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Clustering

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University of Southern California

Mining of Massive Datasets Jure Leskovec, Anand Rajaraman, Jeff Ullman Stanford University

http://www.mmds.org



High Dimensional Data

Given a cloud of data points we want to understand its structure



The Problem of Clustering

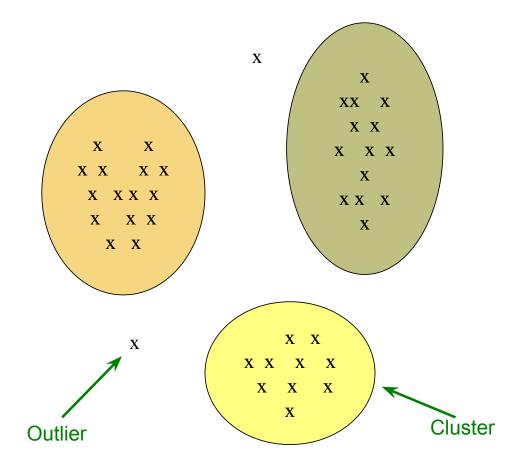
- Given a set of points, with a notion of distance between points, group the points into some number of clusters, so that
 - Members of a cluster are close/similar to each other
 - Members of different clusters are dissimilar

Usually:

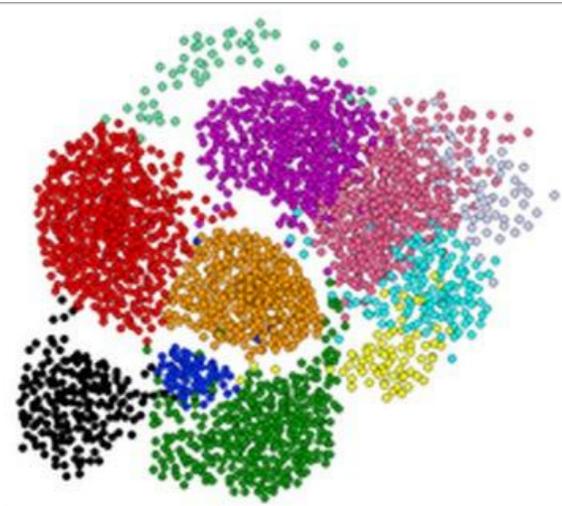
- Points are in a high-dimensional space
- Similarity is defined using a distance measure
 - Euclidean, Cosine, Jaccard, edit distance, ...

so similarity becomes hard to judge.

Example: Clusters & Outliers



Clustering is a hard problem!

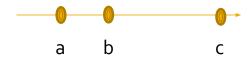


Why is it hard?

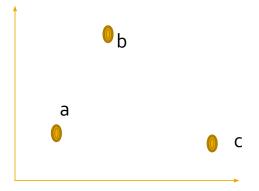
- Clustering in two dimensions looks easy
- Clustering small amounts of data looks easy
- Many applications involve not 2, but 10 or 10,000 dimensions
- High-dimensional spaces look different: Almost all pairs of points are at about the same distance

High Dimension: Euclidean

- Consider a set of data points on a line
 - dist(a, b) < dist(a, c)</p>



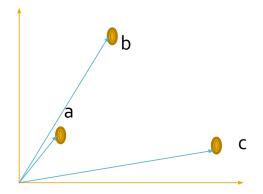
- Consider increasing the dimension by 1
 - dist(a, b) ~ dist(a, c)



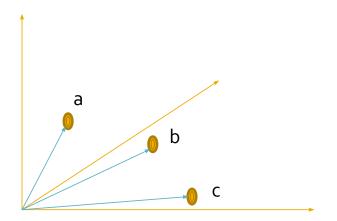
High Dimension: Cosine

Cosine(a, b) > Cosine(a, c)

- Increase d to 3
 - Cosine(a, b) ~ Cosine(a, c)

