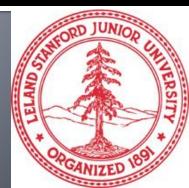
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Clustering

CS246: Mining Massive Datasets
Jure Leskovec, Stanford University
Mina Ghashami, Stanford University
http://cs246.stanford.edu



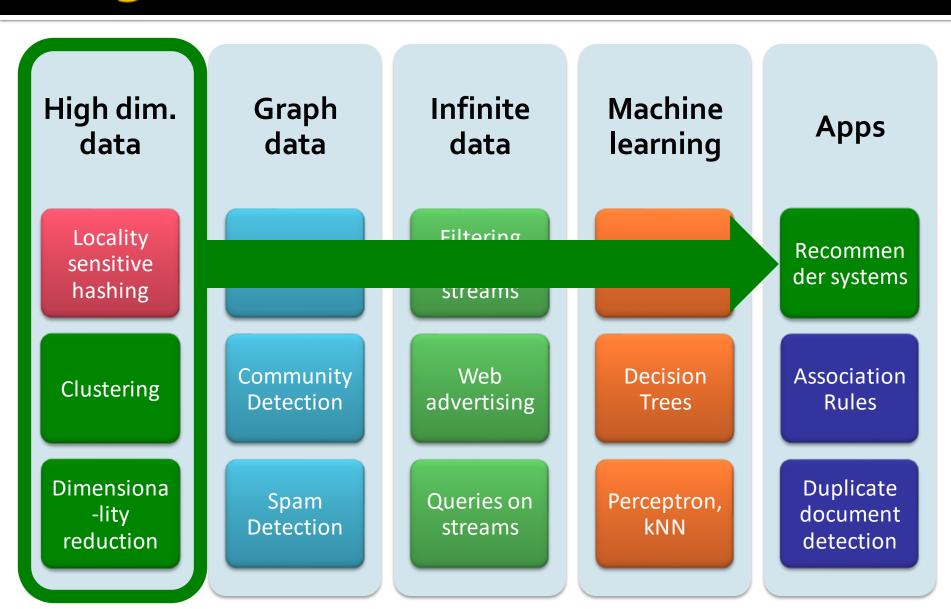
Announcements

- Colab 0 & 1 grades released.
 - Solutions to be released soon.
- Colab 2 and Homework 1 due this Thursday.

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High Dimensional Data



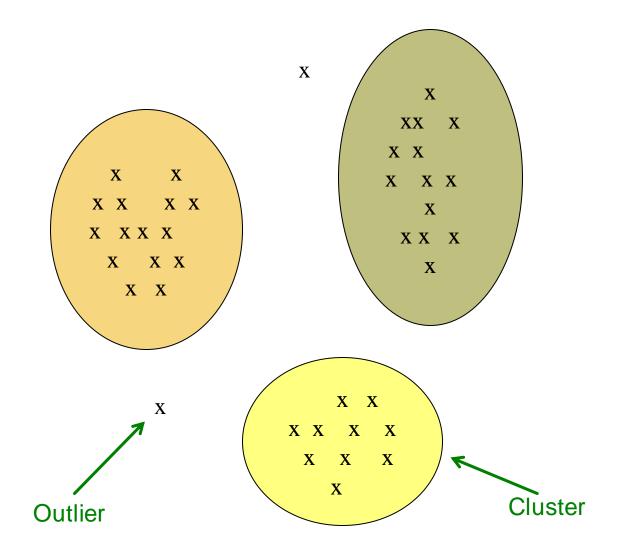
The Problem of Clustering

- 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



Clustering Problem: Galaxies

- A catalog of 2 billion "sky objects" represents objects by their radiation in 7 dimensions (frequency bands)
- Problem: Cluster similar objects, e.g., galaxies, nearby stars, quasars, etc.
- Sloan Digital Sky Survey



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

Clustering Problem: Documents

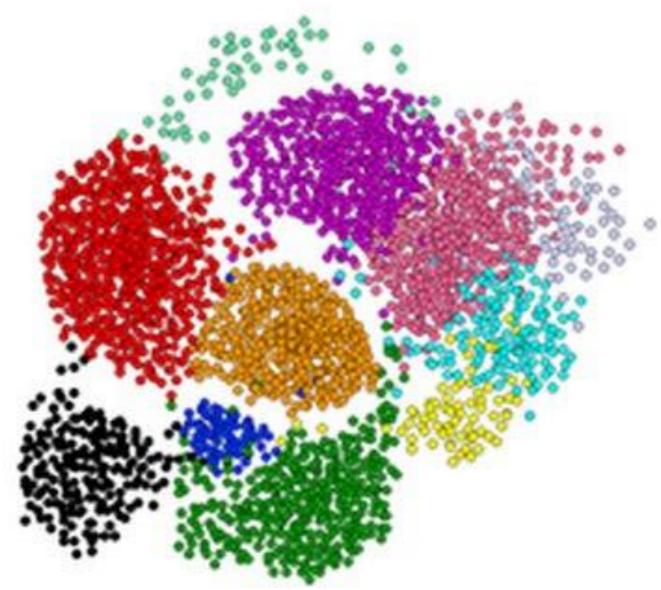
Finding topics:

