

Cluster Analysis

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```
health <- read.csv(file = "C:/Users/LILIAN/Desktop/Cluster Analysis/health outcome.csv")
```

```
sum(is.na(health))
```

```
## [1] 321
```

DATA WRANGLING AND EXPLORATION ANALYSIS

```
#DROP INSIGNIFICANT COLUMNS OF THE DATASET
```

```
health <- subset(health, select = -c(Infants.exclusively.breastfed...ages.0D5.months..2008D2013,Deaths
```

```
#df <- df[-c(209,210),]
```

```
head(health,3)
```

```
##      Country  Infants_DTP  Infants_Measles  Infant_Mortality  UnderFive_Mortality
## 1      Norway            1                7                2.3                2.8
## 2 Netherlands            1                4                3.3                4.0
## 3      Sweden            1                3                2.4                3.0
##  Female_Adult_mortality  Male_Adult_Mortality  Tuberculosis_death
## 1                    47                    73                0.1
## 2                    54                    69                0.2
## 3                    43                    69                0.1
##  Life_expectancy  Physicians  Public_health
## 1             24.0         37.4            9.6
## 2             23.5         31.5           12.9
## 3             24.1         32.7            9.7
```

```
nrow(health)
```

```
## [1] 208
```

```
ncol(health)
```

```
## [1] 11
```

```
# Add row names
rownames(health) = c(health$Country)
head(health)
```

```
##           Country Infants_DTP Infants_Measles Infant_Mortality
## Norway         Norway         1             7             2.3
## Netherlands    Netherlands     1             4             3.3
## Sweden          Sweden          1             3             2.4
## Korea_Republic Korea_Republic  1             1             3.2
## Luxembourg      Luxembourg      1             5             1.6
## Japan           Japan           1             5             2.1
##           UnderFive_Mortality Female_Adult_mortality Male_Adult_Mortality
## Norway         2.8             47             73
## Netherlands    4.0             54             69
## Sweden          3.0             43             69
## Korea_Republic 3.7             38             93
## Luxembourg      2.0             50             79
## Japan           2.9             42             81
##           Tuberculosis_death Life_expectancy Physicians Public_health
## Norway         0.1             24.0          37.4          9.6
## Netherlands    0.2             23.5          31.5         12.9
## Sweden          0.1             24.1          32.7          9.7
## Korea_Republic 5.4             24.0          21.4          7.2
## Luxembourg      0.4             23.4          28.2          7.1
## Japan           1.7             26.1          23.0         10.3
```

```
newhealth <- health[,-1]
head(newhealth,3)
```

```
##           Infants_DTP Infants_Measles Infant_Mortality UnderFive_Mortality
## Norway         1             7             2.3             2.8
## Netherlands    1             4             3.3             4.0
## Sweden          1             3             2.4             3.0
##           Female_Adult_mortality Male_Adult_Mortality Tuberculosis_death
## Norway         47             73             0.1
## Netherlands    54             69             0.2
## Sweden          43             69             0.1
##           Life_expectancy Physicians Public_health
## Norway         24.0          37.4          9.6
## Netherlands    23.5          31.5         12.9
## Sweden          24.1          32.7          9.7
```

```
health <- newhealth
```

```
health <- na.omit(health)
head(health,3)
```

```
##           Infants_DTP Infants_Measles Infant_Mortality UnderFive_Mortality
## Norway         1             7             2.3             2.8
## Netherlands    1             4             3.3             4.0
## Sweden          1             3             2.4             3.0
##           Female_Adult_mortality Male_Adult_Mortality Tuberculosis_death
```

```
## Norway          47          73          0.1
## Netherlands     54          69          0.2
## Sweden          43          69          0.1
##               Life_expectancy Physicians Public_health
## Norway          24.0          37.4          9.6
## Netherlands     23.5          31.5         12.9
## Sweden          24.1          32.7          9.7
```

```
glimpse(health)
```

```
## Rows: 192
## Columns: 10
## $ Infants_DTP      <int> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, ~
## $ Infants_Measles  <int> 7, 4, 3, 1, 5, 5, 8, 11, 3, 10, 1, 1, 14, 1, 2, ~
## $ Infant_Mortality <dbl> 2.3, 3.3, 2.4, 3.2, 1.6, 2.1, 3.5, 3.5, 2.1, 3.~
## $ UnderFive_Mortality <dbl> 2.8, 4.0, 3.0, 3.7, 2.0, 2.9, 4.4, 4.2, 2.6, 3.~
## $ Female_Adult_mortality <int> 47, 54, 43, 38, 50, 42, 57, 52, 51, 38, 57, 41, ~
## $ Male_Adult_Mortality <int> 73, 69, 69, 93, 79, 81, 98, 109, 114, 69, 127, ~
## $ Tuberculosis_death <dbl> 0.1, 0.2, 0.1, 5.4, 0.4, 1.7, 0.4, 0.5, 0.3, 0.~
## $ Life_expectancy  <dbl> 24.0, 23.5, 24.1, 24.0, 23.4, 26.1, 23.6, 25.1, ~
## $ Physicians       <dbl> 37.4, 31.5, 32.7, 21.4, 28.2, 23.0, 29.9, 31.8, ~
## $ Public_health    <dbl> 9.6, 12.9, 9.7, 7.2, 7.1, 10.3, 11.2, 11.7, 9.4~
```

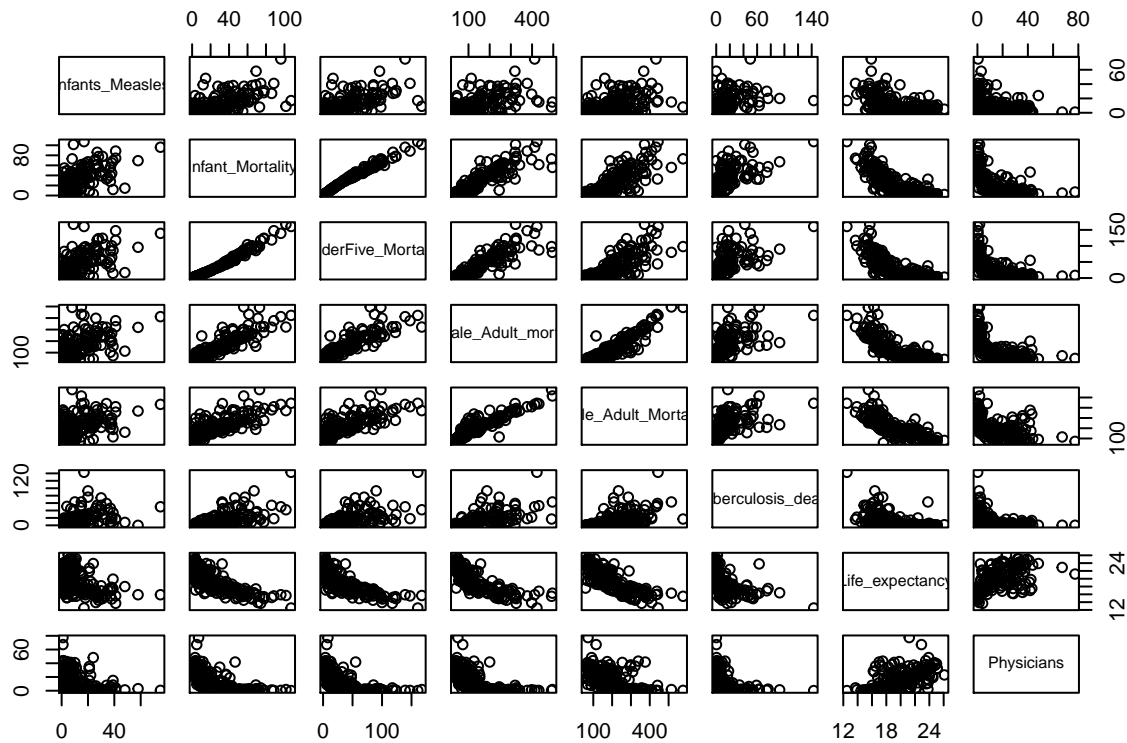
```
summary(health)
```

```
##   Infants_DTP      Infants_Measles Infant_Mortality UnderFive_Mortality
## Min.   : 1.000    Min.   : 1.00    Min.   : 1.600    Min.   : 2.00
## 1st Qu.: 1.000    1st Qu.: 3.00    1st Qu.: 7.175    1st Qu.: 8.35
## Median : 3.000    Median : 7.50    Median : 16.050   Median : 18.85
## Mean   : 6.359    Mean   :11.68    Mean   : 25.921   Mean   : 35.00
## 3rd Qu.: 8.000    3rd Qu.:17.00    3rd Qu.: 39.650   3rd Qu.: 51.85
## Max.   :65.000    Max.   :75.00    Max.   :107.200   Max.   :167.40
## Female_Adult_mortality Male_Adult_Mortality Tuberculosis_death Life_expectancy
## Min.   : 36.0      Min.   : 59.0      Min.   : 0.00     Min.   :12.50
## 1st Qu.: 69.0      1st Qu.:117.8     1st Qu.: 1.00     1st Qu.:17.00
## Median :103.0      Median :187.5     Median : 4.45     Median :19.05
## Mean   :140.7      Mean   :203.2     Mean   : 12.74     Mean   :19.38
## 3rd Qu.:196.5      3rd Qu.:262.2     3rd Qu.: 17.00     3rd Qu.:21.50
## Max.   :496.0      Max.   :577.0     Max.   :143.00     Max.   :26.10
##   Physicians      Public_health
## Min.   : 0.10     Min.   : 1.300
## 1st Qu.: 2.30     1st Qu.: 4.675
## Median :11.70     Median : 6.400
## Mean   :15.43     Mean   : 6.669
## 3rd Qu.:26.93     3rd Qu.: 8.525
## Max.   :77.40     Max.   :17.100
```

```
uplicated(health)%>%
head(n = 20)
```

```
## [1] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
## [13] FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
```

```
pairs(health[2:9])
```



CLUSTER ANALYSIS

```
library(cluster)
```

```
## Warning: package 'cluster' was built under R version 4.0.5
```

```
library(factoextra)
```

```
## Warning: package 'factoextra' was built under R version 4.0.5
```

```
## Loading required package: ggplot2
```

```
## Warning: package 'ggplot2' was built under R version 4.0.5
```

```
## Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
```

```
health_scale <- scale(health)
head(health_scale)
```

```
##           Infants_DTP Infants_Measles Infant_Mortality
## Norway      -0.6059315      -0.3872758      -1.0108556
```

```
## Netherlands      -0.6059315      -0.6354081      -0.9680605
## Sweden            -0.6059315      -0.7181188      -1.0065761
## Korea_Republic    -0.6059315      -0.8835403      -0.9723400
## Luxembourg        -0.6059315      -0.5526973      -1.0408121
## Japan             -0.6059315      -0.5526973      -1.0194146
##                   UnderFive_Mortality Female_Adult_mortality Male_Adult_Mortality
## Norway              -0.9123059              -0.9576841              -1.297255
## Netherlands         -0.8783119              -0.8861452              -1.337107
## Sweden              -0.9066402              -0.9985635              -1.337107
## Korea_Republic      -0.8868104              -1.0496627              -1.097997
## Luxembourg          -0.9349686              -0.9270246              -1.237478
## Japan               -0.9094731              -1.0087833              -1.217552
##                   Tuberculosis_death Life_expectancy Physicians Public_health
## Norway              -0.6541151              1.575400  1.5361327  1.1181616
## Netherlands         -0.6489387              1.405048  1.1235741  2.3769875
## Sweden              -0.6541151              1.609471  1.2074843  1.1563079
## Korea_Republic      -0.3797653              1.575400  0.4173296  0.2026519
## Luxembourg          -0.6385859              1.370978  0.8928209  0.1645057
## Japan               -0.5712925              2.290878  0.5292099  1.3851853
```

