Module 5: Unsupervised Learning

Supervised learning vs. unsupervised learning

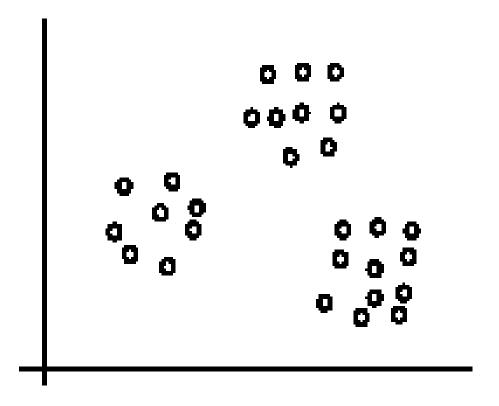
- Supervised learning: discover patterns in the data that relate data attributes with a target (class) attribute.
 - These patterns are then utilized to predict the values of the target attribute in future data instances.
- Unsupervised learning: The data have no target attribute.
 - We want to explore the data to find some intrinsic structures in them.

Clustering

- Clustering is a technique for finding **similarity groups** in data, called **clusters**. i.e.,
 - it groups data instances that are similar to (near) each other in one cluster and data instances that are very different (far away) from each other into different clusters.
- Clustering is often called an **unsupervised learning** task as no class values denoting an *a priori* grouping of the data instances are given, which is the case in supervised learning.
- Due to historical reasons, clustering is often considered synonymous with unsupervised learning.
 - In fact, association rule mining is also unsupervised

An illustration

• The data set has three natural groups of data points, i.e., 3 natural clusters.



What is clustering for?

- Let us see some real-life examples
- Example 1: groups people of similar sizes together to make "small", "medium" and "large" T-Shirts.
 - **Tailor-made for each person: too expensive**
 - One-size-fits-all: does not fit all.
- Example 2: In marketing, segment customers according to their similarities
 - To do targeted marketing.

What is clustering for? (cont...)

- Example 3: Given a collection of text documents, we want to organize them according to their content similarities,
 - To produce a topic hierarchy
- In fact, clustering is one of the most utilized data mining techniques.
 - It has a long history, and used in almost every field, e.g., medicine, psychology, botany, sociology, biology, archeology, marketing, insurance, libraries, etc.
 - In recent years, due to the rapid increase of online documents, text clustering becomes important.

Types of Clustering

- Hierarchical algorithms: these find successive clusters using previously established clusters.
 - 1. Agglomerative ("bottom-up"): Agglomerative algorithms begin with each element as a separate cluster and merge them into successively larger clusters.
 - 2. Divisive ("top-down"): Divisive algorithms begin with the whole set and proceed to divide it into successively smaller clusters.
- Partitional clustering: Partitional algorithms determine all clusters at once. They include:
 - K-means and derivatives
 - Fuzzy c-means clustering
 - QT clustering algorithm

Aspects of clustering

- ♠ A distance (similarity, or dissimilarity) function
- Clustering quality
 - **№** Inter-clusters distance ⇒ maximized
 - Intra-clusters distance ⇒ minimized
- The quality of a clustering result depends on the algorithm, the distance function, and the application.

K-means clustering

- K-means is a partitional clustering algorithm
- Let the set of data points (or instances) D be

$$\{X_1, X_2, ..., X_n\},\$$

- where $x_i = (x_{i1}, x_{i2}, ..., x_{ir})$ is a vector in a real-valued space $X \in \mathbb{R}^r$, and r is the number of attributes (dimensions) in the data.
- The k-means algorithm partitions the given data into k clusters.
 - Each cluster has a cluster center, called centroid.
 - k is specified by the user

K-means algorithm

- Given *k*, the *k-means* algorithm works as follows:
 - 1. Randomly choose *k* data points (seeds) to be the initial centroids, cluster centers
 - 2. Assign each data point to the closest centroid
 - 3. Re-compute the centroids using the current cluster memberships.
 - 4. If a convergence criterion is not met, go to 2).

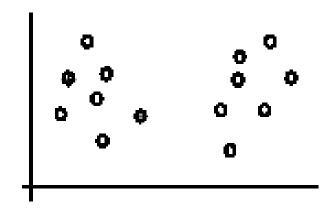
Stopping/convergence criterion

- no (or minimum) re-assignments of data points to different clusters,
- 2. no (or minimum) change of centroids, or
- minimum decrease in the sum of squared error (SSE), k

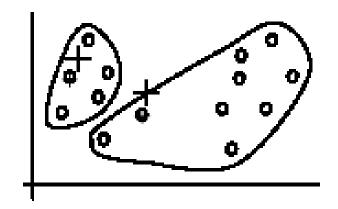
$$SSE = \sum_{j=1}^{\infty} \sum_{j=1}^{\infty} dist(x, m_j)^2$$
 (1)

 C_i is the jth cluster, \mathbf{m}_j is the centroid of cluster C_j (the mean vector of all the data points in C_j), and $dist(\mathbf{x}, \mathbf{m}_j)$ is the distance between data point \mathbf{x} and centroid \mathbf{m}_j .

