Clustering - Agglomerative & K-Means

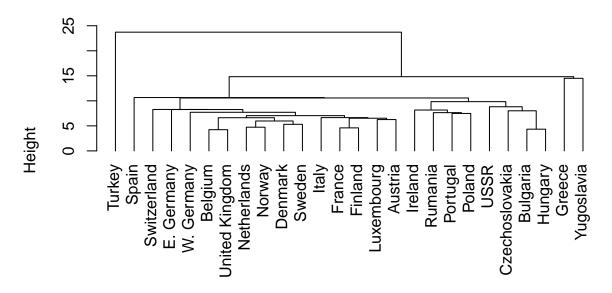
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#Agglomerative clustering of Europen Job Data (single link, complete link, and group average clustering)

```
data <- read.csv("europe", sep = "")

diss <- dist(data, method = "euclidean")
labels<-as.character(data[,1])
sing<-hclust(diss, method = "single")
plot(sing, hang = -1, main = "Single")</pre>
```

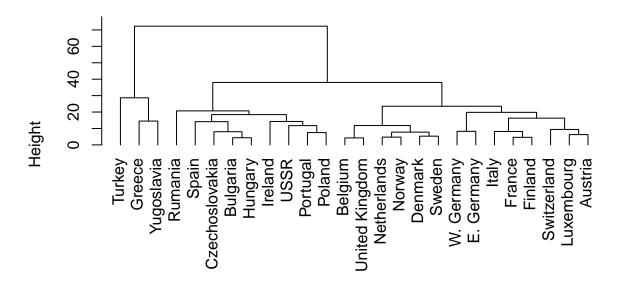
Single



diss hclust (*, "single")

```
comp<-hclust(diss, method = "complete")
plot(comp, hang = -1, main = "Complete")</pre>
```

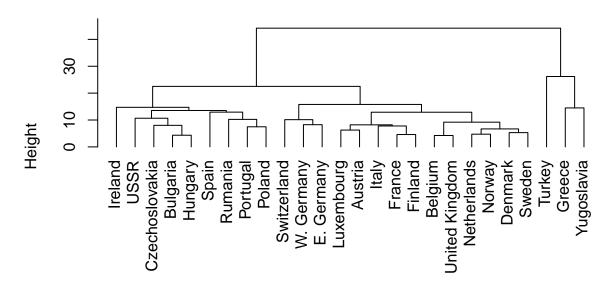
Complete



diss hclust (*, "complete")

```
avg<-hclust(diss, method = "average")
plot(avg, hang = -1, main = "Average")</pre>
```

Average



diss hclust (*, "average")

##Single dendrogram: ##Single dendrogram provides few important information about the clusters in this data set. ##- It is clear that Turkey in terms of occupation was quite different from other countries and may form a separate cluster itself. From the data it is clear that people mostly were occupied in agriculture. ##Similarly Greece and Yugoslavia appear to be clustering together and looks distant from other countries. ##It exposes 3 more major clusters in terms of occupation similarity - West European countries, East European countries and mid European countries. ##So single method looks to form approximately 5 major clusters.

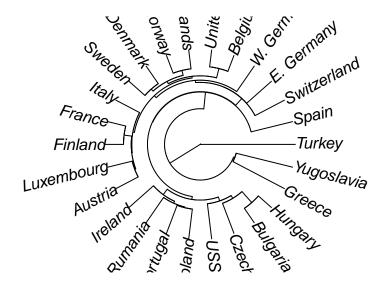
##Complete dendrogram:

##This exposes approximately 4 major clusters. In this looks like cluster Turkey got merged with Greece/Yugoslavia. ##This method does a better job separating countries(sub clusters) among other clusters seen in single dendrogram. ##This exposes some differences(clusters) within other regions.

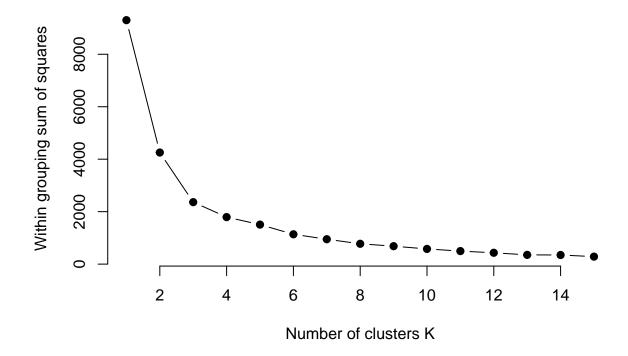
##Average dendrogram:

##This looks to form clusters which are not extremes like single or complete. Clearly it provides cluster separation with granularity more than single but little bit conservative as compared to complete dendrogram.

```
library(ape)
plot(as.phylo(sing), type='fan')
```



#k-means clustering of Europen Job Data



It is clear from the elbow diagram that a good choice of number of cluster for this data set could k=6 or k=7 or k=8 as it seems to represent sufficient/efficient encoding of the data set.

```
kmdata<-kmeans(data, 7, nstart = 10, iter.max = 1000 ,algorithm="MacQueen")
## K-means clustering with 7 clusters of sizes 6, 1, 7, 4, 1, 3, 4
##
## Cluster means:
##
                                         PS
                                                 Con
                                                            SI
                     Min
                              Man
                                                                    Fin
           Agr
## 1 10.250000 1.1833333 31.61667 0.8833333 8.983333 17.016667 4.566667
## 2 66.800000 0.7000000 7.90000 0.1000000 2.800000
                                                     5.200000 1.100000
     7.085714 0.5428571 25.18571 0.9714286 8.071429 16.371429 5.914286
## 4 26.250000 1.1500000 24.85000 0.8750000 8.950000 11.825000 3.725000
     4.200000 2.9000000 41.20000 1.3000000 7.600000 11.200000 1.200000
## 6 41.600000 1.4000000 21.50000 0.7666667 7.233333
                                                      7.933333 5.000000
  7 21.375000 2.3250000 30.80000 1.0750000 8.500000 8.175000 0.750000
##
           SPS
                     TC
## 1 19.400000 6.066667
## 2 11.900000 3.200000
## 3 28.557143 7.328571
## 4 16.350000 6.050000
## 5 22.100000 8.400000
## 6 9.333333 5.233333
```

```
## 7 19.225000 7.750000
##
## Clustering vector:
               Denmark France W. Germany Ireland
##
       Belgium
##
        Italy Luxembourg Netherlands United Kingdom
##
                                                      Austria
##
                            3 3
##
       Finland
                   Greece
                               Norway
                                         Portugal
                                                       Spain
##
                        6
                                3
       Sweden
                Switzerland
##
                               Turkey
                                          Bulgaria Czechoslovakia
##
                               2
##
     E. Germany
                               Poland
                                         Rumania
                                                        USSR
                   Hungary
##
                                  4
                                              6
                                                          7
##
     Yugoslavia
##
##
## Within cluster sum of squares by cluster:
## (between_SS / total_SS = 87.9 %)
## Available components:
## [1] "cluster" "centers"
                            "totss"
                                        "withinss"
## [5] "tot.withinss" "betweenss"
                            "size"
                                        "iter"
## [9] "ifault"
```