```
health_euc <- dist(health_scale, method = "euclidean")
```

```
round(as.matrix(health_euc)[1:3, 1:3], 1)
```

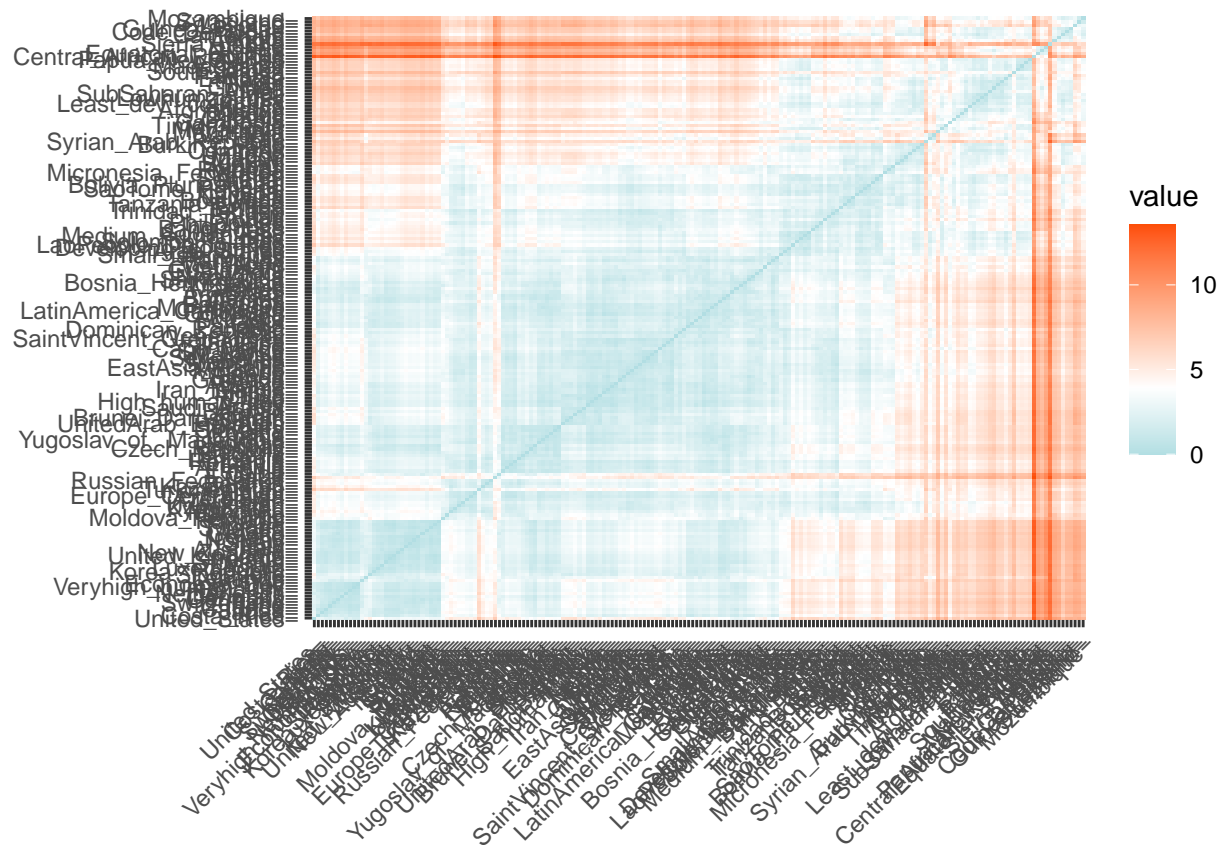
```
##           Norway Netherlands Sweden
## Norway      0.0           1.4     0.5
## Netherlands 1.4           0.0     1.2
## Sweden      0.5           1.2     0.0
```

```
health_cor <- get_dist(health_scale, method = "pearson")
round(as.matrix(health_cor)[1:3, 1:3], 1)
```

```
##           Norway Netherlands Sweden
## Norway      0.0           0.1     0
## Netherlands 0.1           0.0     0
## Sweden      0.0           0.0     0
```

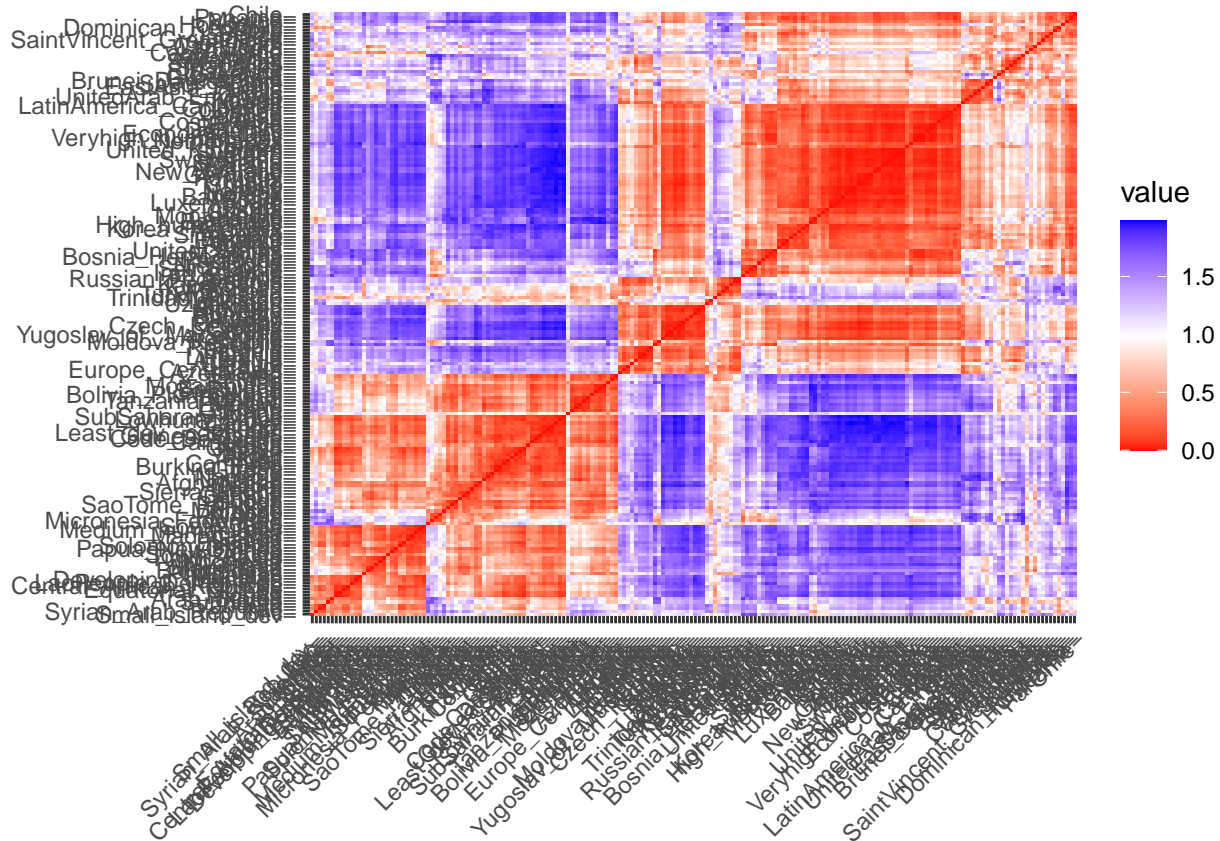
```
#to visualize the distance matrices
#library(factoextra)
```

```
#visualize the euclidean matrix
fviz_dist(health_euc, gradient = list(low = "#00AFBB", mid = "white", high = "#FC4E07"))
```



The red color indicate high similarity while the blue color indicates low similarity. the color level is proportionate to the value of similarity between observations where pure red represents 0 and pure blue represents 1

```
fviz_dist(health_cor)
```



K-MEANS ALGORITHM

```
#Select randomly k objects from the data set as the initial cluster centers or means
health_Kmeans2 <- kmeans(health,centers = 2, nstart = 25)
str(health_Kmeans2)
```

```
## List of 9
## $ cluster      : Named int [1:192] 1 1 1 1 1 1 1 1 1 1 ...
##   .. attr(*, "names")= chr [1:192] "Norway" "Netherlands" "Sweden" "Korea_Republic" ...
## $ centers      : num [1:2, 1:10] 4.29 11.51 8.08 20.65 14.55 ...
##   .. attr(*, "dimnames")=List of 2
##     ..$ : chr [1:2] "1" "2"
##     ..$ : chr [1:10] "Infants_DTP" "Infants_Measles" "Infant_Mortality" "UnderFive_Mortality" ...
## $ totss       : num 4251418
## $ withinss    : num [1:2] 756122 782988
## $ tot.withinss: num 1539110
## $ betweenss   : num 2712308
## $ size        : int [1:2] 137 55
## $ iter        : int 1
## $ ifault      : int 0
## - attr(*, "class")= chr "kmeans"
```

THE AVAILABLE COMPONENTS LISTED AS THE OUTPUT OF KMEANS ARE AS FOLLOWS:

cluster: A vector of integers (from 1:k) indicating the cluster to which each point is allocated.

centers: A matrix of cluster centers.

totss: The total sum of squares.

withinss: Vector of within-cluster sum of squares, one component per cluster.

tot.withinss: Total within-cluster sum of squares, i.e. $\text{sum}(\text{withinss})$.

betweenss: The between-cluster sum of squares, i.e. $\text{totss} - \text{tot.withinss}$.

size: The number of points in each cluster.

health_Kmeans2

