

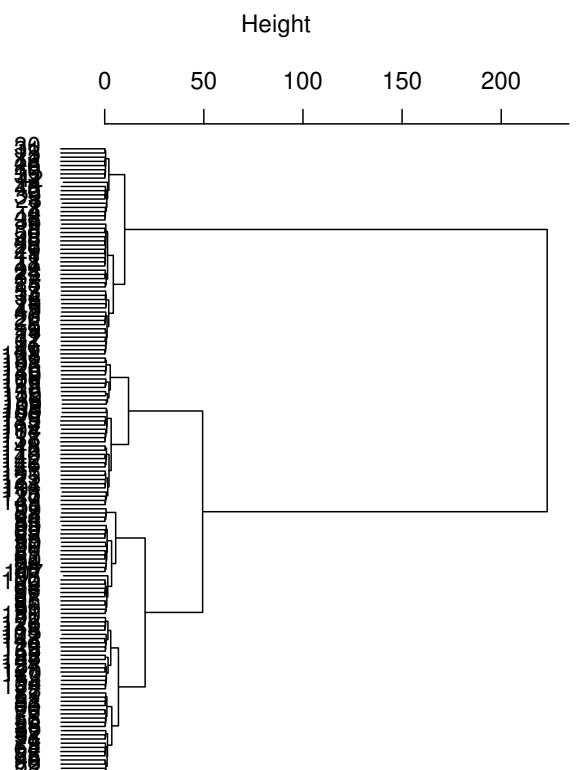
## R Solution Exercise Sheet 3: Cluster Analysis

### Computer Problems:

- (a) 

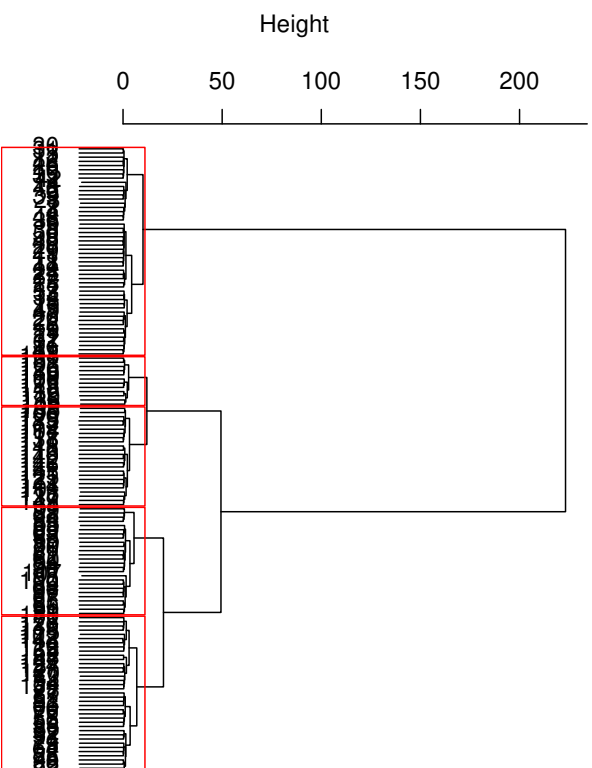
```
# Ward Hierarchical Clustering
d <- dist(iris, method = "euclidean") # distance matrix
fit <- hclust(d, method="ward")
plot(fit) # display dendrogram
groups <- cutree(fit, k=5) # cut tree into 5 clusters
# draw dendrogram with red borders around the 5 clusters
rect.hclust(fit, k=5, border="red")
```

**Cluster Dendrogram**



`d`  
`hclust (*, "ward.D")`

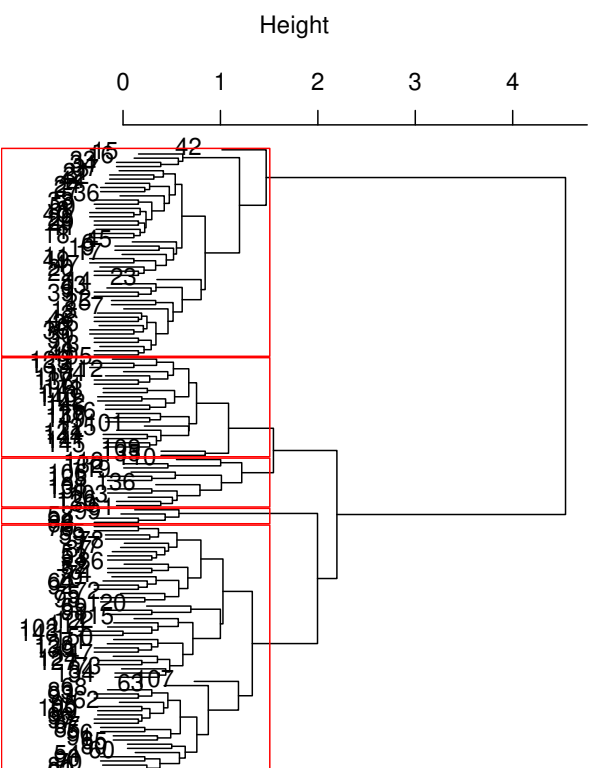
Cluster Dendrogram



d  
hclust(\*, "ward.D")

