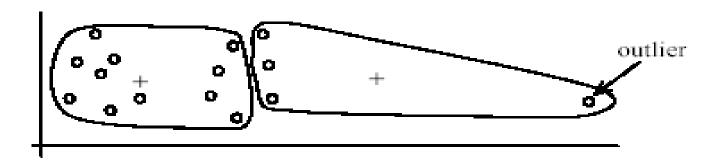
CS 461 Artificial Intelligence

Limitations in K-means Clustering

- K-means has problems when the data contains outliers
- The K-means algorithm is very sensitive to the initial seeds.

- K-means has problems when clusters are of different
 - Sizes
 - Densities
 - Non-globular shapes

K-means has problems when the data contains outliers

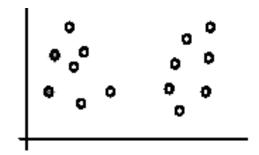


(A): Undesirable clusters

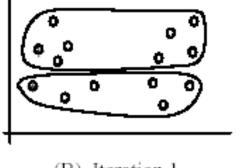


(B): Ideal clusters

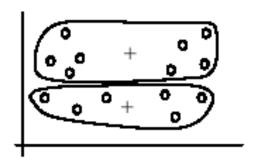
The algorithm is sensitive to initial seeds



(A). Random selection of seeds (centroids)



(B). Iteration 1

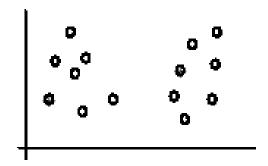


(C). Iteration 2

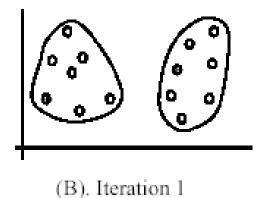
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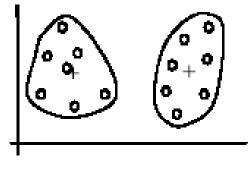
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The algorithm is sensitive to initial seeds



(A). Random selection of k seeds (centroids)



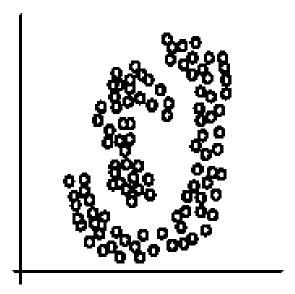


(C). Iteration 2

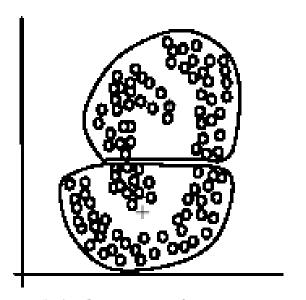
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▶ The *k*-means algorithm is not suitable for discovering clusters that are not hyper-ellipsoids (or hyper-spheres).



(A): Two natural clusters



(B): k-means clusters

- In the Euclidean space, standardization of attributes is recommended so that all attributes can have equal impact on the computation of distances.
- Consider the following pair of data points:

$$x_i$$
: (0.1, 20) and x_j : (0.9, 720)
$$dist(\mathbf{x}_i, \mathbf{x}_j) = \sqrt{(0.9 - 0.1)^2 + (720 - 20)^2} = 700.000457,$$

- The distance is almost completely dominated by (720 20) = 700.
- Standardize attributes: to force the attributes to have a common value range,

Interval-scaled attributes:

- Their values are real numbers following a linear scale.
 - The difference in Age between 10 and 20 is the same as that between 40 and 50.
 - The key idea is that intervals keep the same importance through out the scale
- Two main approaches to standardize interval scaled attributes, range and z-score.

Range:

Consider f is an attribute $range(x_{if}) = \frac{x_{if} - \min(f)}{\max(f) - \min(f)},$

Z-score:

- transforms the attribute values so that they have a mean of zero and a mean absolute deviation of 1.
- The mean and absolute deviation of attribute f, denoted by m_f and s_f respectively is computed as,

$$m_{f} = \frac{1}{n} \left(x_{1f} + x_{2f} + \dots + x_{nf} \right),$$

$$s_{f} = \frac{1}{n} \left(|x_{1f} - m_{f}| + |x_{2f} - m_{f}| + \dots + |x_{nf} - m_{f}| \right),$$

$$z(x_{if}) = \frac{x_{if} - m_{f}}{s_{f}}.$$

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Ratio-scaled attributes:

- Numeric attributes, but unlike interval-scaled attributes, their scales are exponential,
- For example, the total amount of microorganisms that evolve in a time t is approximately given by

$$Ae^{Bt}$$
,

- where A and B are some positive constants.
- Do log transform:

$$\log(x_{if})$$

Then treat it as an interval-scaled attribute

- The k-means algorithm is sensitive to outliers!
 - Since an object with an extremely large value may substantially distort the distribution of the data.

