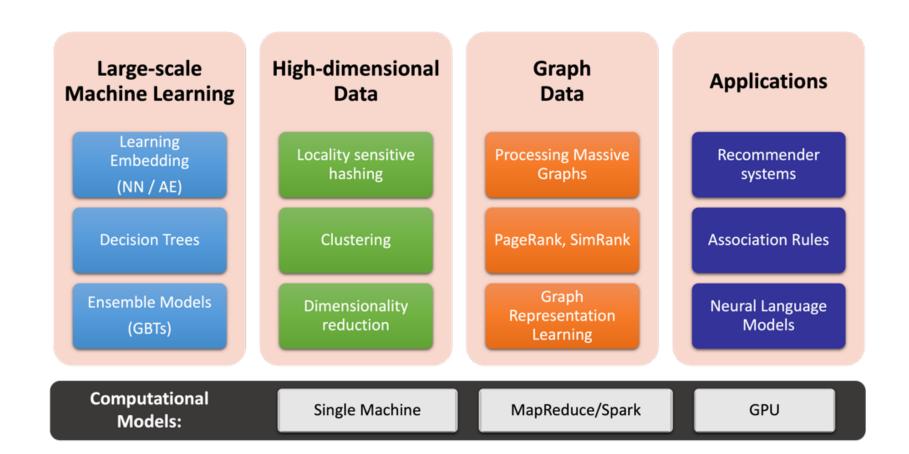
MIE524 Data Mining Clustering

Slides Credits:

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MIE524: Course Topics (Tentative)



The Problem of Clustering

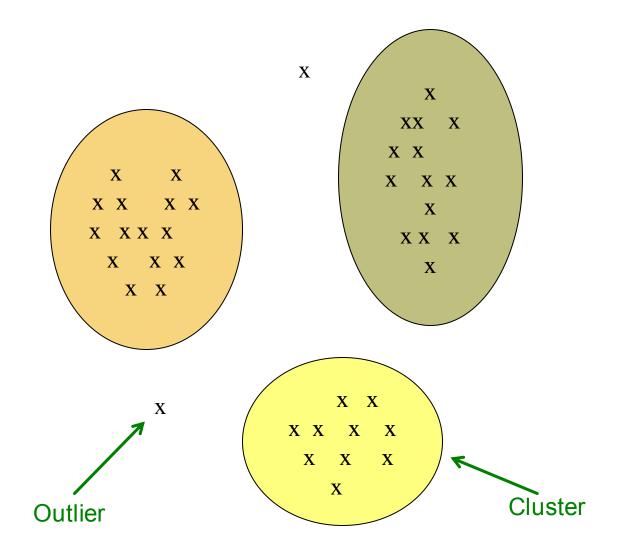
- Given a set of points, with a notion of distance between points, group the points into some number of clusters, so that
 - Members of the same cluster are close/similar to each other
 - Members of different clusters are dissimilar

Usually:

- Points are in a high-dimensional space
- Similarity is defined using a distance measure
 - Euclidean, Cosine, Jaccard, edit distance, ...

Example: Clusters & Outliers

Example: Clusters & Outliers



Examples of clustering tasks

- Identify similar groups of online shoppers based on their browsing and purchasing history
- Identify similar groups of music listeners or movie viewers based on their ratings or recent listening/viewing patterns
- Cluster input variables based on their correlations to remove redundant predictors from consideration
- Cluster hospital patients based on their medical histories
- Determine how to place sensors, broadcasting towers, law enforcement, or emergency-care centers to guarantee that desired coverage criteria are met

Clustering Problem: Music CDs

- Intuitively: Music can be divided into categories, and customers prefer a few genres
 - But what are categories really?
- Represent a CD by a set of customers who bought it
- Similar CDs have similar sets of customers, and vice-versa

Clustering Problem: Music CDs

Space of all CDs:

- Think of a space with one dim. for each customer
 - Values in a dimension may be 0 or 1 only
 - A CD is a "point" in this space $(x_1, x_2, ..., x_d)$, where $x_i = 1$ iff the i th customer bought the CD
- For Amazon, the dimension is tens of millions
- Task: Find clusters of similar CDs

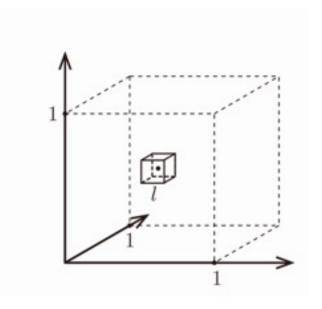
Why is it hard?

