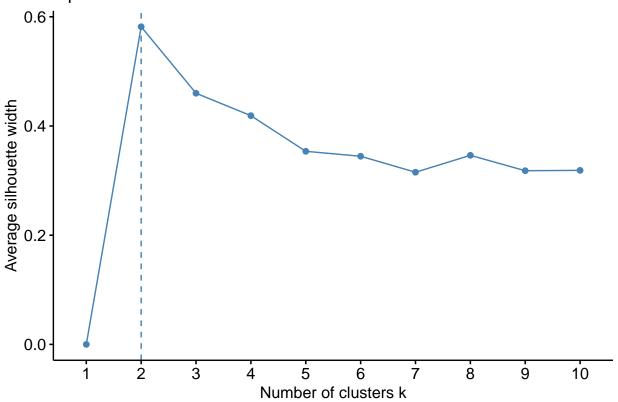
ex_clusters_Machine_Leraning

Identify clusters in the data

```
df <- iris[,1:4]
# find optimal number of clusters
dff <- scale(df) # base R function
factoextra::fviz_nbclust(dff, kmeans, method = "silhouette")</pre>
```

Optimal number of clusters



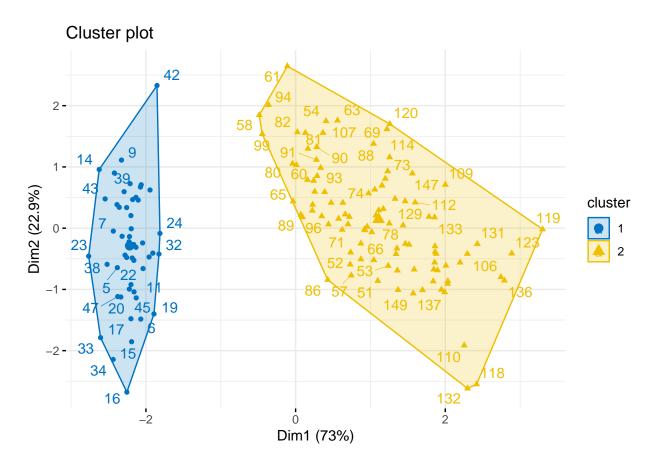
```
# methods may be<. gap_stat, wss, silhouette
#WHICH k? Where sudden "change decay slope"!

# compute and visualise
set.seed(123)
km.res <- kmeans(dff, 2, nstart = 25)
km.res</pre>
```

```
\mbox{\tt \#\#} K-means clustering with 2 clusters of sizes 50, 100 \mbox{\tt \#\#}
```

```
## Cluster means:
## Sepal.Length Sepal.Width Petal.Length Petal.Width
                   -1.300630 -1.2507035
## 1 -1.0111914 0.8504137
     0.5055957 -0.4252069
## 2
                     0.650315 0.6253518
## Clustering vector:
## [149] 2 2
## Within cluster sum of squares by cluster:
## [1] 47.35062 173.52867
## (between_SS / total_SS = 62.9 %)
## Available components:
##
## [1] "cluster"
                                           "tot.withinss"
              "centers"
                       "totss"
                                 "withinss"
                       "iter"
## [6] "betweenss"
              "size"
                                 "ifault"
# visualize
 factoextra::fviz_cluster(km.res, data = dff,
         ellipse.type = "convex",
          # "norm",
           # "convex",
         palette = "jco",
         repel = TRUE,
         ggtheme = ggplot2::theme_minimal())
```

Warning: ggrepel: 81 unlabeled data points (too many overlaps). Consider ## increasing max.overlaps



may adjust max.overlaps to show all labels on plots

... and continue

```
# compare with PAM clustering
# Compute PAM
pam.res <- cluster::pam(dff, 2)
# Visualize
factoextra::fviz_cluster(pam.res)</pre>
```

