



DEPARTMENT OF COMPUTER SCIENCE & ENGINEERING

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WORKSHEET:-7

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Branch:- BE CSE

Section:- 20BCS_DM_716/B

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Subject:- Data Mining Lab

Subject Code:- 20CSP-376

AIM:-

- ❖ To perform the cluster analysis by k-means method using R

THEORY:-

K Means Clustering in R Programming is an Unsupervised Non-linear algorithm that cluster data based on similarity or similar groups. It seeks to partition the observations into a pre-specified number of clusters. Segmentation of data takes place to assign each training example to a segment called a cluster. In the unsupervised algorithm, high reliance on raw data is given with large expenditure on manual review for review of relevance is given. It is used in a variety of fields like Banking, healthcare, retail, Media, etc.

- K-Means clustering groups the data on similar groups. The algorithm is as follows:-
 - Choose the number **K** clusters.
 - Select at random **K** points, the centroids (Not necessarily from the given data).
 - Assign each data point to closest centroid that forms **K** clusters.



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- Compute and place the new centroid of each centroid.
- After final reassignment, name the cluster as Final cluster.

DATASET:-

Iris dataset consists of 50 samples from each of 3 species of Iris (Iris setosa, Iris virginica, Iris versicolor) and a multivariate dataset introduced by British statistician and biologist Ronald Fisher in his 1936 paper The use of multiple measurements in taxonomic problems. Four features were measured from each sample i.e length and width of the sepals and petals and based on the combination of these four features, Fisher developed a linear discriminant model to distinguish the species from each other.

```
# Loading data  
data(iris)  
  
# Structure  
str(iris)
```

Performing K-Means Clustering on Dataset:-

Using K-Means Clustering algorithm on the dataset which includes 11 persons and 6 variables or attributes.

```
# Installing Packages  
install.packages("ClusterR")  
install.packages("cluster")  
  
# Loading package  
library(ClusterR)  
library(cluster)  
  
# Removing  
initial label of #  
Species from  
original
```



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```
dataset iris_1 <-  
iris[, -5]  
  
# Fitting K-Means clustering  
Model  
# to training dataset  
set.seed(240) #  
Setting seed  
kmeans.re <- kmeans(iris_1, centers = 3, nstart =  
20) kmeans.re  
cluster identification for  
  
#each observation  
kmeans.re$cluster  
#confusion matrix  
cm <- table(iris$Species,  
kmeans.re$cluster) cm  
# Model Evaluation and  
visualization  
plot(iris_1[c("Sepal.Length",  
"Sepal.Width")])  
plot(iris_1[c("Sepal.Length",  
"Sepal.Width")], col =  
kmeans.re$cluster)  
plot(iris_1[c("Sepal.Length",  
"Sepal.Width")], col =  
kmeans.re$cluster,  
main = "K-means with 3  
clusters")  
  