- Clustering in two dimensions looks easy
- Clustering small amounts of data looks easy
- And in most cases, looks are not deceiving
- Many applications involve not 2, but 10 or 10,000 dimensions
- High-dimensional spaces look different: Almost all pairs of points are very far from each other --> The Curse of Dimensionality!

Example: Curse of Dimensionality

- Take 10,000 uniform random points on [0,1] line. Assume query point is at the origin
- What fraction of "space" do we need to cover to get 0.1% of data (10 nearest neighbors)
- In 1-dim to get 10 neighbors we must go to distance 10/10,000=0.001 on the average
- In 2-dim we must go $\sqrt{0.001}$ =0.032 to get a square that contains 0.001 volume
- In general, in d-dim we must go $(0.001)^{\frac{1}{d}}$
- So, in 10-dim to capture 0.1% of the data we need 50% of the range.

Curse of dimensionality



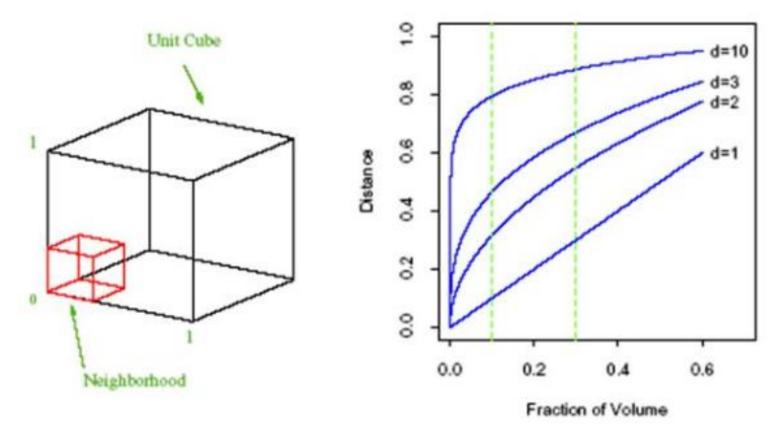
Formally, imagine the unit cube $[0,1]^d$. All training data is sampled *uniformly* within this cube, i.e. $\forall i, x_i \in [0,1]^d$, and we are considering the k=10 nearest neighbors of such a test point.

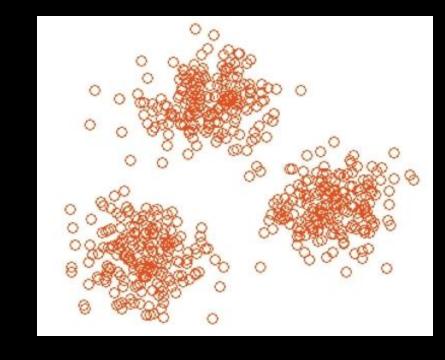
Let ℓ be the edge length of the smallest hyper-cube that contains all k-nearest neighbor of a test point. Then $\ell^d \approx \frac{k}{n}$ and $\ell \approx \left(\frac{k}{n}\right)^{1/d}$. If n=1000, how big is ℓ ?

d	ℓ	
2	0.1	
10	0.63	
100	0.955	
1000	0.9954	

Example: Curse of Dimensionality

Curse of Dimensionality: All points are very far from each other





k-means Clustering

Method: K-mean clustering

- Main idea: A good clustering is one for which the within-cluster variation is as small as possible.
- ullet The within-cluster variation for cluster C_k is some measure of the amount by which the observations within each class differ from one another
- We'll denote it by $WCV(C_k)$
- Goal: Find C_1, \ldots, C_K that minimize

$$\sum_{k=1}^{K} WCV(C_k)$$

ullet This says: Partition the observations into K clusters such that the WCV summed up over all K clusters is as small as possible

How to define within-cluster variation?