```
## K-means clustering with 2 clusters of sizes 137, 55
##
## Cluster means:
##   Infants_DTP Infants_Measles Infant_Mortality UnderFive_Mortality
## 1    4.291971    8.080292    14.54672    17.53431
## 2   11.509091   20.654545    54.25273    78.52182
##   Female_Adult_mortality Male_Adult_Mortality Tuberculosis_death
## 1         89.0365         153.3577         6.249635
## 2        269.4182         327.3818        28.894545
##   Life_expectancy Physicians Public_health
## 1    20.65036  19.967153    6.968613
## 2    16.20182   4.134545    5.921818
##
## Clustering vector:
##           Norway           Netherlands           Sweden
##             1             1             1
##   Korea_Republic           Luxembourg           Japan
##             1             1             1
##           Belgium           France           Finland
##             1             1             1
##             Italy       Czech_Republic           Greece
##             1             1             1
##           Cyprus           Qatar           Slovakia
##             1             1             1
##           Poland           Malta           Portugal
##             1             1             1
##           Hungary           Bahrain           Kuwait
##             1             1             1
##           Belarus           Oman           Bahamas
##             1             1             1
##           Kazakhstan           Malaysia           Mauritius
##             2             1             1
##           Seychelles           Turkey           Sri_Lanka
##             1             1             1
##           Brazil           Georgia           Grenada
##             1             1             1
##           Jordan   Yugoslav_of_ Macedonia           Algeria
##             1             1             1
##           Albania           Ecuador           Saint_Lucia
##             1             1             1
##           China           Fiji           Thailand
##             1             1             1
##           Libya           Tunisia           Tonga
##             1             1             1
```


##	Maldives	Samoa	Turkmenistan
##	1	1	2
##	Uzbekistan	Nicaragua	Morocco
##	1	1	1
##	Bangladesh	SaoTome_Principe	Swaziland
##	1	1	2
##	Tanzania_United	Rwanda	Sudan
##	2	2	2
##	Gambia	Switzerland	Germany
##	2	1	1
##	Ireland	United_States	Canada
##	1	1	1
##	Singapore	United_Kingdom	Slovenia
##	1	1	1
##	Spain	Saudi_Arabia	Croatia
##	1	1	1
##	Montenegro	Uruguay	Serbia
##	1	1	1
##	Cuba	Costa_Rica	Iran_Islamic
##	1	1	1
##	Mongolia	Botswana	Indonesia
##	2	2	1
##	Philippines	Bolivia_Plurinational	Kyrgyzstan
##	1	1	1
##	Micronesia_Federated	Guyana	Tajikistan
##	1	2	1
##	Sierra_Leone	Burundi	High_human_dev
##	2	2	1
##	Veryhigh_human_dev	EastAsia_Pacific	Europe_CentralAsia
##	1	1	1
##	Denmark	Iceland	Lithuania
##	1	1	1
##	Russian_Federation	Peru	Armenia
##	2	1	1
##	Colombia	Jamaica	Egypt
##	1	1	1
##	ElSalvador	Guatemala	Bhutan
##	1	1	2
##	Angola	Lesotho	Liberia
##	2	2	2
##	Eritrea	Economic_Dev	Estonia
##	2	1	1
##	Brunei_Darussalam	Latvia	Romania
##	1	1	1
##	Bulgaria	SaintVincent_Grenadines	Belize
##	1	1	1
##	Moldova_Republic	Senegal	Malawi
##	1	2	2
##	Israel	Azerbaijan	Bosnia_Herzegovina
##	1	1	1
##	Suriname	Kiribati	Cambodia
##	1	1	1
##	Cameroon	Mauritania	LatinAmerica_Caribbean
##	2	2	1

```

##      UnitedArab_Emirates      Trinidad_Tobago      Namibia
##              1              1              2
##              Ghana              Nepal              Yemen
##              2              1              2
##              Togo              Burkina_Faso      New_Zealand
##              2              2              1
##              Austria      Argentina      Barbados
##              1              1              1
##              Panama      Paraguay      Cabo_Verde
##              1              1              1
##      Code_De_Ivoire      Mozambique      Australia
##              2              2              1
##              Chile      Guinea_Bissau      Arab_States
##              1              2              1
##              World      Dominican_Republic      Small_island_dev
##              1              1              1
##      Medium_human_dev      Developing_countries      Venezuela
##              1              1              1
##              Mexico      Ukraine      Congo
##              1              1              2
##              Myanmar      Least_dev_ countries      LaoPeople_Demo_Rep
##              2              2              1
##              Uganda      South_Asia      India
##              2              1              1
##              Honduras      Solomon_Islands      Papua_NewGuinea
##              1              1              2
##              Djibouti      Timor_Leste      Zambia
##              2              1              2
##              Afghanistan      Congo      Comoros
##              2              2              2
##              Benin      Niger      Lebanon
##              2              2              1
##              Ethiopia      Lowhuman_dev      Viet_Nam
##              2              2              1
##      SubSaharan_Africa      Iraq      Kenya
##              2              1              2
##              Mali      Gabon      Madagascar
##              2              2              2
##              Pakistan      Vanuatu      Guinea
##              1              1              2
##              South_Africa      Nigeria      Syrian_Arab_Republic
##              2              2              1
##      Equatorial_Guinea      Chad Central_African_Republic
##              2              2              2
##
## Within cluster sum of squares by cluster:
## [1] 756122.2 782987.9
## (between_SS / total_SS = 63.8 %)
##
## Available components:
##
## [1] "cluster"      "centers"      "totss"        "withinss"     "tot.withinss"
## [6] "betweenss"    "size"         "iter"         "ifault"

```

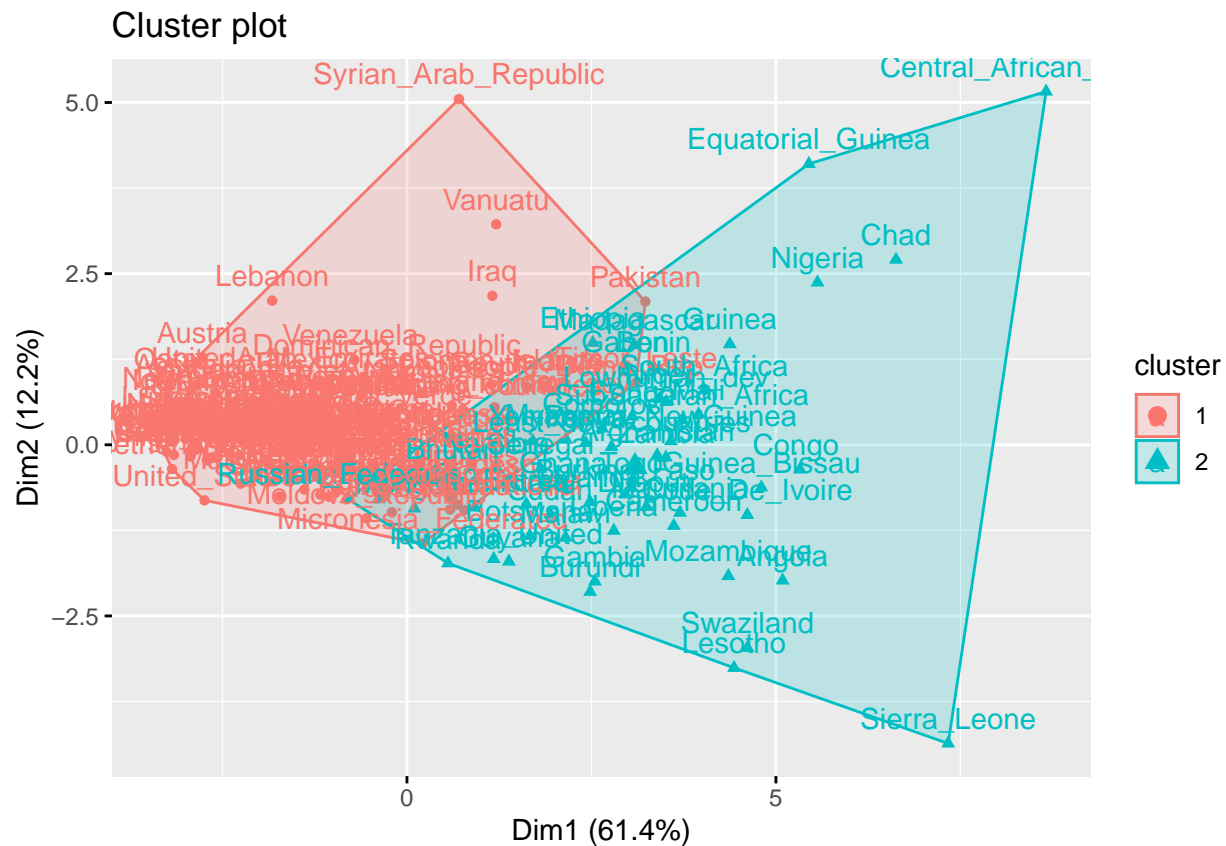
THE RESULT ABOVE SHOWS: That our groupings resulted in 2 cluster sizes of 137 and 55.

Also, the cluster centers (means) for the two groups across the ten variables (Infants_DTP, Infants_Measles, Infant_Mortality, UnderFive_Mortality, Female_Adult_mortality, Male_Adult_Mortality, Tuberculosis_death, Life_expectancy, Physicians, Public_health).

The cluster assignment for each observation (i.e. Nigeria was assigned to cluster 2, Mexico was assigned to cluster 1, etc.). #####

FOR more than two dimensions (variables) fviz_cluster will perform principal component analysis (PCA) and plot the data points according to the first two principal components that explain the majority of the variance.

```
fviz_cluster(health_Kmeans2, data = health)
```



Because the number of clusters (k) must be set before we start the algorithm, it is often advantageous to use several different values of k and examine the differences in the results. We can execute the same process for 3, 4, and 5 clusters, and the results are shown in the figure:

```
#health_Kmeans2 <- kmeans(health, centers = 2, nstart = 25)
#health_Kmeans3 <- kmeans(health, centers = 3, nstart = 25)
#health_Kmeans4 <- kmeans(health, centers = 4, nstart = 25)
#health_Kmeans5 <- kmeans(health, centers = 5, nstart = 25)

# plots to compare
#plot1 <- fviz_cluster(health_Kmeans2, geom = "point", data = health),ggtitle("k = 2")
#plot2 <- fviz_cluster(health_Kmeans3, geom = "point", data = health),ggtitle("k = 3")
#plot3 <- fviz_cluster(health_Kmeans4, geom = "point", data = health),ggtitle("k = 4")
```

```
#plot4 <- fviz_cluster(health_Kmeans5, geom = "point", data = health),ggtitle("k = 5")  
library(gridExtra)
```

```
## Warning: package 'gridExtra' was built under R version 4.0.3
```

```
##
```

```
## Attaching package: 'gridExtra'
```

```
## The following object is masked from 'package:dplyr':
```

```
##
```

```
##      combine
```

```
#grid.arrange(plot1, plot2, plot3, plot4, nrow = 2)
```

```
set.seed(123)
```

```
# function to compute total within-cluster sum of square
```

```
wss <- function(k) {  
  kmeans(health, k, nstart = 10 )$tot.withinss  
}
```

```
# Compute and plot wss for k = 1 to k = 15
```

```
k.values <- 1:15
```

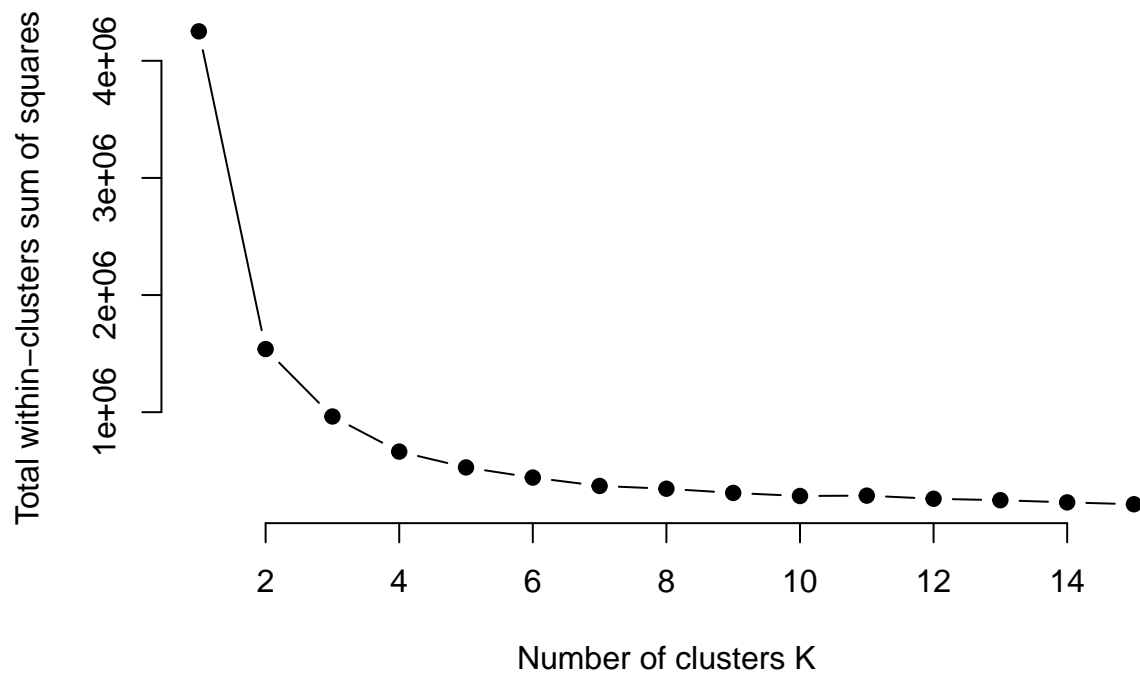
```
library(purrr)
```