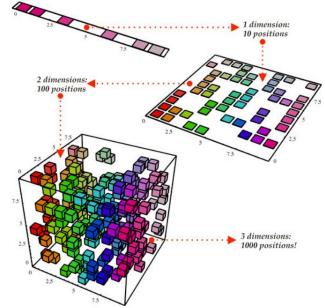


- Higher d
 - Angle -> 90°
 - Cosine -> 0



Curse of Dimensionality

- Data points have similar distance btw each other
 - Euclidean distance breaks
 - almost all pairs of points are equally far away from one another
- Data vectors become orthogonal
 - Cosine function breaks
 - almost any two vectors are orthogonal



https://bigsnarf.wordpress.com/2013/06/14/curse-of-dimensionality/

Clustering Problem: Music CDs

- Intuitively: Music divides into categories, and customers prefer a few categories
 - 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

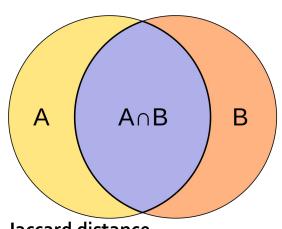
Space of all CDs:

- Think of a space with one dim. for each customer (there are many customers!)
 - Values in a dimension may be 0 or 1 only
 - A CD is a point in this space $(x_1, x_2, ..., x_k)$, where $x_i = 1$ iff the ith customer bought the CD
- For Amazon, the <u>dimension</u> is tens of millions.
- Task: Find clusters of similar CDs

Cosine, Jaccard, and Euclidean

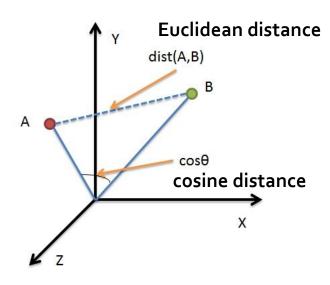
- As with CDs we have a choice when we think of documents as sets of words or shingles:
 - Sets as vectors: Measure similarity by the cosine distance
 - Sets as sets: Measure similarity by the
 Jaccard distance
 - Sets as points: Measure similarity by Euclidean distance

Measure similarity

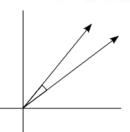


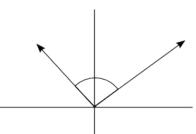
Jaccard distance

$$J(A, B) = \frac{|A \cap B|}{|A \cup B|}$$



cosine distance





Similar scores Score Vectors in same direction Angle between then is near 0 deg. Cosine of angle is near 1 i.e. 100%

Unrelated scores Score Vectors are nearly orthogonal Angle between then is near 90 deg. Cosine of angle is near 0 i.e. 0%

Opposite scores Score Vectors in opposite direction Angle between then is near 180 deg. Cosine of angle is near -1 i.e. -100%

