10 PROVE: CLUSTERING THE STATES

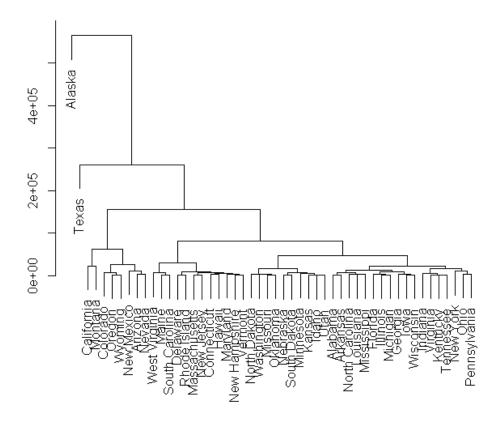
```
library(tidyverse)
library(datasets)
library(cluster)

data <- state.x77

# Computing the distance matrix distance <- dist(as.matrix(data))

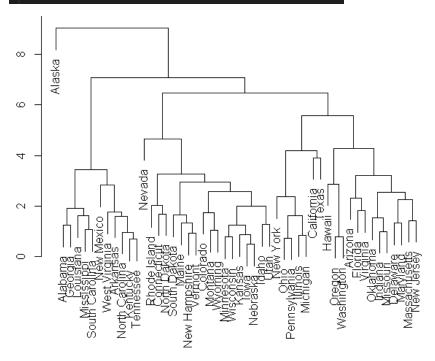
# Clustering hc <- hclust(distance)

# Plotting a dendogram plot(hc)</pre>
```

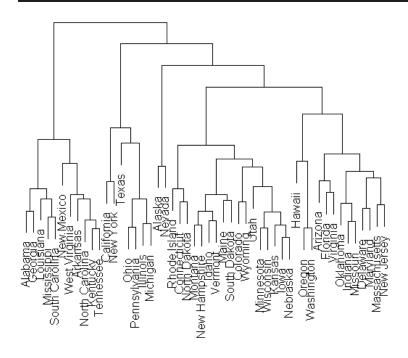


```
# Scaling data (normalization)
data.scaled <- scale(data)

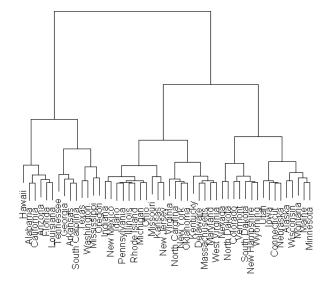
# dendogram with normalized data
distance <- dist(as.matrix(data.scaled))
hc <- hclust(distance)
plot(hc)</pre>
```



```
# dendogram without Area
data.scaled_noArea <- scale(data[,1:7])
distance <- dist(as.matrix(data.scaled_noArea))
hc <- hclust(distance)
plot(hc)</pre>
```



```
# dendogram only with Frost
data.scaled_Frost <- scale(data[,7])
distance <- dist(as.matrix(data.scaled_Frost))
hc <- hclust(distance)
plot(hc)</pre>
```



```
# K-means CLustering
clust <- kmeans(data.scaled, 3) # k = 5
summary(clust)</pre>
```

Centers of the clusters (mean values)
clust\$centers

```
Length Class Mode
cluster
centers
                     24
                                -none- numeric
                              -none- numeric
                           -none- numeric
-none- numeric
betweenss
                              -none- numeric
                               -none- numeric
                                -none- numeric
                               -none- numeric
> clust$centers
Population Income Illiteracy Life Exp Murder HS Grad
1 -0.2269956 -1.3014617 1.391527063 -1.1773136 1.0919809 -1.4157826
2 -0.4873370 0.1329601 -0.641201154 0.7422562 -0.8552439 0.5515044
3 0.9462026 0.741609 0.005468667 -0.3242467 0.5676042 0.1558335
                         Area
          Frost
1 -0.7206500 -0.2340290
2 0.4528591 -0.1729366
3 -0.1960979 0.4483198
```

Clusters clust\$cluster

```
Alaska
                                    Arizona
                                                  Arkansas
                                                               California
    Colorado
                Connecticut
                                   Delaware
                                                   Florida
                                                                  Georgia
                                  Illinois
      Hawaii
                      Idaho
                                                   Indiana
                                                                     Iowa
                                                     Maine
                                                                 Maryland
                                                                 Missouri
Massachusetts
                   Michigan
                                  Minnesota
                                               Mississippi
                   Nebraska
                                    Nevada New Hampshire
     Montana
                                                               New Jersey
  New Mexico
                   New York North Carolina
                                              North Dakota
                                                                     Ohio
    Oklahoma
                     Oregon
                               Pennsylvania
                                              Rhode Island South Carolina
South Dakota
                  Tennessee
                                                                  Vermont
    Virginia
                 Washington West Virginia
                                                                  Wyoming
```

```
# Within-cluster sum of squares
clust$withinss
# Total sum of squares across clusters
clust$tot.withinss
```

```
3 2

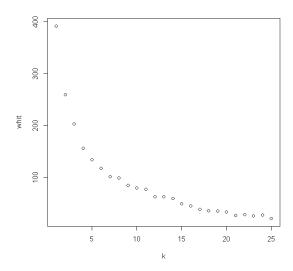
> clust$withinss

[1] 23.62227 67.72742 111.66951

> clust$tot.withinss

[1] 203.0192
```

```
# Plotting k-means clusters
clusplot(data.scaled, clust$cluster, color = T, shaed = T, labels = 2, lines = 0)
# CHOOSING NUMBER OF K USING ELBOW METHOD
whit <- c()
k <- c()
for (i in 1:25) {
    clust <- kmeans(data.scaled, i)
    whit[[i]] <- clust$tot.withinss
    k[[i]] <- i
}
elbow <- cbind(k, whit) %>%
    data.frame()
plot(elbow)
```



```
clust <- kmeans(data.scaled, 5)
clust$cluster
clusplot(data.scaled, clust$cluster, color = T, shaed = T, labels = 2, lines = 0)</pre>
```

