

Individual Assignment 10

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R Markdown

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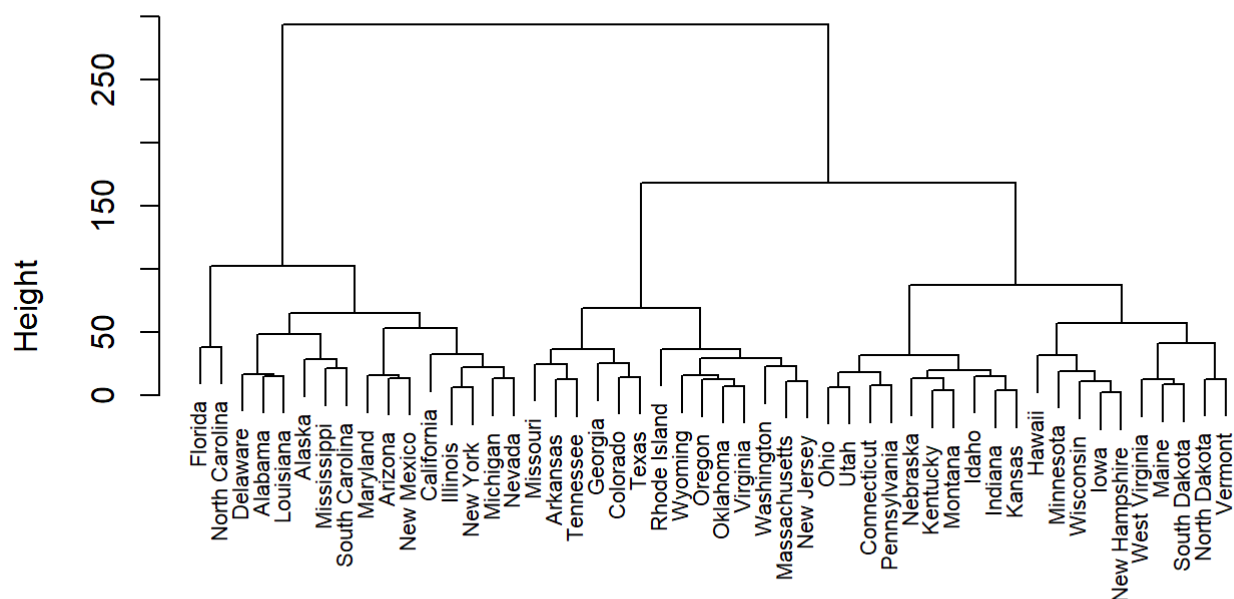
Exercise 10.7: Problem 9

a.

```
set.seed(114514)

hc = hclust(dist(USArrests), method = 'complete')
plot(hc, cex = 0.7)
```

Cluster Dendrogram



```
dist(USArrests)
hclust (*, "complete")
```

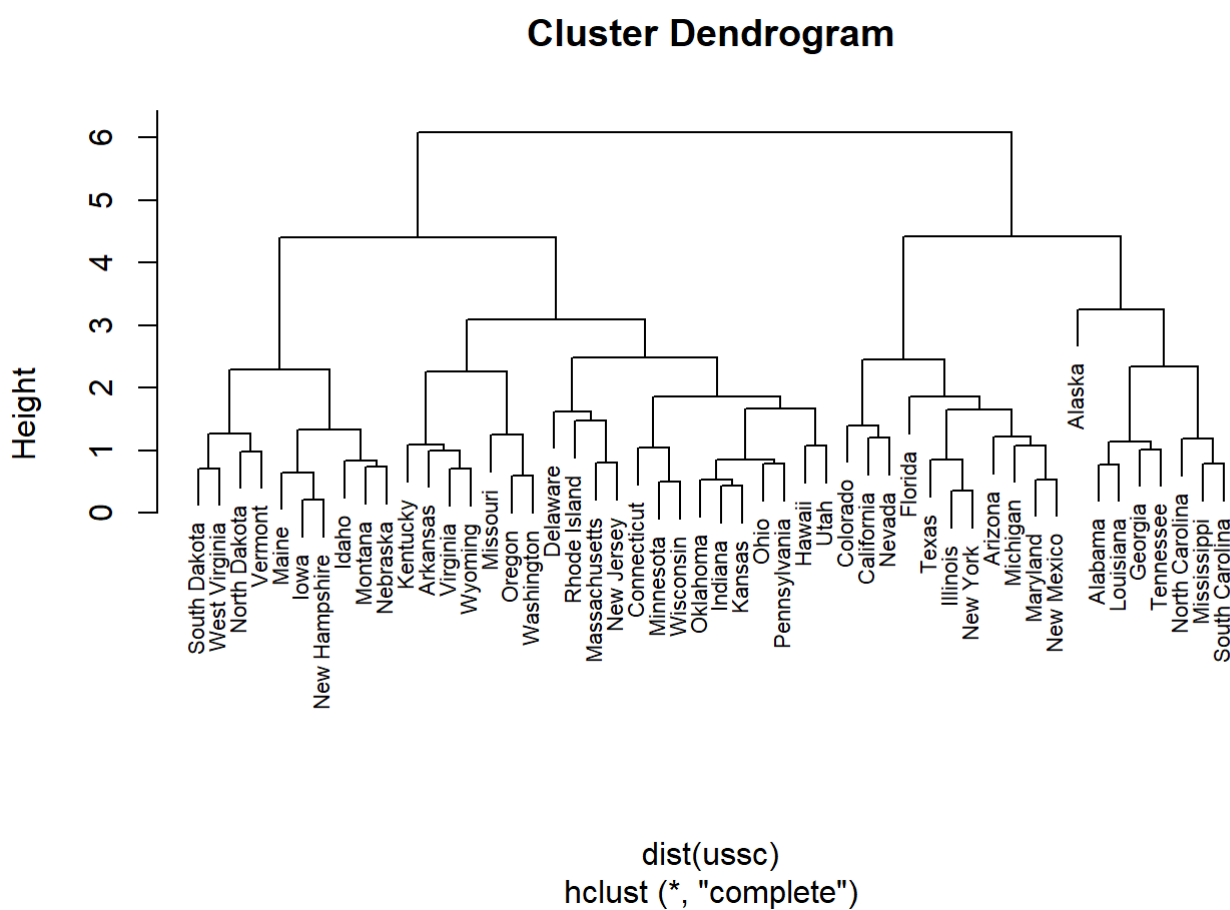
b.

```
hccut = cutree(hc, k=3)
hccut
```

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

c.

```
ussc = scale(USArrests)
hcsc = hclust(dist(ussc), method = 'complete')
plot(hcsc, cex = 0.7)
```



- d. We can find that original model has a height of 10, and the scaled model is 6, means that the clusters are different. In general, variables with different measurement units should be scaled before computing the inter-observation dissimilarities. From this model we can also scale the variable to make every variable has a equal measurement.