```
## Warning: package 'purrr' was built under R version 4.0.5
```

```
# extract wss for 2-15 clusters
```

```
wss_values <- map_dbl(k.values, wss)
```

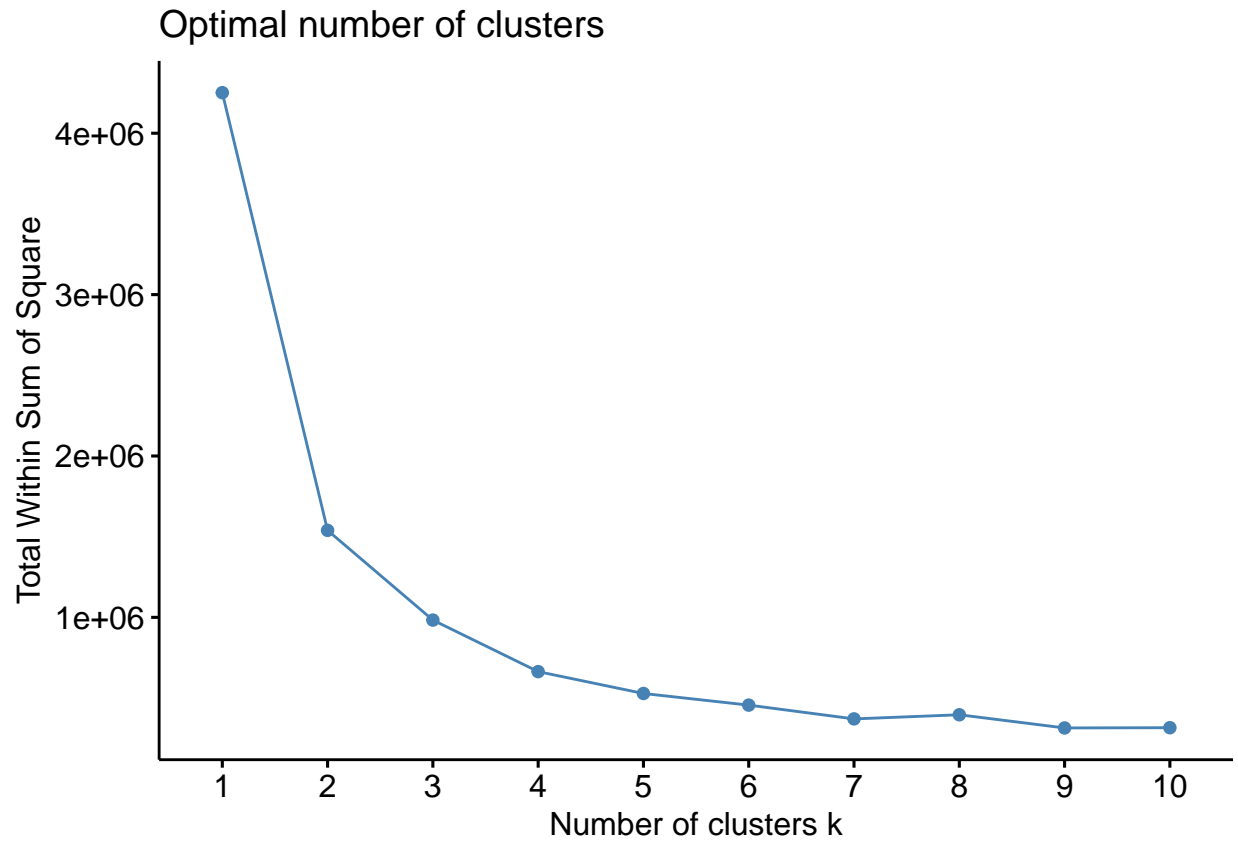
```
plot(k.values, wss_values,  
     type="b", pch = 19, frame = FALSE,  
     xlab="Number of clusters K",  
     ylab="Total within-clusters sum of squares")
```



The results suggest that 4 is the optimal number of clusters as it appears to be the bend in the knee (or elbow).

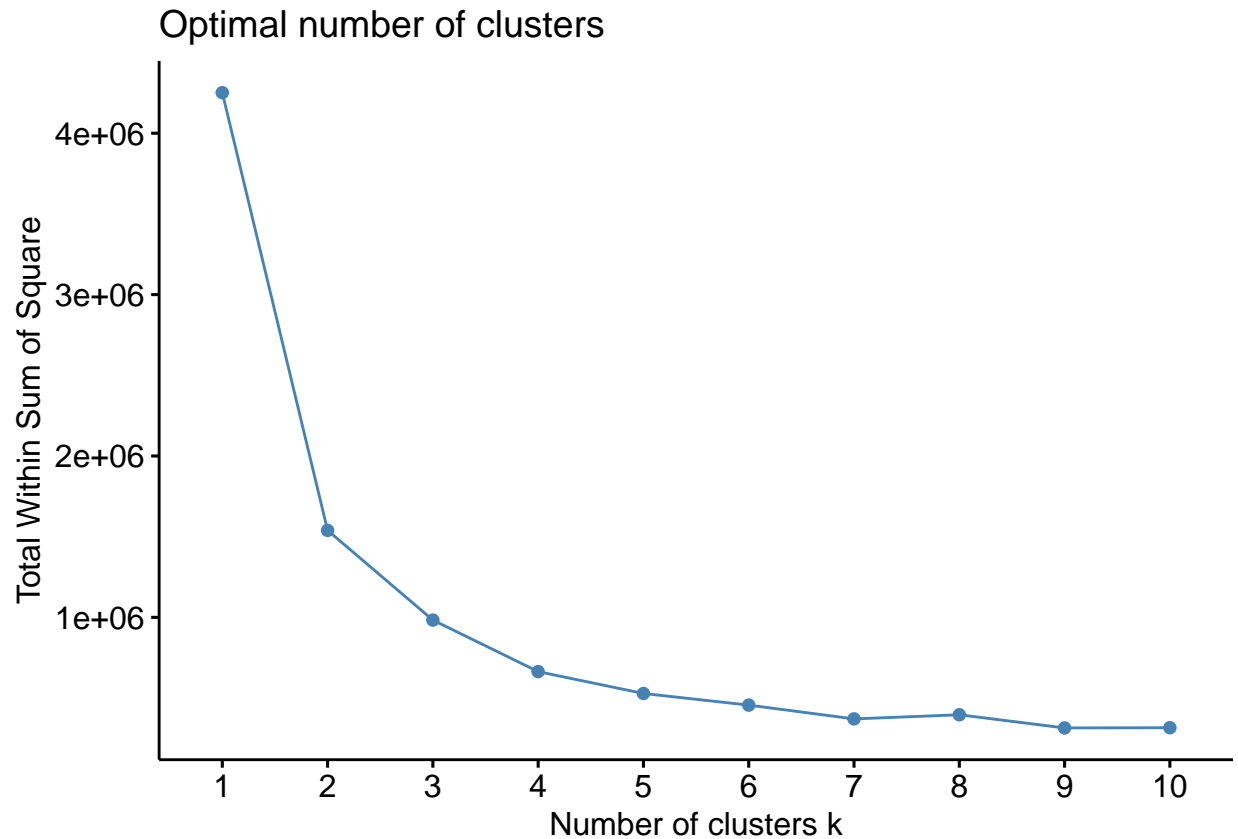
K-MEANS CLUSTERING

```
fviz_nbclust(health, kmeans, method = "wss")
```



The bent is generally considered as an indicator of the optimal number of clusters. the plot shows variances decreases as the k number increases. a bent on the elbow can be seen as $k = 4$, meaning additional clusters beyond 4 has little values. Let's classify our observations into 4 clusters.

```
fviz_nbclust(health, kmeans, method = "wss")
```



```
geom_vline(xintercept = 4, linetype = 2) #the line did not appear bcos of the #sign
```

```
## mapping: xintercept = ~xintercept
## geom_vline: na.rm = FALSE
## stat_identity: na.rm = FALSE
## position_identity
```

```
labs(subtitle = "Elbow method")
```

```
## $subtitle
## [1] "Elbow method"
##
## attr(,"class")
## [1] "labels"
```

To compute k means clustering on a data matrix, We set a seed for random number generator to randomly selected centroids for k means algorithms.

```
# Compute k-means clustering with k = 4
set.seed(123)
final_result <- kmeans(health, 4, nstart = 25)
print(final_result)
```