## Plotting cluster centers  
kmeans.re$centers kmeans.re$centers[  
c("Sepal.Length", "Sepal.Width")]  
  
# cex is font size, pch is symbol  
points(kmeans.re$centers[, c("Sepal.Length",  
"Sepal.Width")], col = 1:3, pch = 8, cex = 3)
```

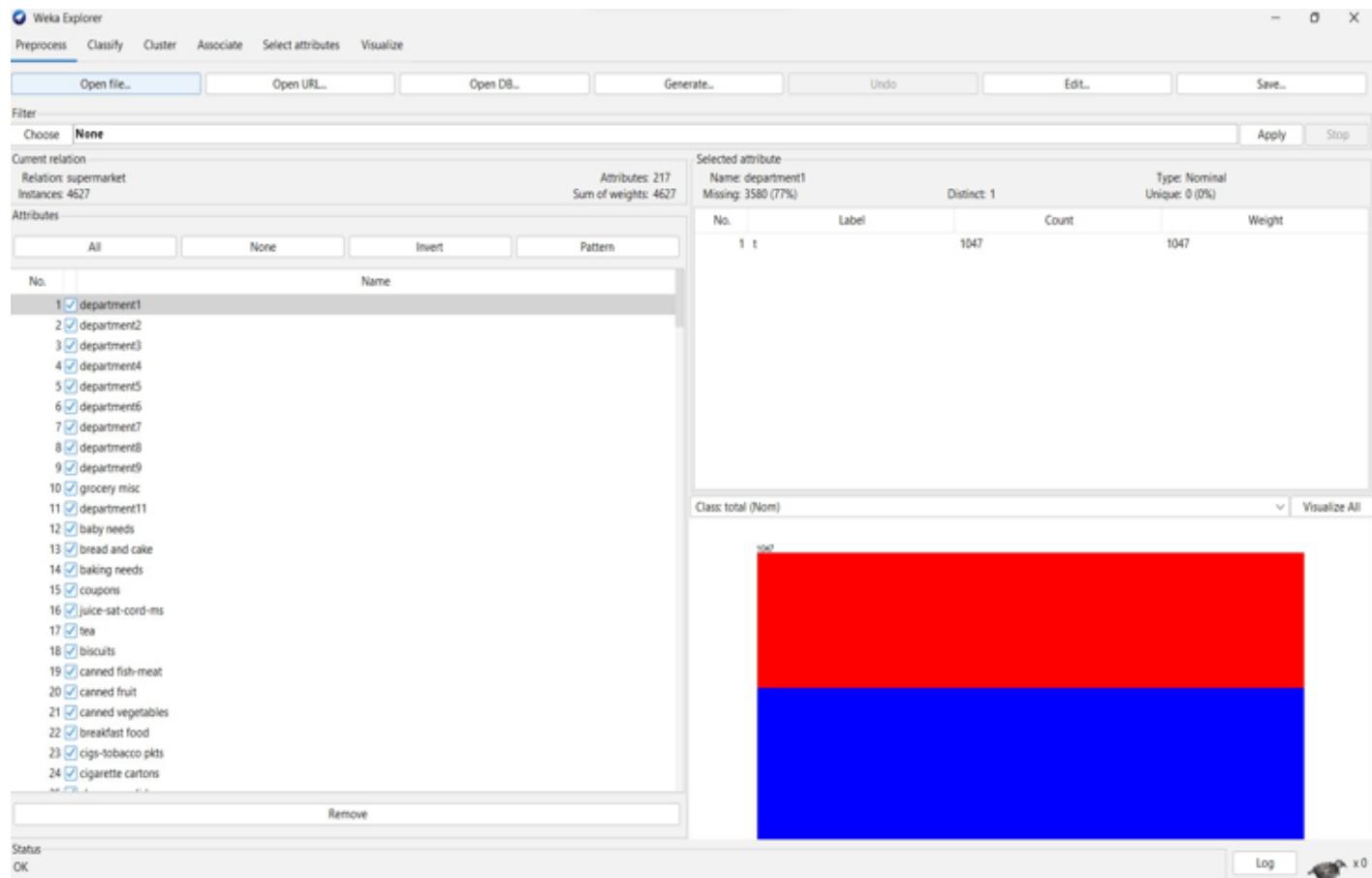


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```
## Visualizing  
clusters  
y_kmeans <-  
  
kmeans.re$clust  
er  
clusplot(iris_1[, c("Sepal.Length",  
"Sepal.Width")], y_kmeans, lines = 0,  
shade = TRUE, color = TRUE, labels = 2,  
plotchar = FALSE, span = TRUE,  
main = paste("Cluster iris"),
```

OUTPUT SCREENSHOT:-





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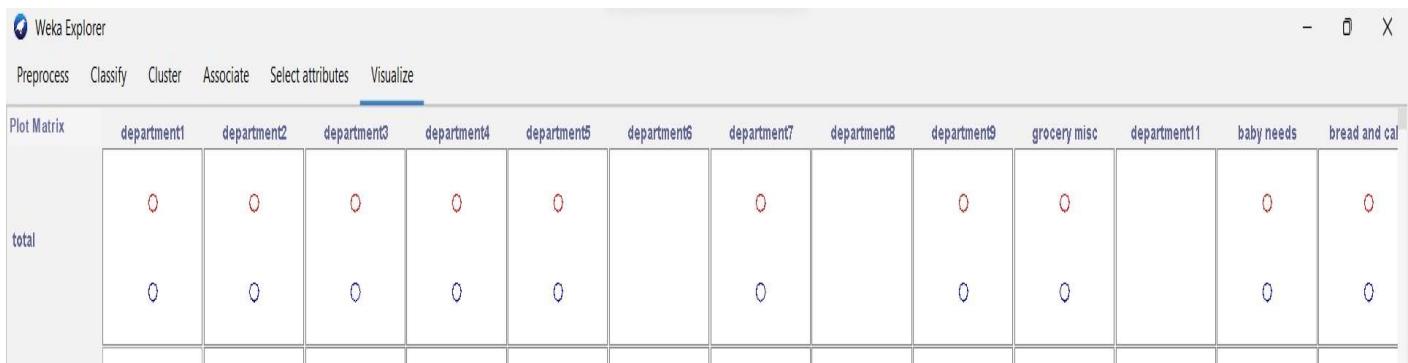
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```
department210          t      t      t
department211          t      t      t
department212          t      t      t
department213          t      t      t
department214          t      t      t
department215          t      t      t
department216          t      t      t
total                  low    low    high

Time taken to build model (percentage split) : 0.12 seconds

Clustered Instances

0      987 ( 63%)
1      587 ( 37%)
```



}