Clustering Algorithms

Hierarchical Clustering

k - Means Algorithms

CURE Algorithm

Methods of Clustering

- Hierarchical (Agglomerative):
 - Initially, each point in cluster by itself.
 - Repeatedly combine the two "nearest" clusters into one.
- Point Assignment:
 - Maintain a set of clusters.
 - Place points into their "nearest" cluster.

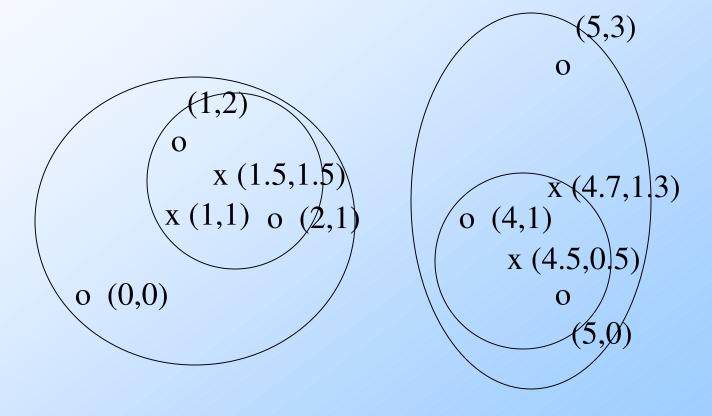
Hierarchical Clustering

- Two important questions:
 - 1. How do you determine the "nearness" of clusters?
 - 2. How do you represent a cluster of more than one point?

Hierarchical Clustering – (2)

- Key problem: as you build clusters, how do you represent the location of each cluster, to tell which pair of clusters is closest?
- Euclidean case: each cluster has a centroid = average of its points.
 - Measure intercluster distances by distances of centroids.

Example



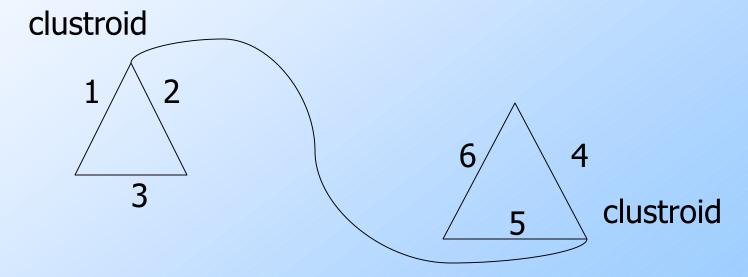
And in the Non-Euclidean Case?

- The only "locations" we can talk about are the points themselves.
 - I.e., there is no "average" of two points.
- Approach 1: clustroid = point "closest" to other points.
 - Treat clustroid as if it were centroid, when computing intercluster distances.

"Closest" Point?

- Possible meanings:
 - 1. Smallest maximum distance to the other points.
 - 2. Smallest average distance to other points.
 - 3. Smallest sum of squares of distances to other points.
 - 4. Etc., etc.

Example



intercluster distance

Other Approaches to Defining "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 1: Use the *diameter* of the merged cluster = maximum distance between points in the cluster.
- Approach 2: Use the average distance between points in the cluster.

Cohesion -(2)

- ◆ Approach 3: Use a density-based approach: take the diameter or average distance, e.g., and divide by the number of points in the cluster.
 - Perhaps raise the number of points to a power first, e.g., square-root.

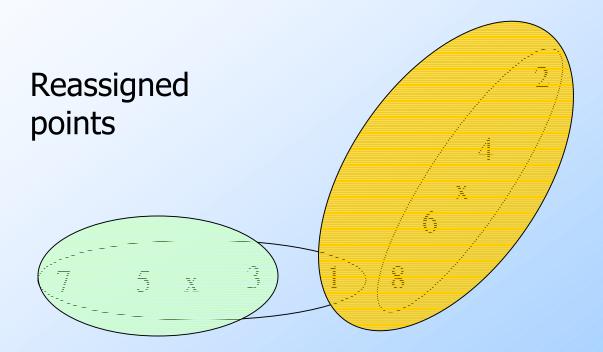
k – Means Algorithm(s)

- Assumes Euclidean space.
- 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.

Populating Clusters

- 1. For each point, place it in the cluster whose current centroid it is nearest.
- 2. After all points are assigned, fix the centroids of the *k* clusters.
- 3. Optional: reassign all points to their closest centroid.
 - Sometimes moves points between clusters.