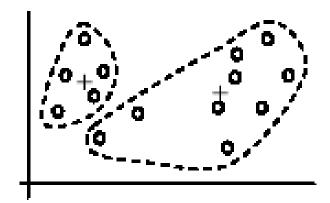
An example



(A). Random selection of k centers

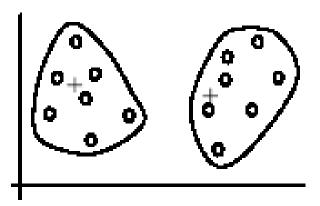


Iteration 1: (B). Cluster assignment

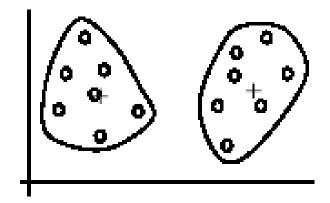


(C). Re-compute centroids

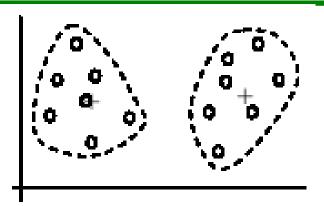
An example (cont ...)



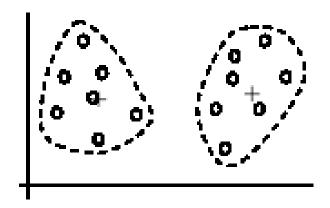
Iteration 2: (D). Cluster assignment



Iteration 3: (F). Cluster assignment



(E). Re-compute centroids



(G). Re-compute centroids

Numerical Example : K- Means Clustering (K = 2)

Individual	Variable 1	Variable 2
1	1.0	1.0
2	1.5	2.0
3	3.0	4.0
4	5.0	7.0
5	3.5	5.0
6	4.5	5.0
7	3.5	4.5

Step 1:

<u>Initialization</u>: Randomly we choose following two centroids (k=2) for two clusters.

In this case the 2 centroid are: m1=(1.0,1.0) and m2=(5.0,7.0).

Individual	Variable 1	Variable 2
1	1.0	1.0
2	1.5	2.0
3	3.0	4.0
4	5.0	7.0
5	3.5	5.0
6	4.5	5.0
7	3.5	4.5

Individual	Mean Vector
1	(1.0, 1.0)
4	(5.0, 7.0)
	Individual 1 4

Step 2:

Thus, we obtain two clusters containing:

 $\{1,2,3\}$ and $\{4,5,6,7\}$.

Their new centroids are:

$$m_1 = (\frac{1}{3}(1.0 + 1.5 + 3.0), \frac{1}{3}(1.0 + 2.0 + 4.0)) = (1.83, 2.33)$$

$$m_2 = (\frac{1}{4}(5.0 + 3.5 + 4.5 + 3.5), \frac{1}{4}(7.0 + 5.0 + 5.0 + 4.5))$$

= (4.12,5.38)

Individual	Centrold 1	Centrold 2
1	0	7.21
2 (1.5, 2.0)	1.12	6.10
3	3.61	3.61
4	7.21	0
5	4.72	2.5
6	5.31	2.06
7	4.30	2.92

$$d(m_1, 2) = \sqrt{|1.0 - 1.5|^2 + |1.0 - 2.0|^2} = 1.12$$

 $d(m_2, 2) = \sqrt{|5.0 - 1.5|^2 + |7.0 - 2.0|^2} = 6.10$

Step 3:

Now using these centroids we compute the Euclidean distance of each object, as shown in table.

Therefore, the new clusters are:

{1,2} and {**3**,4,5,6,7}

Next centroids are: m1=(1.25,1.5) and m2=(3.9,5.1)

Individual	Centroid 1	Centroid 2
1	1.57	5.38
2	0.47	4.28
3	2.04	1.78
4	5.64	1.84
5	3.15	0.73
6	3.78	0.54
7	2.74	1.08

<u>Step 4</u>:

The clusters obtained

are:

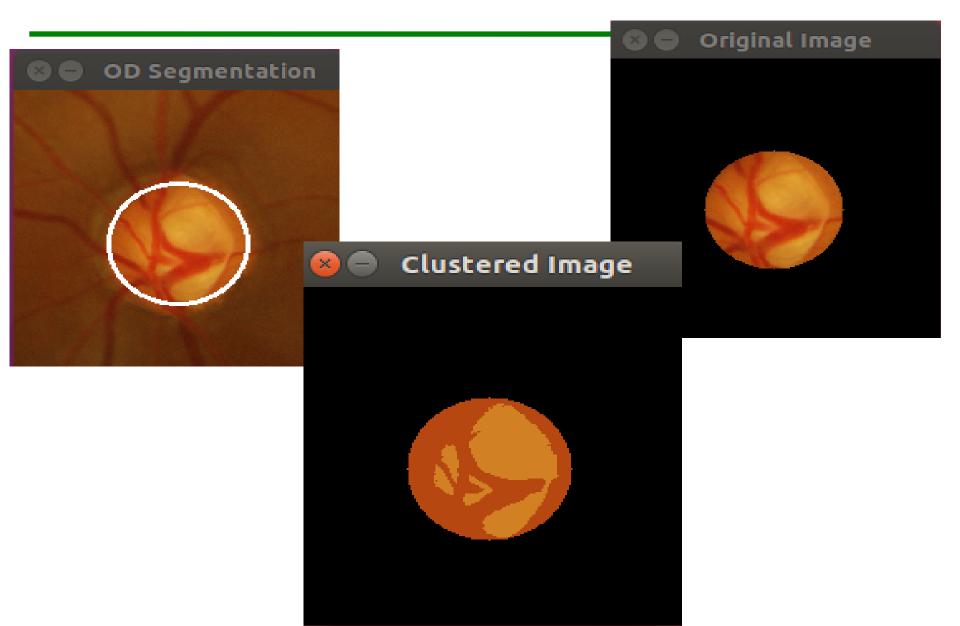
{1,2} and {3,4,5,6,7}

Therefore, there is no change in the cluster.

Thus, the algorithm comes to a halt here and final result consist of 2 clusters {1,2} and {3,4,5,6,7}.

Individual	Centroid 1	Centroid 2
1	0.56	5.02
2	0.56	3.92
3	3.05	1.42
4	6.66	2.20
5	4.16	0.41
6	4.78	0.61
7	3.75	0.72

Example



Strengths of k-means

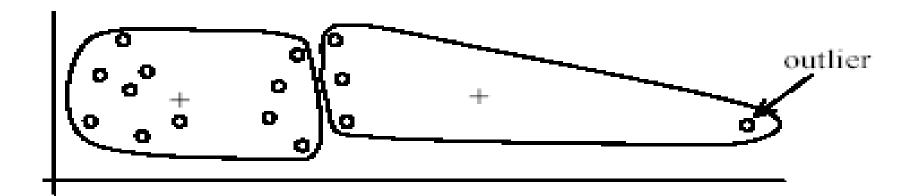
Strengths:

- Simple: easy to understand and to implement
- ❖ Efficient: Time complexity: *O*(*tkn*),
 where *n* is the number of data points, *k* is the number of clusters, and *t* is the number of iterations.
- **♥** Since both *k* and *t* are small. *k*-means is considered a linear algorithm.
- K-means is the most popular clustering algorithm.
- Note that: it terminates at a local optimum if SSE is used. The global optimum is hard to find due to complexity.

Weaknesses of k-means

- The algorithm is only applicable if the mean is defined.
 - For categorical data, *k*-mode the centroid is represented by most frequent values.
- The user needs to specify k.
- The algorithm is sensitive to outliers
 - Outliers are data points that are very far away from other data points.
 - Outliers could be errors in the data recording or some special data points with very different values.

Weaknesses of k-means: Problems with outliers



(A): Undesirable clusters



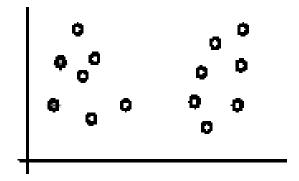
(B): Ideal clusters

Weaknesses of k-means: To deal with outliers

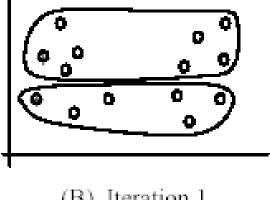
- One method is to remove some data points in the clustering process that are much further away from the centroids than other data points.
 - To be safe, we may want to monitor these possible outliers over a few iterations and then decide to remove them.
- Another method is to perform random sampling. Since in sampling we only choose a small subset of the data points, the chance of selecting an outlier is very small.
 - Assign the rest of the data points to the clusters by distance or similarity comparison, or classification

Weaknesses of k-means (cont ...)

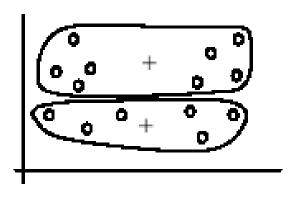
The algorithm is sensitive to initial seeds.



(A). Random selection of seeds (centroids)



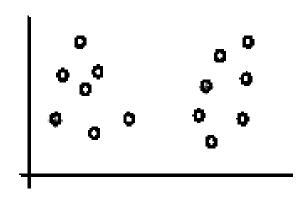
(B). Iteration 1



(C). Iteration 2

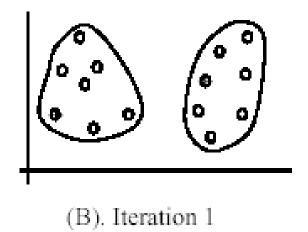
Weaknesses of k-means (cont ...)

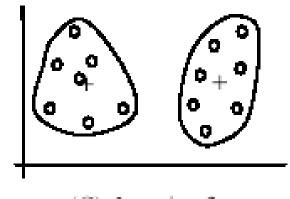
If we use different seeds: good results



There are some methods to help choose good seeds

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





(C). Iteration 2

K-means summary

- Despite weaknesses, k-means is still the most popular algorithm due to its simplicity, efficiency and
 - other clustering algorithms have their own lists of weaknesses.
- No clear evidence that any other clustering algorithm performs better in general
 - although they may be more suitable for some specific types of data or applications.
- Comparing different clustering algorithms is a difficult task. No one knows the correct clusters!