# K-Means Problem

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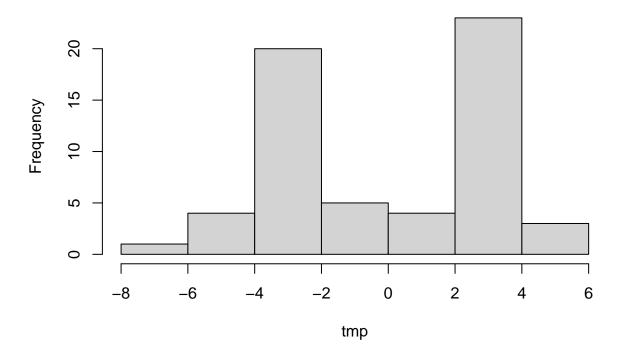
# Try K-Means Clustering

Generate fake data and explore how the method works.

### Generate example data

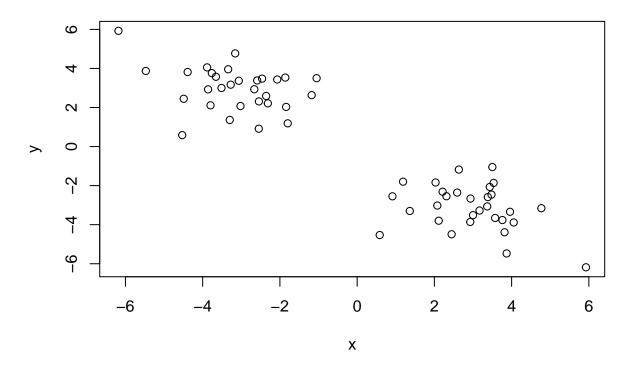
```
tmp <- c(rnorm(30,-3), rnorm(30,3))
hist(tmp)</pre>
```

# Histogram of tmp



## Generate multidimensional example data

```
x <- cbind(x = tmp, y = rev(tmp))
plot(x)</pre>
```



## Use the kmeans() function to explore the fake data

Use it while specifying 2 expected clusters and iterating 20 times.

```
clusters <- kmeans(x, centers = 2, nstart = 20)</pre>
clusters
## K-means clustering with 2 clusters of sizes 30, 30
## Cluster means:
##
          x
## 1 -3.131876 2.964332
## 2 2.964332 -3.131876
##
## Clustering vector:
  ##
## Within cluster sum of squares by cluster:
## [1] 77.62072 77.62072
   (between_SS / total_SS = 87.8 %)
## Available components:
##
## [1] "cluster"
                  "centers"
                               "totss"
                                                        "tot.withinss"
                                           "withinss"
## [6] "betweenss"
                  "size"
                               "iter"
                                           "ifault"
   [Q] How many points are in each cluster?
There are 30 points in each cluster.
```

## [1] 30 30

clusters\$size

[Q] What component of your results object dteails:

Cluster size

```
clusters$size
```

```
## [1] 30 30
```

Cluster assignment

#### clusters\$cluster

Cluster center

#### clusters\$centers

```
## x y
## 1 -3.131876 2.964332
## 2 2.964332 -3.131876
```

#### Plot x colored by the kmeans cluster centers as blue points

Load ggplot2

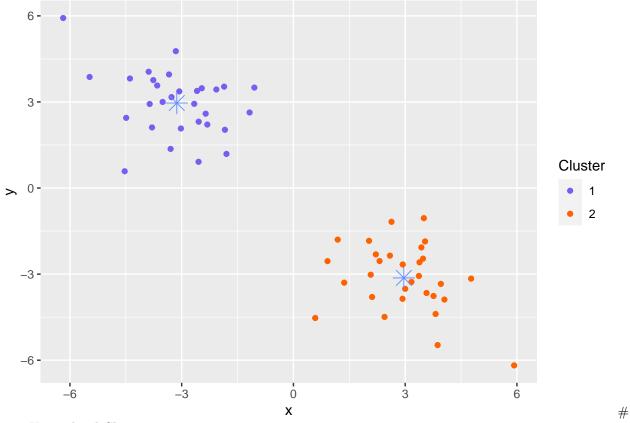
```
library(ggplot2)
```

Convert matrices to be used in ggplot to data frames.

```
df <- data.frame(x)
centroids <- data.frame(clusters$centers)</pre>
```

Plot the original data colored by kmenas clusters and add blue centroids. IBM's colorblind palette is used.

```
ggplot(data = df) +
  aes(x = x, y = y, color = factor(clusters$cluster)) +
  geom_point() +
  scale_color_manual(values = c("#785EF0", "#FE6100"), name = "Cluster") +
  geom_point(data = centroids, aes(x = x, y = y), color = "#648FFF", shape = 8, size = 5)
```



Try Hierarchical Clustering

Using the same example data x.

#### Generate the distance matrix

```
dm <- dist(x)
str(dm)

## 'dist' num [1:1770] 1.58 1.17 1.27 1.47 2.38 ...

## - attr(*, "Size")= int 60

## - attr(*, "Diag")= logi FALSE

## - attr(*, "Upper")= logi FALSE

## - attr(*, "method")= chr "euclidean"

## - attr(*, "call")= language dist(x = x)</pre>
```

## Call hclust() to determine clusters

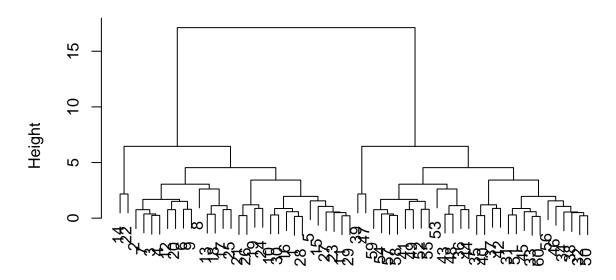
```
hc <- hclust(dm)
hc

##
## Call:
## hclust(d = dm)
##
## Cluster method : complete
## Distance : euclidean</pre>
```

## Plot the hierachical cluster

plot(hc)

# **Cluster Dendrogram**



dm hclust (\*, "complete")