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Vellore Institute of Technology

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19BCE1311

**CSE3506 – ESSENTIALS OF DATA ANALYTICS
LAB-7**

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Tasks for Week-7: Partitioning Based Clustering

Aim: To understand the following operations/functions on 'iris' data and perform similar operations on 'USArrests' dataset based on given instructions.

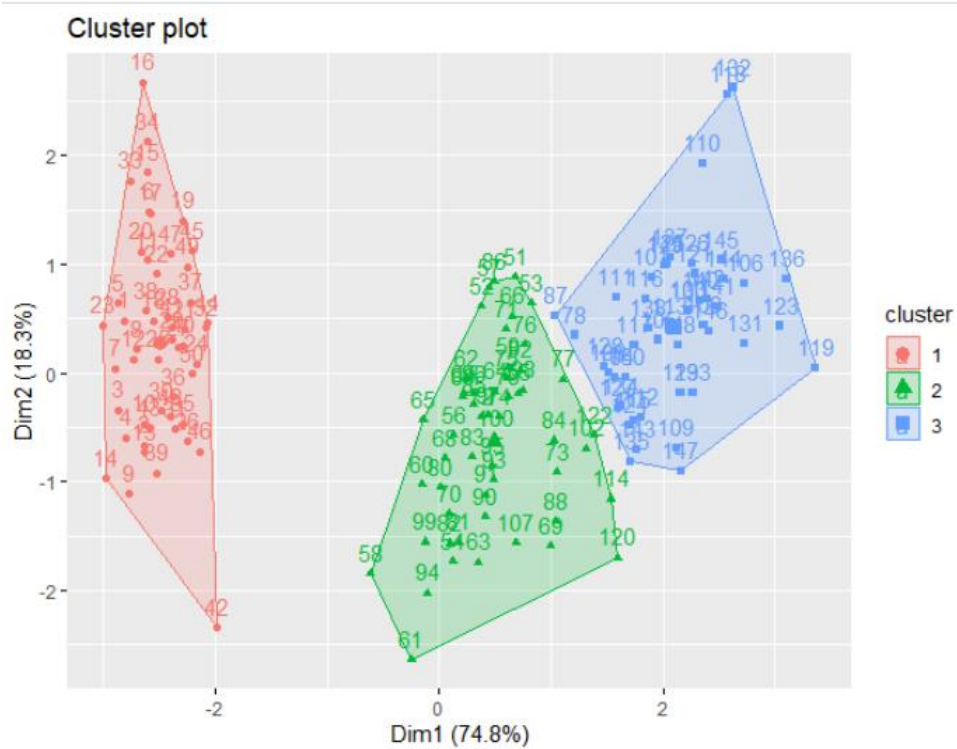
Algorithm:

1. Removing all the values from the global environment
2. Set the working directory to the dataset where we store by using `setwd()`.
3. To see the dataset use `view()` function.
4. By using `scale` function. We scale the data and store it in another variable.
5. Using `kmeans` function we find the kmeans clustering with 2 center at first it can be of any centers and store the result in `fit`.
6. By using `fit$cluster` we can find the cluster values.
7. By using `fit$size` we can find the size of each cluster.
8. By using `fit$withinss` we can find with in cluster sum of squares for each cluster.
9. By using `fit$tot.withinss` we can find with in cluster sum of squares with respective to all clusters.
10. Create the no of iterations we need to find the perfect cluster and size of `wcss` and the `nclust` list.
11. To find the best no of center from 1 to 15 we create a for loop
 1. find the kmeans cluster with each center value in for loop
 2. put to the total with in cluster sum of squares for each iteration in `wcss`
 3. put the size of cluster in `nclust`.
12. Plot the graph between the no of center and the `wcss` values for each center. the place where we find the bend that is our no of cluster should be taken.
13. In other way we can use `factoextra` library.
14. Using `fviz_nbclust` function we can find the graph.
15. Using `fviz_cluster` function we can find the clusters
16. Call `cluster` library
17. We use `pam` function to find the k medoid clusters and store the values in `fitm`.
18. By using the `fitm$medoid` we can find no of medoid.
19. Using `fviz_cluster` function we can find the medoids.

