# K-means method for clustering K-means Examples

STAT 32950-24620

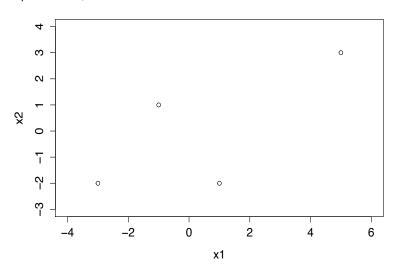
Spring 2025 (wk7)

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# $\underline{\mathsf{Example}\ 1}\ \mathsf{Consider}\ \mathsf{various}\ \mathsf{initialization}\ \mathsf{and}\ \mathsf{clustering}\ \mathsf{methods}$

4 points in  $\mathbb{R}^2$ , to form 2 clusters



# K-means method for clustering

For a given K – number of desired clusters, the optimal clustering minimizes the **total within-cluster sums of squares**:

$$\sum_{i=1}^K \sum_{i=1}^{n_i} d_{ij,c(i)}^2 = \sum_{i=1}^K \sum_{i=1}^{n_i} (x_{ij} - c(i))^2$$

where

K the total number of clusters (usually preassigned in the K-means method)

 $n_i$  the number of members in cluster i (varies at each step of the algorithm)

c(i) is the centroid of cluster i (varies at each step of the algorithm)

 $X_{i_i}$  the jth member of cluster i (varies at each step of the algorithm)

 $d_{i_i,c(i)}$  the distance between  $x_{i_i}$  and c(i) (Euclidean distance most common)

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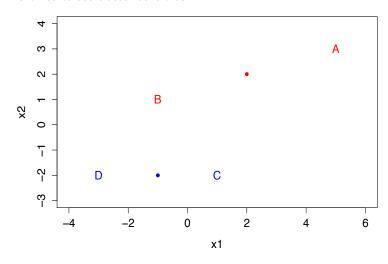
Start with a random assignment:

- Either, assign K cluster centroids randomly:
  - Assign all objects to the cluster of nearest centroid
  - Recalculate the cluster centroids
  - Repeat
- Or, assign cluster membership randomly:
  - Calculate the cluster centroids
  - Re-assign all objects to the cluster of nearest centroid
  - Repeat

Ex1: Start with randomly assigned membership (vs start with centroids)

Start with clusters (AB) and (CD).

Next: Calculate cluster centroids.



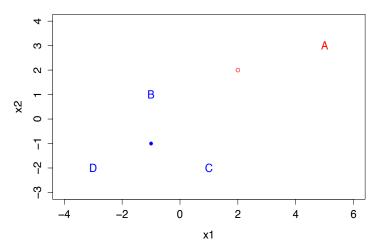
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Re-assign element to nearest centroids.

Reassign A, C, D - rejected.

 $Reassigning \ B \ - \ accepted. \ \ (\textit{Early MacQueen: Update affected cluster centroids right away.})$ 

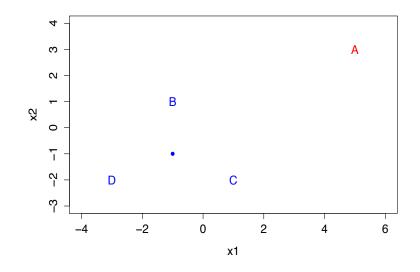


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#### With the new clusters,

- Re-calculate the new cluster centroids.
- Check the distance of each items to the centroids (Early Floyd: update centroids after checking all items)
- Re-assign items to the cluster with the nearest centroid.
- Repeat.
- Stop when no appreciable improvement of total within-cluster sum of squares.

Final clusters: (A) and (BCD)



All possible combinations of k = 2 clusters in the example:

- (A) (BCD)
- (B) (ACD)
- (C) (ABD)
- (D) (ABC)
- (AB) (CD)
- (AC) (BD)
- (AD) (BC)

Of all 2-clusters, the partition  $\{(A), (BCD)\}$  minimizes the total within-cluster sums of squares

$$\sum_{i=1}^{2} \sum_{j=1}^{2} d_{i_{j},c(i)}^{2} = \sum_{i=1}^{2} \sum_{j=1}^{2} (x_{i_{j}} - c(i))^{2}$$

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#### help(kmeans)

kmeans {stats} K-Means Clustering

Description

Perform k-means clustering on a data matrix.