```
## K-means clustering with 4 clusters of sizes 38, 69, 68, 17
```

```

##
## Cluster means:
##   Infants_DTP Infants_Measles Infant_Mortality UnderFive_Mortality
## 1      8.763158      17.947368      46.663158      66.010526
## 2      5.246377       9.898551      21.768116      26.430435
## 3      3.323529       6.235294       7.219118       8.507353
## 4     17.647059     26.705882     71.217647     106.488235
##   Female_Adult_mortality Male_Adult_Mortality Tuberculosis_death
## 1           226.47368           289.5789           24.61053
## 2           117.33333           200.1739           10.65942
## 3            60.32353            105.8529            1.77500
## 4           365.41176           411.8824           38.47059
##   Life_expectancy Physicians Public_health
## 1           16.48158      5.286842      5.710526
## 2           19.20580     14.291304      6.097101
## 3           22.11618     25.726471      7.852941
## 4           15.57647      1.558824      6.394118
##
## Clustering vector:
##           Norway           Netherlands           Sweden
##           3             3             3
##   Korea _Republic           Luxembourg           Japan
##           3             3             3
##           Belgium           France           Finland
##           3             3             3
##           Italy           Czech_Republic           Greece
##           3             3             3
##           Cyprus           Qatar           Slovakia
##           3             3             3
##           Poland           Malta           Portugal
##           2             3             3
##           Hungary           Bahrain           Kuwait
##           2             3             3
##           Belarus           Oman           Bahamas
##           2             3             3
##           Kazakhstan           Malaysia           Mauritius
##           1             2             2
##           Seychelles           Turkey           Sri_Lanka
##           2             3             2
##           Brazil           Georgia           Grenada
##           2             2             2
##           Jordan   Yugoslav_of_ Macedonia           Algeria
##           3             3             2
##           Albania           Ecuador           Saint_Lucia
##           3             2             2
##           China           Fiji           Thailand
##           3             2             2
##           Libya           Tunisia           Tonga
##           3             3             2
##           Maldives           Samoa           Turkmenistan
##           3             2             1
##           Uzbekistan           Nicaragua           Morocco
##           2             2             2
##           Bangladesh           SaoTome_Principe           Swaziland

```

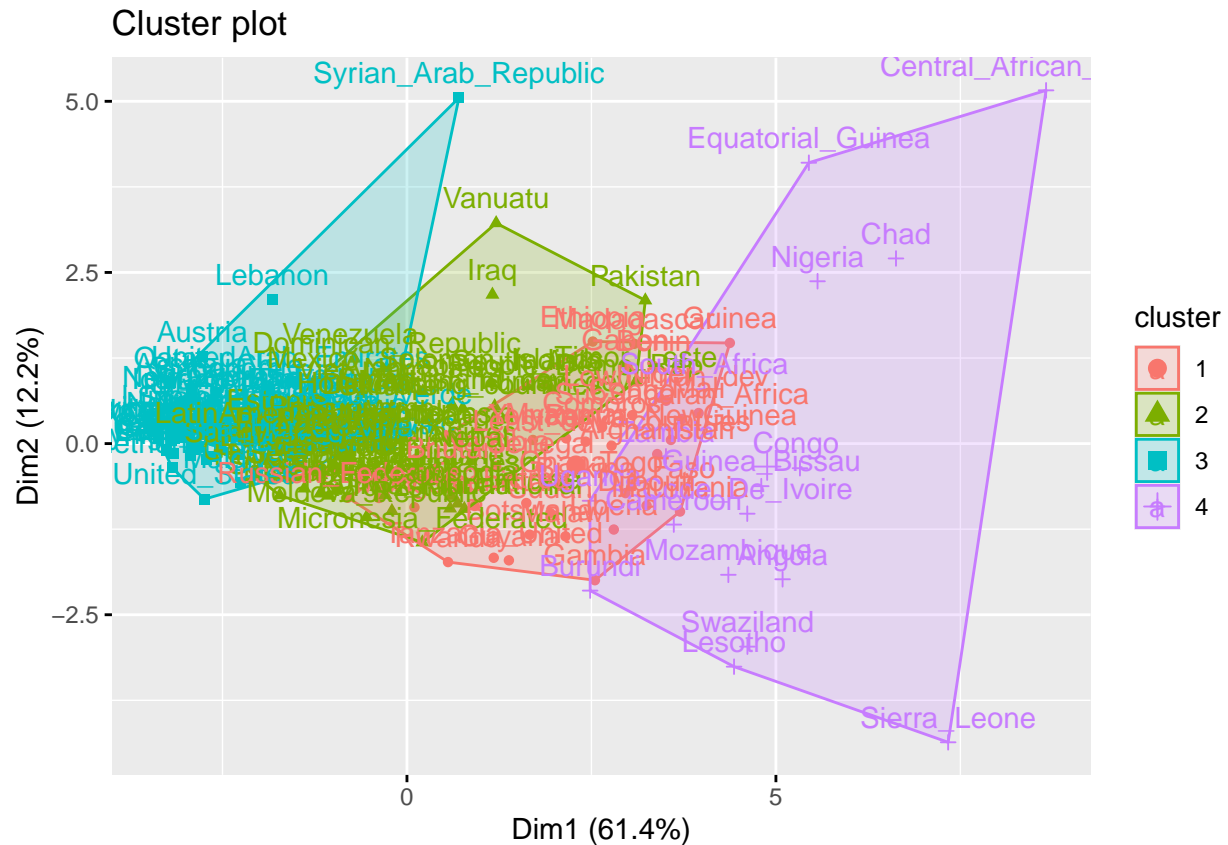

##	2	2	4
##	Tanzania_United	Rwanda	Sudan
##	1	1	1
##	Gambia	Switzerland	Germany
##	1	3	3
##	Ireland	United_States	Canada
##	3	3	3
##	Singapore	United_Kingdom	Slovenia
##	3	3	3
##	Spain	Saudi_Arabia	Croatia
##	3	3	3
##	Montenegro	Uruguay	Serbia
##	3	3	2
##	Cuba	Costa_Rica	Iran_Islamic
##	3	3	3
##	Mongolia	Botswana	Indonesia
##	1	1	2
##	Philippines	Bolivia_Plurinational	Kyrgyzstan
##	2	2	2
##	Micronesia_Federated	Guyana	Tajikistan
##	2	1	2
##	Sierra_Leone	Burundi	High_human_dev
##	4	4	3
##	Veryhigh_human_dev	EastAsia_Pacific	Europe_CentralAsia
##	3	3	2
##	Denmark	Iceland	Lithuania
##	3	3	2
##	Russian_Federation	Peru	Armenia
##	1	3	2
##	Colombia	Jamaica	Egypt
##	3	2	2
##	ElSalvador	Guatemala	Bhutan
##	2	2	1
##	Angola	Lesotho	Liberia
##	4	4	1
##	Eritrea	Economic_Dev	Estonia
##	1	3	2
##	Brunei_Darussalam	Latvia	Romania
##	3	2	2
##	Bulgaria	SaintVincent_Grenadines	Belize
##	2	2	3
##	Moldova_Republic	Senegal	Malawi
##	2	1	1
##	Israel	Azerbaijan	Bosnia_Herzegovina
##	3	2	3
##	Suriname	Kiribati	Cambodia
##	2	2	2
##	Cameroon	Mauritania	LatinAmerica_Caribbean
##	4	1	2
##	UnitedArab_Emirates	Trinidad_Tobago	Namibia
##	3	2	1
##	Ghana	Nepal	Yemen
##	1	2	1
##	Togo	Burkina_Faso	New_Zealand

```

##          1          1          3
##      Austria      Argentina      Barbados
##          3          3          3
##      Panama      Paraguay      Cabo_Verde
##          3          2          3
##      Code_De_Ivoire      Mozambique      Australia
##          4          4          3
##      Chile      Guinea_Bissau      Arab_States
##          3          4          2
##      World      Dominican_Republic      Small_island_dev
##          2          2          2
##      Medium_human_dev      Developing_countries      Venezuela
##          2          2          2
##      Mexico      Ukraine      Congo
##          2          2          1
##      Myanmar      Least_dev_ countries      LaoPeople_Demo_Rep
##          1          1          2
##      Uganda      South_Asia      India
##          4          2          2
##      Honduras      Solomon_Islands      Papua_NewGuinea
##          2          2          1
##      Djibouti      Timor_Leste      Zambia
##          1          2          4
##      Afghanistan      Congo      Comoros
##          1          4          1
##      Benin      Niger      Lebanon
##          1          1          3
##      Ethiopia      Lowhuman_dev      Viet_Nam
##          1          1          2
##      SubSaharan_Africa      Iraq      Kenya
##          1          2          1
##      Mali      Gabon      Madagascar
##          1          1          1
##      Pakistan      Vanuatu      Guinea
##          2          2          1
##      South_Africa      Nigeria      Syrian_Arab_Republic
##          4          4          3
##      Equatorial_Guinea      Chad Central_African_Republic
##          4          4          4
##
## Within cluster sum of squares by cluster:
## [1] 171201.28 220402.40 93307.34 178776.31
## (between_SS / total_SS = 84.4 %)
##
## Available components:
##
## [1] "cluster"      "centers"      "totss"        "withinss"     "tot.withinss"
## [6] "betweenss"   "size"         "iter"         "ifault"

```

```
fviz_cluster(final_result, data = health)
```



```
health %>%
  mutate(Cluster = final_result$cluster) %>%
  group_by(Cluster) %>%
  summarise_all("mean")
```

```
## # A tibble: 4 x 11
##   Cluster Infants_DTP Infants_Measles Infant_Mortality UnderFive_Mortality
##   <int>      <dbl>      <dbl>          <dbl>          <dbl>
## 1     1         8.76        17.9           46.7           66.0
## 2     2         5.25         9.90           21.8           26.4
## 3     3         3.32         6.24            7.22            8.51
## 4     4        17.6        26.7           71.2          106.
## # ... with 6 more variables: Female_Adult_mortality <dbl>,
## #   Male_Adult_Mortality <dbl>, Tuberculosis_death <dbl>,
## #   Life_expectancy <dbl>, Physicians <dbl>, Public_health <dbl>
```

OR SIMPLY PUT

```
aggregate(health, by=list(cluster= final_result$cluster), mean)
```

```
##   cluster Infants_DTP Infants_Measles Infant_Mortality UnderFive_Mortality
## 1     1     8.763158    17.947368     46.663158     66.010526
## 2     2     5.246377     9.898551     21.768116     26.430435
## 3     3     3.323529     6.235294      7.219118      8.507353
```

```
## 4      4  17.647059      26.705882      71.217647      106.488235
##   Female_Adult_mortality Male_Adult_Mortality Tuberculosis_death
## 1      226.47368      289.5789      24.61053
## 2      117.33333      200.1739      10.65942
## 3      60.32353      105.8529      1.77500
## 4      365.41176      411.8824      38.47059
##   Life_expectancy Physicians Public_health
## 1      16.48158   5.286842   5.710526
## 2      19.20580  14.291304   6.097101
## 3      22.11618  25.726471   7.852941
## 4      15.57647   1.558824   6.394118
```

This computes summary statistics of data subsets

```
#combine R objects by rows and columns
```

```
point_class <- cbind(health, cluster = final_result$cluster)
head(point_class)
```

```
##           Infants_DTP Infants_Measles Infant_Mortality
## Norway           1           7           2.3
## Netherlands      1           4           3.3
## Sweden           1           3           2.4
## Korea_Republic   1           1           3.2
## Luxembourg       1           5           1.6
## Japan            1           5           2.1
##           UnderFive_Mortality Female_Adult_mortality Male_Adult_Mortality
## Norway           2.8           47           73
## Netherlands      4.0           54           69
## Sweden           3.0           43           69
## Korea_Republic   3.7           38           93
## Luxembourg       2.0           50           79
## Japan            2.9           42           81
##           Tuberculosis_death Life_expectancy Physicians Public_health
## Norway           0.1           24.0      37.4           9.6
## Netherlands      0.2           23.5      31.5          12.9
## Sweden           0.1           24.1      32.7           9.7
## Korea_Republic   5.4           24.0      21.4           7.2
## Luxembourg       0.4           23.4      28.2           7.1
## Japan            1.7           26.1      23.0          10.3
##           cluster
## Norway           3
## Netherlands      3
## Sweden           3
## Korea_Republic   3
## Luxembourg       3
## Japan            3
```

```
final_result
```

```
## K-means clustering with 4 clusters of sizes 38, 69, 68, 17
##
## Cluster means:
```

