

# Data Science and Machine Learning 2187 & 2087: Unsupervised Learning

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### Goals of this lecture



- Understand the difference between unsupervised- and supervised learning
- Understand the definition of hard clustering
- Understand clustering cost from different similarity measures
- Understand the K-means algorithm
- Understand the K-medeoids algorithm

## Some notation



Feature vectors x, labels y

$$x \in \mathbb{R}^d$$
$$y \in \{-1, 1\}$$

Training set

$$S_n = \{(x^{(i)}, y^{(i)}), i = 1, ..., n\}$$

Classifier

$$h: \mathbb{R}^d \to \{-1, 1\}$$

## Supervised Learning vs. Unsupervised Learning



- ▶ In supervised learning we have labeled data:  $S_n = \{(x^{(i)}, y^{(i)}), i = 1, ..., n\}$  and want to learn to correctly classify unseen data
  - Think of:
  - A gazillion of photos with a "cat" and "not cat" classification.
  - etc.
- ▶ In clustering we only have feature vectors:  $S_n = \{x^{(i)} | i = 1, \dots, n\}$  and want to find structures in unlabeled data
  - ► Think of:
  - Clustering a data set of customer into groups
  - Find spatial patterns, e.g. crime hotspots
  - Find similar news stories
  - Recommend products to customers "like you"
  - Create labels for supervised learning algorithms
  - Exploratory data analysis
- Types
  - Hard clustering
  - Soft clustering
  - ► Hierarchical clustering

## Clustering as Partitioning



Clustering input:  $S_n = \{x^{(i)} | i = 1, \dots, n\}, K$ 

Number of clusters: K

The output of the clustering algorithm are indexes that partition the data:  $C_1, \dots, C_k$ ; where  $C_1 \cup C_2 \cup \dots \cup C_K = \{1, 2, \dots, n\}$  and  $C_i \cap C_i = \emptyset$  {for any  $i \neq j$  in  $\{1, \dots, k\}$ .

In other words: the union of all  $C_j$  's is the original set and the intersection of any  $C_i$  and  $C_j$  is an empty set.

### In plain English:

We want to assign each element of the training data set  $S_n$  into K separate clusters in a way that each element only belongs to one cluster.

## Clustering as selecting representatives



Clustering input: 
$$S_n = \{x^{(i)} | n = 1, \dots, n\}, K$$

Number of clusters: *K* 

Select the best representatives of each cluster:  $z^{(1)}, \dots, z^{(k)}$ .

# Similarity Measures-Cost functions



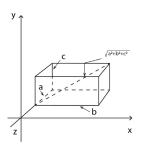
- Cost of partitioning
  - Sum of costs of all individual clusters:  $cost(C_1, \dots, C_k) \sum_{i=1}^k cost(C_i)$ .
- Cost of a single cluster
  - Sum of distances from data points to the representative of the cluster:  $Cost(C, z) = \sum_{i \in C} distance(x^{(i)}, z)$
- Total Cost to be minimized
  - ► Cost $(C_1, ..., C_K) = \sum_{j=1}^K \text{Cost}(C_j) = \sum_{j=1}^K \sum_{i \in C_j} ||x_i z_j||^2$

#### Two common distance measures

- ► Cosine similarity:  $cos(x^{(i)}, x^{(j)}) = \frac{x^{(i)} \cdot x^{(j)}}{||x^{(i)}||||x^{(j)}||}$ 
  - Is not sensitive of magnitude of vector (will not react to length).
- ▶ Euclidean squared distance:  $dist(x^{(i)}, x^{(j)}) = ||x^{(i)} x^{(j)}||^2$ .
  - ▶ Will react to length of the vectors



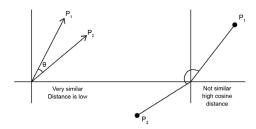




- ➤ The Euclidean distance between any two points is the square root of the sum of squares of differences between the coordinates. Straight line distance between any two points (pythagorean theorem)
- ▶ In two dimensions:  $dist(p,q) = \sqrt{(p_1 q_1)^2 + (p_2 q_2)^2}$ .
- Squared Euclidean distance is the sum of squares:  $dist^2(p,q) = (p_1 q_1)^2 + (p_2 q_2)^2$
- Generalizes to n-dimensions but looses meaning in very high dimensional data

## Cosine distance and Cosine similarity

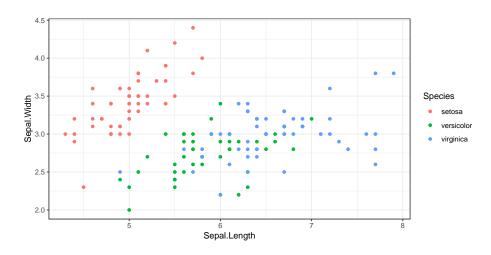




- Cosine similarity between any two points is defined as the cosine of the angle between any two points with the origin as its vertex.
- Cosine distance is defined as: 1 cosine similarity
- ► Cosine distance varies from 0 to 2, whereas cosine similarity varies between -1 to 1.

