More Clustering

CURE Algorithm Non-Euclidean Approaches

The CURE Algorithm

Problem with BFR/k -means:

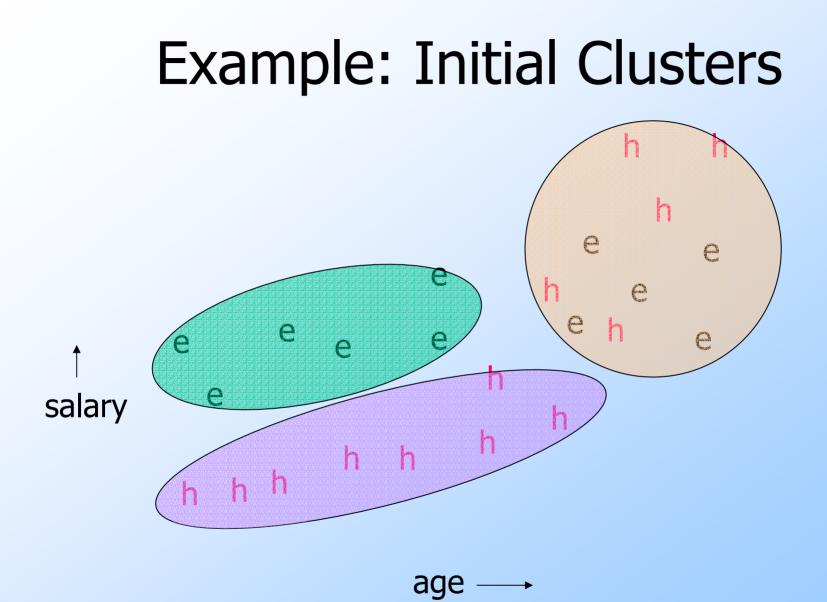
- Assumes clusters are normally distributed in each dimension.
- And axes are fixed --- ellipses at an angle are not OK.
- CURE:
 - Assumes a Euclidean distance.
 - Allows clusters to assume any shape.

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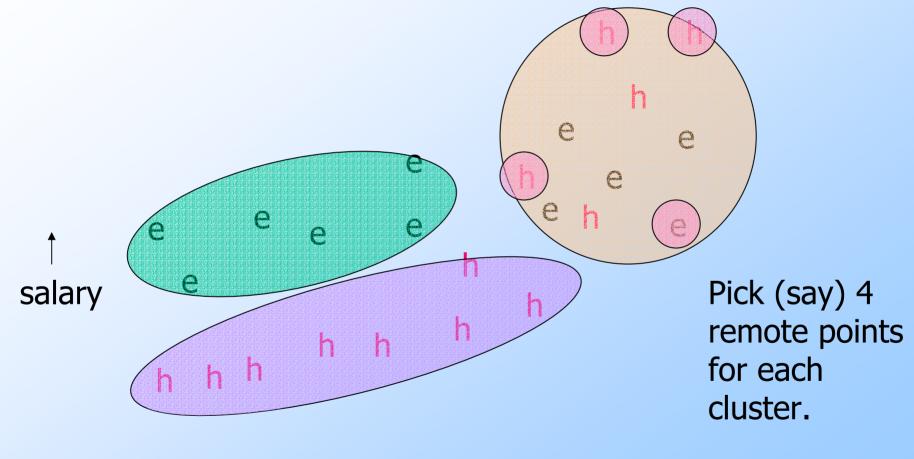
age —