(b) # Average Hierarchical Clustering  
d <- dist(iris, method = "euclidean") # distance matrix  
fit <- hclust(d, method="average")  
plot(fit) # display dendrogram  
groups <- cutree(fit, k=5) # cut tree into 5 clusters  
# draw dendrogram with red borders around the 5 clusters  
rect.hclust(fit, k=5, border="red")

Cluster Dendrogram



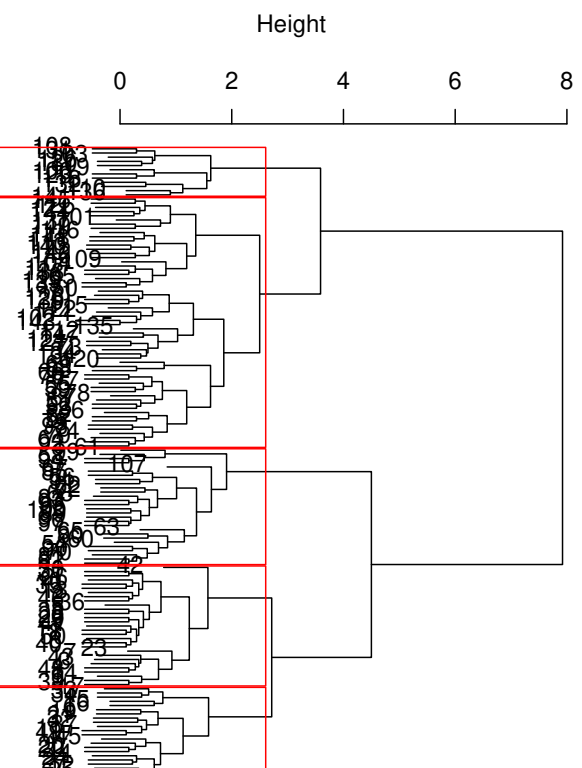
```

d
hclust(*, "average")

(c) # Complete Hierarchical Clustering
d <- dist(iris, method = "euclidean") # distance matrix
fit <- hclust(d, method="complete")
plot(fit) # display dendrogram
groups <- cutree(fit, k=5) # cut tree into 5 clusters
# draw dendrogram with red borders around the 5 clusters
rect.hclust(fit, k=5, border="red")

```

Cluster Dendrogram



```

d
hclust(*, "complete")

(d) # Single Hierarchical Clustering
d <- dist(iris, method = "euclidean") # distance matrix
fit <- hclust(d, method="single")
plot(fit) # display dendrogram
groups <- cutree(fit, k=5) # cut tree into 5 clusters
# draw dendrogram with red borders around the 5 clusters
rect.hclust(fit, k=5, border="red")

```



```

[77] 2 3 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 3 2 3 3 3 3 2 3 3
[115] 2 3 3 3 3 2 3 2 3 2 3 3 2 2 3 3 3 3 3 2 3 3 3 3 2 3 3 3 2 3 3 3 2

```

Within cluster sum of squares by cluster:

```

[1] 15.15100 39.82097 23.87947
(between_SS / total_SS = 88.4 %)

```

Available components:

```

[1] "cluster"      "centers"      "totss"        "withinss"     "tot.wi
[6] "betweenss"    "size"         "iter"         "ifault"
>

```

# K-Means Clustering with 2 clusters

```
fit2 = kmeans(ir, 2)
```

```
fit2
```

K-means clustering with 2 clusters of sizes 97, 53

Cluster means:

```

      Sepal.Length Sepal.Width Petal.Length Petal.Width
1      6.301031      2.886598      4.958763      1.695876
2      5.005660      3.369811      1.560377      0.290566

```

Clustering vector:

```

[1] 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
[39] 2 2 2 2 2 2 2 2 2 2 2 2 1 1 1 1 1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1
[77] 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 2 1 1 1 1 2 1 1 1 1 1 1 1 1 1 1
[115] 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1

```

Within cluster sum of squares by cluster:

```

[1] 123.79588 28.55208
(between_SS / total_SS = 77.6 %)

```

Available components:

```

[1] "cluster"      "centers"      "totss"        "withinss"     "tot.wi
[6] "betweenss"    "size"         "iter"         "ifault"

```

(b)

```
library(cluster)
```

```
clusplot(ir, fit3$cluster, color=TRUE, shade=TRUE,
         labels=2, lines=0)
```

