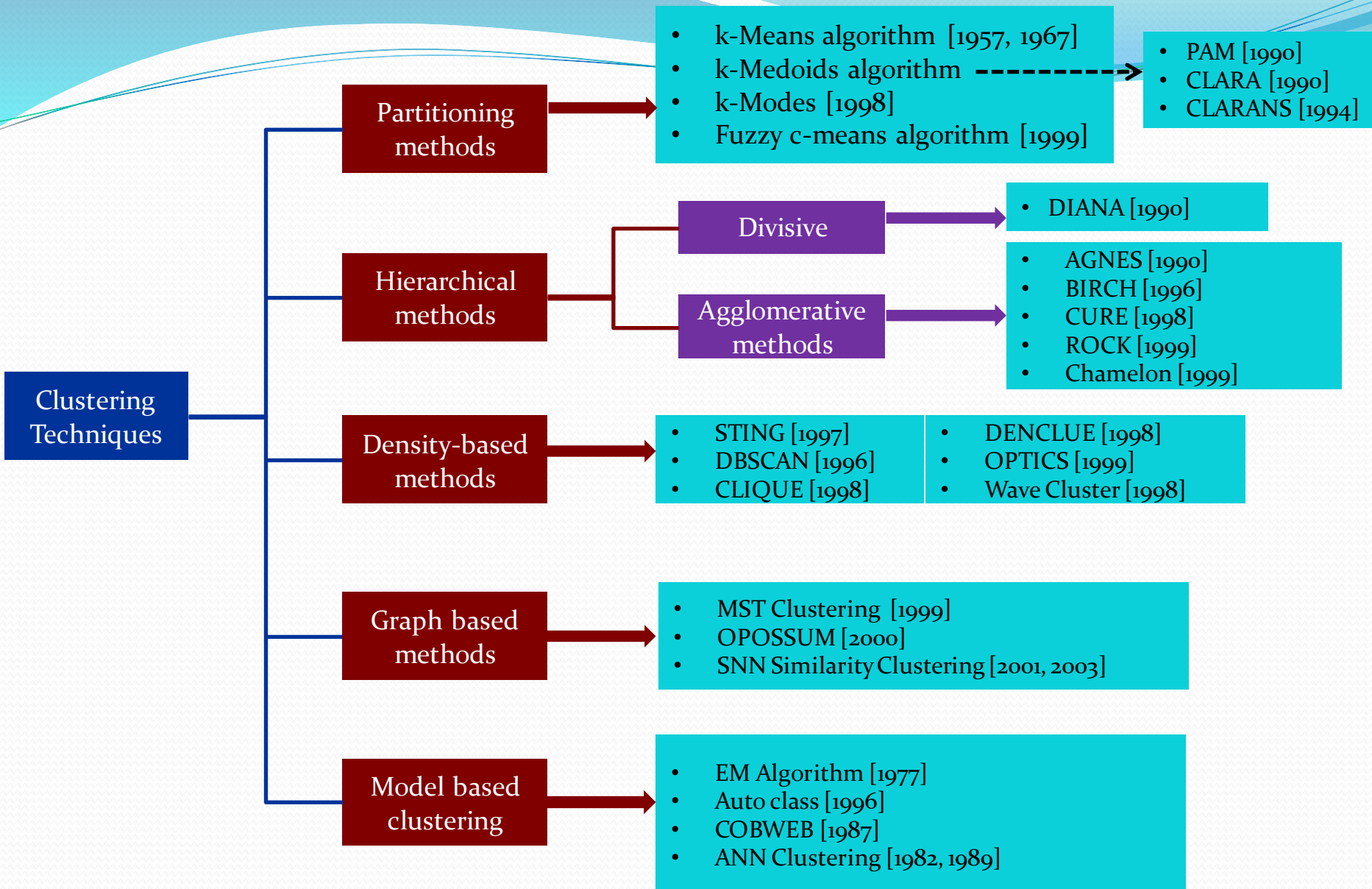


# Clustering Techniques

- Introduction to clustering
- Similarity and dissimilarity measures
- Clustering techniques
- Partitioning algorithms
- Hierarchical algorithms
- Density-based algorithm

# Clustering techniques

- Clustering has been studied extensively for more than 40 years and across many disciplines due to its broad applications.
- As a result, many clustering techniques have been reported in the literature.
- Let us categorize the clustering methods. In fact, it is difficult to provide a crisp categorization because many techniques overlap to each other in terms of clustering paradigms or features.
- A broad taxonomy of existing clustering methods is shown in Fig. 16.1.
- It is not possible to cover all the techniques in this lecture series. We emphasize on major techniques belong to partitioning and hierarchical algorithms.



# Clustering techniques

- In this lecture, we shall cover the following clustering techniques only.
  - Partitioning
    - k-Means algorithm
    - PAM (k-Medoids algorithm)
  - Hierarchical
    - DIANA (divisive algorithm)
    - AGNES } (Agglomerative algorithm)
    - ROCK }
  - Density – Based
    - DBSCAN

# k-Means Algorithm

- k-Means clustering algorithm proposed by J. Hartigan and M. A. Wong [1979].
- Given a set of  $n$  distinct objects, the k-Means clustering algorithm partitions the objects into  $k$  number of clusters such that intracluster similarity is high but the intercluster similarity is low.
- In this algorithm, user has to specify  $k$ , the number of clusters and consider the objects are defined with numeric attributes and thus using any one of the distance metric to demarcate the clusters.

# k-Means Algorithm

The algorithm can be stated as follows.

- First it selects  $k$  number of objects at random from the set of  $n$  objects. These  $k$  objects are treated as the **centroids or center of gravities** of  $k$  clusters.
- For each of the **remaining objects**, it is assigned to one of the **closest centroid**. Thus, it forms a **collection of objects assigned to each centroid** and is called a **cluster**.
- Next, the centroid of each cluster is then updated (by calculating the mean values of attributes of each object).
- The assignment and update procedure is until it reaches some stopping criteria (such as, number of iteration, centroids remain unchanged or no assignment, etc.)