• Goal: Find C_1, \ldots, C_K that minimize

$$\sum_{k=1}^{K} \mathrm{WCV}(C_k)$$

• Typically, we use squared Euclidean distance:

$$WCV(C_k) = \frac{1}{|C_k|} \sum_{i,i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2$$

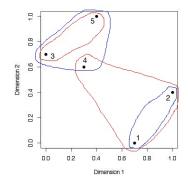
where $|C_k|$ denotes the number of observations in cluster k

To be clear: We're treating K as fixed ahead of time. We are not
optimizing K as part of this objective.

Simple example

Here n=5 and K=2, The full distance matrix for all 5 observations is shown below.

	I	2	3	4	5
ı	0	0.25	0.98	0.52	1.09
2	0.25	0	1.09	0.53	0.72
3	0.98	1.09	0	0.10	0.25
4	0.52	0.53	0.10	0	0.17
5	1.09	0.72	0.25	0.17	0



- Red clustering: $\sum WCV_k = (0.25 + 0.53 + 0.52)/3 + 0.25/2 = 0.56$
- Blue clustering: $\sum WCV_k = 0.25/2 + (0.10 + 0.17 + 0.25)/3 = 0.30$
- ullet It's easy to see that the Blue clustering minimizes the within-cluster variation among all possible partitions of the data into K=2 clusters

How do we minimize WCV?

$$\sum_{k=1}^{K} \text{WCV}(C_k) = \sum_{k=1}^{K} \frac{1}{|C_k|} \sum_{i,i' \in C_k} \sum_{j=1}^{p} (x_{ij} - x_{i'j})^2$$
$$= \sum_{k=1}^{K} \frac{1}{|C_k|} \sum_{i,i' \in C_k} ||x_i - x_{i'}||_2^2$$

- It's computationally infeasible to actually minimize this criterion
- We essentially have to try all possible partitions of n points into K sets.
- When n=10, K=4, there are $34{,}105$ possible partitions
- When n = 25, K = 4, there are 5×10^{13} ...
- We're going to have to settle for an approximate solution

K-means algorithm

• It turns out that we can rewrite WCV_k more conveniently:

$$WCV_k = \frac{1}{|C_k|} \sum_{i,i' \in C_k} ||x_i - x_{i'}||_2^2 = 2 \sum_{i \in C_k} ||x_i - \bar{x}_k||^2$$

where $\bar{x}_k = \frac{1}{|C_k|} \sum_{i \in C_k} x_i$ is just the average of all the points in cluster C_k

k–means Algorithm(s)

Initialize clusters by picking k centers

Until convergence:

- 1) For each point, assign it to the cluster whose current centroid is the closest
- 2) After all points are assigned, update the centroids of the k clusters as average of datapoints within each cluster

Convergence means Points don't move between clusters and centroids stabilize

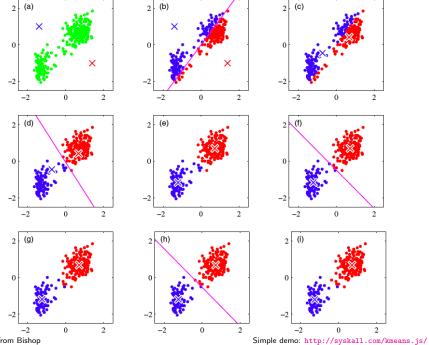
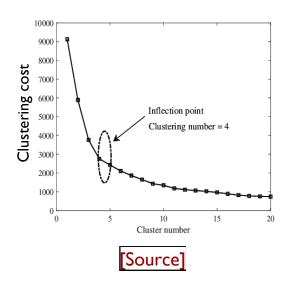


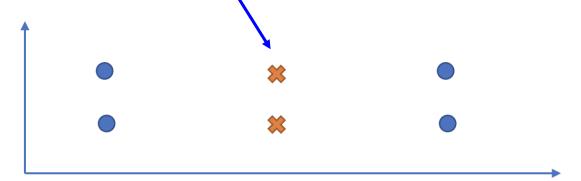
Figure from Bishop

How to choose K?