- Represent a document by a vector $(x_1, x_2, ..., x_k)$, where $x_i = 1$ iff the i th word (in some order) appears in the document
 - It actually doesn't matter if k is infinite; i.e., we don't limit the set of words
- Documents with similar sets of words may be about the same topic

Cosine, Jaccard, and Euclidean

- We have a choice when we think of documents as sets of words or shingles:
 - Sets as vectors: Measure similarity by the cosine distance
 - Sets as sets: Measure similarity by the Jaccard distance
 - Sets as points: Measure similarity by Euclidean distance

Clustering is a hard problem!



Why is it hard?

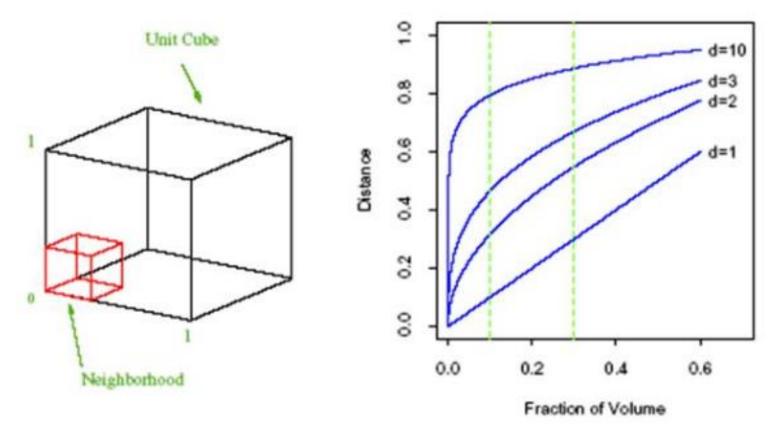
- Clustering in two dimensions looks easy
- Clustering small amounts of data looks easy
- And in most cases, looks are not deceiving
- Many applications involve not 2, but 10 or 10,000 dimensions
- High-dimensional spaces look different: Almost all pairs of points are very far from each other --> The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

Example: Curse of Dimensionality

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



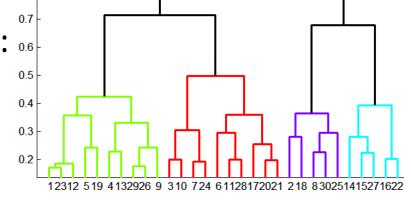
Overview: Clustering Strategies (1)

 $0.9 \, \mathrm{r}$

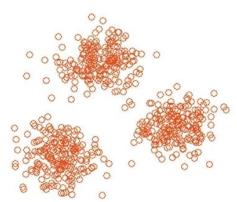
0.8

Two group of methods:

- Hierarchical:
 - Agglomerative (bottom up):
 - Initially, each point is a cluster
 - Repeatedly combine the two "nearest" clusters into one



- Divisive (top down):
 - Start with one cluster and recursively split it
- Point assignment:
 - Maintain a set of clusters
 - Points belong to the "nearest" cluster



Overview: Clustering Strategies (2)

- Is the space Euclidean or non-Euclidean?
- In Euclidean:
 - Points are vectors of real numbers, i.e. coordinates
 - It is possible to summarize a collection of points as their average. We call it centroid.
 - Distance measure: L2 norm, L1 norm
- In non-Euclidean:
 - There is no notion of location, and centroid
 - We summarize a collection of points differently
 - Distance measures: Jaccard, Hamming, cosine

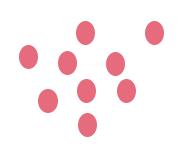
Overview: Clustering Strategies (3)

Does the data fit in memory or it resides on disk?

- In-memory clustering is more straightforward
 - Example: K-means
- Large-data clustering requires loading one batch of data at a time, cluster them in memory and keep summaries of clusters
 - Example: BFR, CURE

Hierarchical vs point-assignment

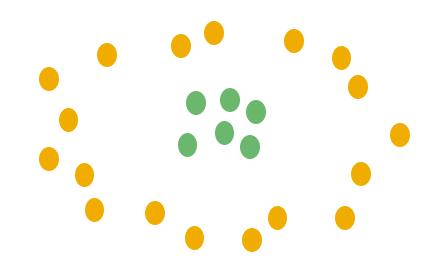
 Point assignment good when clusters are nice, convex shapes:

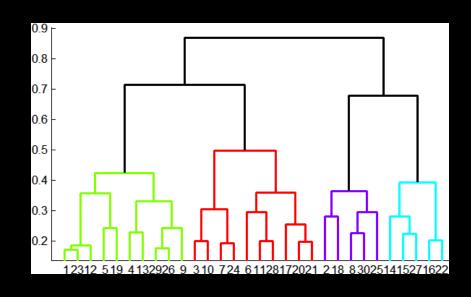




- Hierarchical can win when shapes are weird:
 - Note both clusters have essentially the same centroid.