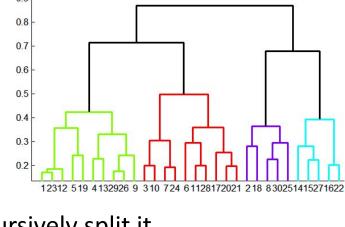
Overview: Methods of Clustering

find one then werge in

- **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

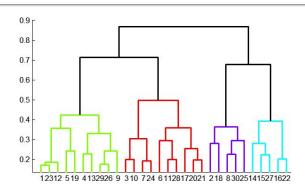
2. Point Assignment

- Maintain a set of clusters
- Points belong to "nearest" cluster



Hierarchical Clustering

Key operation:
Repeatedly combine
two nearest clusters

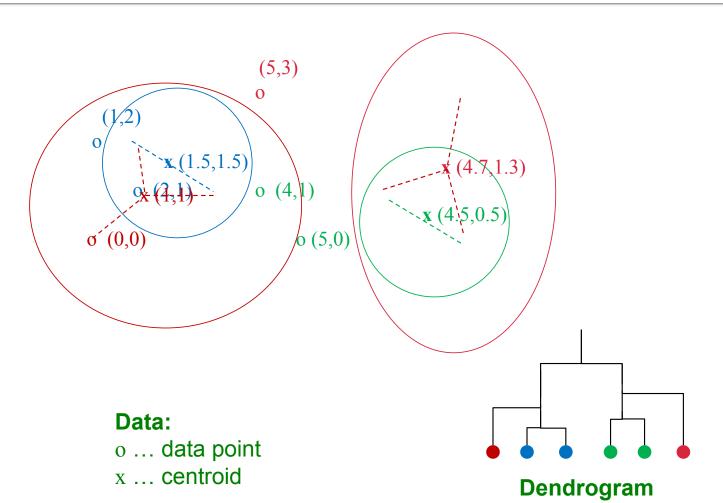


- Three important questions:
 - 1) How do you represent a cluster of more than one point?
 - 2) How do you determine the "nearness" of clusters?
 - 3) When to stop combining clusters?

Euclidean Case?

- Key operation: Repeatedly combine two nearest clusters
- (1) How to represent a cluster of many points?
 - Euclidean case: each cluster has a rest of data points
 centroid = average of its (data)points
- (2) How to determine "nearness" of clusters?
 - Measure cluster distances by distances of centroids

Example: Hierarchical clustering



And in the Non-Euclidean Case?

What about the Non-Euclidean case?

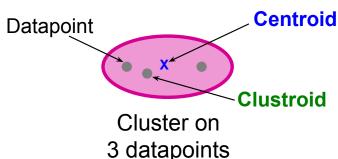
- The only "locations" we can talk about are the points themselves
 - i.e., there is no "average" of two points (e.g., students)

Approach 1:

- (1) How to represent a cluster of many points?
 clustroid = (data)point "closest" to other points
 (2) How do you determine the "nearness" of clusters?
- (2) How do you determine the "nearness" of clusters
 Treat clustroid as if it were centroid, when computing inter-cluster distances

"Closest" Point?

- (1) How to represent a cluster of many points?
 clustroid = 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: $\min_{c} \sum_{x \in C} d(x,c)^2$



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.

Defining "Nearness" of Clusters

- (2) How do you determine the "nearness" of clusters?
 - Approach 2:
 Intercluster distance = minimum of the distances between any two points, one from each cluster
 - Approach 3:
 Pick a notion of "cohesion" of clusters, e.g., maximum distance from the clustroid
 - Merge clusters whose union is most cohesive

Cohesion

- Approach 3.1: Use the diameter of the merged cluster = maximum distance between points in the cluster
- Approach 3.2: Use the average distance between points in the cluster
- Approach 3.3: Use a density-based approach
- Take the diameter or avg. distance, e.g., and divide by the number of points in the cluster

Example

- Consider a cluster of 4 points:
 - abcd, aecdb, abecb, ecdab
- Their edit distances:

aecdbabecbecdababcd335aecdb22abecb4

# 0 eg:	ed to do to change as b about, accords) => { insert: e remone: b WHEA: b	
	ecdab	1 mark: b
	5	edif_d(necdb,aberb)=2
	2	(het: 6
	4	remove: b

Determine Clusteroid

- "aecdb" will be chosen as clusteroid
 - Located in "center" judged by all 3 measures

	aecdb	abecb	ecdab
abcd	3	3	5
aecdb		2	2
abecb			4

Point	Sum	Sum-sq	Max
abcd	11	43	5
aecdb	_7	17	3
abecb	9	29	4
ecdab	11	45	5

Complexity of Hierarchical Clustering

- n data points
- At most n − 1 step of merging
- Naive implementation, e.g., storing pairwise cluster distances in a matrix

	C1	C ₂	C ₃	C ₄
C1	0	2	3	2
C ₂		0	4	5
C ₃			0	3
C ₄				0

Complexity of Naive Implementation

- Initially, O(n²) for creating matrix and finding pair with minimum distance
- Subsequent merge, assuming matrix: k x k
 - Delete columns for old clusters: O(k)
 - Add new column for new cluster C': O(k)
 - Compute dist. of C' with other clusters: O(k)
 - Find new pair of clusters with min. dist: O(k²)
- => Overall complexity: O(n³)

Implementation Summary

- Naïve implementation of hierarchical clustering:
 - At each step, compute pairwise distances between all pairs of clusters, then merge
 - $O(N^3)$
- Careful implementation using priority queue can reduce time to $O(N^2 \log N)$ (read textbook)
 - Still too expensive for really big datasets that do not fit in memory