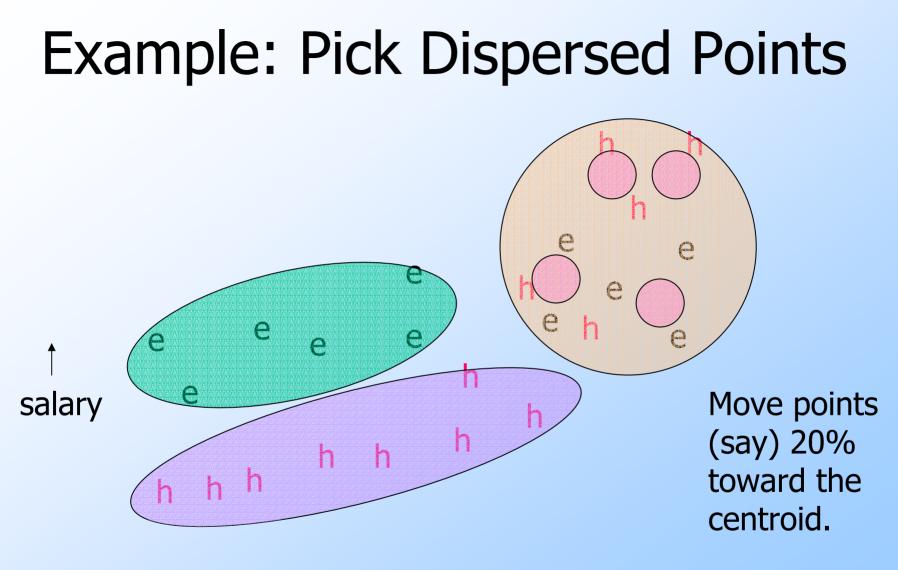
Starting CURE

- 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: 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.

Curse of Dimensionality

- One way to look at it: in largedimension spaces, random vectors are perpendicular. Why?
 - Argument #1: Lots of 2-dim subspaces.
 There must be one where the vectors' projections are almost perpendicular.
 - Argument #2: Expected value of cosine of angle is 0.

Cosine of Angle Between Random Vectors

- Assume vectors emanate from the origin (0,0,...,0).
- Components are random in range [-1,1].
 (a₁, a₂,..., a_n).(b₁, b₂,..., b_n) has expected value 0 and a standard deviation that grows as √n.
- But lengths of both vectors grow as \sqrt{n} .
- So dot product around $\sqrt{n} (\sqrt{n} * \sqrt{n}) = 1/\sqrt{n}$.

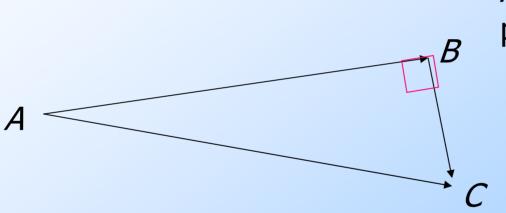
Random Vectors --- Continued

- Thus, a typical pair of vectors has an angle whose cosine is on the order of $1/\sqrt{n}$.
- As n -> ∞, that's 0; i.e., the angle is about 90°.

Interesting Consequence

- Suppose "random vectors are perpendicular," even in non-Euclidean spaces.
- Suppose we know the distance from A to B, say d (A,B), and we also know d (B,C), but we don't know d (A,C).
- Suppose B and C are fairly close, say in the same cluster.
- What is d(A,C)?

Diagram of Situation



Approximately perpendicular

Assuming points lie in a plane: $d(A,B)^2 + d(B,C)^2 = d(A,C)^2$

Important Point

- Why do we assume AB is perpendicular to AC, and not that either of the other two angles are right-angles?
 - 1. AB and AC are not "random vectors"; they each go to points that are far away from A and close to each other.
 - 2. If AB is longer than AC, then it is angle ACB that is right, but both ACB and ABC are approximately right-angles.

Dealing With a Non-Euclidean Space

- Problem: clusters cannot be represented by centroids.
- Why? Because the "average" of "points" might not be a point in the space.
- Best substitute: the *clustroid* = point in the cluster that minimizes the sum of the squares of distances to the points in the cluster.