Aside: if you realized you had concentric clusters, you could map points based on distance from center, and turn the problem into a simple, one-dimensional case.

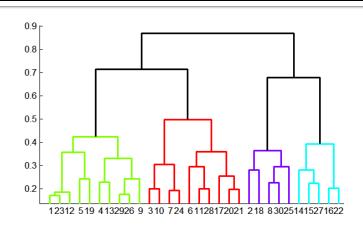




Hierarchical Clustering

Hierarchical Clustering

 Key operation:
 Repeatedly merge two "nearest" clusters



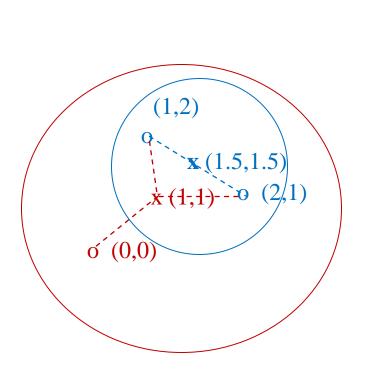
- Three important questions:
 - 1) How to represent a cluster?
 - 2) How to determine the nearness of clusters?
 - **3)** When to stop merging clusters?

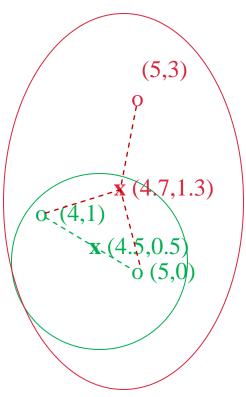
Hierarchical Clustering: Euclidean

In Euclidean case:

- (1) How to represent a cluster of many points?
 - As we merge clusters, we represent the "location" of each cluster by its
 - **centroid** = average of its (data)points
- (2) How to determine the nearness of clusters?
 - Measure cluster distances by distances of centroids
 - Merge two clusters with the shortest distance

Example: Hierarchical clustering

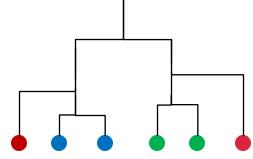




Data:

o ... data point

x ... centroid



Dendrogram

Hierarchical Clustering: Non-Euclidean

In non-Euclidean case:

The only "locations" we can talk about are the points themselves. There are three main approaches:

(1) How to represent a cluster of many points?

- 1. pick a *clustroid* = point "*closest*" to other points
- 2. As the collection of points it is.
- 3. As the collection of points it is.

(2) How to determine the nearness of clusters?

- 1. Treat clustroid as if it were centroid
- 2. Various distance measures between points of two clusters
- 3. Various cohesion measures of the union of two clusters

Hierarchical Clustering: Non-Euclidean

Approach 1:

(1) How to represent a cluster:

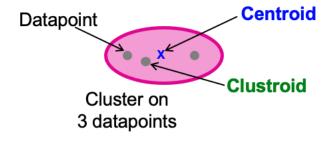
- pick a clustroid = (data)point "closest" to other points
 Possible meanings of "closest":
 - Smallest maximum distance to other points
 - Smallest average distance to other points
 - Smallest sum of squares of distances to other points
 - For distance metric d clustroid c of cluster c is $\arg\min_{c}\sum_{x\in C}d(x,c)^2$

(2) How to determine the nearness of clusters?

Treat clustroid as if it were centroid

Centroid vs Clusteroid

- Centroid is the avg. of all (data)points in the cluster.
 - This means centroid is an "artificial" point.
- Clustroid is an existing (data)point that is "closest" to all other points in the cluster.



Hierarchical Clustering: Non-Euclidean

Approach 2:

(1) How to represent a cluster? As the collection of points

(2) How to determine the nearness of clusters? Define *inter-cluster distance*:

- Minimum of the distances between any two points, one from each cluster
- Average distance of all pairs of points, one from each cluster

Hierarchical Clustering: Non-Euclidean

Approach 3:

- (1) How to represent a cluster? As the collection of points
- (2) How to determine the nearness of clusters? Define a notion of *cohesion*, and merge clusters whose *union* is most cohesive Possible notions of cohesion (the smaller, the more cohesive):
- diameter of the merged cluster = maximum distance between points in the cluster
- average distance between points in the cluster
- Density of the merged cluster = divide by the number of points in the cluster by diameter or avg. distance

When to Stop?

When do we stop merging clusters?