Usage

Arguments

Value

kmeans returns an object of class "kmeans" which has a print and a fitted method. It is a list with at least the following components:

cluster

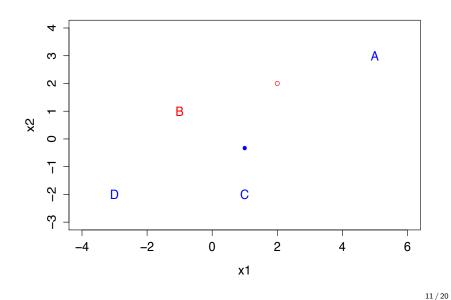
centers

. . .

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The kmeans function in R

#### Discussion: Different algorithms. (Lloyd, vs MacQueen vs. Hartigan-Wong: min ESS update)



#### Using the kmeans function in R for Example 1

```
x1=c(5,-1,1,-3); x2=c(3,1,-2,-2)
M0 = kmeans(cbind(x1,x2),2)
```

MO\$cluster

[1] 1 2 2 2 # (A), (BCD)

MO\$size [1] 1 3

[1] 1 0

MO\$center

x1 x2

1 5 3

2 -1 -1

MO\$withinss

[1] 0 14 MO\$tot.withinss

[1] 14

MO\$betweenss

[1] 39

MO\$totss [1] 53

dist(MO\$center)

2 7.211103

Available components:

-

[1] "cluster" "centers" [7] "size" "iter"

"totss" "ifault" "withinss"

ninss" "tot.withinss" "betweenss"

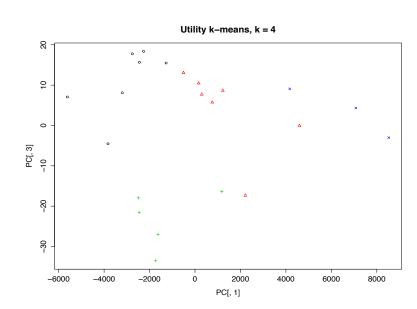
#### Example 2: K-means method applied to the utility data: First, let K=4

```
data = read.table("T12-4.dat"); X = data[,1:8]
NormX = as.matrix(X)%*%solve(diag(sqrt(diag(var(X)))))
Mkm = kmeans(NormX,4)
Mkm$cluster
[1] 3 4 3 1 4 3 4 2 3 1 2 4 1 3 4 2 4 3 3 1 4 1
Mkm$size
[1] 5 3 7 7
print(Mkm$center, digits=3)
  [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8]
1 6.03 5.12 4.30 12.4 1.206 2.11 2.280 1.39
2 5.44 3.95 5.42 12.3 2.031 4.37 0.000 1.02
3 6.54 5.56 3.10 12.4 0.550 2.86 0.191 1.57
4 5.80 4.13 4.34 13.6 0.985 1.65 0.426 3.23
Mkm$withinss
[1] 10.177094 9.533522 26.507769 34.164812
Mkm$betweenss
[1] 87.6168
print(dist(Mkm$center),digits=3)
    1 2 3
2 3.75
3 2.70 3.75
4 3.08 4.07 3.15
```

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#### Ex2: K-means method applied to the utility data, now K=5

```
Mkm5 = kmeans(NormX,5)
Mkm5$cluster
[1] 4 3 4 2 1 4 3 5 4 2 5 3 2 4 3 5 3 4 4 2 3 2
Mkm5$size
[1] 1 5 6 7 3
print(Mkm5$center, digits=3)
  [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8]
1 8.08 3.92 4.66 11.5 0.321 0.93 0.929 3.68
2 6.03 5.12 4.30 12.4 1.206 2.11 2.280 1.39
3 5.42 4.16 4.28 13.9 1.096 1.77 0.342 3.16
4 6.54 5.56 3.10 12.4 0.550 2.86 0.191 1.57
5 5.44 3.95 5.42 12.3 2.031 4.37 0.000 1.02
Mkm5$withinss
[1] 0.000000 10.177094 21.187976 26.507769 9.533522
Mkm5$betweenss
[1] 100.5936
print(dist(Mkm5$center),digits=3)
1 2 3 4
2 3.98
3 3.89 3.25
4 4.14 2.70 3.30
5 5.56 3.75 4.04 3.75
```



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# The effect of initial assignments

#### Ex2: K-means method applied to the utility data with K=4 again.

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# The effect of initial assignments (cont.)

