# Machine Learning

**CSE 142** 

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Friday, November 12, 2021

Clustering (k-means, k-medoids)

# Guest Lecture next Monday

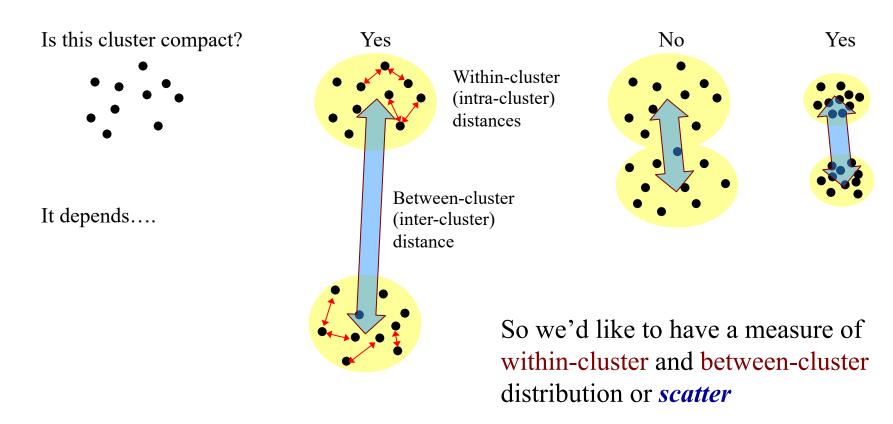
- Predicting behavior of road users, by Vihan Jain
- Bio: Vihan Jain is a Staff Software Engineer at Waymo in Mountain View where he is tech-leading a team which develops and deploys Machine Learned models that predict the behavior of road users as the autonomous driving vehicle navigates through a scene. Previously, he was at Google Research where he worked on multi-modal learning, natural language compositionality and video semantic understanding. He has also worked on long-term value modeling in recommendation systems, designing configurable simulation platforms for studying recommendations, wide and deep learning and TensorFlow infrastructure. Prior to moving to the US, he worked with Ads Infrastructure at Google Canada and did an internship at Google India during the summer of 2012. He graduated from Indian Institute of Technology Roorkee as a gold medalist in 2013. In his free time, he likes to travel and play/follow several sports.

# Clustering vs. classification

- Classification vs. clustering
  - In a classifier, possible class labels are provided
    - { dog, cat, elephant, mouse, ...}, { spam, ham }, etc.
    - Given in the training data (for supervised classification)
  - In a clustering problem, possible labels are the cluster labels learned from the training set
    - { cluster 1, cluster 2, cluster 3, ...}
    - Not given in the training data
- Terminology: In both cases, people often refer to the assigning of labels or clusters to data points (during the learning/training process, or afterwards in testing) as classification
  - Even if it's a clustering problem!

# Clustering

- The goal of clustering is to find clusters (groupings) that are compact with respect to the distance metric
- What do we mean by compactness?



#### Covariance matrix:

Sample covariance: 
$$\hat{\Sigma}_{ij} = \frac{1}{k} \sum_{k} (x_{ik} - \hat{\mu}_i) (x_{jk} - \hat{\mu}_j) = \frac{1}{k} S_{ij}$$

If  $X_z$  is a matrix that holds all the zero-centered samples as <u>column</u> vectors, then

$$\hat{\Sigma} = \frac{1}{k} X_z X_z^T = \frac{1}{k} S$$

S is the Scatter matrix

Alternatively, if  $X_z$  is a matrix that holds all the zero-centered samples as row vectors, then

$$S = X_z^T X_z$$

It depends on how we define  $X_z$ !

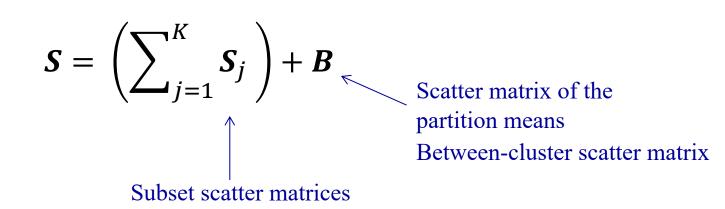
For the scatter matrix (and thus the covariance matrix),  $X_z$  is zero-mean

– That is, the mean data point  $\overline{x}$  (or  $\mu$  or  $\mu_x$ ) is first subtracted from every data point  $x_i$ 

By the way, the Gram matrix is not zero-mean...