Representing Clusters in Non-Euclidean Spaces

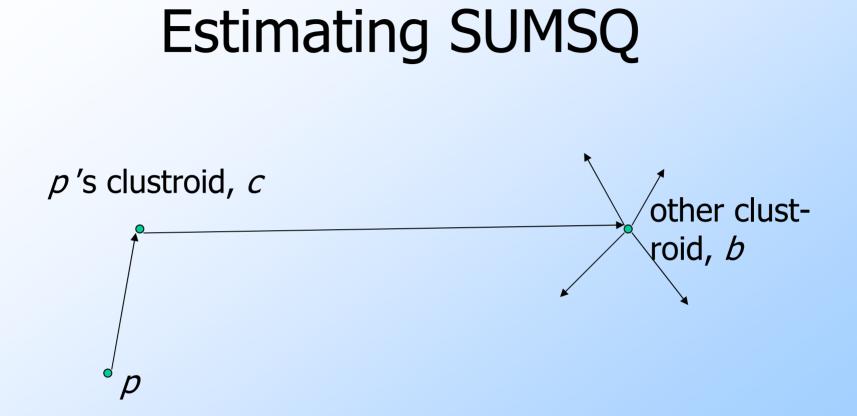
- Recall BFR represents a Euclidean cluster by *N*, SUM, and SUMSQ.
- A non-Euclidean cluster is represented by:
 - /V.
 - The clustroid.
 - Sum of the squares of the distances from clustroid to all points in the cluster.

Example of CoD Use

 Problem: in non-Euclidean space, we want to decide whether to merge two clusters.

- Each cluster represented by *N*, clustroid, and "SUMSQ."
- Also, SUMSQ for each point in the cluster, even if it is not the clustroid.

Merge if SUMSQ for new cluster is "low."



Suppose *p* Were the Clustroid of Combined Cluster

- It's SUMSQ would be the sum of:
 - 1. Old SUMSQ(*p*) [for old cluster containing *p*].
 - 2. SUMSQ(*b*) plus *d* (*p*,*b*)² times number of points in *b* 's cluster.
- Critical point: vector p -> b assumed perpendicular to vectors from b to all other points in its cluster --- justifies (2).

Combining Clusters --- Continued

We can thus estimate SUMSQ for each point in the combined cluster. Take the point with the least SUMSQ as the clustroid of the new cluster ---- provided that SUMSQ is small enough.

The GRGPF Algorithm

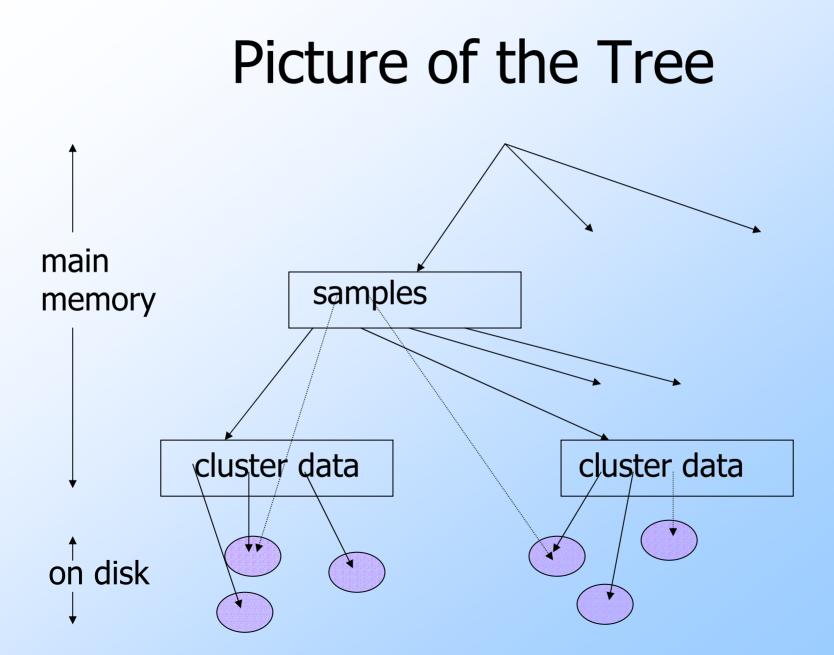
From Ganti et al. --- see reading list.
Works for non-Euclidean distances.
Works for massive (disk-resident) data.
Hierarchical clustering.
Clusters are grouped into a tree of disk blocks (like a B-tree or R-tree).