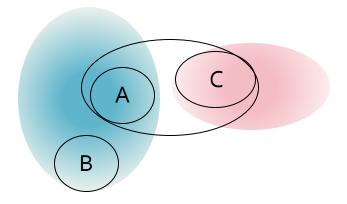
- When some number k of clusters are found (assumes we know the number of clusters)
- When stopping criterion is met
 - Stop if diameter exceeds threshold
 - Stop if density is below some threshold
 - Stop if merging clusters yields a bad cluster
 - E.g., diameter suddenly jumps
- Keep merging until there is only 1 cluster left

Which design choice is the best?

- It really depends on the shape of clusters.
 - Which you may not know in advance.
- Example: we'll compare two approaches:
 - Merge clusters with smallest distance between centroids (or clustroids for non-Euclidean)
 - 2. Merge clusters with the smallest distance between two points, one from each cluster

Case 1: Convex Clusters

- Centroid-based merging works well.
- But merger based on closest members might accidentally merge incorrectly.

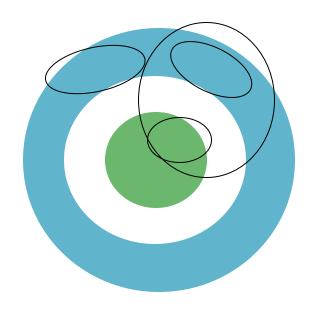


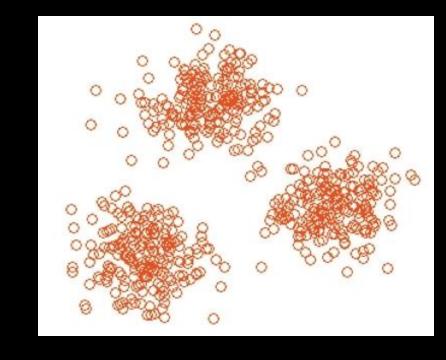
A and B have closer centroids than A and C, but closest points are from A and C.



Case 2: Concentric Clusters

- Linking based on closest members works well
- But Centroid-based linking might cause errors





k-means Clustering

k–means Algorithm(s)

- It is a problem formulation, not an algorithm.
- Problem: Given Euclidean space/distance and k = number of clusters, find cluster centers that minimizes sum of squared distances from each point to its cluster center
- Finding an exact solution is NP-hard.
- The approximate solution is Lloyd's algorithm or the k-means algorithm.

k–means Algorithm(s)

Initialize clusters by picking k centers

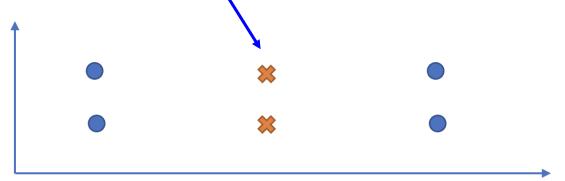
Until convergence:

- 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

Shortcoming of k-means

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



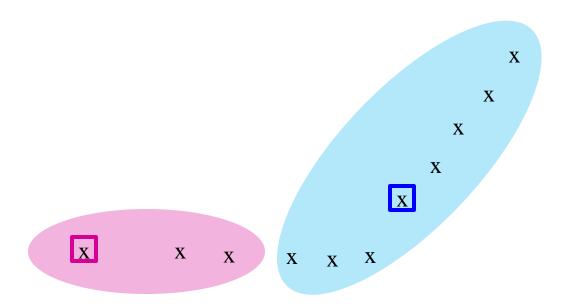
Different strategies for picking k centers:

- Pick k datapoints at random
- k-means ++

k-means++

- Basic idea: Pick a small sample of points S, cluster them by any algorithm, and use the centroids as a seed
- In **k-means++**, sample size |S| = k times a factor that is logarithmic in the total number of points
- How to pick sample points: Visit points in random order, but the probability of adding a point p to the sample is proportional to $D(p)^2$.
 - D(p) = distance between p and the nearest already picked point.

