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Vellore Institute of Technology

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19BCE1311

CSE3506 – ESSENTIALS OF DATA ANALYTICS
LAB-8

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Tasks for Week-8: Hierarchical Clustering

Aim: To understand the following operations/functions on 'USArrests' data and perform similar operations on 'iris' 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 `dist` function we find the Euclidean distances for the scaled data.
6. By using the Euclidean distances and `hclust` function we can create and then plot the hierarchical clustering dendrogram.
7. By using `cutree` we divide the elements of the dendrogram into k number of clusters ($k=4$ in our case).
8. Then, by using `rect.hclust` function we can divide the dendrogram into k clusters ($k=4$ in our case) , i.e. create k rectangular divisions/borders in the dendrogram

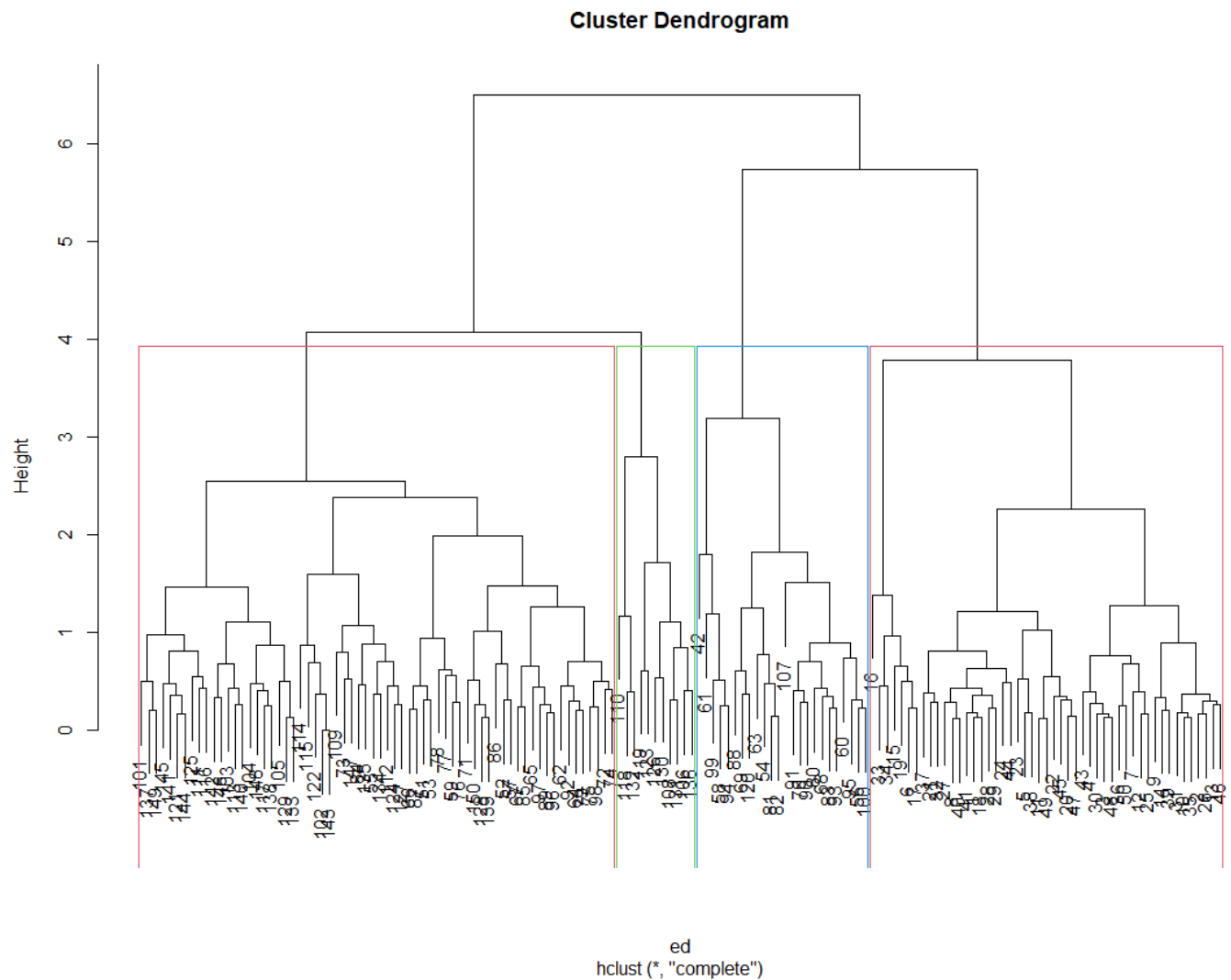
Result:

For iris.csv

Cluster:

[illegible]

Cluster Dendrogram:



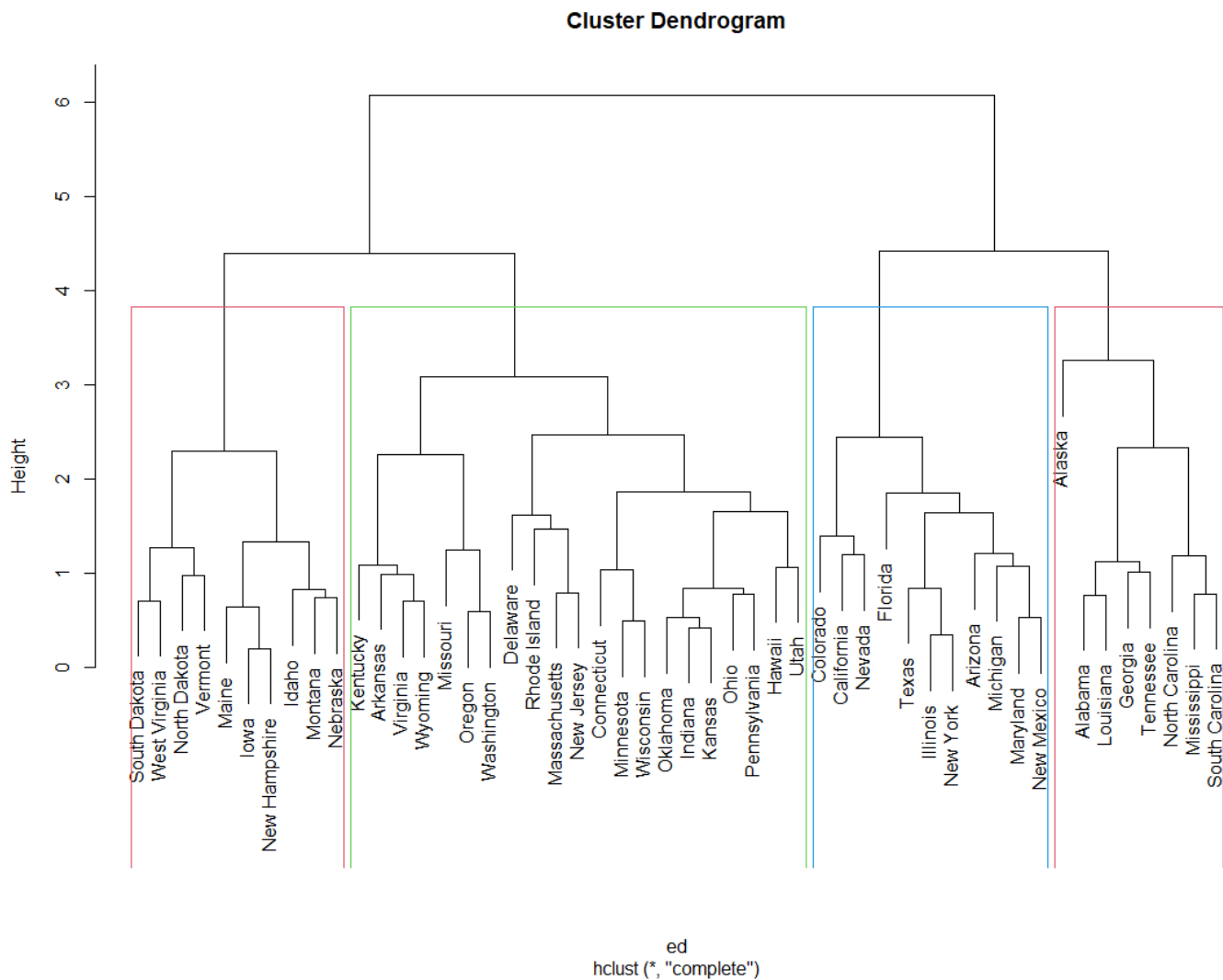
For USArrests.csv

Cluster:

```
> cluster
```

Alabama	Alaska	Arizona	Arkansas	California	Colorado	Connecticut	Delaware
1	1	2	3	2	2	3	3
Florida	Georgia	Hawaii	Idaho	Illinois	Indiana	Iowa	Kansas
2	1	3	4	2	3	4	3
Kentucky	Louisiana	Maine	Maryland	Massachusetts	Michigan	Minnesota	Mississippi
3	1	4	2	3	2	3	1
Missouri	Montana	Nebraska	Nevada	New Hampshire	New Jersey	New Mexico	New York
3	4	4	2	4	3	2	2
North Carolina	North Dakota	Ohio	Oklahoma	Oregon	Pennsylvania	Rhode Island	South Carolina
1	4	3	3	3	3	3	1
South Dakota	Tennessee	Texas	Utah	Vermont	Virginia	Washington	West Virginia
4	1	2	3	4	3	3	4
Wisconsin	Wyoming						
3	3						

Cluster Dendrogram:



Program:

For iris.csv

```
rm(list=ls())  
setwd("C:/Users/Abhinav Vijayakumar/Desktop/VIT Academics/Sem 6/EDA/LAB/LAB 8")  
  
data <- read.csv("iris.csv", row.names=1)  
View(data)  
  
df <- scale(data)  
View(df)  
  
ed <- dist(df, method = 'euclidean')
```

```
hierClust <- hclust(ed, method = 'complete')
plot(hierClust)

cluster <- cutree(hierClust, k = 4)
cluster

rect.hclust(hierClust, k = 4, border = 2:4)
```

For USArrests.csv

```
rm(list=ls())
setwd("C:/Users/Abhinav Vijayakumar/Desktop/VIT Academics/Sem 6/EDA/LAB/LAB 8")

data <- read.csv("USArrests.csv", row.names=1)
View(data)

df <- scale(data)
View(df)

ed <- dist(df, method = 'euclidean')
hierClust <- hclust(ed, method = 'complete')
plot(hierClust)

cluster <- cutree(hierClust, k = 4)
cluster

rect.hclust(hierClust, k = 4, border = 2:4)
```