### Scatter matrix

• If the data D is partitioned into K subsets  $\{D_1, D_2, ... D_K\}$  then the scatter matrix can be written as



To compute B, replace every point in D with the mean of its partition  $D_i$  and compute the scatter matrix

Within-cluster scatter matrices

# Example: Scatter matrix

$$x_{3} = (0, 4)$$
 $x_{4} = (3, 4)$ 
 $x_{5} = (5, 2)$ 
 $x_{1} = (2, 0)$ 

$$X = [x_1 \quad x_2 \quad x_3 \quad x_4 \quad x_5] = \begin{bmatrix} 2 & 4 & 0 & 3 & 5 \\ 0 & 1 & 4 & 4 & 2 \end{bmatrix}$$

The Gram matrix is ...

$$\mathbf{G} = \mathbf{X}^T \mathbf{X} = \begin{bmatrix} 4 & 8 & 0 & 6 & 10 \\ 8 & 17 & 4 & 16 & 22 \\ 0 & 4 & 16 & 16 & 8 \\ 6 & 16 & 16 & 25 & 23 \\ 10 & 22 & 8 & 23 & 29 \end{bmatrix}$$

$$\overline{x} = \frac{1}{5} \sum_{i=1}^{5} x_i = \frac{1}{5} \begin{bmatrix} 14\\11 \end{bmatrix} = \begin{bmatrix} 2.8\\2.2 \end{bmatrix}$$

 $(k \times k)$ , where k is the number of data points

$$X_z = [x_1 - \overline{x} \quad x_2 - \overline{x} \quad x_3 - \overline{x} \quad x_4 - \overline{x} \quad x_5 - \overline{x}] = \begin{bmatrix} -0.8 & 1.2 & -2.8 & 0.2 & 2.2 \\ -2.2 & -1.2 & 1.8 & 1.8 & -0.2 \end{bmatrix}$$

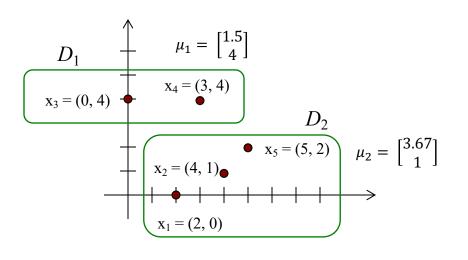
The Scatter matrix is ...

$$S = X_z X_z^T = \begin{bmatrix} 14.8 & -4.8 \\ -4.8 & 12.8 \end{bmatrix}$$

 $(N \times N)$ , where N is the dimensionality of the data points

# Example: Scatter matrix via partitions

$$S = \left(\sum_{j=1}^K S_j\right) + B$$



$$\begin{bmatrix} 1.5 & 1.5 & 3.67 & 3.67 & 3.67 \\ 4 & 4 & 1 & 1 & 1 \end{bmatrix} \qquad \mu_B = \begin{bmatrix} 2.8 \\ 2.2 \end{bmatrix}$$

Zero-mean partition means

$$B_z = \begin{bmatrix} -1.3 & -1.3 & 13/15 & 13/15 & 13/15 \\ 1.8 & 1.8 & -1.2 & -1.2 & -1.2 \end{bmatrix}$$

Between-cluster scatter matrix

$$\mathbf{B} = \mathbf{B}_z \mathbf{B}_z^T = \begin{bmatrix} 5.633 & -7.8 \\ -7.8 & 10.8 \end{bmatrix}$$

Scatter matrix of  $D_1$ 

$$S_1 = \begin{bmatrix} x_3 - \begin{bmatrix} 1.5 \\ 4 \end{bmatrix} & x_4 - \begin{bmatrix} 1.5 \\ 4 \end{bmatrix} \end{bmatrix} \begin{bmatrix} x_3 - \begin{bmatrix} 1.5 \\ 4 \end{bmatrix} & x_4 - \begin{bmatrix} 1.5 \\ 4 \end{bmatrix} \end{bmatrix}^T = \begin{bmatrix} -1.5 & 1.5 \\ 0 & 0 \end{bmatrix} \begin{bmatrix} -1.5 & 1.5 \\ 0 & 0 \end{bmatrix}^T = \begin{bmatrix} 4.5 & 0 \\ 0 & 0 \end{bmatrix}$$

Scatter matrix of  $D_2$ 

$$\mathbf{S}_2 = \begin{bmatrix} -5/3 & 1/3 & 4/3 \\ -1 & 0 & 1 \end{bmatrix} \begin{bmatrix} -5/3 & 1/3 & 4/3 \\ -1 & 0 & 1 \end{bmatrix}^T = \begin{bmatrix} 4.67 & 3 \\ 3 & 2 \end{bmatrix}$$

$$S = S_1 + S_1 + B = \begin{bmatrix} 4.5 & 0 \\ 0 & 0 \end{bmatrix} + \begin{bmatrix} 4.67 & 3 \\ 3 & 2 \end{bmatrix} + \begin{bmatrix} 5.633 & -7.8 \\ -7.8 & 10.8 \end{bmatrix} = \begin{bmatrix} 14.8 & -4.8 \\ -4.8 & 12.8 \end{bmatrix}$$

