## Homework 5

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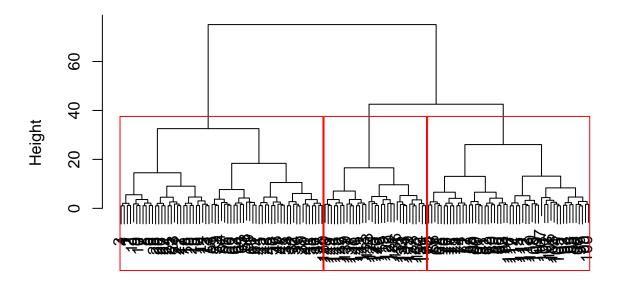
#### Problem 1

```
library(stats)
data = read.csv(file = "~/Documents/Math189/iris.csv", header = TRUE, fill = TRUE)
iris = data[,2:6]

#subset of data that is purely integer
pureIris = data[,1:4]

d1 = dist(pureIris, method = "euclidean")
hc1 = hclust(d1, method = "average")
clusterCut1 = cutree(hc1, k = 3)
plot(hc1, main = "Cluster Dendogram of IRIS Data")
rect.hclust(hc1, k=3, border="red")
```

### **Cluster Dendogram of IRIS Data**

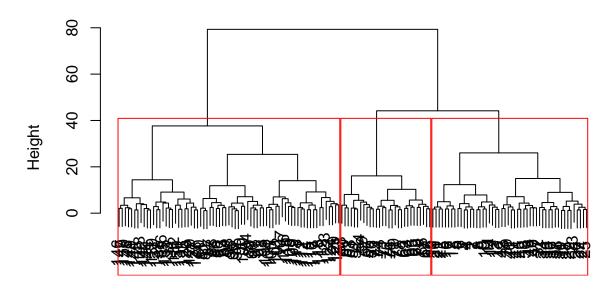


d1 hclust (\*, "average")

```
#Correct clustering rate is
table1 = table(clusterCut1,iris$Species)
paste("Correct rate is ", sum(diag(table1))/150*100 , "%")
```

```
########----- Part (1) -----######
#(1) Retaining all the settings in part 1, except using the Taxicab
#distance as the similarity measure.
d2 = dist(pureIris, method = "manhattan")
hc2 = hclust(d2, method = "average")
clusterCut2 = cutree(hc2, k = 3)
plot(hc2, main = "Cluster Dendogram of IRIS Data")
rect.hclust(hc2, k=3, border="red")
```

#### **Cluster Dendogram of IRIS Data**

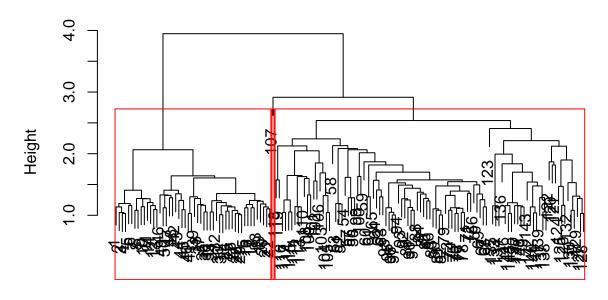


d2 hclust (\*, "average")

```
table(clusterCut2,iris$Species)
##
## clusterCut2 setosa versicolor virginica
##
             1
                   50
                               0
             2
                    0
                              29
##
                                         0
##
             3
                              21
                                        50
#Correct clustering rate is
table2 = table(clusterCut2,iris$Species)
paste("Correct rate is ", sum(diag(table2))/150*100 , "%")
## [1] "Correct rate is 86 %"
#######----- Part (2) -----#####
#(2) Retaining all the settings in part 1, except using the single
#linkage as the linkage function.
d3 = dist(pureIris, method = "euclidean")
hc3 = hclust(d3, method = "single")
```

```
clusterCut3 = cutree(hc3, k = 3)
plot(hc3, main = "Cluster Dendogram of IRIS Data")
rect.hclust(hc3, k=3, border="red")
```

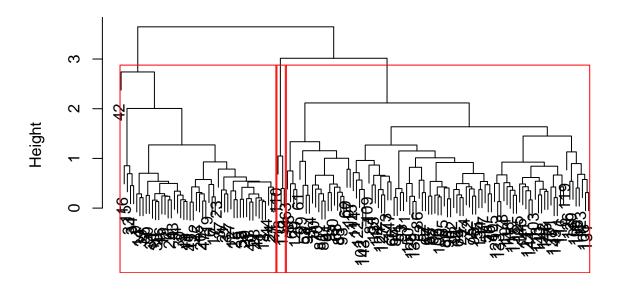
#### **Cluster Dendogram of IRIS Data**



d3 hclust (\*, "single")

```
table(clusterCut3,iris$Species)
## clusterCut3 setosa versicolor virginica
                  50
##
            1
                             0
            2
                            50
##
                  0
                                      49
#Correct clustering rate is
table3 = table(clusterCut3,iris$Species)
paste("Correct rate is ", sum(diag(table3))/150*100 , "%")
#######----- Part (3) -----#####
#(3) Retaining all the settings in part 1, except standardizing
#each variable before applying the clustering. (Standardize each
#column in the dataset to have zero mean and unit variance).
#standardize variables
standardizIris = scale(iris[,1:4])
d4 = dist(standardizIris, method = "euclidean")
hc4 = hclust(d4, method = "average")
clusterCut4 = cutree(hc4, k = 3)
plot(hc4, main = "Cluster Dendogram of IRIS Data")
```

# **Cluster Dendogram of IRIS Data**



d4 hclust (\*, "average")

```
table(clusterCut4,iris$Species)
##
```

```
## clusterCut4 setosa versicolor virginica
## 1 50 0 0
## 2 0 50 47
## 3 0 0 3
```

```
#Correct clustering rate is
table4 = table(clusterCut4,iris$Species)
paste("Correct rate is ", sum(diag(table4))/150*100 , "%")
```