Information Retained About a Cluster

- 1. *N*, clustroid, SUMSQ.
- 2. The *p* points closest to the clustroid, and their values of SUMSQ.
- 3. The *p* points of the cluster that are furthest away from the clustroid, and their SUMSQ's.

At Interior Nodes of the Tree

- Interior nodes have samples of the clustroids of the clusters found at descendant leaves of this node.
- Try to keep clusters on one leaf block close, descendants of a level-1 node close, etc.
- Interior part of tree kept in main memory.



Initialization

Take a main-memory sample of points.
Organize them into clusters hierarchically.
Build the initial tree, with level-1 interior nodes representing clusters of clusters, and so on.

All other points are inserted into this tree.

Inserting Points

Start at the root.

- At each interior node, visit one or more children that have sample clustroids near the inserted point.
- At the leaves, insert the point into the cluster with the nearest clustroid.

Updating Cluster Data

Suppose we add point X to a cluster.
Increase count N by 1.
For each of the 2p + 1 points Y whose SUMSQ is stored, add d (X,Y)².
Estimate SUMSQ for X.

Estimating SUMSQ(X)

If C is the clustroid, SUMSQ(X) is, by the CoD assumption: Nd (X,C)² + SUMSQ(C)

- Based on assumption that vector from X to C is perpendicular to vectors from C to all the other nodes of the cluster.
- This value may allow X to replace one of the closest or furthest nodes.

Possible Modification to Cluster Data

There may be a new clustroid --- one of the p closest points --- because of the addition of X.

Eventually, the clustroid may migrate out of the p closest points, and the entire representation of the cluster needs to be recomputed.

Splitting and Merging Clusters

- Maintain a threshold for the *radius* of a cluster = $\sqrt{(SUMSQ/N)}$.
- Split a cluster whose radius is too large.
- Adding clusters may overflow leaf blocks, and require splits of blocks up the tree.
 - Splitting is similar to a B-tree.
 - But try to keep locality of clusters.

Splitting and Merging --- (2)

- The problem case is when we have split so much that the tree no longer fits in main memory.
- Raise the threshold on radius and merge clusters that are sufficiently close.

Merging Clusters

Suppose there are nearby clusters with clustroids *C* and *D*, and we want to consider merging them.

Assume that the clustroid of the combined cluster will be one of the *p* furthest points from the clustroid of one of those clusters.

Merging --- (2)

- Compute SUMSQ(X) [from the cluster of C] for the combined cluster by summing:
 - 1. SUMSQ(X) from its own cluster.
 - 2. SUMSQ(D) + $N[d(X,C)^2 + d(C,D)^2]$.
 - Uses the CoD to reason that the distance from X to each point in the other cluster goes to C, makes a right angle to D, and another right angle to the point.

Merging --- Concluded

 Pick as the clustroid for the combined cluster that point with the least SUMSQ.

- But if this SUMSQ is too large, do not merge clusters.
- Hope you get enough mergers to fit the tree in main memory.

Fastmap

Not a clustering algorithm --- rather, a method for applying *multidimensional scaling*.

 That is, mapping the points onto a smalldimension space, so the CoD does not apply.