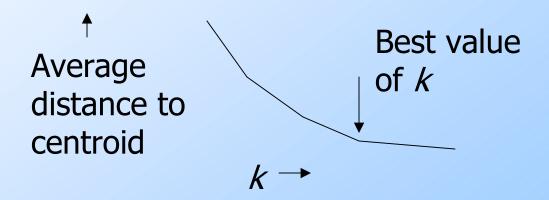
Example: Assigning Clusters



Clusters after first round

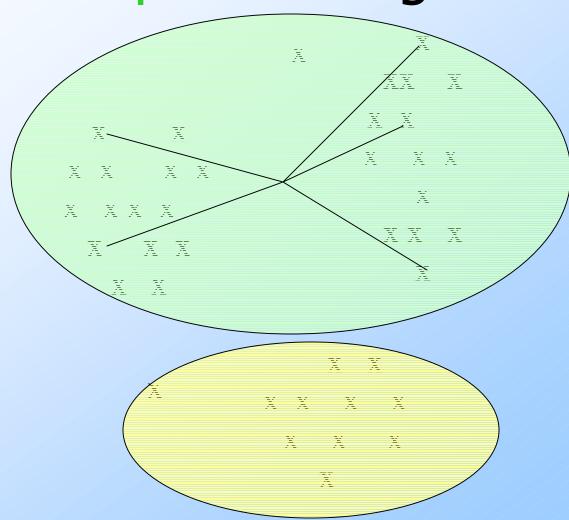
Getting k Right

- 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.



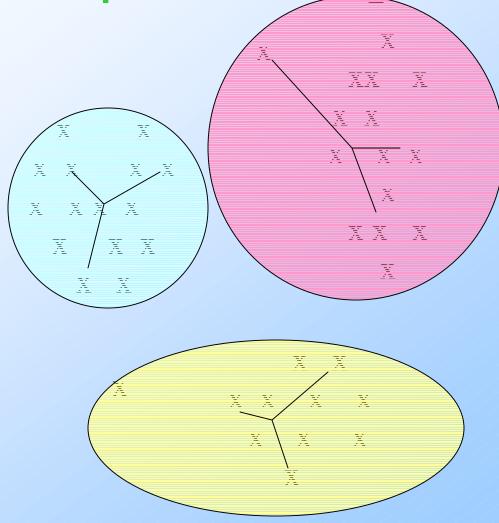
Example: Picking k

Too few; many long distances to centroid.



Example: Picking k

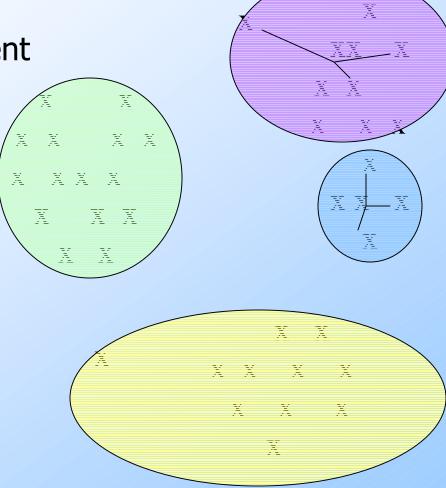
Just right; distances rather short.



Example: Picking k

Too many; little improvement

in average distance.



BFR Algorithm

- ◆BFR (Bradley-Fayyad-Reina) is a variant of k-means designed to handle very large (disk-resident) data sets.
- It assumes that clusters are normally distributed around a centroid in a Euclidean space.
 - Standard deviations in different dimensions may vary.

BFR - (2)

- Points are read one main-memory-full 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.

Initialization: k-Means

- Possibilities include:
 - Take a small random sample and cluster optimally.
 - 2. 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

- 1. The *discard set*: points close enough to a centroid to be summarized.
- 2. The *compression set*: groups of points that are close together but not close to any centroid. They are summarized, but not assigned to a cluster.
- 3. The *retained set*: isolated points.

Summarizing Sets of Points

- For each cluster, the discard set is summarized by:
 - 1. The number of points, N.
 - 2. The vector SUM, whose *i*th component is the sum of the coordinates of the points in the *i*th dimension.
 - 3. The vector SUMSQ: i^{th} component = sum of squares of coordinates in i^{th} dimension.

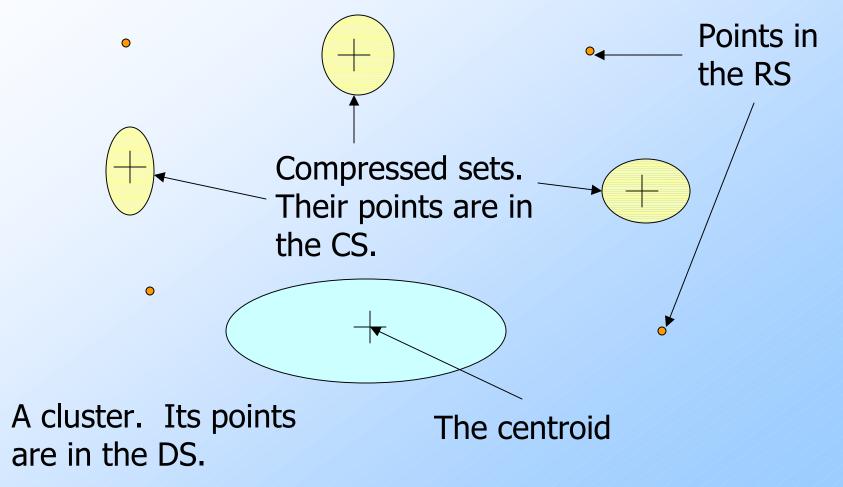
Comments

- ◆2d + 1 values represent any number of points.
 - d = number of dimensions.
- Averages in each dimension (centroid coordinates) can be calculated easily as SUM_i/N.
 - $SUM_i = i^{th}$ component of SUM.