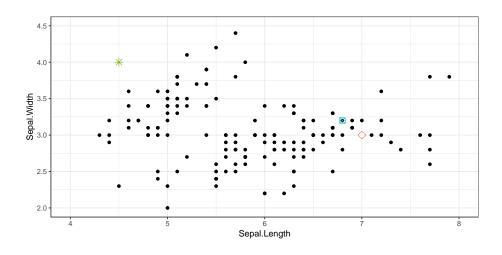
## Iris Dataset





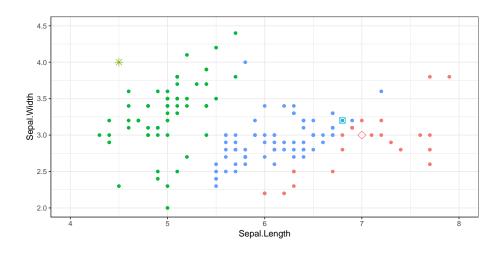
# Assign random initialization points (representatives $z_j$ )





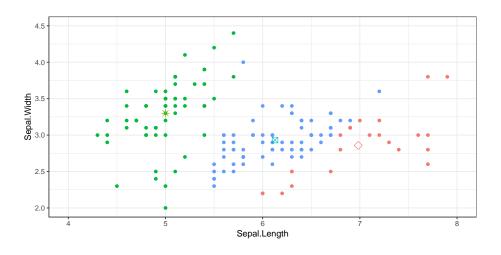






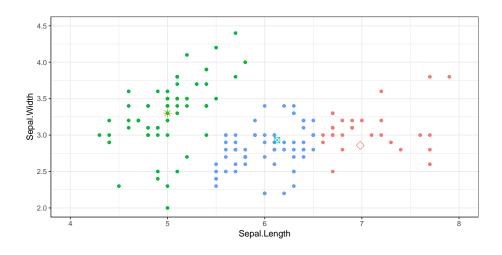
# Recalculate representatives $z_j$ as centroids of the new cluster (1st round)





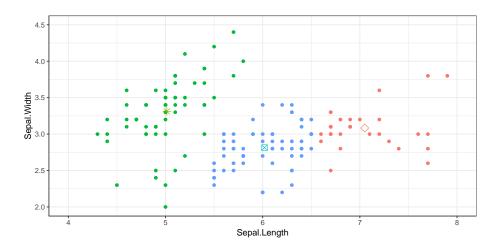






# Recalculate representatives $z_j$ as centroids of the new cluster (2nd round)





## K-means Algorithm



Given a set of feature vectors  $S_n = \{x^{(i)} | i = 1, ..., n\}$  and the number of clusters K we can find cluster assignments  $C_1, \dots, C_K$  and the representatives of each of the K clusters  $z_1, \dots, z_K$ :

- 1. Randomly select  $z_1, \dots, z_K$
- 2. Iterate until no change in cost
  - 2.1 Given  $z_1, \dots, z_K$ , assign each data point  $x^{(i)}$  to the closest  $z_j$ , such that  $Cost(z_1, \dots z_K) = \sum_{i=1}^n \min_{j=1,\dots,K} ||x^{(i)} z_j||^2$
  - 2.2 Given  $C_1, \dots, C_K$  find the best representatives  $z_1, \dots, z_K$ , i.e. find  $z_1, \dots, z_K$  such that  $z_j = \operatorname{argmin}_z \sum_{i \in C_j} \|x^{(i)} z\|^2$

## Minimizing the cost in K-means



K-means only works with Euclidean square distance!

The best representative is found by optimization (gradient with respect to  $z^{(j)}$ , setting to zero and solving for  $z^{(j)}$ ).

$$\nabla_{z_j} \left( \sum_{i \in \mathbb{C}_j} \|x^{(i)} - z_j\|^2 \right) = 0$$

$$\sum_{i\in\mathbb{C}_j}-2(x^{(i)}-z_j)=0$$

$$z^{(j)} = \frac{\sum_{i \in C_j} x^{(i)}}{|C_j|}$$

It is the centroid of the cluster, where  $C_j$  is the size of the respective cluster.

The clustering output that the K-Means algorithm converges to depends on the intialization! Suboptimal initializations are possible.

## K-means in pseudo code for two clusters



- 1. Choose any two random coordinates,  $z_1$  and  $z_2$ , on the scatter plot as initial cluster centers.
- 2. Calculate the distance of each data point in the scatter plot from coordinates  $z_1$  and  $z_2$
- 3. Assign each data point to a cluster based on whether it is closer to  $z_1$  or  $z_2$
- 4. Find the mean coordinates of all points in each cluster and update the values of  $z_1$  and  $z_2$  to those coordinates respectively.
- 5. Start again from Step 2 until the coordinates of  $z_1$  and  $z_2$  stop moving significantly, or after a certain pre-determined number of iterations of the process.

## Limitations of the K-Means Algorithm



- ▶ Algorithm is only guaranteed to converge to local minimum
- Initialization matters
  - ▶ Bad initalization can lead to suboptimal clusters in pathological cases
- Unlclear how many cluster we should plug into the algo (more about that next lecture)
- Only works with eucledian distance

### K-Medoids



Any distance measure possible!

Gives actual data points as representatives.

Finds the cost-minimizing representatives  $z_1, \dots, z_K$  for any distance measure. Uses real data points for initialization.

- 1. Randomly select  $\{z_1, ..., z_K\} \subseteq \{x_1, ..., x_n\}$
- 2. Iterate until no change in cost
  - 2.1 Given  $z_1, \dots, z_K$ , assign each data point  $x^{(i)}$  to the closest  $z_j$ , so that  $Cost(z_1, \dots z_K) = \sum_{i=1}^n \min_{j=1,\dots,K} ||x^{(i)} z_j||^2$
  - 2.2 Given  $C_j \in \{C_1, ..., C_K\}$  find the best representative  $z_j \in \{x_1, ..., x_n\}$  such that  $\sum_{x^{(i)} \in C_j} \operatorname{dist}(x^{(i)}, z_j)$  is minimal

## K-Mediods pseudocode



- 1. Choose k data points from the scatter plot as starting points for cluster centers.
- 2. Calculate their distance from all the points in the scatter plot.
- 3. Classify each point into the cluster whose center it is closest to.
- 4. Select a new point in each cluster that minimizes the sum of distances of all points in that cluster from itself.
- 5. Repeat Step 2 until the centers stop changing.

## Literature