9 measure

Sum Squared Error (SSE)

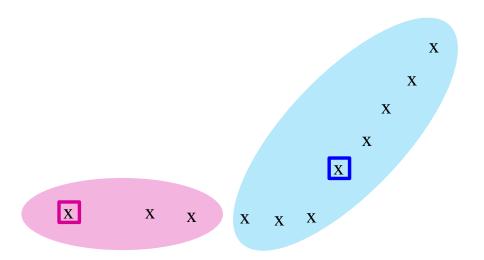
- SSE (sum squared error) is a common measure of the quality of a cluster.
 - It is the <u>sum of the squares of the distances between each of the points of the cluster and the centroid.</u>
- Sometimes, we decide to split a cluster in order to reduce the SSE.
 - Calculate the SSE before split
 - Decide the split (you can choose)
 - Calculate the sum of SSE(s) for the clusters after split
 - If SSE after < SSE before, then split
- Example: before split: (9,5), (2,2), and (4,8)
- how to split?
- should it be split?

k-means clustering

k-means Algorithm(s)

- Assumes Euclidean space/distance
- \blacksquare Start by picking k, the number of clusters
- Initialize clusters by picking one point per cluster
 - Example: Pick one point at random, then k-1 other points, each as far away as possible from the previous points

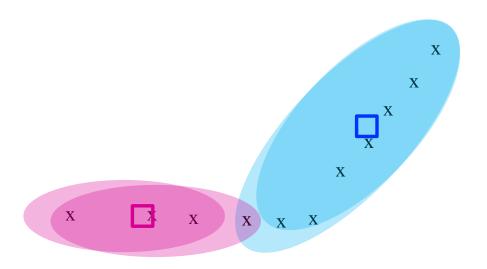
Example: Assigning Clusters



x ... data point ... centroid

Clusters after round 1

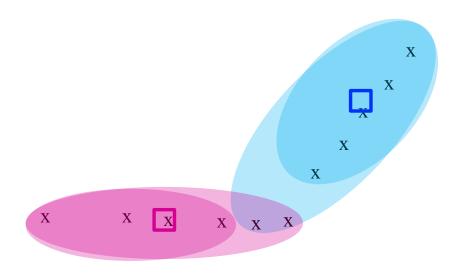
Example: Assigning Clusters



x ... data point ... centroid

Clusters after round 2

Example: Assigning Clusters



x ... data point ... centroid

Clusters at the end

Populating Clusters

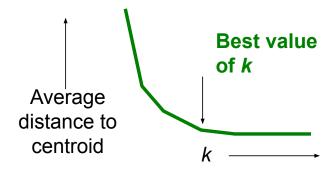
- 3. Step.
 - **1)** For each point, place it in the cluster whose current centroid it is nearest
 - 2) After all points are assigned, update the locations of centroids of the k clusters
 - 3) Reassign all points to their closest centroid
 - Sometimes moves points between clusters
 - Repeat 2 and 3 until convergence
 - Convergence: Points don't move between clusters and centroids stabilize

Getting the k right

3

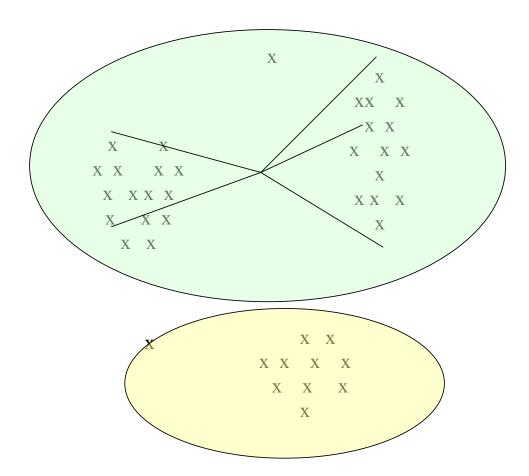
How to select k?