```

##   Infants_DTP Infants_Measles Infant_Mortality UnderFive_Mortality
## 1      8.763158      17.947368      46.663158      66.010526
## 2      5.246377      9.898551      21.768116      26.430435
## 3      3.323529      6.235294      7.219118      8.507353
## 4     17.647059     26.705882     71.217647     106.488235
##   Female_Adult_mortality Male_Adult_Mortality Tuberculosis_death
## 1           226.47368           289.5789           24.61053
## 2           117.33333           200.1739           10.65942
## 3            60.32353           105.8529            1.77500
## 4           365.41176           411.8824           38.47059
##   Life_expectancy Physicians Public_health
## 1           16.48158      5.286842      5.710526
## 2           19.20580     14.291304      6.097101
## 3           22.11618     25.726471      7.852941
## 4           15.57647      1.558824      6.394118
##
## Clustering vector:
##           Norway           Netherlands           Sweden
##           3             3             3
##   Korea _Republic           Luxembourg           Japan
##           3             3             3
##           Belgium           France           Finland
##           3             3             3
##           Italy           Czech_Republic           Greece
##           3             3             3
##           Cyprus           Qatar           Slovakia
##           3             3             3
##           Poland           Malta           Portugal
##           2             3             3
##           Hungary           Bahrain           Kuwait
##           2             3             3
##           Belarus           Oman           Bahamas
##           2             3             3
##           Kazakhstan           Malaysia           Mauritius
##           1             2             2
##           Seychelles           Turkey           Sri_Lanka
##           2             3             2
##           Brazil           Georgia           Grenada
##           2             2             2
##           Jordan   Yugoslav_of_ Macedonia           Algeria
##           3             3             2
##           Albania           Ecuador           Saint_Lucia
##           3             2             2
##           China           Fiji           Thailand
##           3             2             2
##           Libya           Tunisia           Tonga
##           3             3             2
##           Maldives           Samoa           Turkmenistan
##           3             2             1
##           Uzbekistan           Nicaragua           Morocco
##           2             2             2
##           Bangladesh           SaoTome_Principe           Swaziland
##           2             2             4
##           Tanzania_United           Rwanda           Sudan

```

##	1	1	1
##	Gambia	Switzerland	Germany
##	1	3	3
##	Ireland	United_States	Canada
##	3	3	3
##	Singapore	United_Kingdom	Slovenia
##	3	3	3
##	Spain	Saudi_Arabia	Croatia
##	3	3	3
##	Montenegro	Uruguay	Serbia
##	3	3	2
##	Cuba	Costa_Rica	Iran_Islamic
##	3	3	3
##	Mongolia	Botswana	Indonesia
##	1	1	2
##	Philippines	Bolivia_Plurinational	Kyrgyzstan
##	2	2	2
##	Micronesia_Federated	Guyana	Tajikistan
##	2	1	2
##	Sierra_Leone	Burundi	High_human_dev
##	4	4	3
##	Veryhigh_human_dev	EastAsia_Pacific	Europe_CentralAsia
##	3	3	2
##	Denmark	Iceland	Lithuania
##	3	3	2
##	Russian_Federation	Peru	Armenia
##	1	3	2
##	Colombia	Jamaica	Egypt
##	3	2	2
##	ElSalvador	Guatemala	Bhutan
##	2	2	1
##	Angola	Lesotho	Liberia
##	4	4	1
##	Eritrea	Economic_Dev	Estonia
##	1	3	2
##	Brunei_Darussalam	Latvia	Romania
##	3	2	2
##	Bulgaria	SaintVincent_Grenadines	Belize
##	2	2	3
##	Moldova_Republic	Senegal	Malawi
##	2	1	1
##	Israel	Azerbaijan	Bosnia_Herzegovina
##	3	2	3
##	Suriname	Kiribati	Cambodia
##	2	2	2
##	Cameroon	Mauritania	LatinAmerica_Caribbean
##	4	1	2
##	UnitedArab_Emirates	Trinidad_Tobago	Namibia
##	3	2	1
##	Ghana	Nepal	Yemen
##	1	2	1
##	Togo	Burkina_Faso	New_Zealand
##	1	1	3
##	Austria	Argentina	Barbados

```

##          3          3          3
##      Panama      Paraguay      Cabo_Verde
##          3          2          3
##      Code_De_Ivoire      Mozambique      Australia
##          4          4          3
##          Chile      Guinea_Bissau      Arab_States
##          3          4          2
##          World      Dominican_Republic      Small_island_dev
##          2          2          2
##      Medium_human_dev      Developing_countries      Venezuela
##          2          2          2
##          Mexico      Ukraine      Congo
##          2          2          1
##          Myanmar      Least_dev_ countries      LaoPeople_Demo_Rep
##          1          1          2
##          Uganda      South_Asia      India
##          4          2          2
##          Honduras      Solomon_Islands      Papua_NewGuinea
##          2          2          1
##          Djibouti      Timor_Leste      Zambia
##          1          2          4
##          Afghanistan      Congo      Comoros
##          1          4          1
##          Benin      Niger      Lebanon
##          1          1          3
##          Ethiopia      Lowhuman_dev      Viet_Nam
##          1          1          2
##      SubSaharan_Africa      Iraq      Kenya
##          1          2          1
##          Mali      Gabon      Madagascar
##          1          1          1
##          Pakistan      Vanuatu      Guinea
##          2          2          1
##          South_Africa      Nigeria      Syrian_Arab_Republic
##          4          4          3
##      Equatorial_Guinea      Chad Central_African_Republic
##          4          4          4
##
## Within cluster sum of squares by cluster:
## [1] 171201.28 220402.40 93307.34 178776.31
## (between_SS / total_SS =  84.4 %)
##
## Available components:
##
## [1] "cluster"      "centers"      "totss"        "withinss"     "tot.withinss"
## [6] "betweenss"    "size"         "iter"         "ifault"

```

```
final_result$size
```

```
## [1] 38 69 68 17
```

HIERARCHICAL CLUSTERING

```
#df_scale <- scale(df)
#head(df_scale)
```

```
require(stats)
health_dist <- dist(x = health_scale, method = "euclidean")
```

```
x <- as.matrix(health_dist)[1:6, 1:6]
x
```

```
##              Norway Netherlands      Sweden Korea _Republic Luxembourg
## Norway      0.0000000      1.362041 0.4726441      1.5689876 1.1826361
## Netherlands 1.3620413      0.000000 1.2492717      2.3395646 2.2308142
## Sweden      0.4726441      1.249272 0.0000000      1.3034367 1.0882037
## Korea _Republic 1.5689876 2.339565 1.3034367      0.0000000 0.6980569
## Luxembourg   1.1826361 2.230814 1.0882037      0.6980569 0.0000000
## Japan        1.2807704 1.472225 1.0126954      1.4448686 1.5751845
##              Japan
## Norway      1.280770
## Netherlands 1.472225
## Sweden      1.012695
## Korea _Republic 1.444869
## Luxembourg   1.575184
## Japan        0.000000
```

```
round(x, digits = 3)
```

```
##              Norway Netherlands Sweden Korea _Republic Luxembourg Japan
## Norway      0.000      1.362 0.473      1.569      1.183 1.281
## Netherlands 1.362      0.000 1.249      2.340      2.231 1.472
## Sweden      0.473      1.249 0.000      1.303      1.088 1.013
## Korea _Republic 1.569 2.340 1.303      0.000      0.698 1.445
## Luxembourg   1.183 2.231 1.088      0.698      0.000 1.575
## Japan        1.281 1.472 1.013      1.445      1.575 0.000
```

HIERARCHICAL CLUSTER ANALYSIS

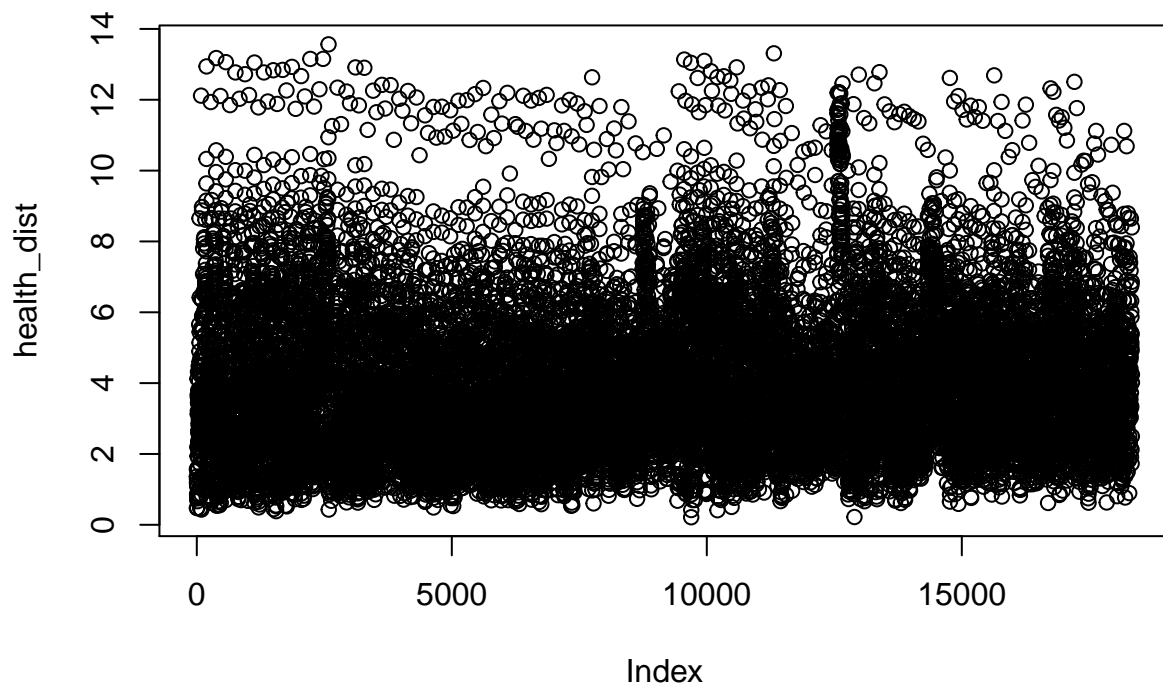
```
#LOAD STATS LIBRARY
```

```
require(stats)
```

```
#FOR HIERARCHICAL CLUSTERING
```

```
health_hc <- hclust(d = health_dist, method = "complete")
```

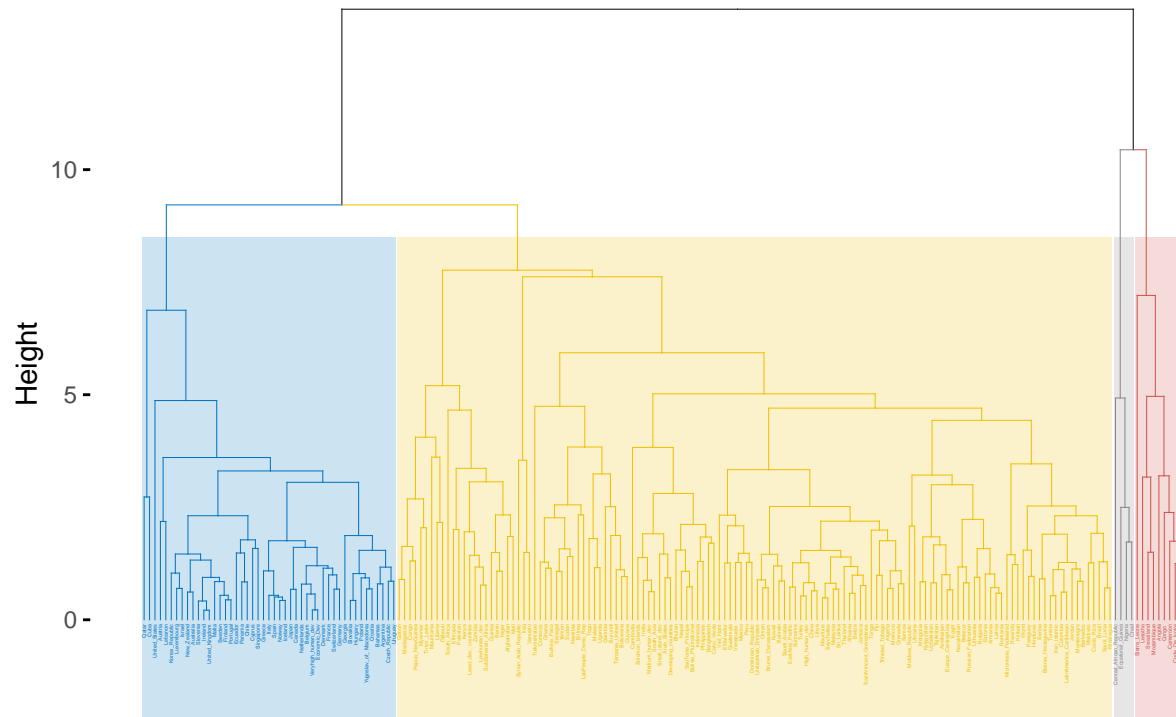
```
plot(x = health_dist)
```