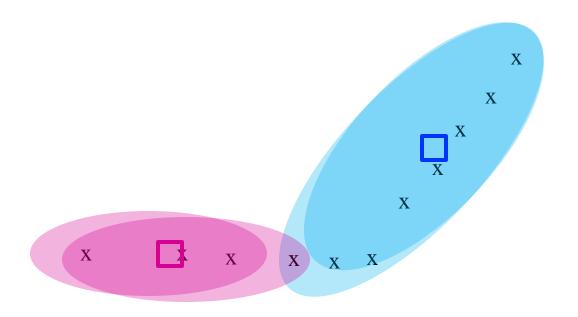
Example: Assigning Clusters



x ... data point ... centroid

Clusters after round 1

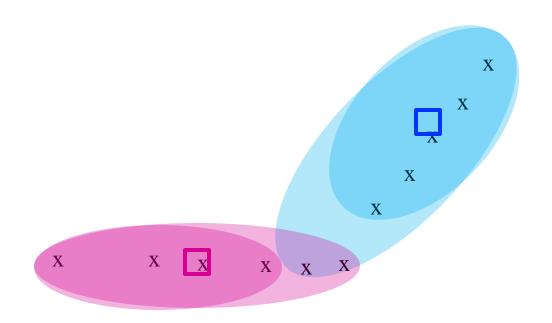
Example: Assigning Clusters



x ... data point ... centroid

Clusters after round 2

Example: Assigning Clusters



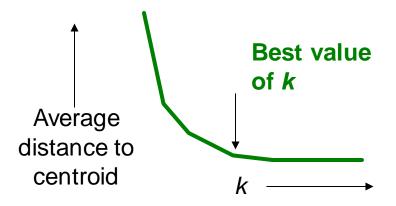
x ... data point ... centroid

Clusters at the end

Getting the k right

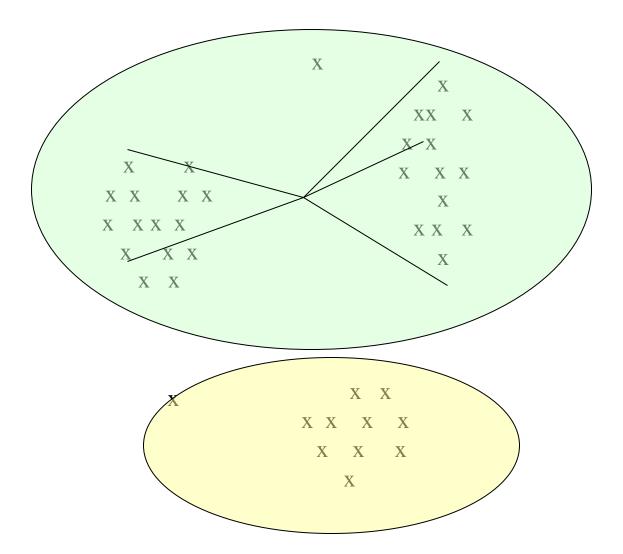
How to select k?

- Try different k, looking at the change in the average distance to centroid as k increases
- Average falls rapidly until right k, then changes little