# k-Means Algorithm

## Algorithm 16.1: k-Means clustering

**Input:**  $D$  is a dataset containing  $n$  objects,  $k$  is the number of cluster

**Output:** A set of  $k$  clusters

**Steps:**

1. Randomly choose  $k$  objects from  $D$  as the initial cluster centroids.
2. **For** each of the objects in  $D$  **do**
  - Compute distance between the current objects and  $k$  cluster centroids
  - Assign the current object to that cluster to which it is closest.
3. Compute the “cluster centers” of each cluster. These become the new cluster centroids.
4. Repeat step 2-3 until the convergence criterion is satisfied
5. Stop

# k-Means Algorithm

## Note:

- 1) Objects are defined in terms of set of attributes.  $A = \{A_1, A_2, \dots, A_m\}$  where each  $A_i$  is continuous data type.
- 2) **Distance computation:** Any distance such as  $L_1, L_2, L_3$  or cosine similarity.
- 3) **Minimum distance** is the measure of closeness between an object and centroid.
- 4) **Mean Calculation:** It is the mean value of each attribute values of all objects.
- 5) **Convergence criteria:** Any one of the following are termination condition of the algorithm.
  - Number of maximum iteration permissible.
  - No change of centroid values in any cluster.
  - Zero (or no significant) movement(s) of object from one cluster to another.
  - Cluster quality reaches to a certain level of acceptance.

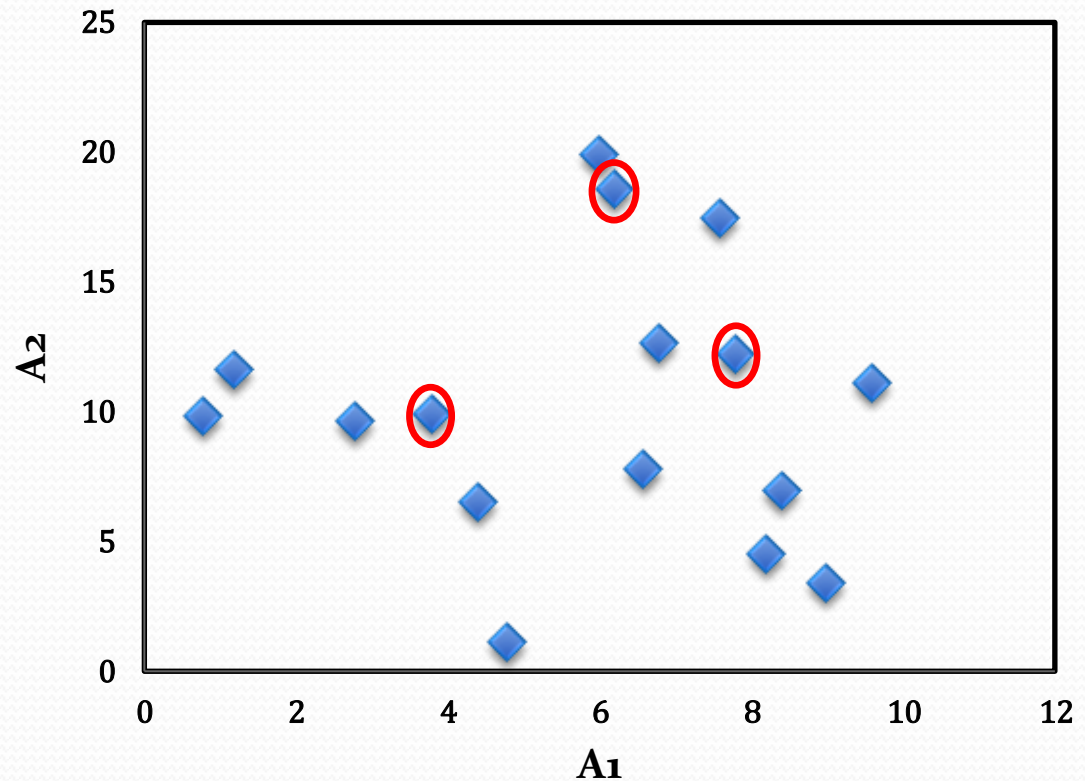


# Illustration of k-Means clustering algorithms

Table 16.1: 16 objects with two attributes  $A_1$  and  $A_2$ .

$A_1$	$A_2$
6.8	12.6
0.8	9.8
1.2	11.6
2.8	9.6
3.8	9.9
4.4	6.5
4.8	1.1
6.0	19.9
6.2	18.5
7.6	17.4
7.8	12.2
6.6	7.7
8.2	4.5
8.4	6.9
9.0	3.4
9.6	11.1

Fig 16.1: Plotting data of Table 16.1



# Illustration of k-Means clustering algorithms

- Suppose,  $k=3$ . Three objects are chosen at random shown as circled (see Fig 16.1). These three centroids are shown below.

