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Lab: CS602-Lab

Machine Learning for Cyber

Security

Assignment 1: To explore R for understanding programming fundamentals for data analysis applications.

Snapshot:

1.

```
> data(iris)
> head(iris)
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species
                       3.5
                                    1.4
                                                0.2 setosa
          4.9
                       3.0
                                    1.4
                                                0.2 setosa
           4.7
                       3.2
                                    1.3
                                                0.2 setosa
           4.6
                       3.1
                                    1.5
                                                0.2 setosa
           5.0
                       3.6
                                    1.4
                                                0.2 setosa
           5.4
                       3.9
                                    1.7
                                                0.4 setosa
```

2.

```
> col = names(iris)
> col
[1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width" "Species"
> class(col)
[1] "character"
> typeof(col)
[1] "character"
```

```
> 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 0.4 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 ...
>
```

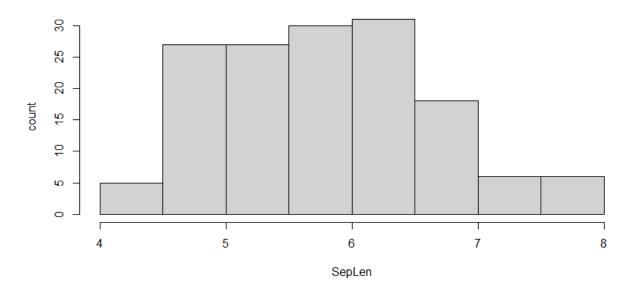
```
> summary(iris)
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
Min. :4.300 Min. :2.000 Min. :1.000 Min. :0.100 setosa :50
1st Qu.:5.100 1st Qu.:2.800 1st Qu.:1.600 1st Qu.:0.300 versicolor:50
Median :5.800 Median :3.000 Median :4.350 Median :1.300 virginica :50
Mean :5.843 Mean :3.057 Mean :3.758 Mean :1.199
3rd Qu.:6.400 3rd Qu.:3.300 3rd Qu.:5.100 3rd Qu.:1.800
Max. :7.900 Max. :4.400 Max. :6.900 Max. :2.500
>
```

5.

```
> unique(iris[, 5])
[1] setosa versicolor virginica
Levels: setosa versicolor virginica
>
>
>
> colnames((iris))
[1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width" "Species"
>
```

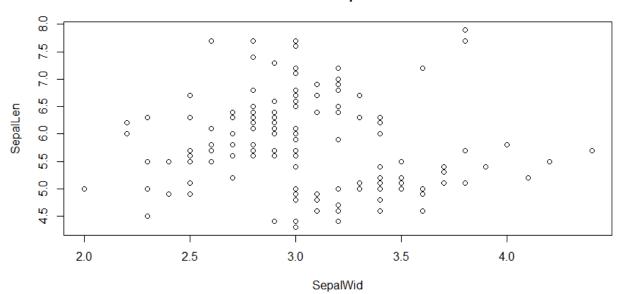
```
> hist(iris$Sepal.Length,
+ main='Histogram',
+ xlab='SepLen',
+ ylab='count')
```





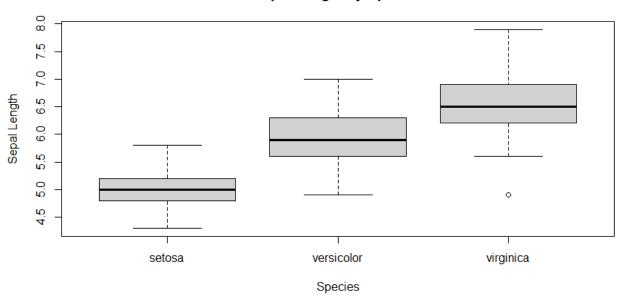
```
> plot(iris$Sepal.width, iris$Sepal.Length,
+ main='Scatterplot',
+ xlab='Sepalwid',
+ ylab='SepalLen')
>
```

Scatterplot

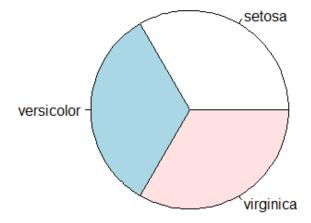


```
> boxplot(Sepal.Length~Species,
+ data=iris,
+ main='Sepal Length -Species',
+ xlab='Species',
+ ylab='Sepal Length')
>
```

Sepal Length by Species



```
> counts <- table(iris$Species)
> pie(counts)
>
```



```
> summary(iris)
 Sepal.Length
                             Petal.Length Petal.Width
              Sepal.Width
                                                          Species
Min. :4.300
              Min. :2.000
                             Min. :1.000 Min. :0.100
1st Qu.:5.100
              1st Qu.:2.800
                             1st Qu.:1.600
                                          1st Qu.:0.300
                                                          versicolor:50
Median :5.800
              Median:3.000
                             Median :4.350
                                           Median :1.300
                                                          virginica:50
Mean :5.843
              Mean :3.057
                             Mean :3.758
                                           Mean :1.199
                             3rd Qu.:5.100
3rd Qu.:6.400
              3rd Qu.:3.300
                                           3rd Qu.:1.800
      :7.900
Max.
              Max.
                    :4.400
                             Max.
                                   :6.900
                                           Max. :2.500
```

```
> A <- matrix(c(2, 4, 6, 8), nrow = 2, ncol = 2)
> B <- matrix(c(1, 3, 5, 7), nrow = 2, ncol = 2)
>
> dot_product <- A * B</pre>
```

```
> dot_product <- A * B
> dot_product
    [,1] [,2]
[1,] 2 30
[2,] 12 56
```

```
> C %*% D

[,1] [,2]

[1,] 76 103

[2,] 100 136
```

Conclusion:

- 1. Using R's built-in functions is more efficient than creating your functions and loops especially while working on some datasets.
- 2. Checking out the shape, dimension, and categorical data per column, as well as the min, max, and median per column, can give a better understanding of the data.
- 3. It is crucial to take care of nrow/ncol kind of params while dealing with matrices.