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### Midterm Admin

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### **Unsupervised Learning**

#### Motivation

- Supervised learning:  $(x_1, y_1), (x_2, y_2), ..., (x_n, y_n)$ .
- Unsupervised learning:  $x_1, x_2, ..., x_n$ .
- There are a few common tasks without labels.
- Olustering: separate instances into groups.
- Novelty (outlier) detection: find instances that are different.
- Dimensionality reduction: represent each instance with a lower dimensional feature vector while maintaining key characteristics.

### Unsupervised Learning Applications Motivation

- Google News
- Google Photo
- Image Segmentation
- Text Processing

### Hierarchical Clustering Description

- Start with each instance as a cluster.
- Merge clusters that are closest to each other.
- Result in a binary tree with close clusters as children.

### Hierarchical Clustering Diagram Description

### Single Linkage Distance

 Usually, the distance between two clusters is measured by the single-linkage distance.

$$d(C_k, C_{k'}) = \min \{d(x_i, x_{i'}) : x_i \in C_k, x_{i'} \in C_{k'}\}$$

• It is the shortest distance from any instance in one cluster to any instance in the other cluster.

### Complete Linkage Distance

• Another measure is complete-linkage distance,

$$d(C_k, C_{k'}) = \max\{d(x_i, x_{i'}) : x_i \in C_k, x_{i'} \in C_{k'}\}$$

• It is the longest distance from any instance in one cluster to any instance in the other cluster.

### Average Linkage Distance Diagram

• Another measure is average-linkage distance.

$$d(C_k, C_{k'}) = \frac{1}{|C_k| |C_{k'}|} \sum_{x_i \in C_k, x_{i'} \in C_{k'}} d(x_i, x_{i'})$$

• It is the average distance from any instance in one cluster to any instance in the other cluster.

## Hierarchical Clustering 4, Diagram

#### Discussion

- K can be chosen using prior knowledge about X.
- The algorithm can stop merging as soon as all the between-cluster distances are larger than some fixed R.
- The binary tree generated in the process is often called dendrogram, or taxonomy, or a hierarchy of data points.
- An example of a dendrogram is the tree of life in biology.

### K Means Clustering Description

- This is not K Nearest Neighbor.
- Start with random cluster centers.
- Assign each point to its closest center.
- Update all cluster centers as the center of its points.

### Distortion Distortion

- Distortion for a point is the distance from the point to its cluster center.
- Total distortion is the sum of distortion for all points.

$$D_{K} = \sum_{i=1}^{n} d\left(x_{i}, c_{k^{*}(x_{i})}\left(x_{i}\right)\right)$$
$$k^{*}\left(x\right) = \operatorname*{argmin}_{k=1,2,...K} d\left(x, c_{k}\right)$$

### Objective Function Counterexample Definition

### **Gradient Descent**

### Definition

 When d is the Euclidean distance. K Means algorithm is the gradient descent when distortion is the objective (cost) function.

$$\frac{\partial}{\partial c_k} \sum_{k=1}^K \sum_{x \in C_k} \|x - c_k\|_2^2 = 0$$

$$\Rightarrow -2 \sum_{x \in C_k} (x - c_k) = 0$$

$$\Rightarrow c_k = \frac{1}{|C_k|} \sum_{x \in C_k} x$$

# K Means Clustering 2

### Number of Clusters

#### Discussion

- There are a few ways to pick the number of clusters K.
- ullet K can be chosen using prior knowledge about X.
- ② K can be the one that minimizes distortion? No, when K = n, distortion = 0.
- ullet K can be the one that minimizes distortion + regularizer.

$$K^{\star} = \underset{k}{\operatorname{argmin}} (D_k + \lambda \cdot m \cdot k \cdot \log n)$$

ullet  $\lambda$  is a fixed constant chosen arbitrarily.

### **Initial Clusters**

#### Discussion

- There are a few ways to initialize the clusters.
- **1** *K* uniform random points in  $\{x_i\}_{i=1}^n$ .
- ② 1 uniform random point in  $\{x_i\}_{i=1}^n$  as  $c_1^{(0)}$ , then find the farthest point in  $\{x_i\}_{i=1}^n$  from  $c_1^{(0)}$  as  $c_2^{(0)}$ , and find the farthest point in  $\{x_i\}_{i=1}^n$  from the closer of  $c_1^{(0)}$  and  $c_2^{(0)}$  as  $c_3^{(0)}$ , and repeat this K times.

### Gaussian Mixture Model

#### Discussion

- In K means, each instance belong to one cluster with certainty.
- One continuous version is called the Gaussian mixture model: each instance belongs to one of the clusters with a positive probability.
- The model can be trained using Expectation Maximization Algorithm (EM Algorithm).

# Gaussian Mixture Model Demo