TP5 : Classification Non Supervisée (Unsupervised Machine Learning) : Méthode des centres mobiles (K-MEANS CLUSTERING)

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Méthode des centres mobiles (K-MEANS)

Packages R: stats, factoextra

L'algorithme de classification par méthode des centres mobiles (KMEANS) de MacQueen, variante de l'algorithme de Forgy/Lloyd est l'une des plus connues et des plus utilisées.

Principe de l'algorithme K-means clustering :

On a des données d'individus qu'on souhaite classer en K groupes, tel que les individus dans un même groupe soient les plus similaires (forte similarité intra-classe), et les individus de groupes différents soient les plus dissimilaires (faible similarité inter-classe). Chaque groupe sera représentés par son centre(centroïd) qui correspond à la moyenne des points assignés au groupe.

On veut définir les groupes tel que la variance intra-groupe (total within-cluster variation) soit minimale.

variation intra-groupe : $W(C_k) = \sum_{x_i \in C_k} (x_i - \mu_k)^2$ où x_i point représentant un individu appartenant au groupe C_k et μ_k moyenne assigné au groupe C_k

Chaque individu (x_i) est assigné à un groupe tel que la somme des variances intra-groupes (total within-cluster variation) soit minimale.

$$tot.withinss = \sum_{k=1}^{K} W(C_k) = \sum_{k=1}^{K} \sum_{x_i \in C_k} (x_i - \mu_k)^2$$

K-means algorithm can be summarized as follow:

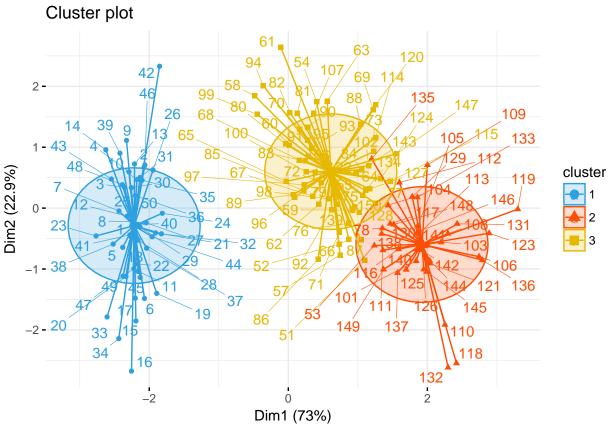
- 1) Specify the number of clusters (K) to be created (by the analyst)
- 2) Select randomly k objects from the dataset as the initial cluster centers or means
- 3) Assigns each observation to their closest centroid, based on the Euclidean distance between the object and the centroid
- 4)For each of the k clusters update the cluster centroid by calculating the new mean values of all the data points in the cluster. The centoid of a Kth cluster is a vector of length p containing the means of all variables for the observations in the kth cluster; p is the number of variables.
- 5) Iteratively minimize the total within sum of square. That is, iterate steps 3 and 4 until the cluster assignments stop changing or the maximum number of iterations is reached. By default, the R software uses 10 as the default value for the maximum number of iterations.

```
ir<-iris[,-5]
species = iris$Species

# Choix de K pour le clustering
fviz_nbclust(ir, kmeans, method = "wss") +
    geom_vline(xintercept = 3, linetype = 2)</pre>
```

