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**Vellore Institute of Technology**  
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# **Essentials of Data Analytics - (CSE3506)**

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## **Lab-8**

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**L21-22 Slot**

## **Tasks for Week-8: Hierarchical Clustering**

Understand the following operations/functions on 'USArrests' data and perform similar operations on 'iris' dataset based on given instructions.

## **AIM**

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

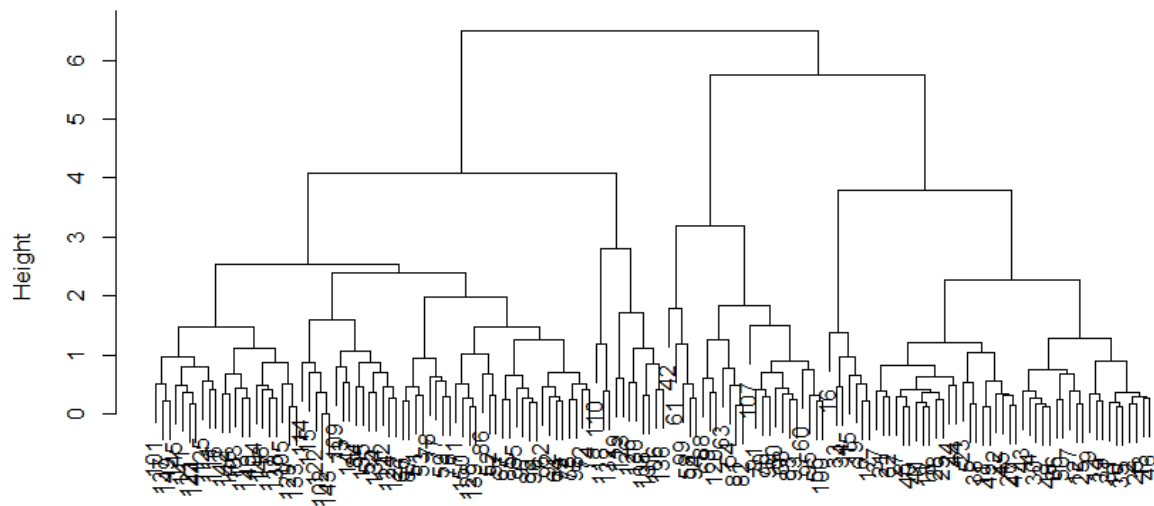
## **Algorithm**

1. Start
2. Use `rm(list=ls())` to clear commands.
3. Read the dataset.
4. Make a scaled version of the original data.
5. Find the euclidean distance between each row
6. Perform hierarchical clustering with 'complete' as the method.
7. Perform hierarchical clustering using the `hclust` method with the 'complete' method.
8. Plot 'heirClust' to get the dendrogram.
9. `Cutree` cuts the data into several groups either by specifying the desired number(s) of groups or the cut height(s).
10. `rect.cluster` is used to get the dendrogram
11. Stop.