# Clustering

- The goal of clustering is to find clusters (groupings) that are compact with respect to the distance metric
- Good clustering is characterized by low within-cluster variance and high between-cluster variance
- The scatter matrix,  $S = X_z X_z^T$ , gives us a measure of variance
- If the data D is partitioned into K subsets/partitions  $\{D_1, D_2, ... D_K\}$ , then the scatter matrix can be written as

$$S = \left(\sum_{j=1}^{K} S_{j}\right) + B$$
Between-cluster scatter matrix (maximize!)

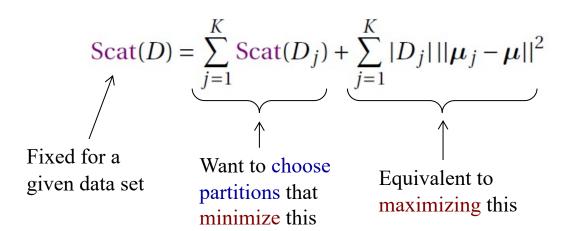
Within-cluster scatter matrices (minimize!)

### Scatter

- The scatter of D is defined as the trace of the scatter matrix
  - The *trace* is the sum of the diagonal elements of a square matrix

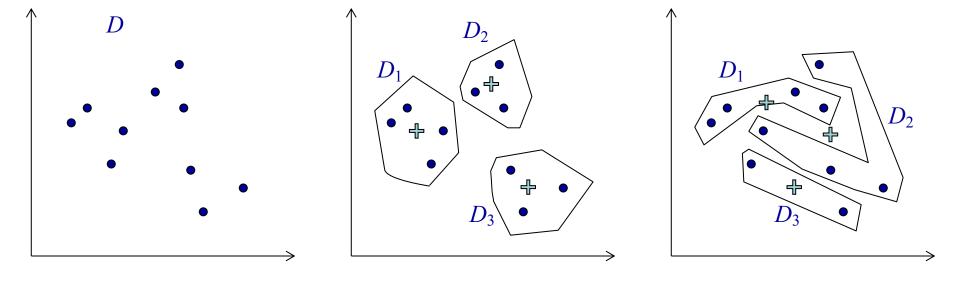
Scat(D) = 
$$Tr(S) = Tr(X_Z X_Z^T)$$
  
=  $Tr(\begin{bmatrix} 14.8 & -4.8 \\ -4.8 & 12.8 \end{bmatrix}) = 14.8 + 12.8 = 27.6$ 

• Since S can be decomposed into partitions, so can Scat(D)



This is the goal of *k*-means clustering

### Scatter



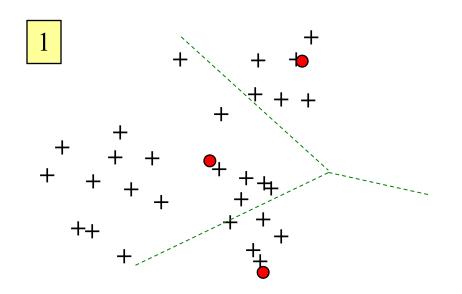
$$\operatorname{Scat}(D) = \sum_{j=1}^K \operatorname{Scat}(D_j) + \sum_{j=1}^K |D_j| \left| |\boldsymbol{\mu}_j - \boldsymbol{\mu}| \right|^2$$

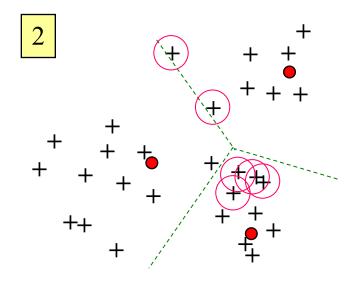
## K-means clustering

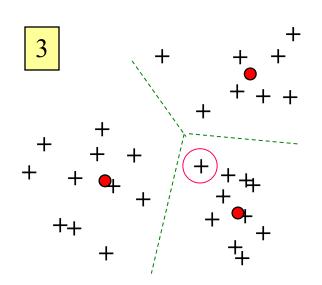
- The general K-means clustering problem is NP-complete, so there is no efficient solution to find the optimal clustering (data partition)
- A widely-used heuristic algorithm for clustering is also known as the K-means algorithm, but it is not optimal
  - It will converge to a solution, but there is no guarantee that the solution is the best one (the global minimum of scatter)
  - But it works quite well in most cases!
- Typically, the K-means algorithm would be run several times (with a random starting point) and then the best solution is selected
  - I.e., the solution with the smallest within-cluster scatter