Optimal number of clusters 600 Total Within Sum of Square 400 200 0 5 1 2 3 4 6 8 9 10 Number of clusters k # K-means avec K=3 set.seed(123) km.res <- kmeans(ir, 3, nstart = 25) # Print the results of kmeans print(km.res) ## K-means clustering with 3 clusters of sizes 50, 38, 62 ## ## Cluster means: ## Sepal.Length Sepal.Width Petal.Length Petal.Width ## 1 5.006000 3.428000 1.462000 0.246000 ## 2 6.850000 3.073684 5.742105 2.071053 ## 3 5.901613 2.748387 4.393548 1.433871 ## ## Clustering vector: ## [141] 2 2 3 2 2 2 3 2 2 3 ## ## Within cluster sum of squares by cluster: ## [1] 15.15100 23.87947 39.82097 (between_SS / total_SS = 88.4 %)

```
## Available components:
##
## [1] "cluster"
                                           "withinss"
                  "centers"
                               "totss"
## [5] "tot.withinss" "betweenss"
                               "size"
                                           "iter"
## [9] "ifault"
# Moyennes des variables dans les 3 groupes
aggregate(ir, by=list(cluster=km.res$cluster), mean)
    cluster Sepal.Length Sepal.Width Petal.Length Petal.Width
## 1
              5.006000
                        3.428000
                                   1.462000
                                             0.246000
         1
## 2
         2
              6.850000
                        3.073684
                                   5.742105
                                             2.071053
## 3
         3
              5.901613
                        2.748387
                                   4.393548
                                             1.433871
# Tableau des individus avec leur moyenne par variable et leur groupe
dd <- cbind(ir, cluster = km.res$cluster)</pre>
head(dd)
##
   Sepal.Length Sepal.Width Petal.Length Petal.Width cluster
                                1.4
           5.1
                     3.5
                                          0.2
## 2
                     3.0
                                          0.2
           4.9
                                1.4
                                                   1
## 3
           4.7
                     3.2
                                1.3
                                          0.2
                                                   1
## 4
           4.6
                     3.1
                                1.5
                                          0.2
                                                   1
## 5
           5.0
                     3.6
                                1.4
                                          0.2
                                                   1
## 6
           5.4
                     3.9
                                1.7
                                          0.4
                                                   1
# Groupe de chaque observations
km.res$cluster
    ## [141] 2 2 3 2 2 2 3 2 2 3
# Effectif des groupes
km.res$size
## [1] 50 38 62
# Centres des groupes
km.res$centers
    Sepal.Length Sepal.Width Petal.Length Petal.Width
## 1
       5.006000
                 3.428000
                            1.462000
                                      0.246000
## 2
       6.850000
                 3.073684
                            5.742105
                                      2.071053
## 3
       5.901613
                 2.748387
                            4.393548
                                      1.433871
# Visualisation des individus en fonction de leur groupes
fviz cluster(km.res, data = ir,
          palette = c("#2E9FDF", "#FC4E07", "#E7B800"),
          ellipse.type = "euclid", # Concentration ellipse
          star.plot = TRUE, # Add segments from centroids to items
          repel = TRUE, # Avoid label overplotting (slow)
           ggtheme = theme_minimal()
```



```
# Etude des pétales
kmoy3 < -kmeans(ir[,c(3,4)],3,nstart=4)
kmoy3
## K-means clustering with 3 clusters of sizes 50, 52, 48
##
## Cluster means:
##
   Petal.Length Petal.Width
## 1
     1.462000
            0.246000
## 2
     4.269231
             1.342308
## 3
     5.595833
             2.037500
##
## Clustering vector:
   ##
  [141] 3 3 3 3 3 3 3 3 3 3 3
##
## Within cluster sum of squares by cluster:
## [1] 2.02200 13.05769 16.29167
  (between_SS / total_SS = 94.3 %)
##
##
## Available components:
##
```

```
## [1] "cluster"
                      "centers"
                                      "totss"
                                                      "withinss"
## [5] "tot.withinss" "betweenss"
                                      "size"
                                                     "iter"
## [9] "ifault"
table(kmoy3$cluster,species)
##
      species
##
       setosa versicolor virginica
           50
##
                       0
     1
##
            0
                      48
                                  4
     2
            0
                       2
                                 46
     3
##
par(mfrow=c(1,2))
plot(ir[c("Petal.Length", "Petal.Width")], col=kmoy3$cluster)
points(kmoy3$centers[,c("Petal.Length", "Petal.Width")],
       col=1:3, pch=23, cex=3)
plot(ir[c("Petal.Length", "Petal.Width")],col=iris$Species)
```