Comments -(2)

- ◆Variance of a cluster's discard set in dimension i can be computed by: (SUMSQ_i/N) – (SUM_i/N)²
- And the standard deviation is the square root of that.
- The same statistics can represent any compression set.

"Galaxies" Picture



Processing a "Memory-Load" of Points

- 1. Find those points that are "sufficiently close" to a cluster centroid; add those points to that cluster and the DS.
- 2. 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.

Processing -(2)

- 3. Adjust statistics of the clusters to account for the new points.
- 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.

A Few Details . . .

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

How Close is Close Enough?

- We need a way to decide whether to put a new point into a cluster.
- BFR suggest two ways:
 - 1. The *Mahalanobis distance* is less than a threshold.
 - 2. Low likelihood of the currently nearest centroid changing.

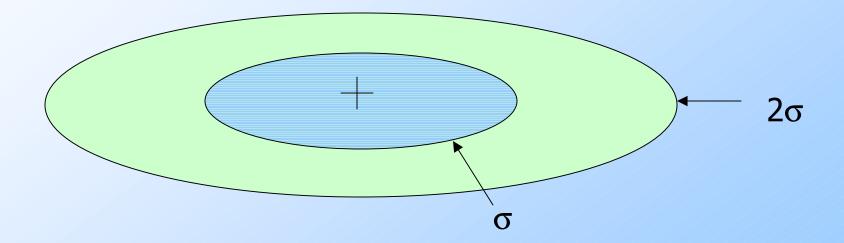
Mahalanobis Distance

- Normalized Euclidean distance.
- For point $(x_1,...,x_k)$ and centroid $(c_1,...,c_k)$:
 - 1. Normalize in each dimension: $y_i = (x_i c_i)/\sigma_i$
 - 2. Take sum of the squares of the y_i 's.
 - 3. Take the square root.

Mahalanobis Distance – (2)

- If clusters are normally distributed in d dimensions, then after transformation, one standard deviation = \sqrt{d} .
 - I.e., 70% 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. 4 standard deviations.

Picture: Equal M.D. Regions



Should Two CS Subclusters Be Combined?

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

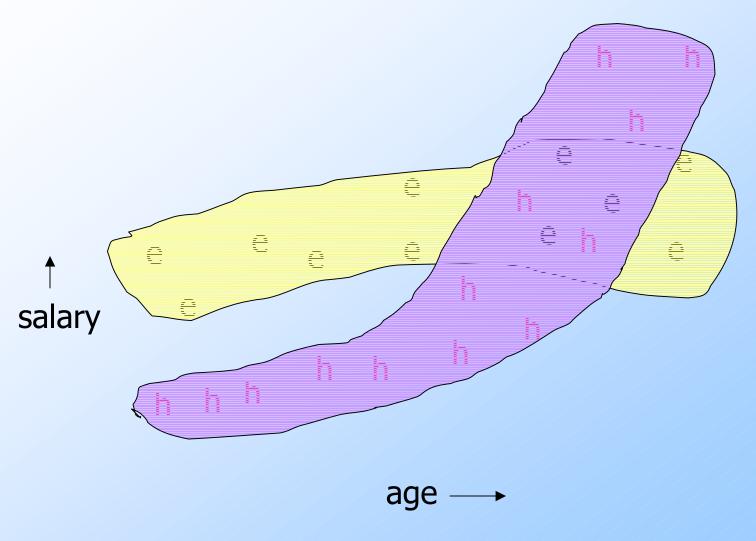
The CURE Algorithm

- Problem with BFR/k-means:
 - Assumes clusters are normally distributed in each dimension.
 - And axes are fixed ellipses at an angle are *not* OK.

CURE:

- Assumes a Euclidean distance.
- Allows clusters to assume any shape.