- 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



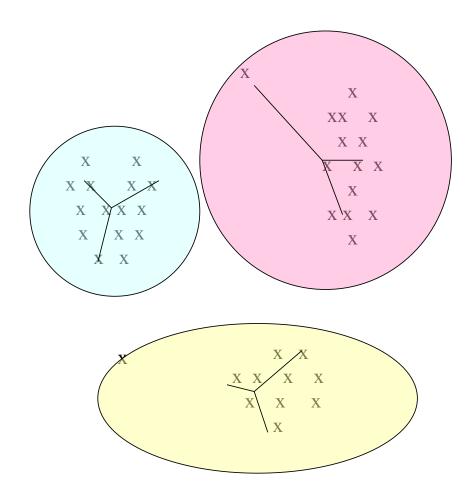
Example: Picking k

Too few; many long distances to centroid.



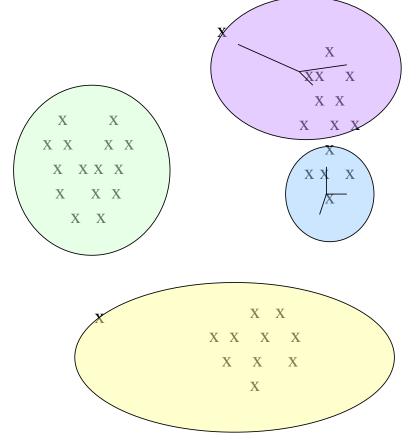
Example: Picking k

Just right; distances rather short.



Example: Picking k

Too many; little improvement in average distance.

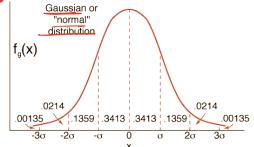


Extension of k-means to large dataBFR Algorithm

1- assumption: ().

- for uniform-distributed axis-aligned shapes

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 (2, X) of wormal distribution is aligned with

About 68% of values are within σ away from the mean; about 95% 2σ away; and about 99.7% 3σ away.

- Standard deviations in different dimensions may vary
 - Clusters are <u>axis-aligned</u> ellipses
- Efficient way to summarize clusters (want memory required O(clusters) and not O(data))

Not OK

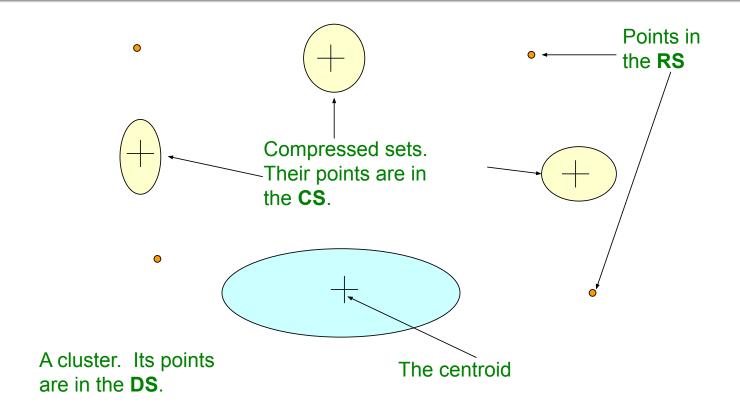
BFR Algorithm

- 2. 5tep:
 - Points are read from disk one "main-memory-full" or "memory-load" at a time
 - Most points from previous memory loads are summarized by simple statistics
 - To begin, from the initial load we select the initial k centroids by some sensible approach:
 - Take k random points band > k cluster > remain h and o > second band > update h and o > repent...
 - 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): don't wed to remember
 - Points close enough to a centroid to be summarized
 - Compression 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



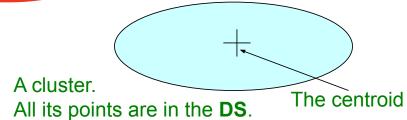
Discard set (DS): Close enough to a centroid to be summarized **Compression set (CS):** Summarized, but not assigned to a cluster **Retained set (RS):** Isolated points

Summarizing Sets of Points

4

For each cluster, the <u>discard set (DS)</u> is <u>summarized</u> by:

- The number of points, N
- The vector \overline{SUM} , whose i^{th} component is the sum of the coordinates of the points in the i^{th} dimension
- The vector \underline{SUMSQ} \underline{i}^{th} component = sum of squares of coordinates in \underline{i}^{th} dimension



Variance

$$Var(X) = E\left[\left(X - E(X)\right)^{2}\right]$$

$$= E\left[X^{2} - 2XE\left[X\right] + \left(E\left[X\right]\right)^{2}\right]$$

$$= E\left[X^{2}\right] - 2E\left[X\right]E\left[X\right] + \left(E\left[X\right]\right)^{2}$$

$$= E\left[X^{2}\right] - \left(E\left[X\right]\right)^{2}$$

$$= E\left[X^{2}\right] - \left(E\left[X\right]\right)^{2}$$

$$Var(X) = \frac{1}{n}\sum_{i=1}^{n}(x_{i} - \mu)^{2}$$

$$\mu = \frac{1}{n}\sum_{i=1}^{n}x_{i}.$$

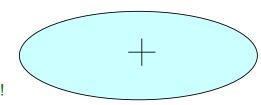
Summarizing Points: Comments

- 2d + 1 values represent any size cluster
 - d = number of dimensions
- Average in each dimension (the centroid) can be calculated as SUM, N
 - **SUM**_i = i^{th} component of SUM

$$\sigma^2 = \frac{\sum X^2}{N} - \mu^2$$

- 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

Processing the "Memory-Load" of points (1):

- 1) 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
- Use any main-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

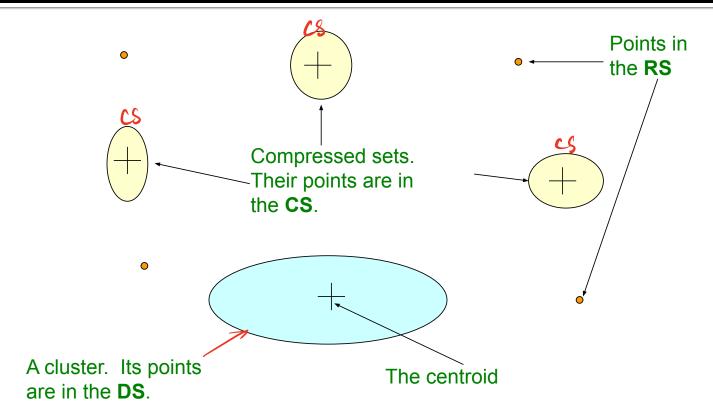
The "Memory-Load" of Points

Processing the "Memory-Load" of points (2):

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

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

A Few Details...

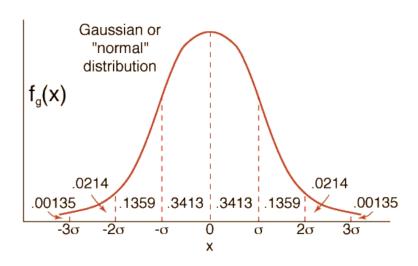
b

- 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?

(D

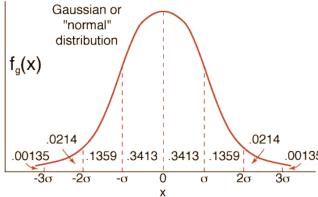
Define "Sufficiently Close"

- ~68% of points: 1 σ away from mean
- ~95% of points: 2 \sigma away
- ~99% of points: 3 σ away



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 like lihood of the point belonging to currently nearest centroid



Mahalanobis Distance

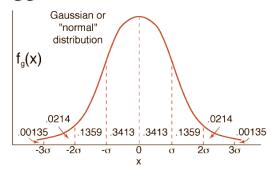
- Normalized Euclidean distance from centroid
- For point $(x_1, ..., x_d)$ and centroid $(c_1, ..., c_d)$
 - 1. Normalize by σ_i in each dimension: $y_i = (x_i c_i) / \sigma_i$
 - Take sum of the squares of the y_i
 - Take the square root