Initial Centroids chosen randomly

Centroid	Objects	
	A1	A2
$c_1$	3.8	9.9
$c_2$	7.8	12.2
$c_3$	6.2	18.5

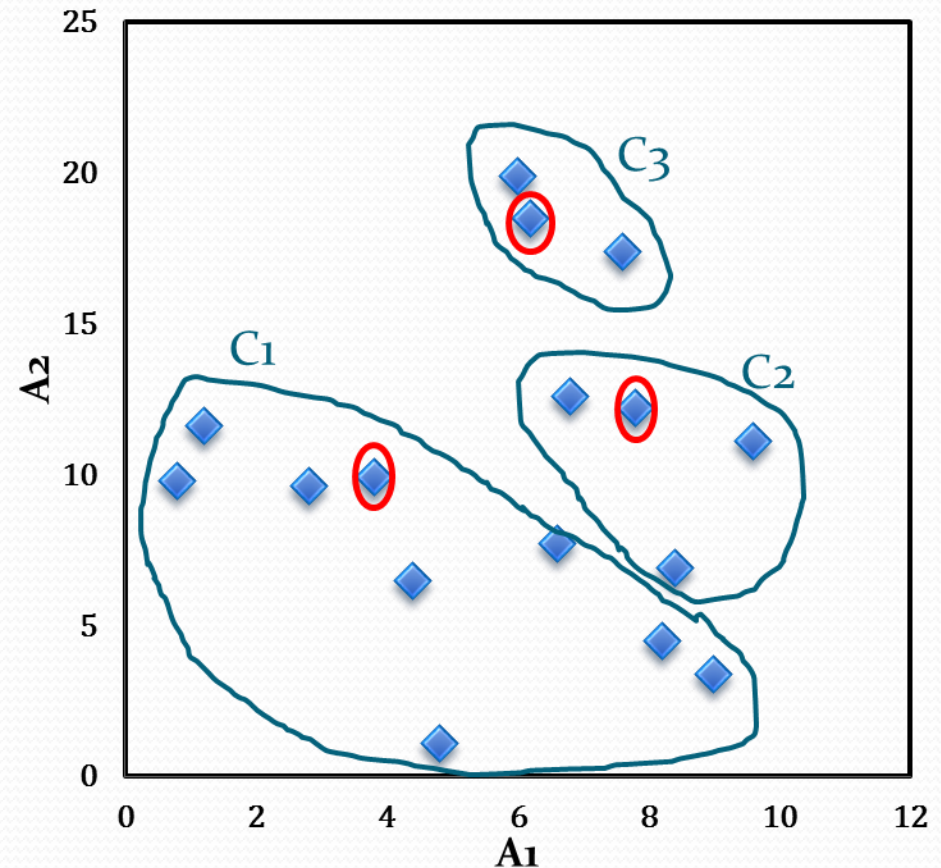
- Let us consider the Euclidean distance measure ( $L_2$  Norm) as the distance measurement in our illustration.
- Let  $d_1$ ,  $d_2$  and  $d_3$  denote the distance from an object to  $c_1$ ,  $c_2$  and  $c_3$  respectively. The distance calculations are shown in Table 16.2.
- Assignment of each object to the respective centroid is shown in the right-most column and the clustering so obtained is shown in Fig 16.2.

# Illustration of k-Means clustering algorithms

Table 16.2: Distance calculation

$A_1$	$A_2$	$d_1$	$d_2$	$d_3$	cluster
6.8	12.6	4.0	1.1	5.9	2
0.8	9.8	3.0	7.4	10.2	1
1.2	11.6	3.1	6.6	8.5	1
2.8	9.6	1.0	5.6	9.5	1
3.8	9.9	0.0	4.6	8.9	1
4.4	6.5	3.5	6.6	12.1	1
4.8	1.1	8.9	11.5	17.5	1
6.0	19.9	10.2	7.9	1.4	3
6.2	18.5	8.9	6.5	0.0	3
7.6	17.4	8.4	5.2	1.8	3
7.8	12.2	4.6	0.0	6.5	2
6.6	7.7	3.6	4.7	10.8	1
8.2	4.5	7.0	7.7	14.1	1
8.4	6.9	5.5	5.3	11.8	2
9.0	3.4	8.3	8.9	15.4	1
9.6	11.1	5.9	2.1	8.1	2

Fig 16.2: Initial cluster with respect to Table 16.2



# Illustration of k-Means clustering algorithms

The calculation new centroids of the three cluster using the mean of attribute values of  $A_1$  and  $A_2$  is shown in the Table below. The cluster with new centroids are shown in Fig 16.3.

## Calculation of new centroids

New Centroid	Objects	
	A1	A2
$c_1$	4.6	7.1
$c_2$	8.2	10.7
$c_3$	6.6	18.6

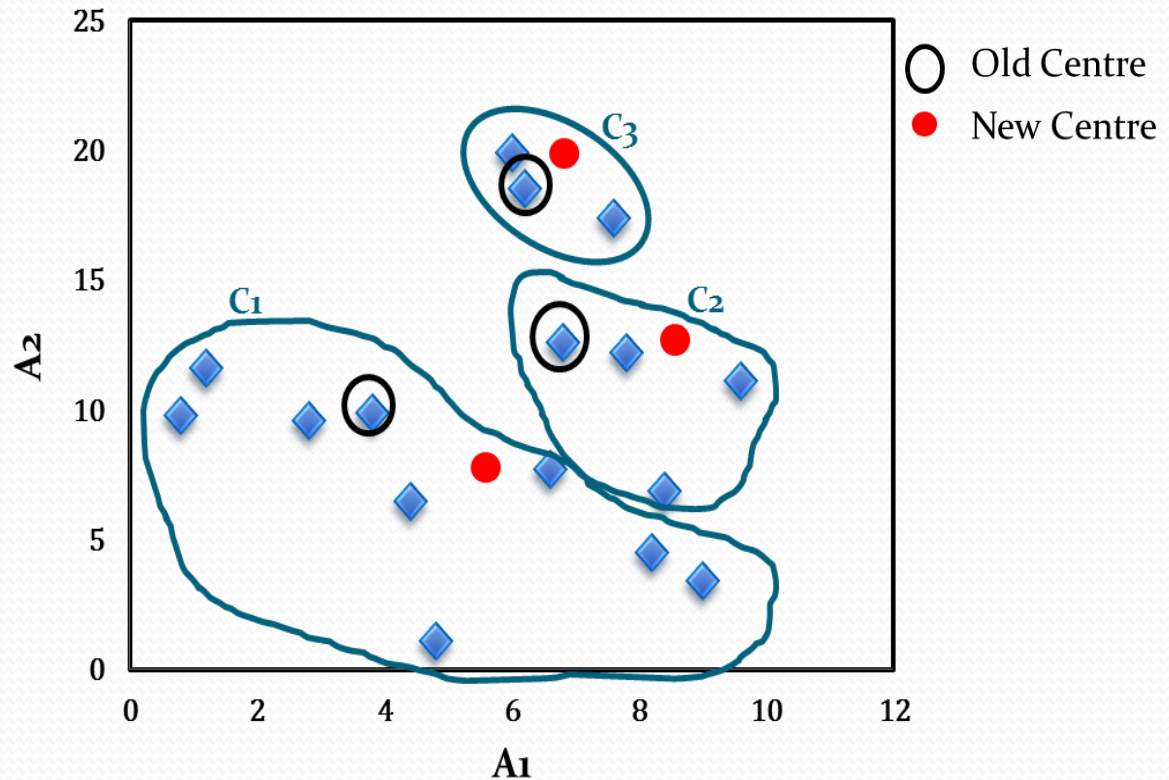


Fig 16.3: Initial cluster with new centroids

# Illustration of k-Means clustering algorithms

We next reassign the 16 objects to three clusters by determining which centroid is closest to each one. This gives the revised set of clusters shown in Fig 16.4.