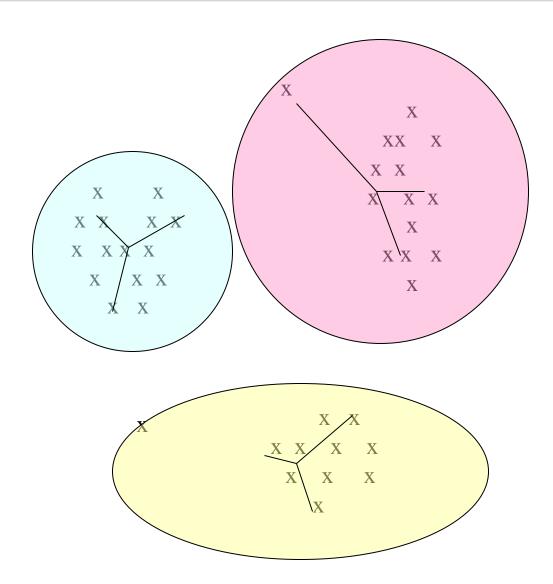
Example: Picking k

Too few; many long distances to centroid



Example: Picking k

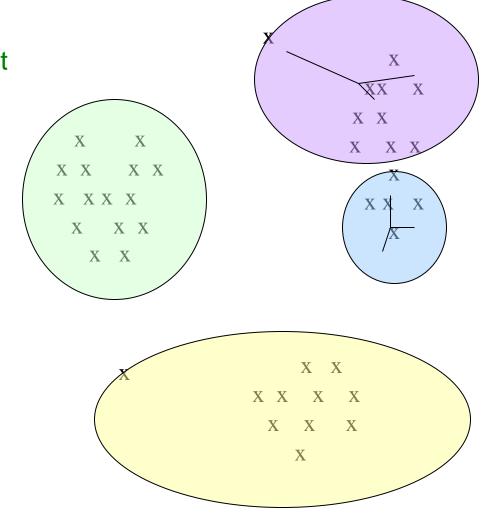
Just right; distances rather short



Example: Picking k

Too many; little improvement

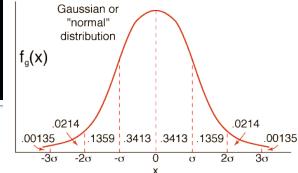
in average distance



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

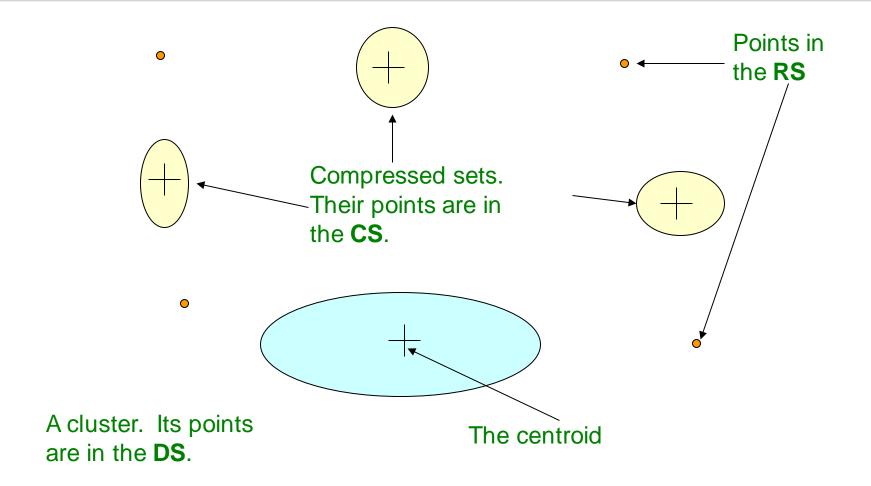
- Points are read from disk one main-memoryfull at a time
- Most points from previous memory loads are summarized by simple statistics
- Step 1) From the initial load we select the initial k centroids by some sensible approach:
 - Take k random points
 - Take a small random sample and cluster optimally
 - 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

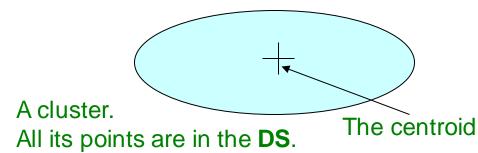


Discard set (DS): Close enough to a centroid to be summarized **Compression set (CS):** Summarized, but not assigned to a cluster **Retained set (RS):** Isolated points, we store them as they are

Summarizing Sets of Points

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

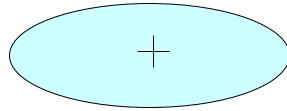
- The number of points, N
- The vector *SUM*, whose *i*th component is the sum of the coordinates of the points in the *i*th dimension
- The vector SUMSQ: ith component = sum of squares of coordinates in ith dimension



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:

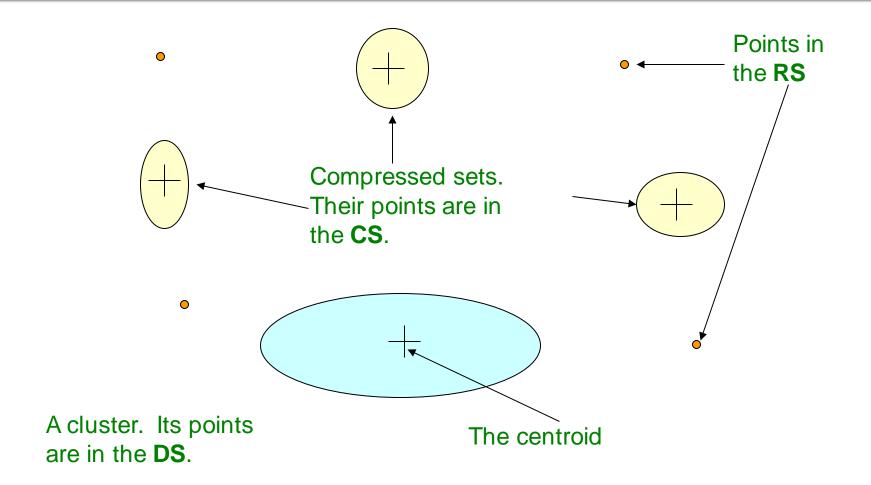
- Step 5) DS set: Adjust statistics of the clusters to account for the new points
 - Add Ns, SUMs, SUMSQs
 - Consider merging compressed sets in the DS

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 If this is the last round, merge all compressed sets in the CS and all RS points into their nearest cluster

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

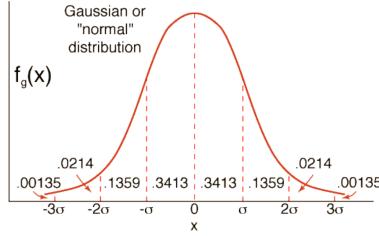
A Few Details...

- 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)
- BFR suggests two ways:
 - 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

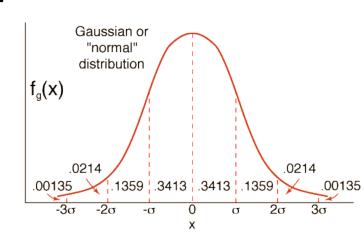
- For a given point $(x_1, ..., x_d)$ and a given centroid $(c_1, ..., c_d)$
 - 1. Normalize in each dimension: $y_i = (x_i c_i) / \sigma_i$
 - 2. Take sum of the squares of the y_i
 - 3. Take the square root