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

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

Mahalanobis Distance

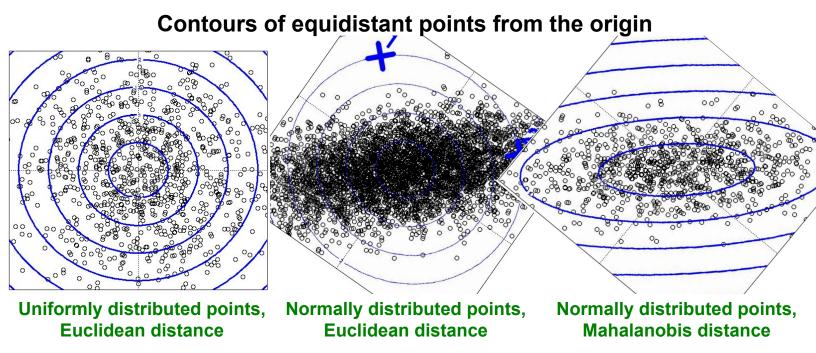
- If clusters are normally distributed in d dimensions, then after transformation, one standard deviation = \sqrt{d}
 - i.e., 68% of the points of the cluster will have a Mahalanobis distance $<\sqrt{d}$
- Accept a point for a cluster if its M.D. is < some threshold, e.g. 2 standard deviations



About 68% of values are within σ away from the mean;
 about 95% 2σ away; and about 99.7% 3σ away.

Picture: Equal M.D. Regions

Euclidean vs. Mahalanobis distance



Should two CS clusters be combined?

Q2) Should two CS subclusters be combined?

- Compute the variance of the combined subclusters
 - 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



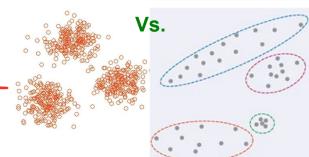




The CURE Algorithm - for arbitrary shapes

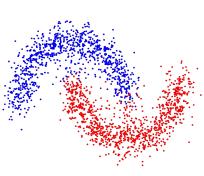


- Assumes clusters are normally distributed in each dimension
- And axes are fixed ellipses at an angle are **not OK**

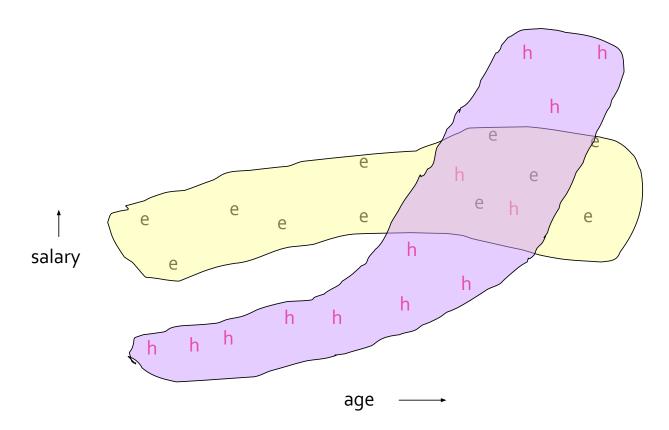


CURE (Clustering Using REpresentatives):

- Assumes a Euclidean distance
- Allows clusters to assume any shape
- Uses a collection of representative points to represent clusters