Ex2: And again, K-means method applied to the utility data with K=4.

```
> Mkm = kmeans(NormX,4)
> Mkm

K-means clustering with 4 clusters of sizes 11, 4, 4, 3

Cluster means:
    [,1]    [,2]    [,3]    [,4]    [,5]    [,6]    [,7]    [,8]
1 6.242539 5.39374 3.555468 12.322459 0.6880315 2.565261 1.1580245 1.490417
2 4.945500 3.665249 4.284880 13.61757 1.1705282 1.684515 0.5136399 3.142881
3 7.018545 5.002118 4.327365 13.51670 1.1304417 1.795924 0.2322546 2.901916
4 5.437792 3.951191 5.421850 12.29131 2.0310536 4.367531 0.0000000 1.017207

Clustering vector:
    [1] 1 2 1 1 3 1 3 4 3 1 4 3 1 1 2 4 2 1 1 1 2 1

Within cluster sum of squares by cluster:
    [1] 51.590389 12.574083 18.083154 9.533522
    (between, SS / total_SS = 45.4 %)
```

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# The effect of initial assignments (cont.) - the worst?

Ex2: K-means method applied to the utility data with K=4. The worst case?

```
> kmeans(NormX,4)
```

K-means clustering with 4 clusters of sizes 5, 4, 4, 9

(between\_SS / total\_SS = 42.8 %)

Cluster means:
[,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8]
1 6.557869 5.810924 2.786993 12.26142 0.2052433 3.023056 0.2679860 1.665534
2 7.018545 5.002118 4.327365 13.51670 1.1304417 1.795924 0.2322546 2.901916
3 4.945500 3.665249 4.284880 13.61757 1.1705282 1.684515 0.5136399 3.142881
4 5.799107 4.614674 4.604527 12.34859 1.4039212 2.911688 1.2664821 1.235394

Clustering vector:
[1] 4 3 1 4 2 1 2 4 2 4 4 2 4 1 3 4 3 1 1 4 3 4

Within cluster sum of squares by cluster:
[1] 15.15613 18.08315 12.57408 50.29653

# The effect of initial assignments (cont.)

Ex2: One more time, K-means method applied to the utility data with K=4.

```
> Mkm = kmeans(NormX,4)
Mkm
K-means clustering with 4 clusters of sizes 5, 6, 1, 10
     [,1] [,2]
                     [,3]
                              [,4]
                                         [,5]
                                                   [,6]
                                                            [,7]
1 6.557869 5.810924 2.786993 12.26142 0.2052433 3.0230559 0.2679860 1.665534
2 5.419726 4.159148 4.284880 13.92018 1.0956999 1.7707122 0.3424266 3.158915
3 8.075392 3.921482 4.661173 11.47687 0.3206927 0.9295816 0.9290183 3.675611
4 5.945439 4.732516 4.551927 12.46764 1.4944278 2.8573086 1.1398339 1.266863
Clustering vector:
[1] 4 2 1 4 3 1 2 4 4 4 4 2 4 1 2 4 2 1 1 4 2 4
Within cluster sum of squares by cluster:
[1] 15.15613 21.18798 0.00000 57.53424
 (between_SS / total_SS = 44.1 %)
```

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# The effect of initial assignments (cont.) - the best?

Ex2: K-means method applied to the utility data with K = 4. The best case?

```
K-means clustering with 4 clusters of sizes 7, 5, 3, 7

Cluster means:
    [,1]    [,2]    [,3]    [,4]    [,5]    [,6]    [,7]    [,8] 
1 6.542384 5.563921 3.097044 12.43434 0.5497589 2.862870 0.1914186 1.572436 
2 6.026735 5.115752 4.301874 12.41385 1.2058044 2.109136 2.2796679 1.387525 
3 5.437792 3.951191 5.421850 12.29131 2.0310536 4.367531 0.0000000 1.017207 
4 5.799107 4.125196 4.338636 13.57114 0.9849846 1.650551 0.4262254 3.232729 
Clustering vector:
    [1] 1 4 1 2 4 1 4 3 1 2 3 4 2 1 4 3 4 1 1 2 4 2

Within cluster sum of squares by cluster:
    [1] 26.507769 10.177094 9.533522 34.164812 
    (between, SS / total_SS = 52.2 %)
```

Remarks: There is always some output, even if the results are not optimal.

Example: kmeans(distance-matrix, k)

> kmeans(NormX,4)