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

 σ_i ... standard deviation of points in the cluster in the i^{th} dimension

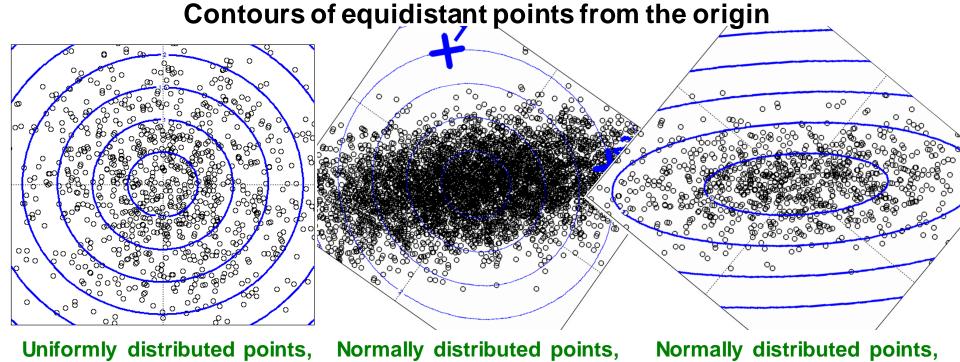
Mahalanobis Distance

- If clusters are normally distributed in d dimensions, then after transformation, one standard deviation = \sqrt{d}
 - i.e., 68% of the points of the cluster will have a Mahalanobis distance $< \sqrt{d}$
- Accept a point for a cluster if its M.D. is < some threshold, e.g. 2 standard deviations



Picture: Equal M.D. Regions

Euclidean vs. Mahalanobis distance



Euclidean distance

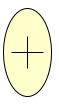
Euclidean distance

Mahalanobis distance

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





The CURE Algorithm

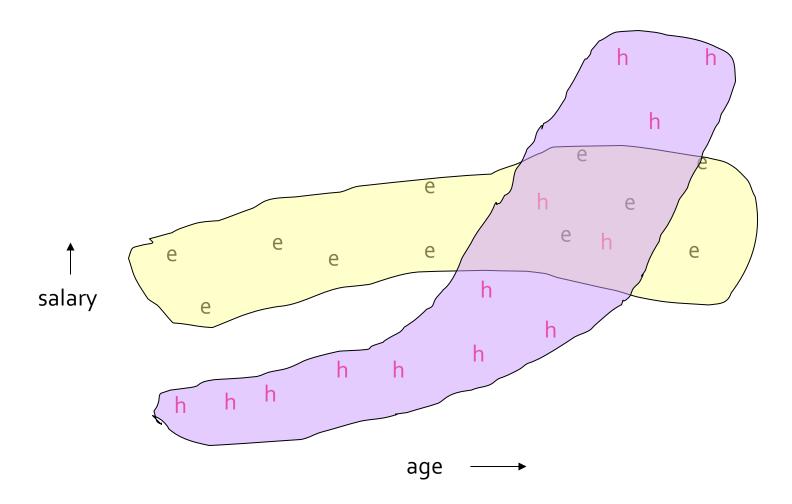
Extension of *k*-means to clusters of arbitrary shapes

CURE Algorithm

CURE (Clustering Using REpresentatives):

- Assumes a Euclidean distance
- No assumption about shape of clusters
 - No need to be normally distributed in each dim
 - No need to have fixed axis
- Instead of centroid, uses a collection of representative points to represent clusters
- Assumes k=number of clusters is given
- In contrast, BFR and k-means assume:
 - clusters are normally distributed in each dimension
 - Axes are fixed ellipses at an angle are not OK