Note that point  $p$  moves from cluster  $C_2$  to cluster  $C_1$ .

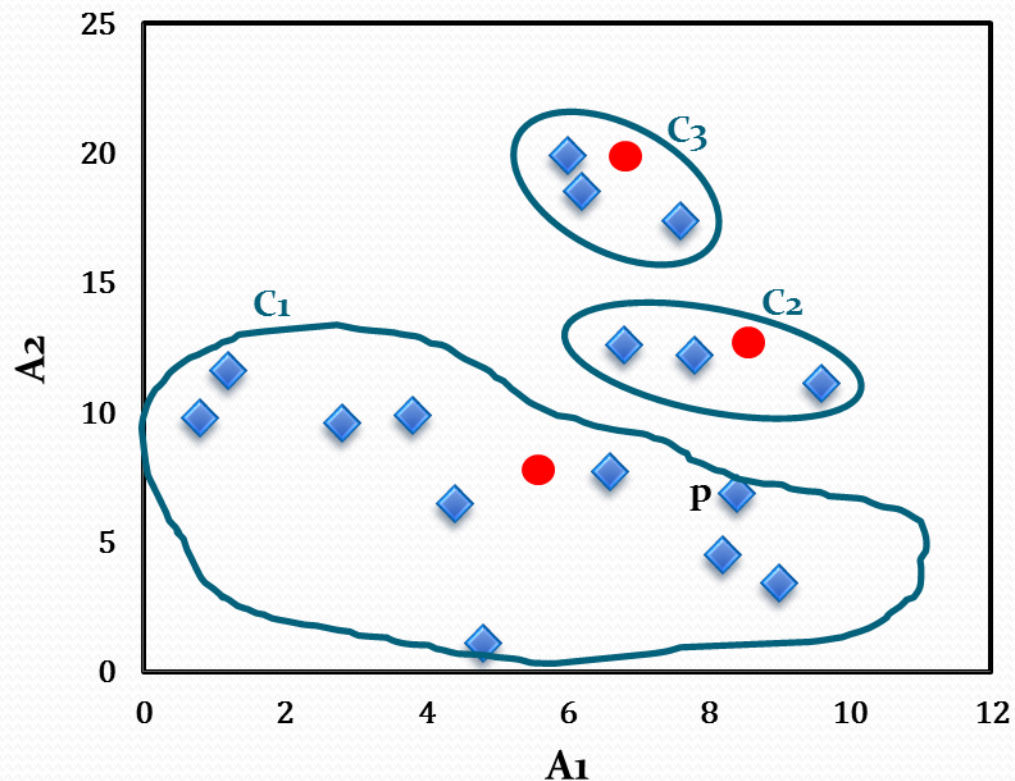


Fig 16.4: Cluster after first iteration

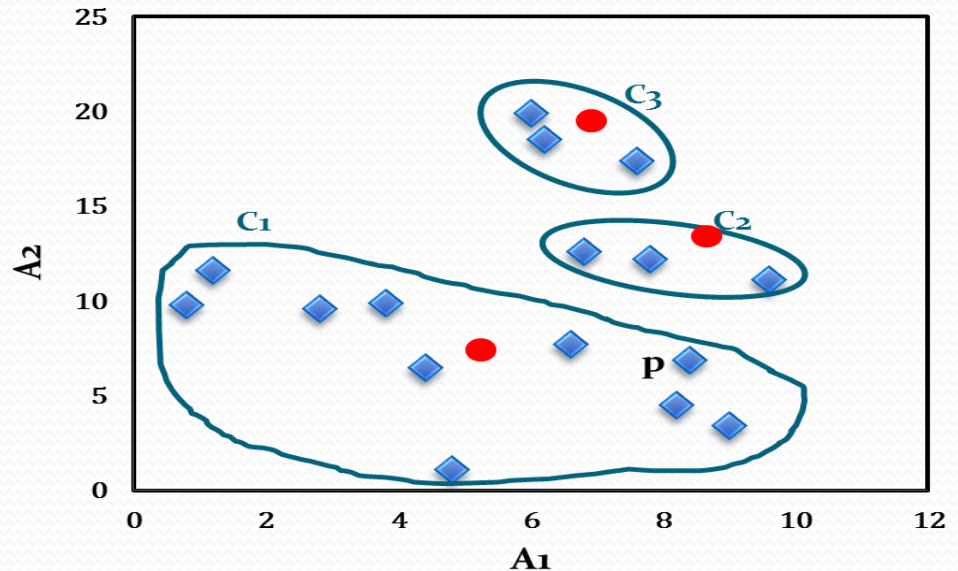
# Illustration of k-Means clustering algorithms

- The newly obtained centroids after second iteration are given in the table below. Note that the centroid  $c_3$  remains unchanged, where  $c_2$  and  $c_1$  changed a little.
- With respect to newly obtained cluster centres, 16 points are reassigned again. These are the same clusters as before. Hence, their centroids also remain unchanged.
- Considering this as the termination criteria, the k-means algorithm stops here. Hence, the final cluster in Fig 16.5 is same as Fig 16.4.