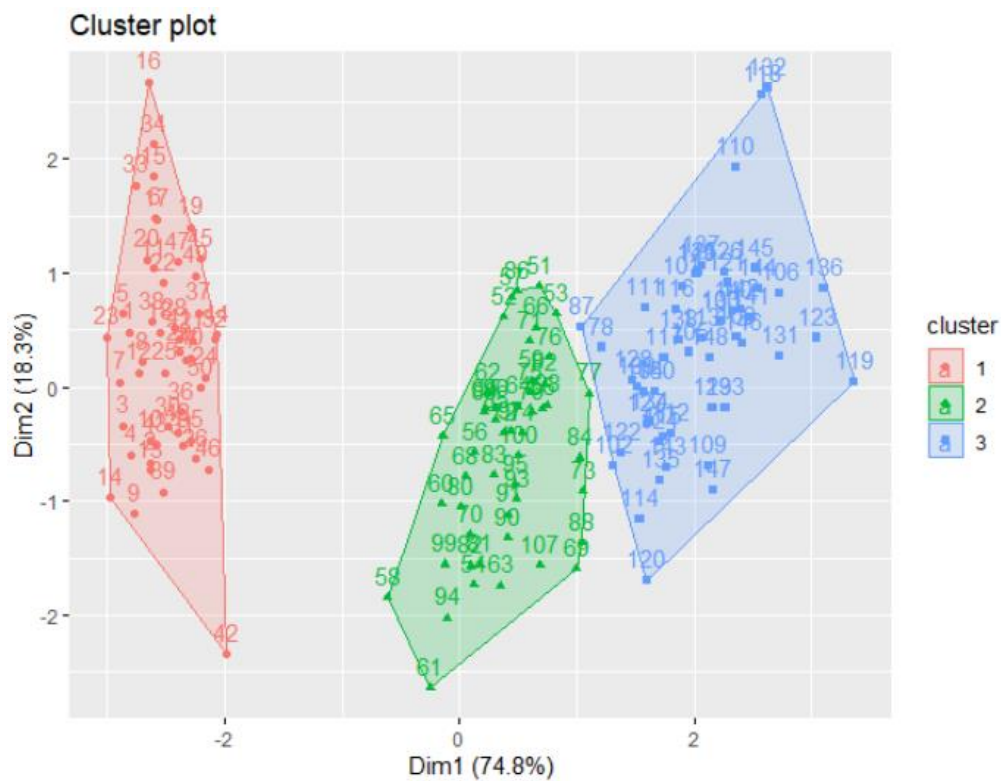
Result:

For iris.csv

K-means centers:

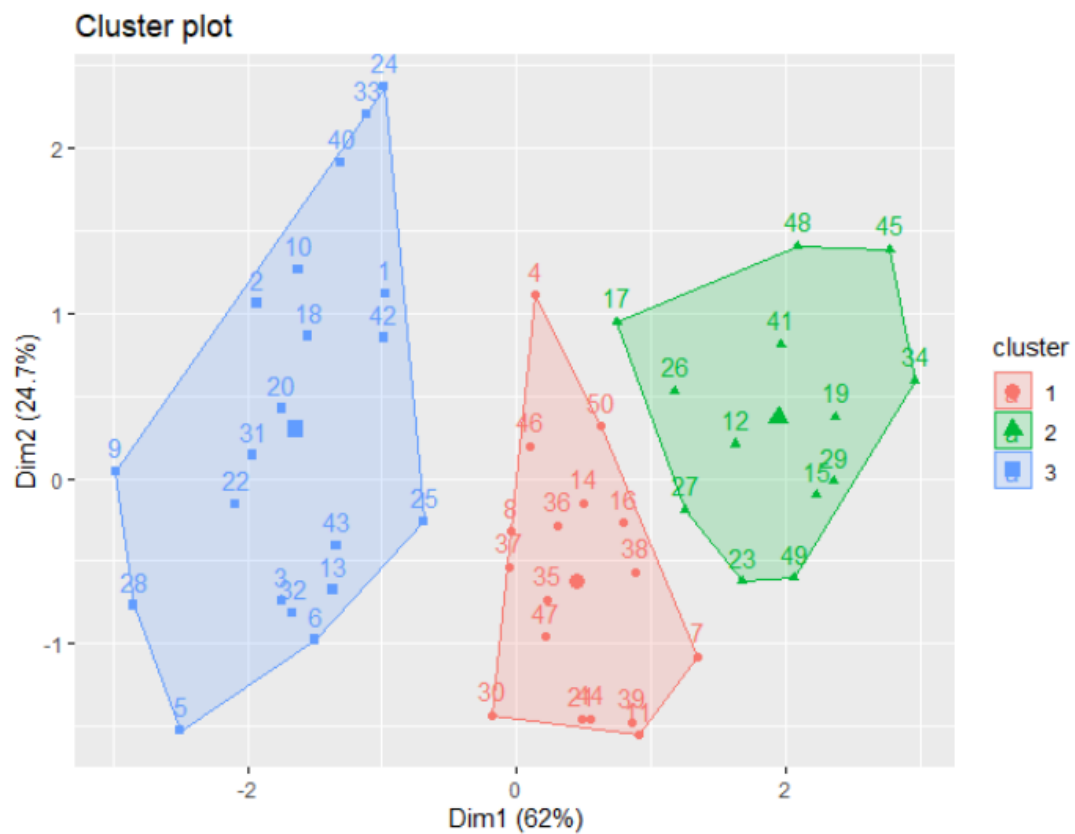


K-medoid centers:

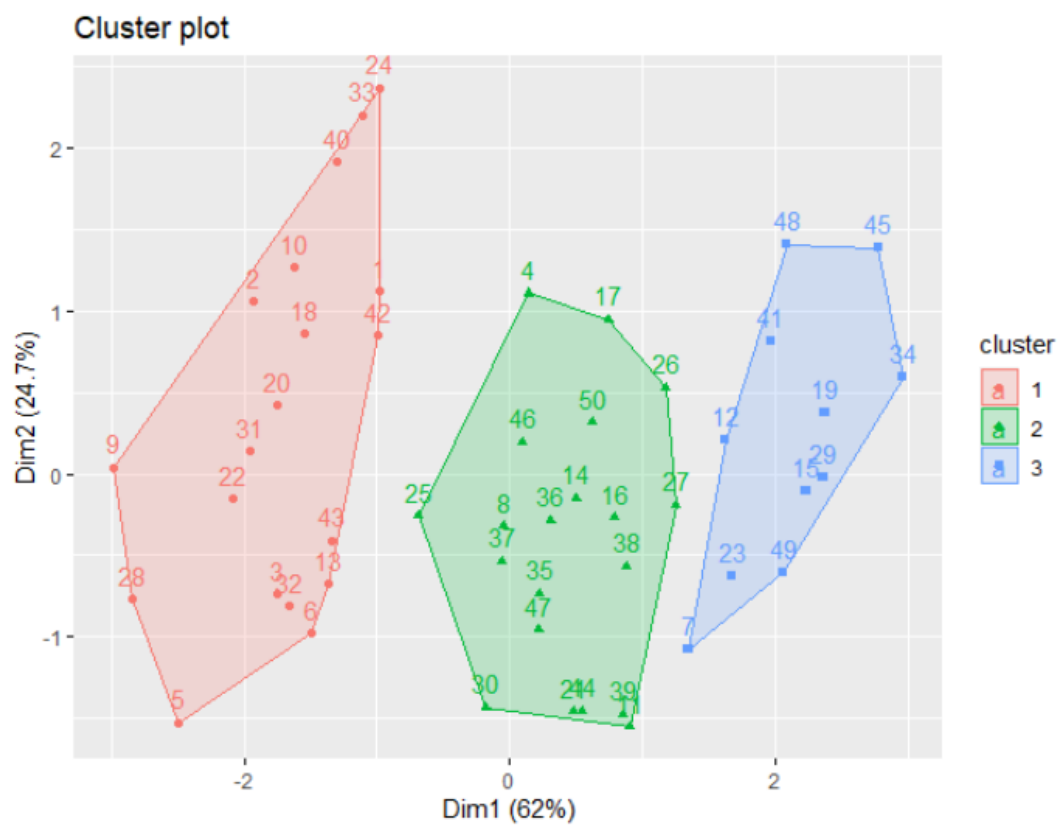


For USArrests.csv

K-means centers:



K-medoid centers:



Statistics:

For iris.csv

- K-means centers

X	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
-1.15087068	-1.01119138	0.85041372	-1.3006301	-1.2507035
0.07534946	0.03881135	-0.73324663	0.3059615	0.2137533
1.13936197	1.03196952	-0.07784286	1.0386287	1.0894947

- K-medoid centers

X	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
-1.07030973	-0.7769106	0.7861738	-1.3357516	-1.3110521
-0.08056095	0.3099591	-0.5903951	0.1370873	0.1320673
0.95522266	0.7930124	-0.1315388	0.9868021	0.7880307

For USArrests.csv

- K-means centers

Murder	Assault	UrbanPop	Rape
-0.4469795	-0.3465138	0.4788049	-0.2571398
-0.9615407	-1.1066010	-0.9301069	-0.9667633
1.0049340	1.0138274	0.1975853	0.8469650

- K-medoid centers

Murder	Assault	UrbanPop	Rape
0.8292944	1.3708088	0.3081225	1.1603196
-0.2727580	-0.2371077	0.1699510	-0.1315342
-1.2829727	-1.3770485	-0.5899924	-1.0603878

Program:

For iris.csv

```
rm(list=ls())
setwd("C:/Users/Abhinav Vijayakumar/Desktop/VIT Academics/Sem 6/EDA/LAB/LAB 7")
data1<-read.csv("iris.csv")
View(data1)
df<-scale(data1)
fit<-kmeans(df,centers=2)
fit$cluster
fit$size
fit$withinss
fit$tot.withinss
Kmax<-15
wcsc<-rep(NA,Kmax)
nClust<- list()
for(i in 1:Kmax){
  fit<-kmeans(df,i)
  wcsc[i]<-fit$tot.withinss
  nClust[[i]]<-fit$size
}
plot(1:Kmax,wcsc,type="b",pch=19)
fit<-kmeans(df,centers=3)
fit$cluster
fit$size
fit$center
library(factoextra)
fviz_nbclust(df, kmeans, method = "wss")
fviz_cluster(fit, data1)
library(cluster)
fitm <- pam(df, 3, metric = "manhattan")
fitm
fitm$medoids
fviz_cluster(fitm, data1)
```

For USArrests.csv

```
rm(list=ls())
setwd("C:/Users/Abhinav Vijayakumar/Desktop/VIT Academics/Sem 6/EDA/LAB/LAB 7")
data2<-read.csv("USArrests.csv")
view(data2)
data2<-data2[,-1]
df1<-scale(data2)
fit1<-kmeans(df1,centers=2)
fit1$cluster
fit1$size
fit1$withinss
fit1$tot.withinss
Kmax1<-15
wcss1<-rep(NA,Kmax1)
nClust1<- list()
for(i in 1:Kmax1){
  fit1<-kmeans(df1,i)
  wcss1[i]<-fit1$tot.withinss
  nClust1[[i]]<-fit1$size
}
plot(1:Kmax1,wcss1,type="b",pch=19)
fit1<-kmeans(df1,centers=3)
fit1$cluster
fit1$size
fit1$center
library(factoextra)
fviz_nbclust(df1, kmeans, method = "wss")
fviz_cluster(fit1, data2)
library(cluster)
fitm1 <- pam(df1, 3, metric = "manhattan")
fitm1
fitm1$medoids
fviz_cluster(fitm1, data2)
```