```
fviz_dend(x = health_hc, cex = 0.15, lwd = 0.18, k = 4,  
  rect = TRUE,  
    k_colors = "jco",  
  rect_border = "jco",  
  rect_fill = TRUE)
```

```
## Warning: 'guides(<scale> = FALSE)' is deprecated. Please use 'guides(<scale> =  
## "none")' instead.
```

Cluster Dendrogram



```
require(grDevices)
require(scales)
```

```
## Loading required package: scales
```

```
##
```

```
## Attaching package: 'scales'
```

```
## The following object is masked from 'package:purrr':
```

```
##
```

```
##   discard
```

```
## The following object is masked from 'package:readr':
```

```
##
```

```
##   col_factor
```

```
require("ggsci")
```

```
## Loading required package: ggsci
```

```
## Warning: package 'ggsci' was built under R version 4.0.5
```

```
require(igraph)
```

```
## Loading required package: igraph
```

```
## Warning: package 'igraph' was built under R version 4.0.5
```

```
##
```

```
## Attaching package: 'igraph'
```

```
## The following objects are masked from 'package:purrr':
```

```
##
```

```
##   compose, simplify
```

```
## The following objects are masked from 'package:dplyr':
```

```
##
```

```
##   as_data_frame, groups, union
```

```
## The following objects are masked from 'package:stats':
```

```
##
```

```
##   decompose, spectrum
```

```
## The following object is masked from 'package:base':
```

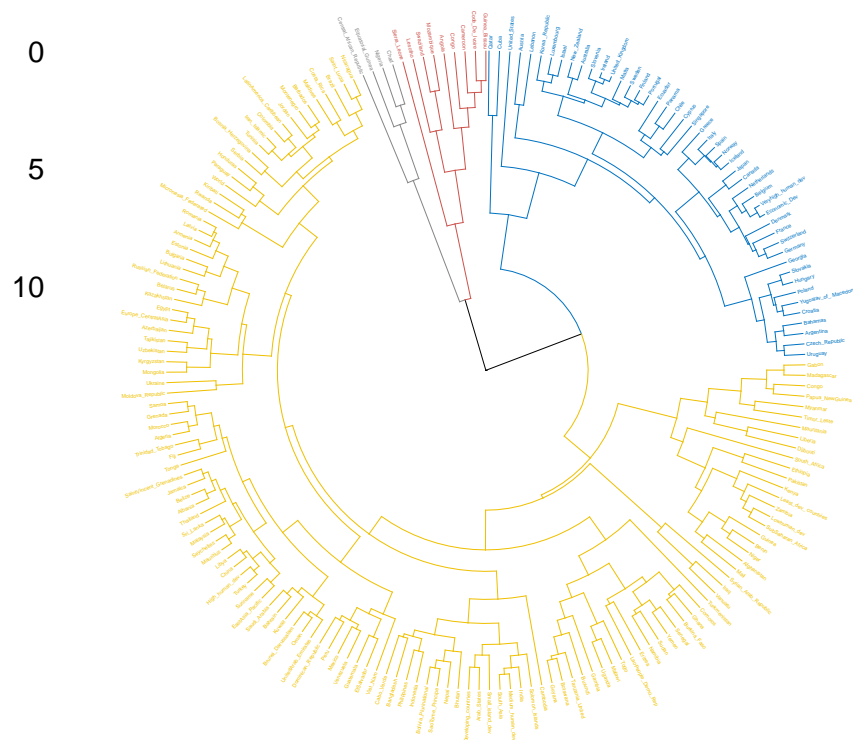
```
##
```

```
##   union
```

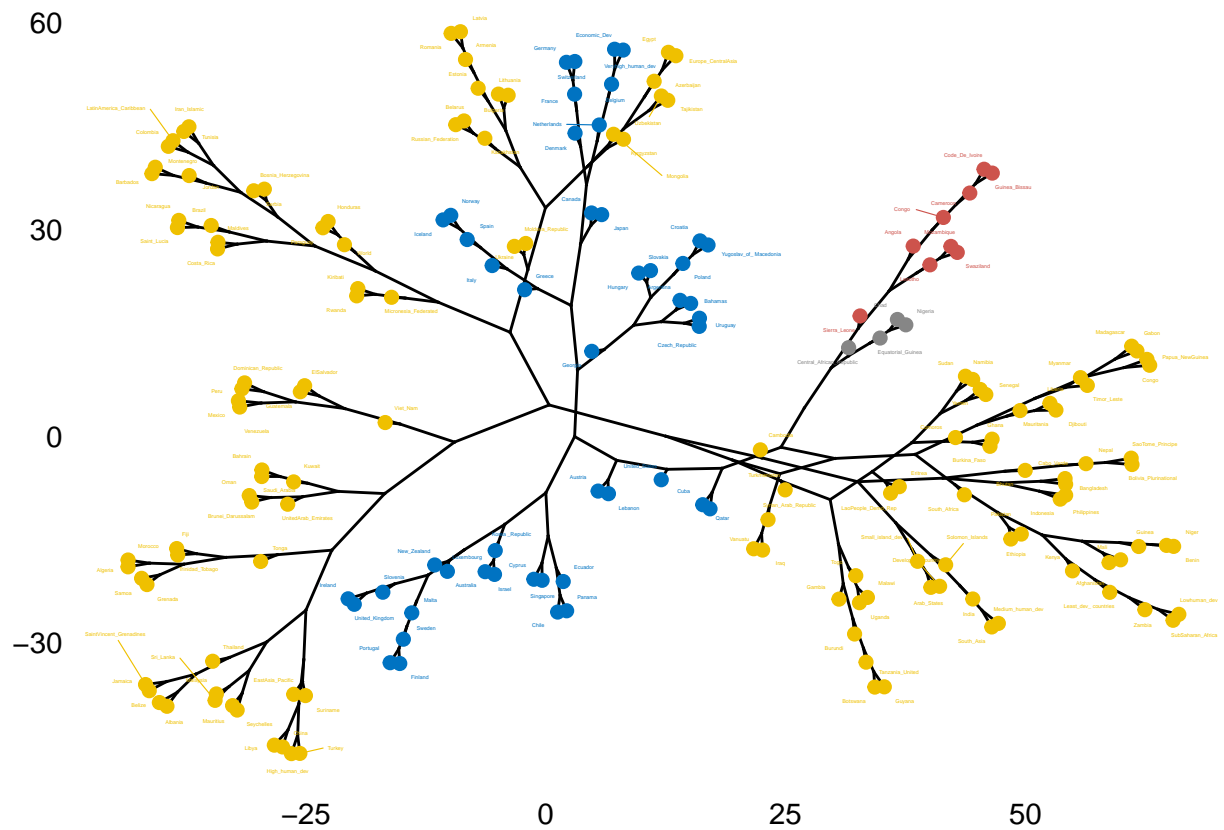
TO ASSIGN COLORS TO THE DENDOGRAM PLOT

```
fviz_dend(x = health_hc, cex = 0.15, lwd = 0.18, k = 4,  
rect = TRUE,  
  k_colors = "jco",  
rect_border = "jco",  
  rect_fill = TRUE,  
  type = "circular")
```

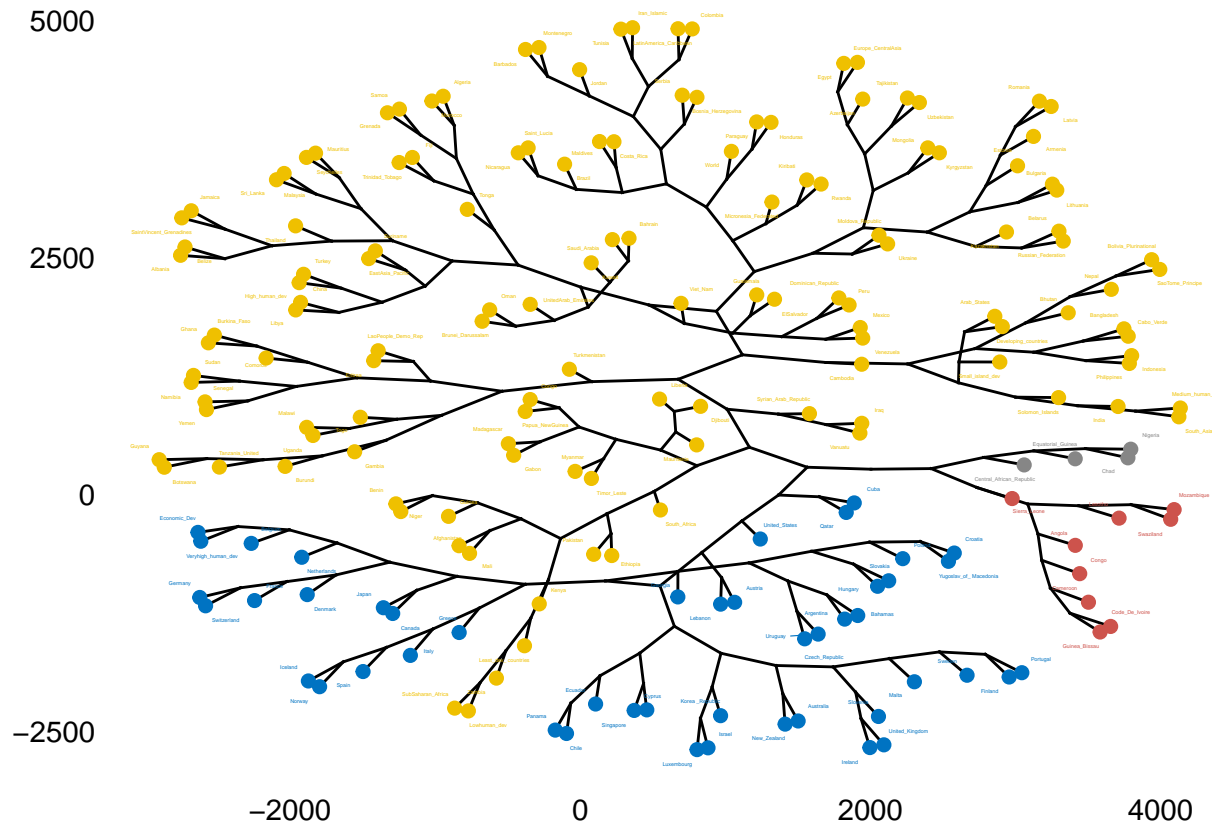
```
## Warning: 'guides(<scale> = FALSE)' is deprecated. Please use 'guides(<scale> =  
## "none")' instead.
```