Cluster centres after second iteration

Centroid	Revised Centroids	
	A1	A2
$c_1$	5.0	7.1
$c_2$	8.1	12.0
$c_3$	6.6	18.6

Fig 16.5: Cluster after Second iteration



# Comments on k-Means algorithm

Let us analyse the k-Means algorithm and discuss the pros and cons of the algorithm. We shall refer to the following notations in our discussion.

- **Notations:**
  - $x$  : an object under clustering
  - $n$  : number of objects under clustering
  - $C_i$  : the  $i$ -th cluster
  - $c_i$  : the centroid of cluster  $C_i$
  - $n_i$  : number of objects in the cluster  $C_i$
  - $c$  : denotes the centroid of all objects
  - $k$  : number of clusters

# Comments on k-Means algorithm

## 1. Value of k:

- The k-means algorithm produces only one set of clusters, for which, user must specify the desired number,  $k$  of clusters.
- In fact,  $k$  should be the best guess on the number of clusters present in the given data. Choosing the best value of  $k$  for a given dataset is, therefore, an issue.
- We may not have an idea about the possible number of clusters for high dimensional data, and for data that are not scatter-plotted.
- Further, possible number of clusters is hidden or ambiguous in image, audio, video and multimedia clustering applications etc.
- There is no principled way to know what the value of  $k$  ought to be. We may try with successive value of  $k$  starting with 2.
- The process is stopped when two consecutive  $k$  values produce more-or-less identical results (with respect to some cluster quality estimation).
- Normally  $k \ll n$  and there is heuristic to follow  $k \approx \sqrt{n}$ .



# Comments on k-Means algorithm

## Example 16.1: k versus cluster quality

- Usually, there is some objective function to be met as a goal of clustering. One such objective function is **sum-square-error** denoted by **SSE** and defined as

$$SSE = \sum_{i=1}^k \sum_{x \in C_i} (x - c_i)^2$$

- Here,  $x - c_i$  denotes the error, if  $x$  is in cluster  $C_i$  with cluster centroid  $c_i$ .
- Usually, this error is measured as distance norms like  $L_1$ ,  $L_2$ ,  $L_3$  or Cosine similarity, etc.

# Comments on k-Means algorithm

## Example 16.1: k versus cluster quality

- With reference to an arbitrary experiment, suppose the following results are obtained.

k	SSE
1	62.8
2	12.3
3	9.4
4	9.3
5	9.2
6	9.1
7	9.05
8	9.0

- With respect to this observation, we can choose the value of  $k \approx 3$ , as with this smallest value of k it gives reasonably good result.
- Note: If  $k = n$ , then  $SSE=0$ ; However, the cluster is useless! This is another example of overfitting.

# Comments on k-Means algorithm

## 2. Choosing initial centroids:

- Another requirement in the k-Means algorithm to choose initial cluster centroid for each  $k$  would be clusters.
- It is observed that the k-Means algorithm terminate whatever be the initial choice of the cluster centroids.
- It is also observed that initial choice influences the ultimate cluster quality. In other words, the result may be trapped into local optima, if initial centroids are chosen properly.
- One technique that is usually followed to avoid the above problem is to choose initial centroids in multiple runs, each with a different set of randomly chosen initial centroids, and then select the best cluster (with respect to some quality measurement criterion, e.g. SSE).
- However, this strategy suffers from the combinational explosion problem due to the number of all possible solutions.

# Comments on k-Means algorithm

## 2. Choosing initial centroids:

- A detail calculation reveals that there are  $c(n, k)$  possible combinations to examine the search of global optima.

$$c(n, k) = \frac{1}{k!} \sum_{i=1}^k (-1)^{k-i} \binom{k}{i} (i)^n$$

- For example, there are  $o(10^{10})$  different ways to cluster 20 items into 4 clusters!
- Thus, the strategy having its own limitation is practical only if
  - 1) The sample is negatively small ( $\sim 100-1000$ ), and
  - 2)  $k$  is relatively small compared to  $n$  (i.e..  $k \ll n$ ).

# Comments on k-Means algorithm

## 3. Distance Measurement:

- To assign a point to the closest centroid, we need a proximity measure that should quantify the notion of “closest” for the objects under clustering.
- Usually Euclidean distance ( $L_2$  norm) is the best measure when object points are defined in n-dimensional Euclidean space.
- Other measure namely cosine similarity is more appropriate when objects are of document type.
- Further, there may be other type of proximity measures that appropriate in the context of applications.
- For example, Manhattan distance ( $L_1$  norm), Jaccard measure, etc.

# Comments on k-Means algorithm

## 3. Distance Measurement:

Thus, in the context of different measures, the **sum-of-squared error** (i.e., objective function/convergence criteria) of a clustering can be stated as under.

Data in Euclidean space ( $L_2$  norm):

$$SSE = \sum_{i=1}^k \sum_{x \in C_i} (c_i - x)^2$$

Data in Euclidean space ( $L_1$  norm):

The Manhattan distance ( $L_1$  norm) is used as a proximity measure, where the objective is to minimize the **sum-of-absolute error** denoted as **SAE** and defined as

$$SAE = \sum_{i=1}^k \sum_{x \in C_i} |c_i - x|$$

# Comments on k-Means algorithm

## Distance with document objects

Suppose a set of  $n$  document objects is defined as  $d$  document term matrix (DTM) (a typical look is shown in the below form).

Document	Term			
	$t_1$	$t_2$	... ..	$t_n$
$D_1$	$f_{11}$	$f_{12}$		$f_{1n}$
$D_2$	$f_{21}$	$f_{22}$		$f_{2n}$
$\vdots$				
$D_n$	$f_{n1}$	$f_{n2}$		$f_{nn}$

Here, the objective function, which is called Total cohesion denoted as TC and defined as

$$TC = \sum_{i=1}^k \sum_{x \in C_i} \cos(x, c_i)$$

where  $\cos(x, c_i) = \frac{x \cdot c_i}{\|x\| \|c_i\|}$

$$x \cdot c_i = \sum_j x_j c_{ij} \quad \text{and} \quad \|x\| = \sqrt{\sum_j^p x_j^2}$$

$$\hat{x} = \sum_{j=1}^p \hat{x}_j \quad \hat{c}_i = \sum_{j=1}^p \hat{c}_{ij} \quad \|\|c_{ij}\|\| = \sqrt{\sum_j^p c_{ij}^2}$$

# Comments on k-Means algorithm

Note: The criteria of objective function with different proximity measures

1. SSE (using  $L_2$  norm) : To **minimize** the SSE.
2. SAE (using  $L_1$  norm) : To **minimize** the SAE.
3. TC(using cosine similarity) : To **maximize** the TC.