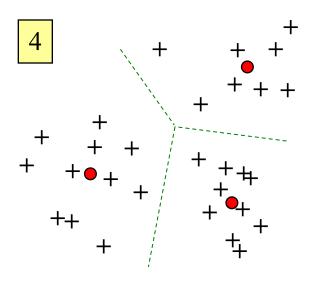
## **Algorithm** KMeans(D(K) - K-means clustering using Euclidean distance $(Dis_2)$

```
: data D \subseteq \mathbb{R}^d; number of clusters K \in \mathbb{N}.
Output: K cluster means \mu_1, \ldots, \mu_K \in \mathbb{R}^d.
randomly initialise K vectors \mu_1, \ldots, \mu_K \in \mathbb{R}^d;
repeat
      assign each \mathbf{x} \in D to \operatorname{argmin}_{i} \operatorname{Dis}_{2}(\mathbf{x}, \mu_{j}); \longleftarrow 1-Nearest neighbor assignment
      for j = 1 to K do
             D_i \leftarrow \{\mathbf{x} \in D | \mathbf{x} \text{ assigned to cluster } j\}; \leftarrow Partition defined by assignment
            \mu_j = \frac{1}{|D_i|} \sum_{\mathbf{x} \in D_i} \mathbf{x}; \quad \longleftarrow \text{Re-compute the cluster mean}
      end
until no change in \mu_1, ..., \mu_K;
return \mu_1, \ldots, \mu_K;
```









### K-means demo

**Demo**: <a href="https://www.naftaliharris.com/blog/visualizing-k-means-clustering/">https://www.naftaliharris.com/blog/visualizing-k-means-clustering/</a>

# K-means applications

In K-means, we simply take a cluster centroid (exemplar) to be the mean of points in the cluster.

- Document clustering
- House clustering based on price, square footage, #bedrooms, etc.
- Image segmentation (pixel clustering based on RGB values and location)

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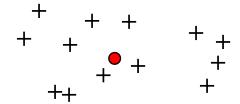


# K-medoids algorithm

- In some problems, the cluster exemplars are required to be data points (from the training data)
  - As opposed to using the mean of the cluster points, for example, since the mean is most likely not a point in the data set
- The concept of *medoid* is useful here the medoid of a set of points is the point with the minimal average dissimilarity (distance) to all other points in the set
  - Using some distance metric: Euclidian, L1, etc.
  - This is a generalization of the concept of median to multiple dimensions
- K-means can be modified to use data points as exemplars rather than means, by instead computing the cluster medoids

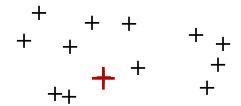
# K-medoids algorithm

#### Cluster mean



Location that minimizes the sum of squared distances to points

#### Cluster medoid



Point that minimizes the sum of squared distances to points

# K-medoids algorithm

**Algorithm** KMedoids(D, K, Dis) – K-medoids clustering using arbitrary distance metric Dis.

```
: data D \subseteq \mathcal{X}; number of clusters K \in \mathbb{N};
                 distance metric Dis: \mathcal{X} \times \mathcal{X} \to \mathbb{R}.
Output: K medoids \mu_1, \ldots, \mu_K \in D, representing a predictive clustering of \mathscr{X}.
randomly pick K data points \mu_1, \ldots, \mu_K \in D;
repeat
      assign each \mathbf{x} \in D to \operatorname{argmin}_{i} \operatorname{Dis}(\mathbf{x}, \mu_{j});
      for j = 1 to K do
             D_j \leftarrow \{\mathbf{x} \in D | \mathbf{x} \text{ assigned to cluster } j\};
            \mu_i = \operatorname{argmin}_{\mathbf{x} \in D_i} \sum_{\mathbf{x}' \in D_i} \operatorname{Dis}(\mathbf{x}, \mathbf{x}'); \leftarrow Re\text{-compute the cluster medoid}
      end
until no change in \mu_1, \ldots, \mu_K;
return \mu_1, \ldots, \mu_K;
```

# Summary: Distance methods and clustering

- Similarity is a function of distance
- Euclidian distance may not always be the right choice of distance metric
- Nearest neighbor methods assign classes/clusters based on distances to points or exemplars, not based on computed boundaries
- For good clustering, we want high within-class (intra-class) similarity and low between-class (inter-class) similarity
- The scatter matrix is an important structure in clustering
- The K-means algorithm (and variations) is widely used