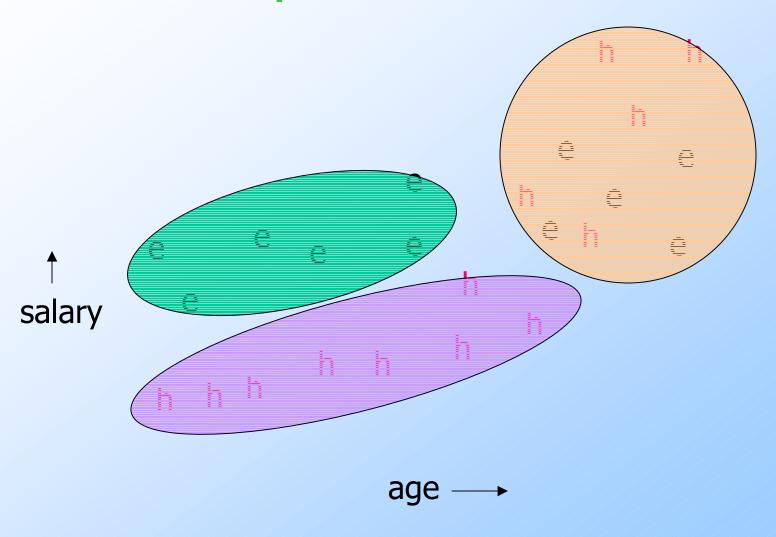
Example: Stanford Faculty Salaries



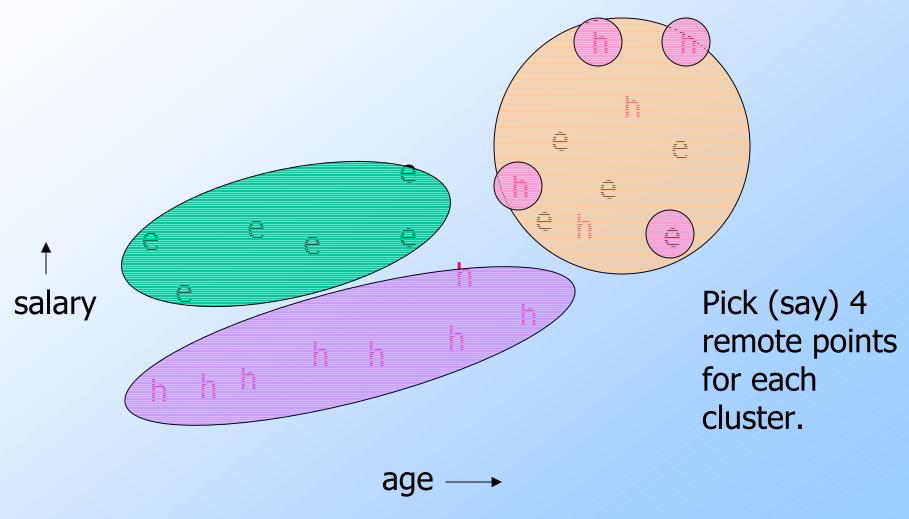
Starting CURE

- 1. Pick a random sample of points that fit in main memory.
- 2. Cluster these points hierarchically group nearest points/clusters.
- 3. For each cluster, pick a sample of points, as dispersed as possible.
- 4. From the sample, pick representatives by moving them (say) 20% toward the centroid of the cluster.

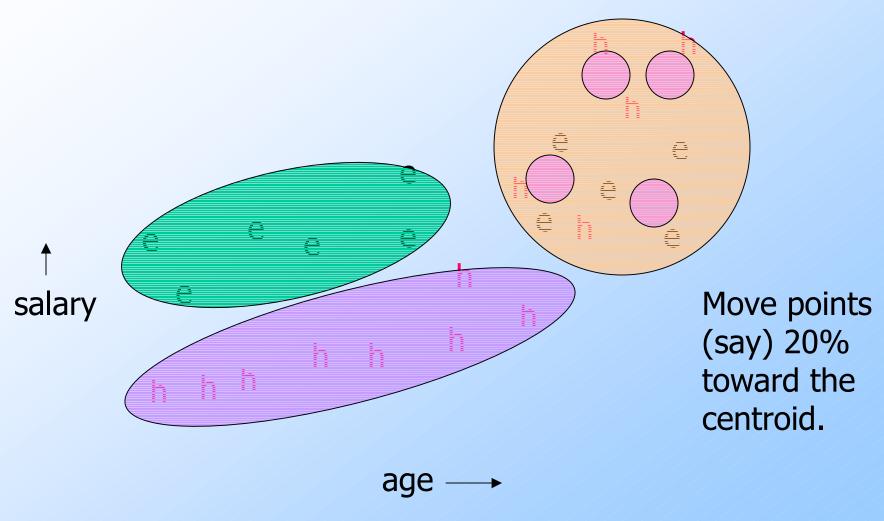
Example: Initial Clusters



Example: Pick Dispersed Points



Example: Pick Dispersed Points



Finishing CURE

- ◆Now, visit each point p in the data set.
- Place it in the "closest cluster."
 - Normal definition of "closest": that cluster with the closest (to p) among all the sample points of all the clusters.