# Comments on k-Means algorithm

## 4. Type of objects under clustering:

- The k-Means algorithm can be applied only when the mean of the cluster is defined (hence it named k-Means). The cluster mean (also called centroid) of a cluster  $C_i$  is defined as

$$c_i = \frac{1}{n_i} \sum_{x \in C_i} x$$

- In other words, the mean calculation assumed that each object is defined with numerical attribute(s). Thus, we cannot apply the k-Means to objects which are defined with categorical attributes.
- More precisely, the k-means algorithm require some definition of cluster mean exists, but not necessarily it does have as defined in the above equation.
- In fact, the k-Means is a very general clustering algorithm and can be used with a wide variety of data types, such as documents, time series, etc.



How to find the mean of objects with composite attributes?

# Comments on k-Means algorithm

## Note:

- 1) When SSE ( $L_2$  norm) is used as objective function and the objective is to minimize, then the cluster centroid (i.e. mean) is the mean value of the objects in the cluster.
- 2) When the objective function is defined as SAE ( $L_1$  norm), minimizing the objective function implies the cluster centroid as the median of the cluster.

The above two interpretations can be readily verified as given in the next slide.

# Comments on k-Means algorithm

## Case 1: SSE

We know,

$$SSE = \sum_{i=1}^k \sum_{x \in C_i} (c_i - x)^2$$

To minimize SSE means,  $\frac{\partial(SSE)}{\partial c_i} = 0$

Thus,

$$\frac{\partial}{\partial c_i} \left( \sum_{i=1}^k \sum_{x \in C_i} (c_i - x)^2 \right) = 0$$

Or,

$$\sum_{i=1}^k \sum_{x \in C_i} \frac{\partial}{\partial c_i} (c_i - x)^2 = 0$$

# Comments on k-Means algorithm

Or,

$$\sum_{x \in \mathcal{C}_i} 2(c_i - x) = 0$$

Or,

$$n_i \cdot c_i = \sum_{x \in \mathcal{C}_i} x$$

Or,

$$c_i = \frac{1}{n_i} \sum_{x \in \mathcal{C}_i} x$$

- Thus, **the best centroid for minimizing SSE of a cluster is the mean of the objects in the cluster.**

# Comments on k-Means algorithm

## Case 2: SAE

We know,

$$SAE = \sum_{i=1}^k \sum_{x \in C_i} |c_i - x|$$

To minimize SAE means,  $\frac{\partial(SAE)}{\partial c_i} = 0$

Thus,

$$\frac{\partial}{\partial c_i} \left( \sum_{i=1}^k \sum_{x \in C_i} |c_i - x| \right) = 0$$

Or,

$$\sum_{i=1}^k \sum_{x \in C_i} \frac{\partial}{\partial c_i} |c_i - x| = 0$$

# Comments on k-Means algorithm

Or,

$$\sum_{x \in C_i} \left\{ (x - c_i) \Big|_{\text{if } x > c_i} + (c_i - x) \Big|_{\text{if } c_i > x} \right\} = 0$$

Solving the above equation, we get

$$c_i = \text{median} \{x | x \in C_i\}$$

- Thus, the best centroid for minimizing SAE of a cluster is the median of the objects in the cluster.



Interpret the best centroid for maximizing TC (with Cosine similarity measure) of a cluster.

The above mentioned discussion is quite sufficient for the validation of k-Means algorithm.

# Comments on k-Means algorithm

## 5. Complexity analysis of k-Means algorithm

Let us analyse the time and space complexities of k-Means algorithm.

### Time complexity:

The time complexity of the k-Means algorithm can be expressed as

$$T(n) = O(n \times m \times k \times l)$$

where  $n$  = number of objects

$m$  = number of attributes in the object definition

$k$  = number of clusters

$l$  = number of iterations.

Thus, time requirement is a linear order of number of objects and the algorithm runs in a modest time if  $k \ll n$  and  $l \ll n$  (the iteration can be moderately controlled to check the value of  $l$ ).

# Comments on k-Means algorithm

## 5. Complexity analysis of k-Means algorithm

**Space complexity:** The storage complexity can be expressed as follows.

It requires  $n \times m$  space to store the objects and  $n \times k$  space to store the proximity measure from  $n$  objects to the centroids of  $k$  clusters.

Thus the total storage complexity is

$$S(n) = O(n \times (m + k))$$

That is, space requirement is in the linear order of  $n$  if  $k \ll n$ .



# Comments on k-Means algorithm

## 6. Final comments:

### Advantages:

- k-Means is simple and can be used for a wide variety of object types.
- It is also efficient both from storage requirement and execution time point of views. By saving distance information from one iteration to the next, the actual number of distance calculations, that must be made can be reduced (specially, as it reaches towards the termination).



How similarity metric can be utilized to run k-Means faster? What is the updation in each iteration?

### Limitations:

- The k-Means is not suitable for all types of data. For example, k-Means does not work on categorical data because mean cannot be defined.
- k-means finds a local optima and may actually minimize the global optimum.