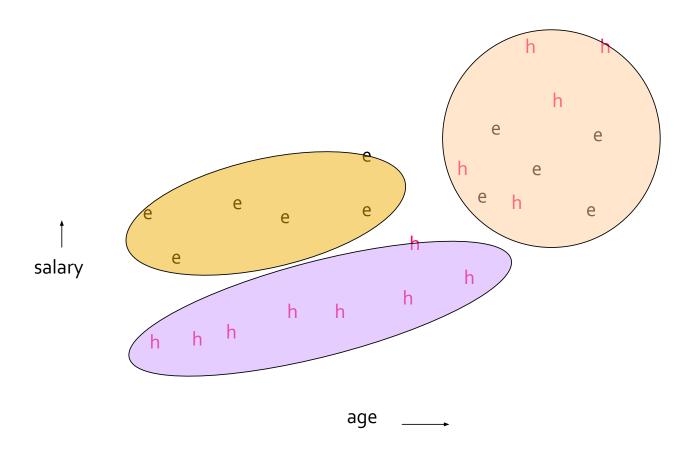
Example: Stanford Salaries



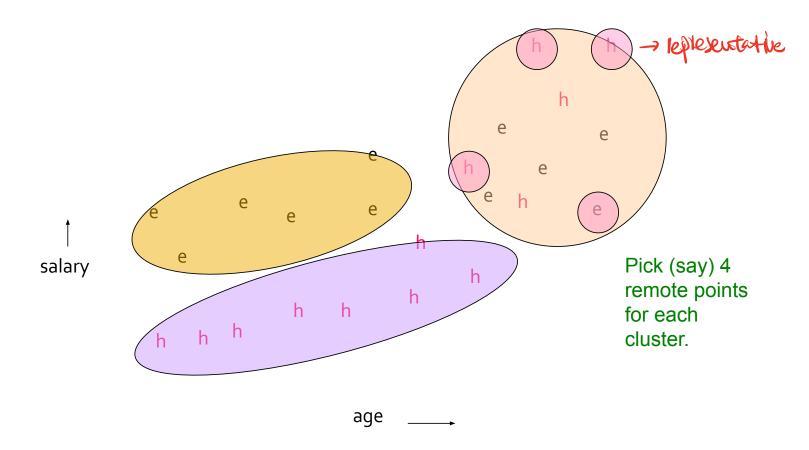
Starting CURE

- Two-Pass Algorithm: Pass 1
 - 0) Pick <u>a random sample</u> of points that fit in main memory
 - 1) Initial clusters:
 - Cluster these points hierarchically group nearest points/clusters
 - 2) Pick representative points:
 - For each cluster, pick a sample of points, as dispersed as possible
 - From the sample, pick representatives by moving them (say) 20% toward the centroid of the cluster

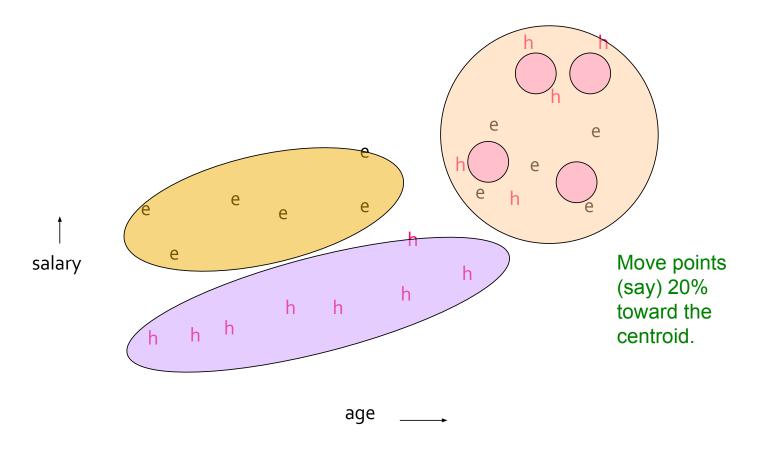
Example: Initial Clusters



Example: Pick Dispersed Points



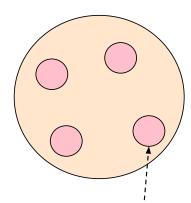
Example: Pick Dispersed Points



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": we is tention but the closest representative to p and assign it to representative's cluster.

Compare CURE with BFR

- Distribution of data
 - CURE: do not assume any particular distribution
 - BFR: data should be normally distributed
- Representation of cluster
 - CURE: a set of representatives
 - BFR: centroid
- Common: both assume data in Euclidean space

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

Standardization vs. Normalization

- Standardization vs. Normalization
- Incomparable units
 - "Age" vs. "GPA"
- Same units, irrelevant features
 - commuting miles vs. maximum jogging miles
- Same units, similar features
 - air quality values today vs. air quality tomorrow
 - length of right vs. left arms
- Others