```
fviz_dend(x = health_hc, cex = 0.15, lwd = 0.18, k = 4,
rect = TRUE,
  k_colors = "jco",
  rect_border = "jco",
  rect_fill = TRUE,
  type = "phylogenetic",
  repel = TRUE)
```

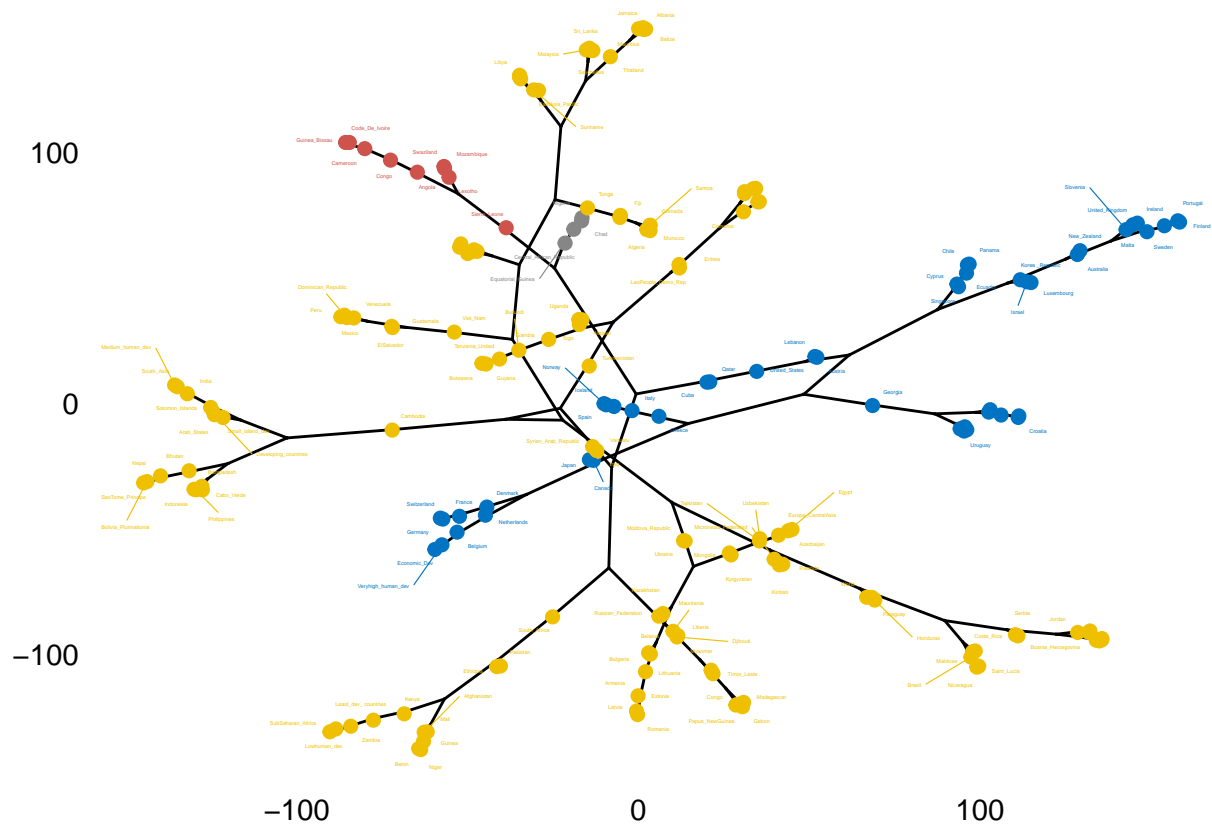


```
fviz_dend(health_hc, cex = 0.15, lwd = 0.18, k = 4,
  rect = TRUE,
  k_colors = "jco",
  rect_border = "jco",
  rect_fill = TRUE,
  type = "phylogenetic",
  repel = TRUE,
  phylo_layout = "layout.gem")
```

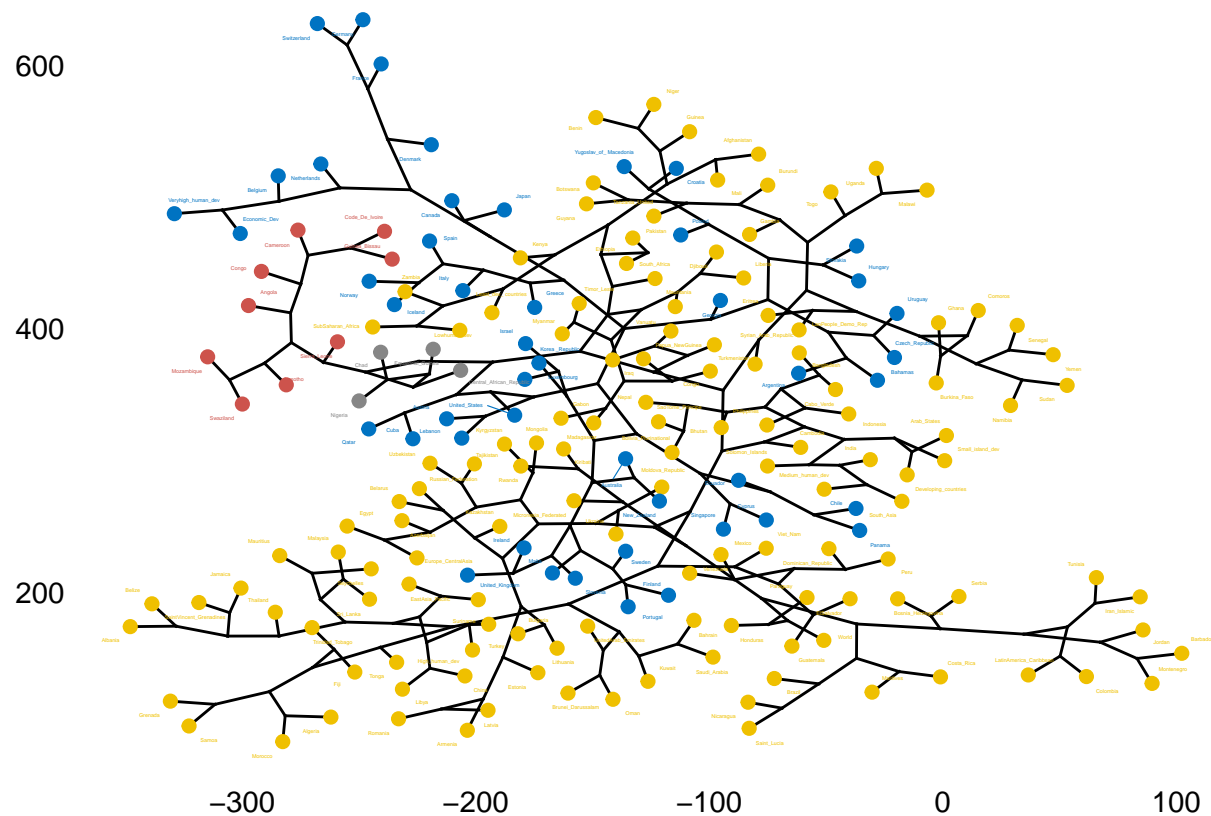


```
fviz_dend(health_hc, cex = 0.15, lwd = 0.18, k = 4,
  rect = TRUE,
  k_colors = "jco",
  rect_border = "jco",
  rect_fill = TRUE,
  type = "phylogenetic",
  repel = TRUE,
  phylo_layout = "layout.mds")
```

```
## Warning: ggrepel: 134 unlabeled data points (too many overlaps). Consider
## increasing max.overlaps
```

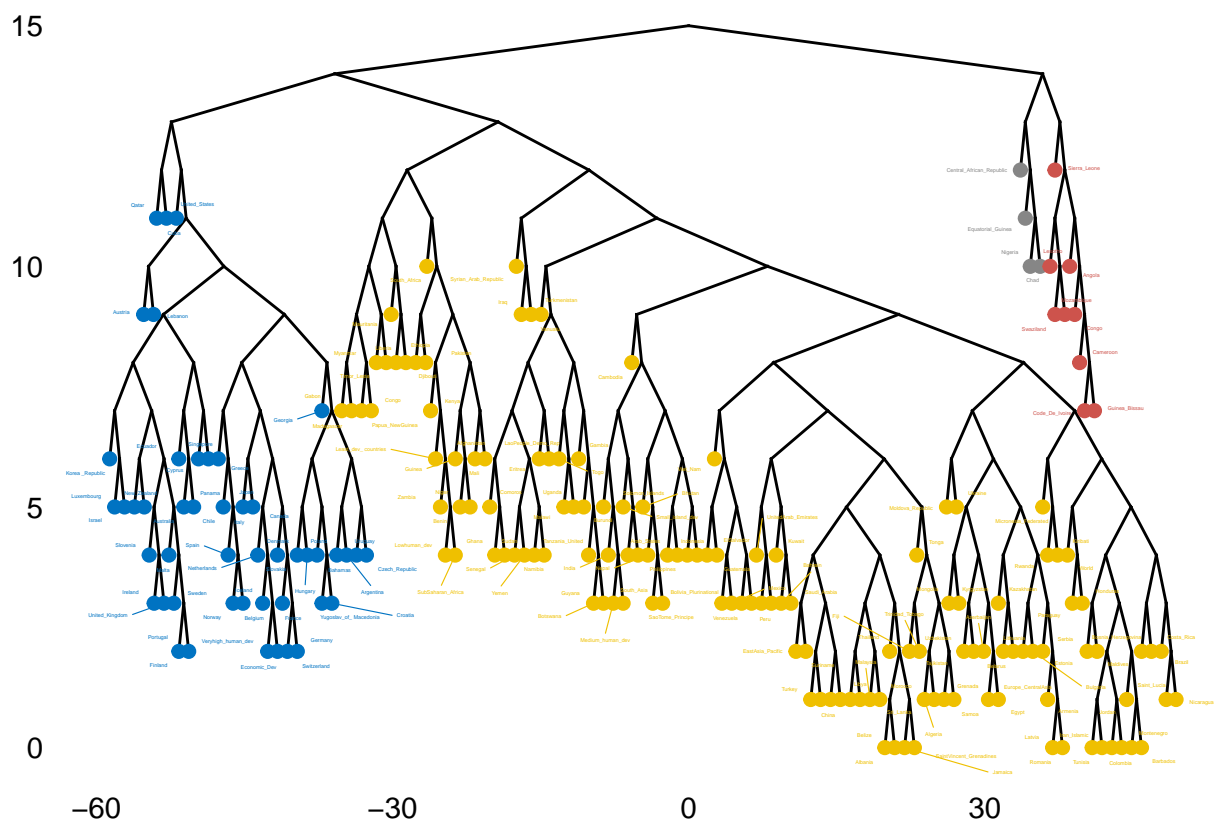



```
fviz_dend(health_hc, cex = 0.15, lwd = 0.18, k = 4,
  rect = TRUE,
  k_colors = "jco",
  rect_border = "jco",
  rect_fill = TRUE,
  type = "phylogenetic",
  repel = TRUE,
  phylo_layout = "layout_with_lgl")
```

```
fviz_dend(health_hc, cex = 0.15, lwd = 0.18, k = 4,
  rect = TRUE,
  k_colors = "jco",
  rect_border = "jco",
  rect_fill = TRUE,
  type = "phylogenetic",
  repel = TRUE,
  phylo_layout = "layout_as_tree")
```

```
## Warning: ggrepel: 12 unlabeled data points (too many overlaps). Consider
## increasing max.overlaps
```



```
fviz_dend(health_hc, cex = 0.15, lwd = 0.18, k = 4,
  rect = TRUE,
  k_colors = "jco",
  rect_border = "jco",
  rect_fill = FALSE)
```

```
## Warning: 'guides(<scale> = FALSE)' is deprecated. Please use 'guides(<scale> =
## "none")' instead.
```

Cluster Dendrogram