K-Medoids:

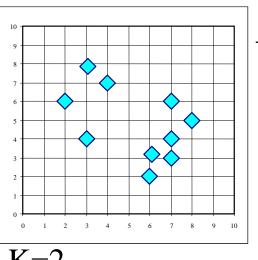
Instead of taking the mean value of the object in a cluster as a reference point, medoids can be used, which is the most centrally located object in a cluster.

Find representative objects, called medoids, in the clusters

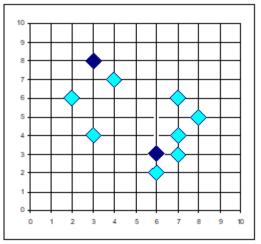
PAM (Partitioning Around Medoids, 1987)

- starts from an initial set of medoids and iteratively replaces one of the medoids by one of the non-medoids if it improves the total distance of the resulting clustering
- PAM works effectively for small data sets, but does not scale well for large data sets

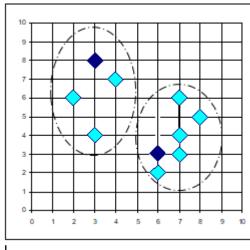




Arbitrary choose k object as initial medoids



Assign each remaining object to nearest medoids



K=2

Total Cost = 26

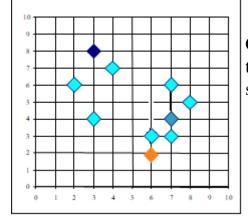
Randomly select a nonmedoid object,O_{ramdom}

Do loop

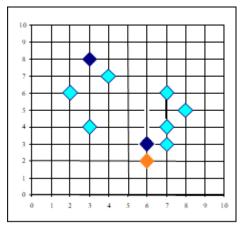
Until no change

Swapping O and O_{ramdom}

If quality is improved.



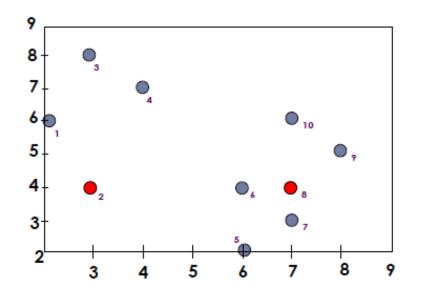
Compute total cost of swapping



- Use real object to represent the cluster
 - 1. Select k representative objects arbitrarily
 - For each pair of non-selected object h and selected object
 i, calculate the total swapping cost TC_{ih}
 - 3. For each pair of i and h,
 - If $TC_{ih} < 0$, \boldsymbol{i} is replaced by \boldsymbol{h}
 - Then assign each non-selected object to the most similar representative object
 - 4. repeat steps 2-3 until there is no change

Data Objects

	A ₁	A ₂
O ₁	2	6
02	3	4
O_3	3	8
O_4	4	7
O_5	6	2
O_6	6	4
O ₇	7	3
O ₈	7	4
O_9	8	5
O ₁₀	7	6



Goal: create two clusters

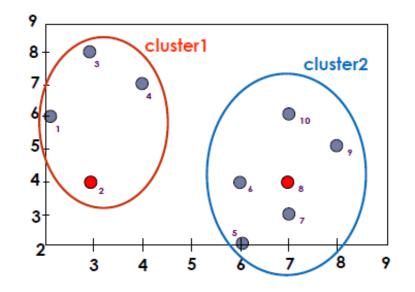
Choose randmly two medoids

$$O_2 = (3,4)$$

 $O_8 = (7,4)$

Data Objects

	A_1	A_2
01	2	6
02	3	4
O_3	3	8
O_4	4	7
O_5	6	2
O_6	6	4
O ₇	7	3
O ₈	7	4
O_9	8	5
O ₁₀	7	6



- →Assign each object to the closest representative object
- →Using L1 Metric (Manhattan), we form the following clusters

