



# DEPARTMENT OF COMPUTER SCIENCE & ENGINEERING

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## Experiment-2.3

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**Subject Name:** Data Mining

**Subject Code:** 20CSP- 351

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**Aim** – To perform the cluster analysis by k-means method using R.

### **Objective-**

- ◆ Represent the reading of file using R studio
- ◆ Displaying the graph using clusterR, caret and cluster.
- ◆ Demonstration of Cluster analysis by K-means method.

### **Script and Output-**

```
str(iris)
# Installing Packages
install.packages("ClusterR")
install.packages("cluster")
# Loading package
library(ClusterR)
library(cluster)
library(caret)

# Removing initial label of Species from original dataset
iris_data <- iris[, -5]
iris_data

# Fitting K-Means clustering Model to training dataset
set.seed(240) # Setting seed
```



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```
km <- kmeans(iris_data, centers = 3, nstart = 20)
km

#Cluster identification for each observation
km$cluster

# Confusion Matrix
cm <- table(iris$Species, kmeans.re$cluster)
cm

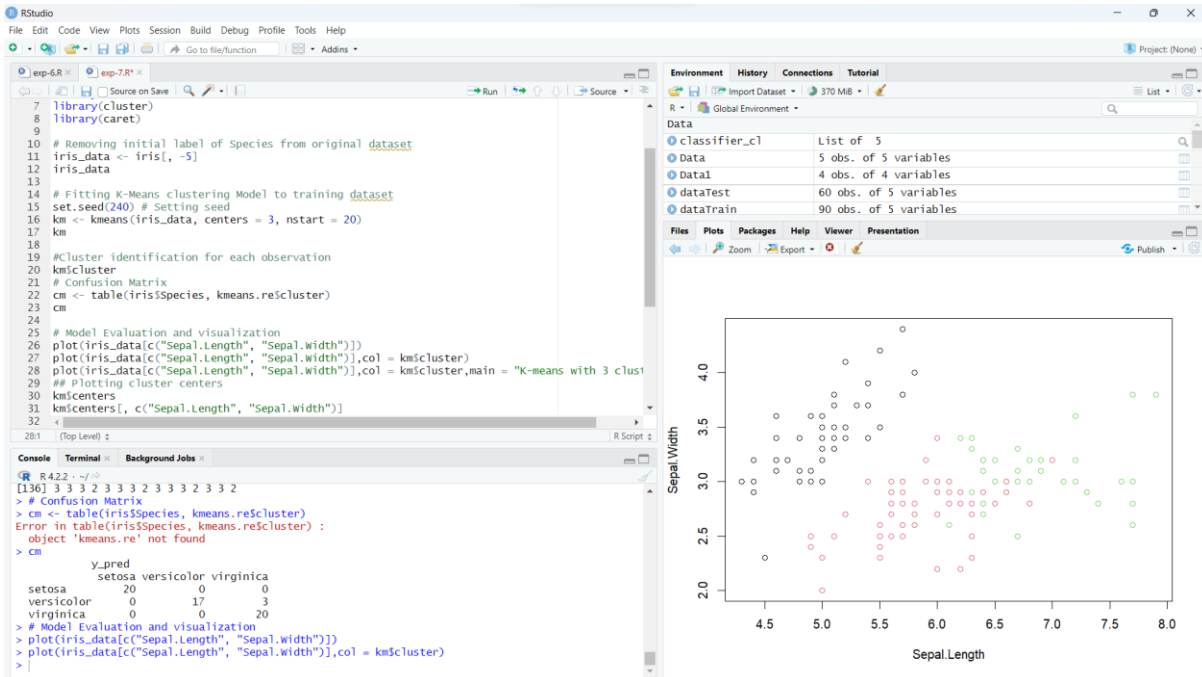
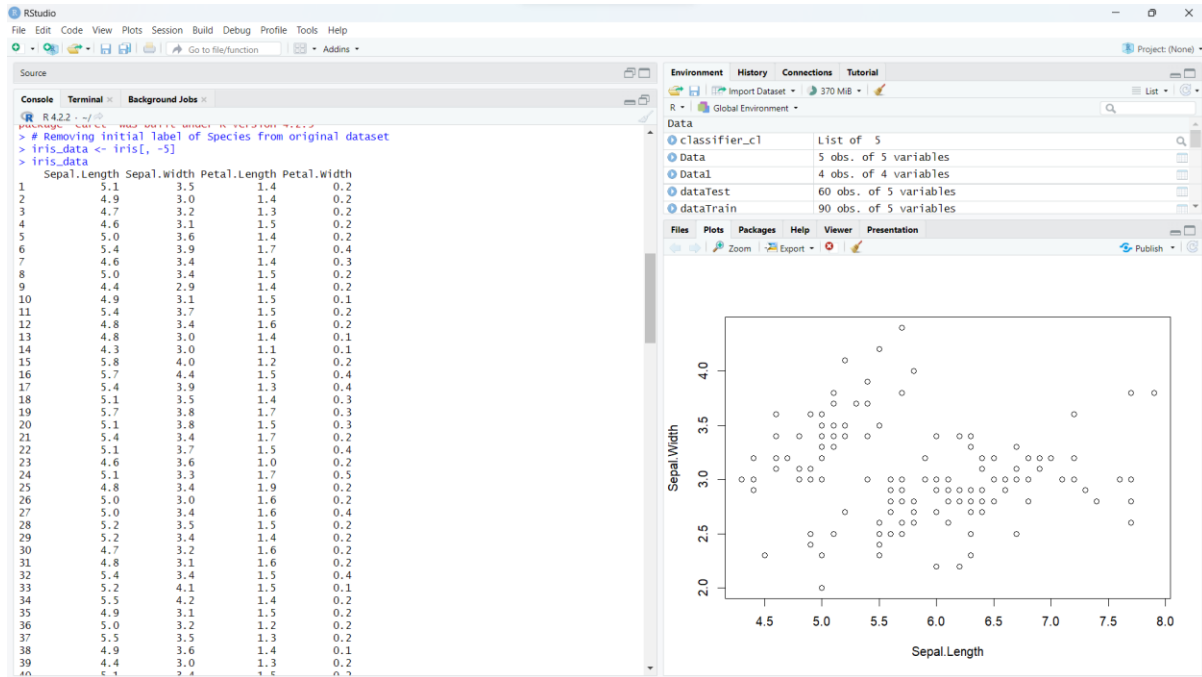
# Model Evaluation and visualization
plot(iris_data[c("Sepal.Length", "Sepal.Width")])
plot(iris_data[c("Sepal.Length", "Sepal.Width")], col = km$cluster)
plot(iris_data[c("Sepal.Length", "Sepal.Width")], col = km$cluster, main = "K-
means with 3 clusters")
## Plotting cluster centers
km$centers
km$centers[, c("Sepal.Length", "Sepal.Width")]
# cex is font size, pch is symbol
points(km$centers[, c("Sepal.Length", "Sepal.Width")], col = 1:3, pch = 8, cex =
3)
## Visualizing clusters
y_kmeans <- km$cluster
clusplot(iris_data[, c("Sepal.Length", "Sepal.Width")],
        y_kmeans,
        lines = 0,
        shade = TRUE,
        color = TRUE,
        labels = 2,
        plotchar = FALSE,
        span = TRUE,
        main = paste("Cluster iris"),
        xlab = 'Sepal.Length',
        ylab = 'Sepal.Width')
```



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## Output-





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