Shortcoming of k-means

Convergence of k-means heavily depends on the initial pick of centroids. It can perform arbitrarily badly:



Summary of K-means

We'd love to minimize

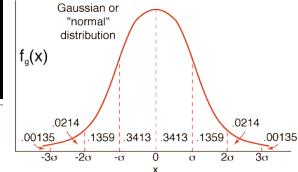
$$\sum_{k=1}^{K} \frac{1}{|C_k|} \sum_{i,i' \in C_k} ||x_i - x_{i'}||_2^2$$

- It's infeasible to actually optimize this in practice, but K-means at least gives us a so-called local optimum of this objective
- The result we get depends both on K, and also on the *random* initialization that we wind up with
- It's a good idea to try different random starts and pick the best result among them
- There's a method called K-means++ that improves how the clusters are initialized

The BFR Algorithm

Extension of k-means to large data

BFR Algorithm



- BFR [Bradley-Fayyad-Reina] is a variant of k-means designed to handle very large (disk-resident) data sets
- Assumes that clusters are normally distributed around a centroid in a Euclidean space
 - Standard deviations in different dimensions may vary
 - Clusters are axis-aligned ellipses
- Goal is to find cluster centroids; point assignment can be done in a second pass through the data.

BFR Overview

- Efficient way to summarize clusters: Want memory required O(clusters) and not O(data)
- IDEA: Rather than keeping points, BFR keeps summary statistics of groups of points
 - 3 sets: Discard set, Compressed set, Retained set
- Overview of the algorithm:
 - **1.** Initialize *K* clusters/centroids
 - 2. Load in a bag of points from disk
 - 3. Assign new points to one of the K original clusters, if they are within some distance threshold of the cluster
 - 4. Cluster the remaining points, and create new clusters
 - 5. Try to merge new clusters from step 4 with any of the existing clusters
 - 6. Repeat steps 2-5 until all points are examined

BFR Algorithm

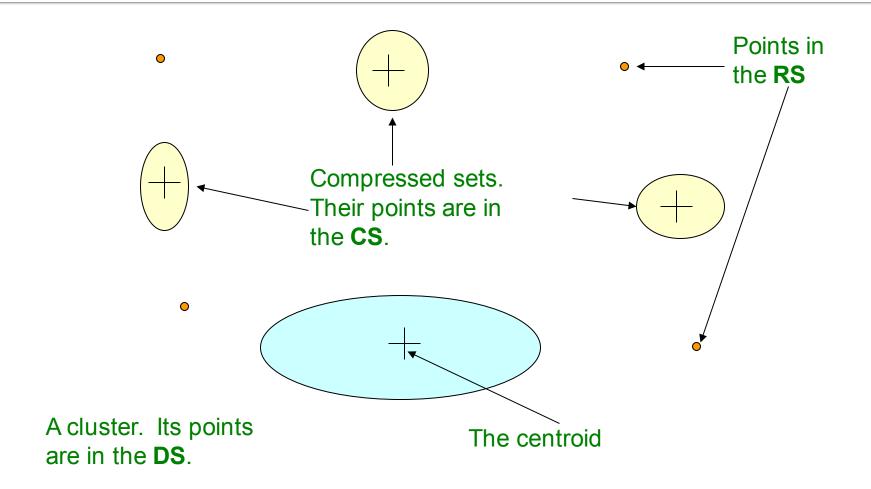
- Points are read from disk one main-memoryfull at a time
- Most points from previous memory loads are summarized by simple statistics
- Step 1) From the initial load we select the initial k centroids by some sensible approach:
 - Take k random points
 - Take a small random sample and cluster optimally
 - Take a sample; pick a random point, and then
 k-1 more points, each as far from the previously selected points as possible

Three Classes of Points

3 sets of points which we keep track of:

- Discard set (DS):
 - Points close enough to a centroid to be summarized
- Compressed set (CS):
 - Groups of points that are close together but not close to any existing centroid
 - These points are summarized, but not assigned to a cluster
- Retained set (RS):
 - Isolated points waiting to be assigned to a compression set

BFR: "Galaxies" Picture

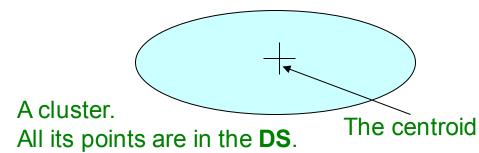


Discard set (DS): Close enough to a centroid to be summarized **Compression set (CS):** Summarized, but not assigned to a cluster **Retained set (RS):** Isolated points, we store them as they are

Summarizing Sets of Points

For each cluster, the discard set (DS) is summarized by:

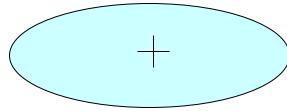
- The number of points, N
- The vector *SUM*, whose *i*th component is the sum of the coordinates of the points in the *i*th dimension
- The vector SUMSQ: ith component = sum of squares of coordinates in ith dimension



Summarizing Points: Comments

- 2d + 1 values represent any size cluster
 - \mathbf{d} = number of dimensions
- Average in each dimension (the centroid)
 can be calculated as SUM; / N
 - **SUM**_i = ith component of SUM
- Variance of a cluster's discard set in dimension i is: (SUMSQ_i / N) – (SUM_i / N)²
 - And standard deviation is the square root of that
- Next step: Actual clustering

Note: Dropping the "axis-aligned" clusters assumption would require storing full covariance matrix to summarize the cluster. So, instead of **SUMSQ** being a *d*-dim vector, it would be a *d* x *d* matrix, which is too big!



The "Memory-Load" of Points

Steps 3-5) Processing "Memory-Load" of points:

- **Step 3)** Find those points that are "sufficiently close" to a cluster centroid and add those points to that cluster and the **DS**
 - These points are so close to the centroid that they can be summarized and then discarded
- Step 4) Use any in-memory clustering algorithm to cluster the remaining points and the old **RS**
 - Clusters go to the CS; outlying points to the RS

Discard set (DS): Close enough to a centroid to be summarized. Compression set (CS): Summarized, but not assigned to a cluster Retained set (RS): Isolated points Jure Les kovec & Mina Ghashami, Stanford CS246: Mining Massive Datasets

The "Memory-Load" of Points

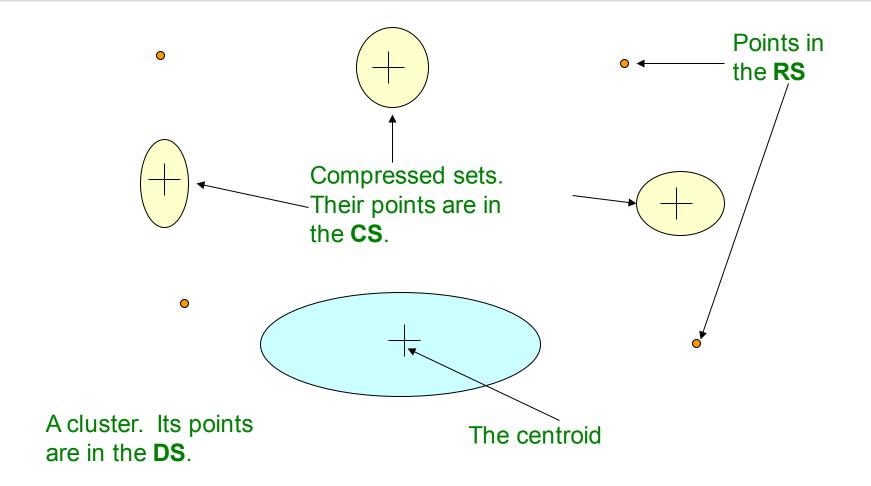
Steps 3-5) Processing "Memory-Load" of points:

- Step 5) DS set: Adjust statistics of the clusters to account for the new points
 - Add Ns, SUMs, SUMSQs
- Consider merging compressed sets in the CS
- If this is the last round, merge all compressed sets in the CS and all RS points into their nearest cluster

Jure Les kovec & Mina Ghashami, Stanford CS246: Mining Massive Datasets

Discard set (DS): Close enough to a centroid to be summarized. **Compression set (CS):** Summarized, but not assigned to a cluster **Retained set (RS):** Isolated points

Summary: BFR



Discard set (DS): Close enough to a centroid to be summarized **Compression set (CS):** Summarized, but not assigned to a cluster **Retained set (RS):** Isolated points

A Few Details...

- Q1) How do we decide if a point is "close enough" to a cluster that we will add the point to that cluster?
- Q2) How do we decide whether two compressed sets (CS) deserve to be combined into one?

How Close is Close Enough?

 Q1) We need a way to decide whether to put a new point into a cluster (and discard)

- The Mahalanobis distance is less than a threshold
- High likelihood of the point belonging to currently nearest centroid

Mahalanobis Distance

Normalized Euclidean distance from centroid

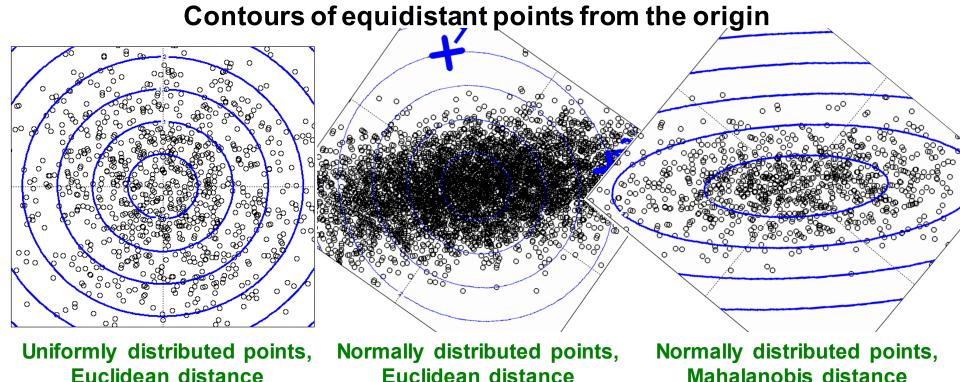
- For a given point $(x_1, ..., x_d)$ and a given centroid $(c_1, ..., c_d)$
 - 1. Normalize in each dimension: $y_i = (x_i c_i) / \sigma_i$
 - 2. Take sum of the squares of the y_i
 - 3. Take the square root

$$d(x,c) = \sqrt{\sum_{i=1}^{d} \left(\frac{x_i - c_i}{\sigma_i}\right)^2}$$

 σ_i ... standard deviation of points in the cluster in the ith dimension

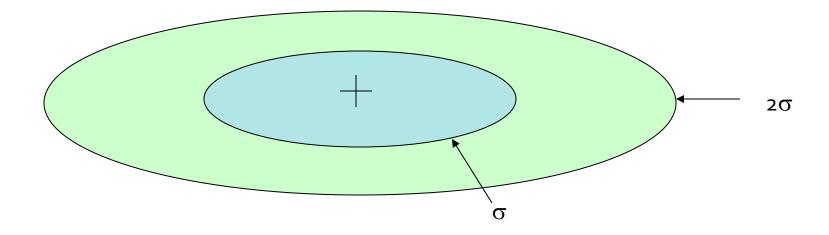
Picture: Equal M.D. Regions

Euclidean vs. Mahalanobis distance



Euclidean distance

Picture: Equal M.D. Regions



Accept a point for a cluster if its M.D. is < some threshold, e.g., 2 standard deviations

Should 2 CS clusters be combined?

Q2) Should 2 CS clusters be combined?

- Compute the variance of the combined subcluster
 - N, SUM, and SUMSQ allow us to make that calculation quickly
- Combine if the combined variance is below some threshold
- Many alternatives: Treat dimensions differently, consider density