Cluster1 =
$$\{O_1, O_2, O_3, O_4\}$$

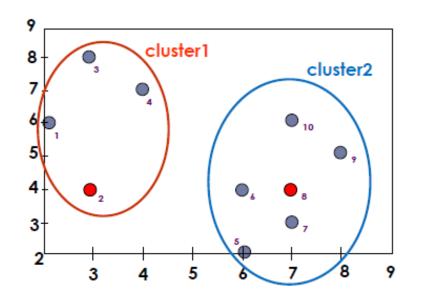
Cluster2 =
$$\{O_5, O_6, O_7, O_8, O_9, O_{10}\}$$

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Data Objects

	A_1	A_2
O ₁	2	6
02	3	4
O_3	3	8
O_4	4	7
O_5	6	2
O_6	6	4
O ₇	7	3
O ₈	7	4
O_9	8	5

O₁₀

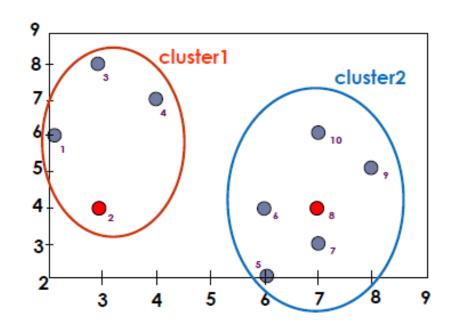


→Compute the absolute error criterion [for the set of Medoids (O2,O8)]

$$\begin{split} E = & \sum_{i=1}^{k} \sum_{p \in C_i} p - o_i \mid = \mid o_1 - o_2 \mid + \mid o_3 - o_2 \mid + \mid o_4 - o_2 \mid \\ & + \mid o_5 - o_8 \mid + \mid o_6 - o_8 \mid + \mid o_7 - o_8 \mid + \mid o_9 - o_8 \mid + \mid o_{10} - o_8 \mid \end{split}$$

Data Objects

	A_1	A_2
O ₁	2	6
02	3	4
O_3	3	8
O_4	4	7
O ₅	6	2
O_6	6	4
O ₇	7	3
O ₈	7	4
O_9	8	5

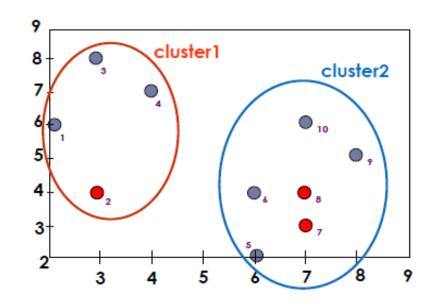


→The absolute error criterion [for the set of Medoids (O2,O8)]

$$E = (3+4+4)+(3+1+1+2+2) = 20$$

Data Objects

	A ₁	A_2
01	2	6
02	3	4
O_3	3	8
O_4	4	7
O_5	6	2
O_6	6	4
O ₇	7	3
O ₈	7	4
O_9	8	5

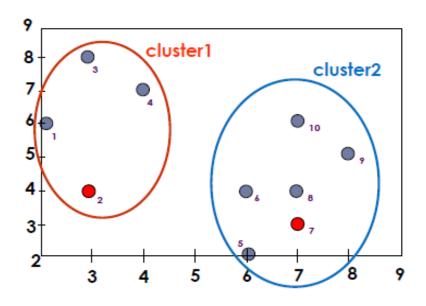


- →Choose a random object O₇
- →Swap O8 and O7
- →Compute the absolute error criterion [for the set of Medoids (O2,O7)]

$$E = (3+4+4)+(2+2+1+3+3)=22$$

Data Objects

	A_1	A_2
01	2	6
02	3	4
O_3	3	8
O_4	4	7
O ₅	6	2
O_6	6	4
O ₇	7	3
O ₈	7	4
O_9	8	5
O ₁₀	7	6



→Compute the cost function

Absolute error [for $O_{2'}O_7$] – Absolute error $[O_{2'}O_8]$

$$S = 22 - 20$$

 $S>0 \Rightarrow$ it is a bad idea to replace O_8 by O_7

- PAM is more robust than k-means in the presence of noise and outliers because a medoid is less influenced by outliers or other extreme values than a mean
- PAM works efficiently for small data sets but does not scale well for large data sets.
- $O(k(n-k)^2)$ for each iteration
 - where n is # of data points,
 - k is # of clusters