensions because they have at least
#similar length and width of sepal, and in order to distinguish their difference, they need
# additional features, like petal measurements, in order to give us an accurate classification
# of these two species.
```

7.

```
library(readxl)
alexadata <- read.table("alexa_file.xlsx", sep = ",", header = TRUE)

## Warning in read.table("alexa_file.xlsx", sep = ",", header = TRUE): line 1
## appears to contain embedded nulls

## Warning in read.table("alexa_file.xlsx", sep = ",", header = TRUE): incomplete
## final line found by readTableHeader on 'alexa_file.xlsx'

alexadata

## [1] PK...
## <0 rows> (or 0-length row.names)
```