# Result

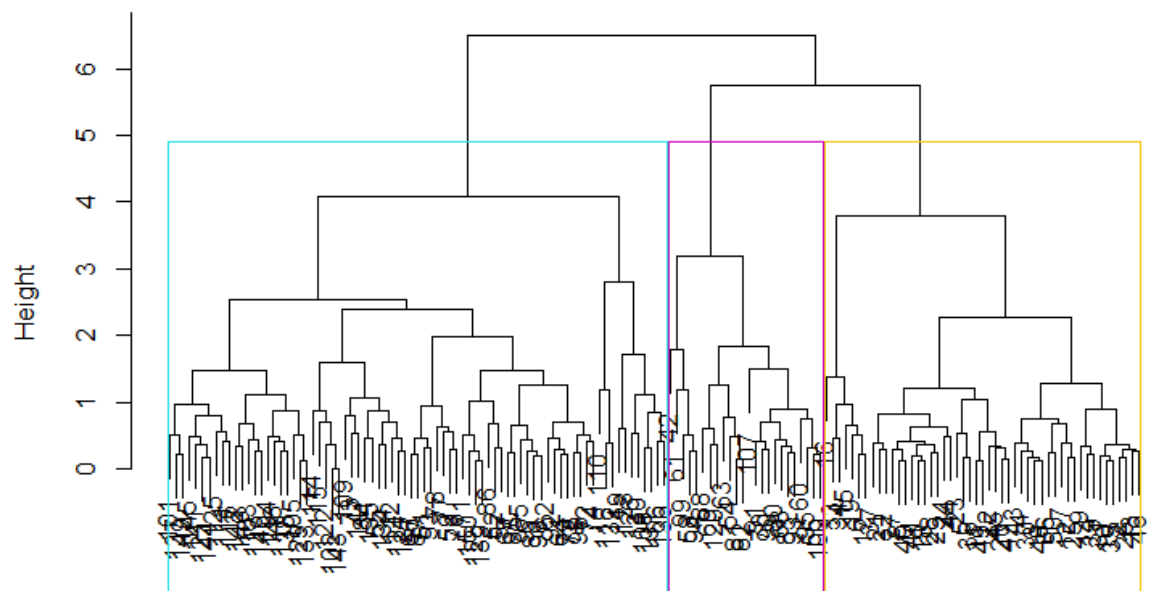
## Case 1: Iris dataset

Cluster Dendrogram



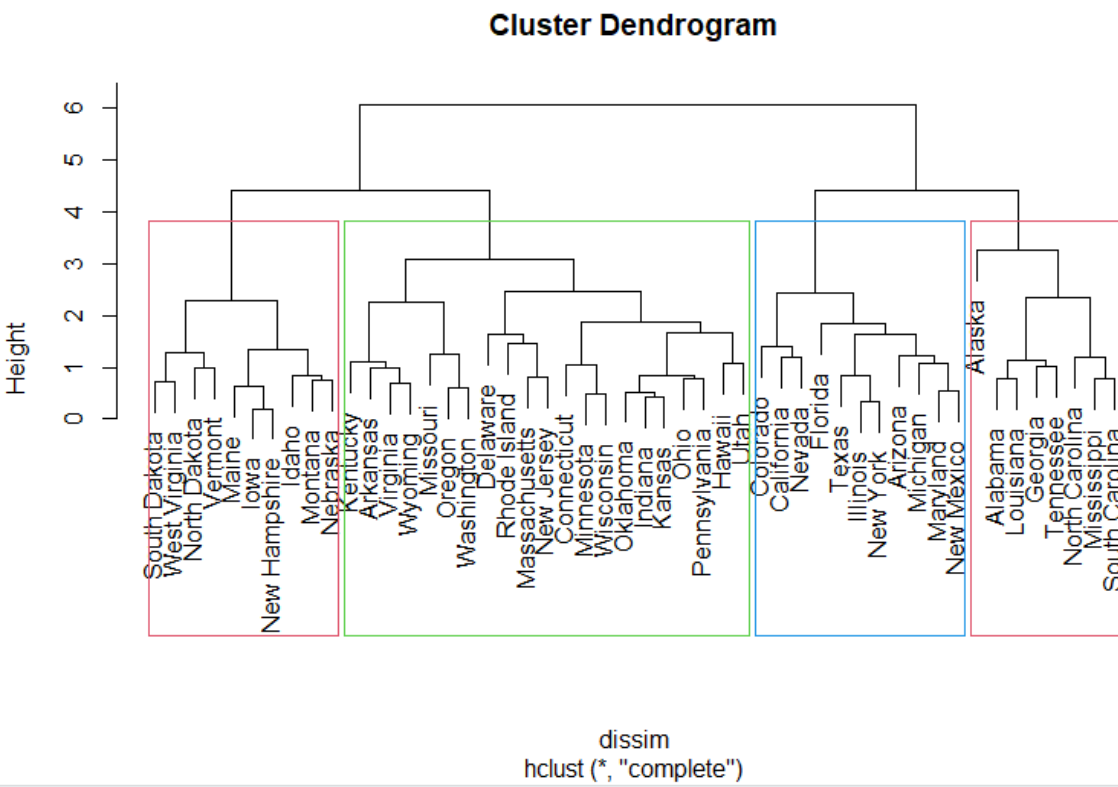
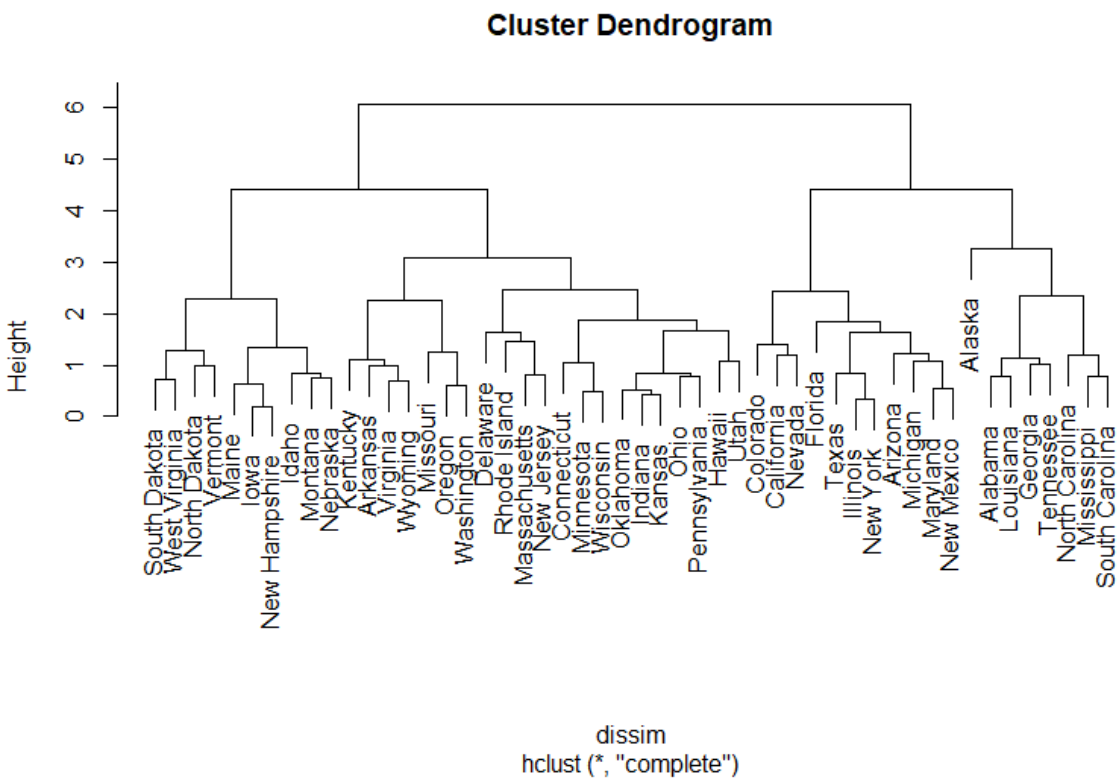
ds  
hclust (\*, "complete")

Cluster Dendrogram



ds  
hclust (\*, "complete")

# Case II: USArrests Dataset.



# Program

## Case 1: Iris dataset

```
rm(list=ls())  
setwd("D:/data analytics/")  
data <- read.csv("D:/6th Sem Works/A2- EDA/LAB/Lab8/iris.csv",row.names=1)  
View(data)  
df <- scale(data)  
View(df)  
ds <- dist(df, method = 'euclidean')  
ds  
hierClust <- hclust(ds, method = 'complete')  
plot(hierClust)  
cluster <- cutree(hierClust, k = 3)  
cluster  
rect.hclust(hierClust,k=3,border = 5:7)
```

## Case II: USArrests Dataset.

```
rm(list=ls())  
data <- read.csv("D:/6th Sem Works/A2- EDA/LAB/Lab8/USArrests.csv",row.names=1)  
df <- scale(data)  
dissim <- dist(df, method = 'euclidean')  
hierClust <- hclust(dissim, method = 'complete')  
plot(hierClust)  
cluster <- cutree(hierClust, k = 4)  
rect.hclust(hierClust, k = 4, border = 2:4)
```