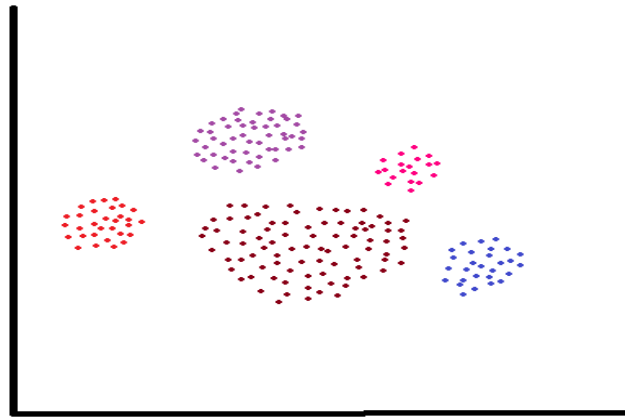
# Comments on k-Means algorithm

## 6. Final comments:

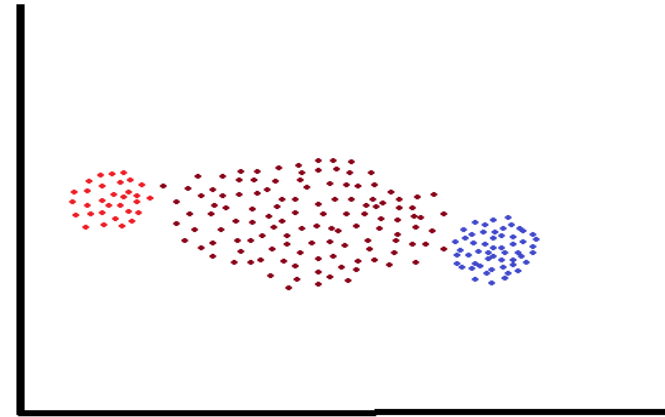
### Limitations :

- k-means has trouble clustering data that contains outliers. When the SSE is used as objective function, outliers can unduly influence the cluster that are produced. More precisely, in the presence of outliers, the cluster centroids, in fact, not truly as representative as they would be otherwise. It also influence the SSE measure as well.
- k-Means algorithm cannot handle non-globular clusters, clusters of different sizes and densities (see Fig 16.6 in the next slide).
- k-Means algorithm not really beyond the scalability issue (and not so practical for large databases).

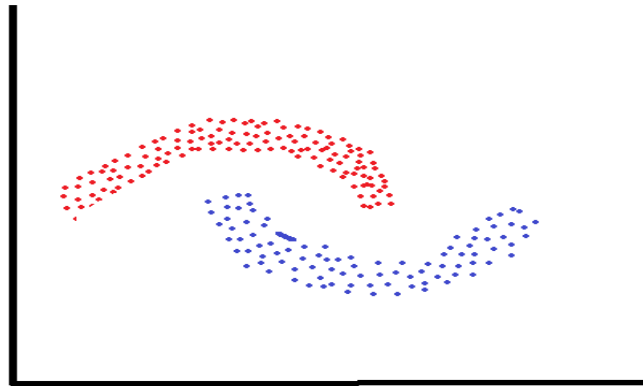
# Comments on k-Means algorithm



Cluster with different sizes



Cluster with different densities



Non-convex shaped clusters

**Fig 16.6: Some failure instance of k-Means algorithm**

# Different variants of k-means algorithm

There are a quite few variants of the k-Means algorithm. These can differ in the procedure of selecting the initial  $k$  means, the calculation of proximity and strategy for calculating cluster means. Another variants of k-means to cluster categorical data.

Few variant of k-Means algorithm includes

- Bisecting k-Means (addressing the issue of initial choice of cluster means).
  1. M. Steinbach, G. Karypis and V. Kumar “A comparison of document clustering techniques”, *Proceedings of KDD workshop on Text mining*, 2000.
- Mean of clusters (Proposing various strategies to define means and variants of means).
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# Different variants of k-means algorithm

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# The k-Medoids algorithm

Now, we shall study a variant of partitioning algorithm called k-Medoids algorithm.

**Motivation:** We have learnt that the k-Means algorithm is sensitive to outliers because an object with an “extremely large value” may substantially distort the distribution. The effect is particularly exacerbated due to the use of the SSE (sum-of-squared error) objective function. The k-Medoids algorithm aims to diminish the effect of outliers.

## Basic concepts:

- The basic concepts of this algorithm is to **select an object as a cluster center** (one representative object per cluster) instead of taking the mean value of the objects in a cluster (as in k-Means algorithm).
- We call this cluster representative as a **cluster medoid** or simply **medoid**.
  1. Initially, it selects a random set of  $k$  objects as the set of medoids.
  2. Then at each step, all objects from the set of objects, which are not currently medoids are examined one by one to see if they should be medoids.

# The k-Medoids algorithm

- That is, the k-Medoids algorithm determines whether there is an object that should replace one of the current medoids.
- This is accomplished by looking all pair of medoid, non-medoid objects, and then choosing a pair that improves the objective function of clustering the best and exchange them.
- The sum-of-absolute error (SAE) function is used as the objective function.

$$SAE = \sum_{i=1}^k \sum_{x \in C_i, x \notin M \text{ and } c_m \in M} |x - c_m|$$

Where  $c_m$  denotes a medoid

$M$  is the set of all medoids at any instant

$x$  is an object belongs to set of non-medoid object, that is,  $x$  belongs to some cluster and is not a medoid. i.e.  $x \in C_i, x \notin M$



# PAM (Partitioning around Medoids)

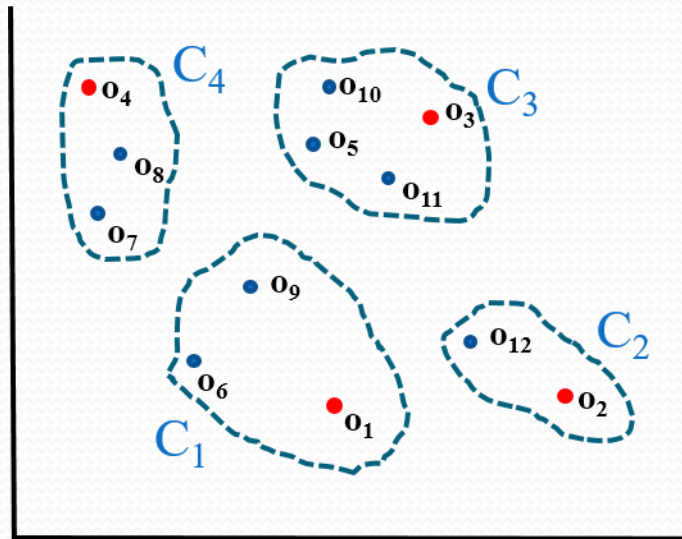
- For a given set of medoids, at any iteration, it select that exchange which has minimum SAE.
- The procedure terminates, if there is no any change in SAE in successive iteration (i.e. there is no change in medoid).
- This k-Medoids algorithm is also known as PAM (Partitioning around Medoids).