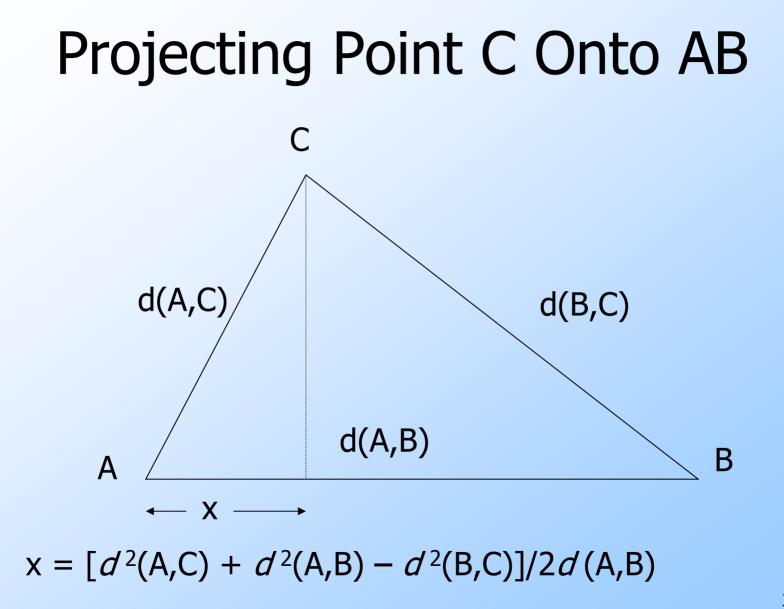
Fastmap --- (2)

Assumes non-Euclidean space.

- But like GRGFP pretends it is working in 2dimensional Euclidean space when it is convenient to do so.
- Goal: map *n* points in much less than $O(n^2)$ time.
 - I.e., you cannot compute distances between each pair of points and place points in k-dim. space to minimize error.

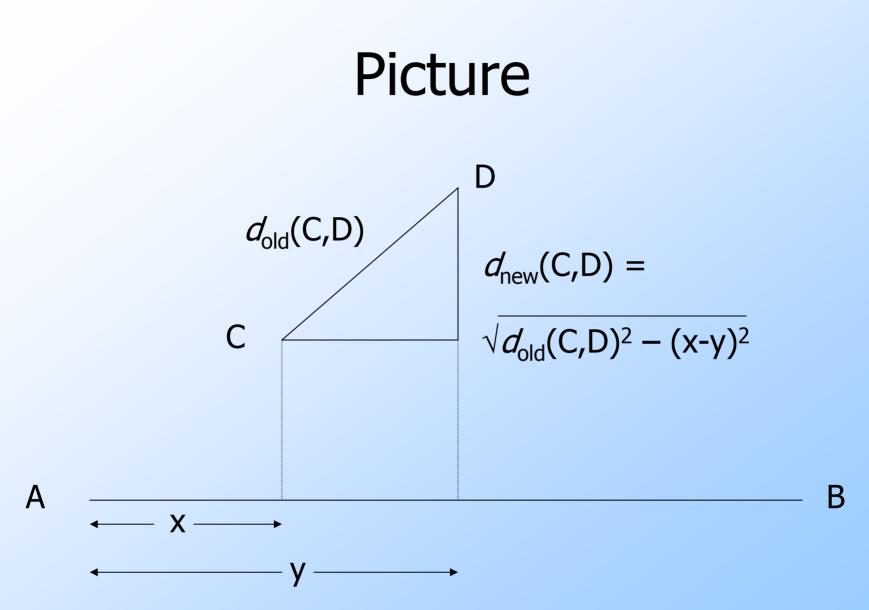
Fastmap --- Key Idea

- Create a "dimension" in non-Euclidean space by:
 - 1. Pick a pair of points A and B that are far apart.
 - Start with random A; pick most distant B.
 - 2. Treat AB as an "axis" and project all points onto AB, using the law of cosines.



Revising Distances

Having computed the position of every point along the *pseudo-axis* AB, we need to lower the distances between points in the "other dimensions."



But ...

- We can't afford to compute new distances for each pseudo-dimension.
 It would take O(n²) time.
- Rather, for each pseudo-dimension, store the position along the pseudo-axis for each point, and adjust the distance between points by square-subtract-sqrt only when needed.
 - I.e., one of the points is an axis-end.

Fastmap --- Summary

Pick a number of dimensions k. FOR i = 1 TO k DO BEGIN Pick a pseudo-axis $A_i B_i$; Compute projection of each point onto this pseudo-axis; END;

• Each step is O(ni); total $O(nk^2)$.