Example: Stanford Salaries

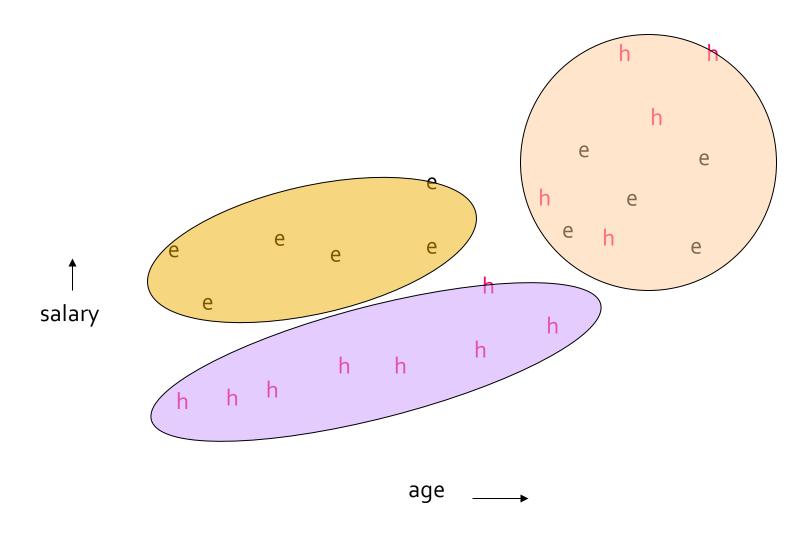


Starting CURE

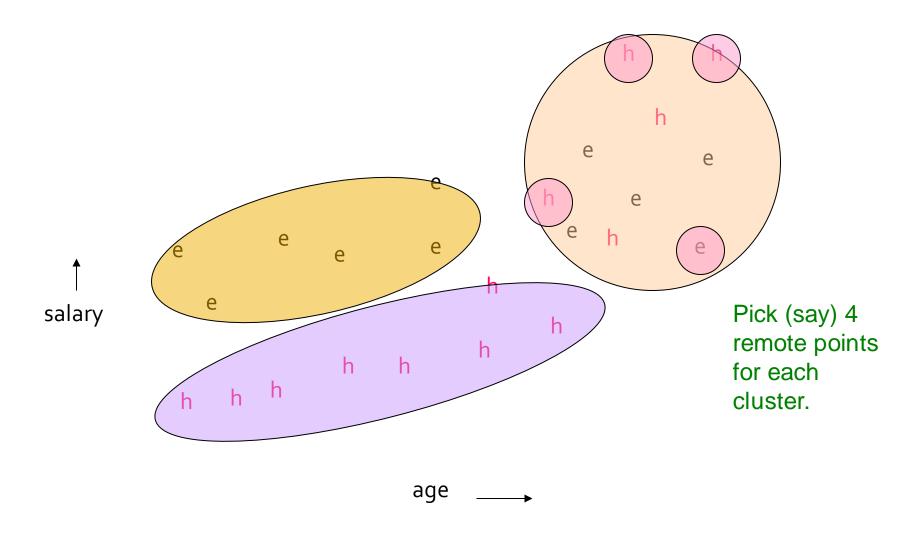
2 Pass algorithm. Pass 1:

- Pick a random sample of data and cluster them in main memory using hierarchical clustering
 - merge two clusters when they have close pair of points
- Pick representative points from each cluster
 - For each cluster, pick a sample of points, as dispersed as possible
 - move representatives a fraction of distance e.g. 20% toward the centroid of the cluster
 - Merge clusters with the closest pair of representatives

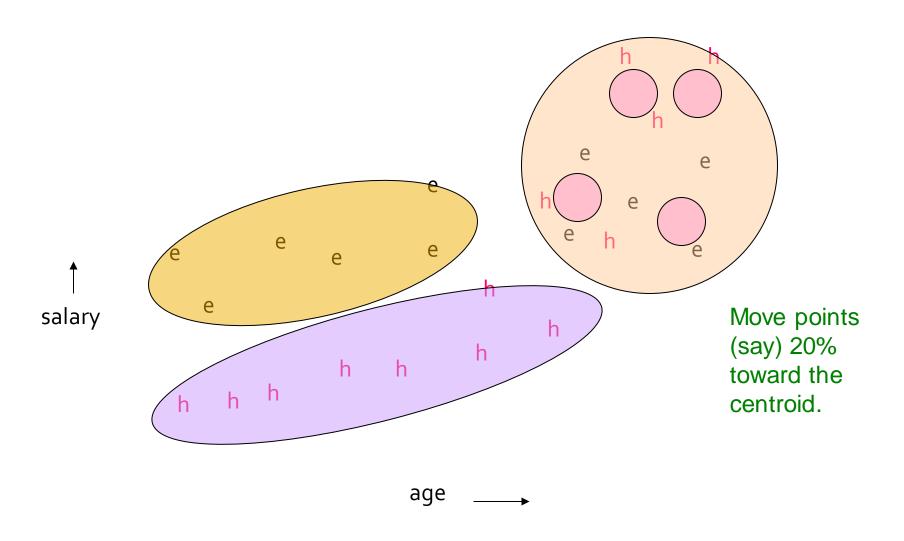
Example: Initial Clusters



Example: Pick Dispersed Points



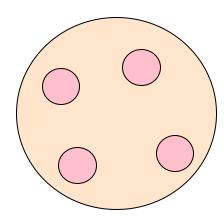
Example: Pick Dispersed Points



Finishing CURE

Pass 2:

 Now, rescan the whole dataset and visit each point p in the data set



- Place it in the "closest cluster"
 - Normal definition of "closest":
 Find the closest representative point to p and assign it to representative's cluster

n

Why the 20% Move Inward?

Intuition:

- If initial sample is large enough, some of the representatives will be on the boundary of clusters
 - Moving them towards centroid, move them inside
- A large, dispersed cluster will have larger moves as opposed to a small, dense cluster
 - Favors a small, dense cluster that is near a larger dispersed cluster

Summary

- Clustering: Given a set of points, with a notion of distance between points, group the points into some number of clusters
- Algorithms:
 - Agglomerative hierarchical clustering:
 - Centroid and clustroid
 - k-means:
 - Initialization, picking k
 - BFR
 - CURE