## Illustration of PAM

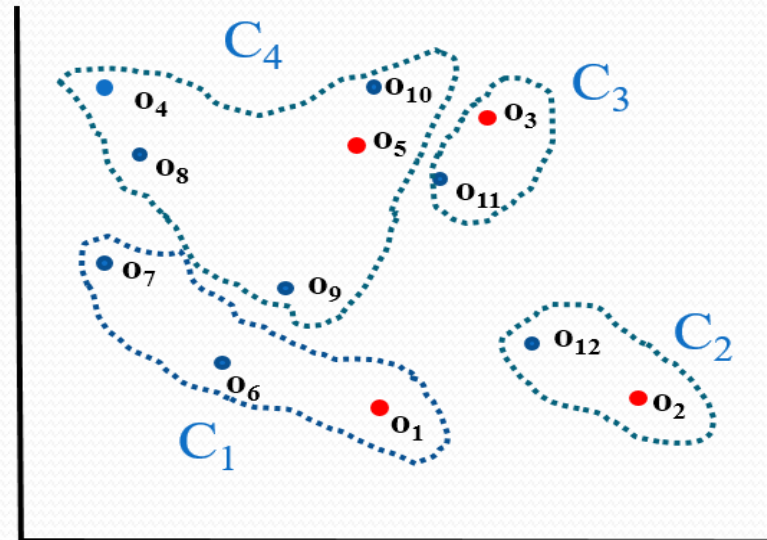
- Suppose, there are set of 12 objects  $O(o_1, o_2, \dots, o_{12})$  and we are to cluster them into four clusters. At any instant, the four clusters  $C_1, C_2, C_3$  and  $C_4$  are shown in Fig. 16.7 (a). Also assume that  $o_1, o_2, o_3$ , and  $o_4$  are the medoids in the clusters  $C_1, C_2, C_3$  and  $C_4$ , respectively. For this clustering we can calculate SAE.
- There are many ways to choose a non-medoid object to be replaced any one medoid object. Out of these, suppose, if  $o_5$  is considered as candidate medoid instead of  $o_4$ , then it gives the lowest SAE. Thus, the new set of medoids would be  $o_1, o_2, o_3$ , and  $o_5$ . The new cluster is shown in Fig 16.7 (b).



# PAM (Partitioning around Medoids)



(a) Cluster with  $o_1, o_2, o_3$ , and  $o_4$  as medoids



(b) Cluster after swapping  $o_4$  and  $o_5$  ( $o_5$  becomes the new medoid).

Fig 16.7: Illustration of PAM

# PAM (Partitioning around Medoids)

PAM algorithm is thus a procedure of iterative selection of medoids and it is precisely stated in Algorithm 16.2.

## Algorithm 16.2: PAM

**Input:** Database of objects D.

k, the number of desired clusters.

**Output:** Set of k clusters

**Steps:**

1. Arbitrarily select k medoids from D.
2. **For** each object  $o_i$  not a medoid **do**
3.     **For** each medoid  $o_j$  **do**
4.     Let  $M = \{o_1, o_2, \dots, o_{i-1}, o_i, o_{i+1}, o_k\}$      //Set of current medoids  
        $M' = \{o_1, o_2, \dots, o_{j-1}, o_j, o_{j+1}, o_k\}$      //set of medoids but swap with non-medoids  $o_j$
5.     Calculate  $cost(o_i, o_j) = SAE|_M - SAE_{M'}$
6.     **End** of 2 for loop

# PAM (Partitioning around Medoids)

## Algorithm 16.2: PAM

7. Find  $o_i, o_j$  for which the  $\text{cost}(o_i, o_j)$  is the smallest.
8. Replace  $o_i$  with  $o_j$  and accordingly update the set  $M$ .
9. Repeat step 2 - step 8 until  $\text{cost}(o_i, o_i) \leq 0$ .
10. Return the cluster with  $M$  as the set of cluster centers.
11. Stop

# Comments on PAM

## 1. Comparing k-Means with k-Medoids:

- Both algorithms need to fix  $k$ , the number of clusters prior to the algorithms. Also, both algorithms arbitrarily choose the initial cluster centroids.
- The k-Medoid method is more robust than k-Means in the presence of outliers, because a medoid is less influenced by outliers than a mean.

## 2. Time complexity of PAM:

- For each iteration, PAM considers  $k(n - k)$  pairs of objects  $o_i, o_j$  for which a cost  $cost(o_i, o_j)$  is determined. Calculating the cost during each iteration requires that the cost be calculated for all other non-medoids  $o_j$ . There are  $n - k$  of these. Thus, the total time complexity per iteration is  $n(n - k)^2$ . The total number of iterations may be quite large.

## 3. Applicability of PAM:

- PAM does not scale well to large databases because of its computational complexity.

# Other variants of k-Medoids algorithms

- There are some variants of PAM that are targeted mainly large datasets are CLARA (Clustering LARge Applications) and CLARANS (Clustering Large Applications based upon RANdomized Search), it is an improvement of CLARA.

## References:

For PAM and CLARA:

- L. kaufman and P. J. Rousseew, “Finding Groups in Data: An introduction to cluster analysis”, John and Wiley, 1990.

For CLARANS:

- R. Ng and J. Han, “Efficient and effective clustering method for spatial Data mining”, Proceeding very large databases [VLDB-94], 1994.



# Any question?

You may post your question(s) at the “Discussion Forum”  
maintained in the course Web page!