Lab 12: Clustering

Donggyun Kim 27008257 4/16/2018

K-means clustering

```
my_kmeans <- function(X, k) {</pre>
  X <- as.matrix(X)</pre>
  N \leftarrow nrow(X)
  P \leftarrow ncol(X)
  K <- k
  cluster_sizes <- numeric(K)</pre>
  cluster_means <- matrix(0, nrow = K, ncol = P)</pre>
  clustering_vector <- numeric(N)</pre>
  wss_cluster <- numeric(K)</pre>
  bss_over_tss <- numeric(1)</pre>
  index <- sample(N, K)</pre>
  centroids <- X[index, ]</pre>
  centroids_new <- X[index, ]</pre>
  distant2 <- matrix(0, nrow = N, ncol = K)</pre>
  1 <- 1
  iteration <- 100
  while (1 <= iteration) {</pre>
    for (i in 1:N) {
      for(j in 1:K) {
         distant2[i, j] <- as.numeric(t(X[i, ] - centroids[j, ]) %*% (X[i, ] - centroids[j, ]))</pre>
       min_k <- which.min(distant2[i, ])</pre>
       clustering_vector[i] <- min_k</pre>
    for (k in 1:K) {
       index <- clustering_vector == k</pre>
       centroids_new[k, ] <- apply(X[index, ], 2, mean)</pre>
    if (sum(!centroids_new == centroids) == 0) {
       break
    centroids <- centroids_new</pre>
    1 <- 1 + 1
  }
  cluster_means <- centroids_new</pre>
  for (k in 1:K) {
    cluster_sizes[k] <- sum(clustering_vector == k)</pre>
```

```
for (k in 1:K) {
  index <- clustering_vector == k</pre>
  wss_cluster[k] <- sum(apply(sweep(X[index, ], 2, centroids[k, ], "-"), 1, function(x) sum(x^2)))
 }
 TSS <- sum(apply(sweep(X, 2, apply(X, 2, mean), "-"), 1, function(x) sum(x^2)))
 BSS <- TSS - sum(wss_cluster)</pre>
 bss over tss <- BSS / TSS
 list(cluster_sizes = cluster_sizes,
     cluster_means = cluster_means,
     clustering_vector = clustering_vector,
     wss_cluster = wss_cluster,
     bss_over_tss = bss_over_tss)
}
set.seed(1991)
my_kmeans(iris[, 1:4], k = 3)
## $cluster_sizes
## [1] 32 96 22
##
## $cluster means
     Sepal.Length Sepal.Width Petal.Length Petal.Width
## [1,]
        5.193750
                 3.631250
                           1.475000
                                    0.271875
## [2,]
        6.314583
                 2.895833
                           4.973958
                                    1.703125
## [3,]
        4.731818
                 2.927273
                           1.772727
                                    0.350000
##
## $clustering_vector
  ## [141] 2 2 2 2 2 2 2 2 2 2 2
##
## $wss cluster
     6.032188 118.651875 18.070000
## [1]
## $bss over tss
## [1] 0.7904898
set.seed(1991)
kmeans(iris[, 1:4], centers = 3)
## K-means clustering with 3 clusters of sizes 33, 96, 21
##
## Cluster means:
   Sepal.Length Sepal.Width Petal.Length Petal.Width
## 1
      5.175758
               3.624242
                         1.472727
                                 0.2727273
## 2
               2.895833
      6.314583
                         4.973958
                                 1.7031250
## 3
      4.738095
               2.904762
                         1.790476
                                 0.3523810
##
## Clustering vector:
```

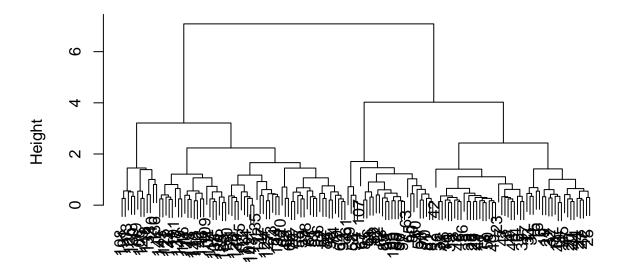
```
## [141] 2 2 2 2 2 2 2 2 2 2 2
##
## Within cluster sum of squares by cluster:
     6.432121 118.651875 17.669524
  (between_SS / total_SS = 79.0 %)
##
##
## Available components:
##
## [1] "cluster"
              "centers"
                        "totss"
                                  "withinss"
## [5] "tot.withinss" "betweenss"
                        "size"
                                  "iter"
## [9] "ifault"
```

Hierarchical clustering

```
d <- dist(iris[, 1:4], method = "euclidean")
hc.complete <- hclust(d, method = "complete")
hc.average <- hclust(d, method = "average")
hc.single <- hclust(d, method = "single")

plot(hc.complete)</pre>
```

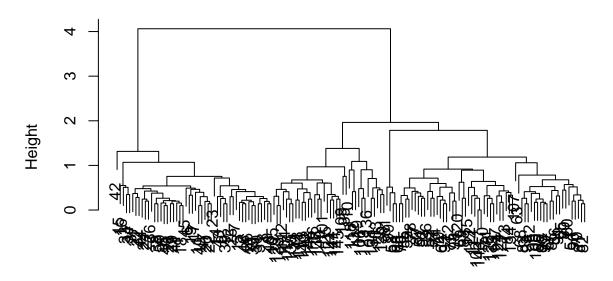
Cluster Dendrogram



d hclust (*, "complete")

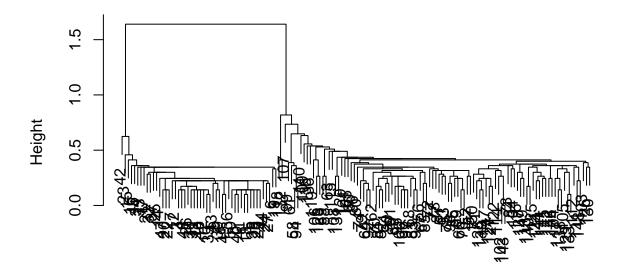
plot(hc.average)

Cluster Dendrogram



d hclust (*, "average")

Cluster Dendrogram



d hclust (*, "single")