# Clustering

Advanced Search Techniques for Large Scale Data Analytics

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### **High Dimensional Data**

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



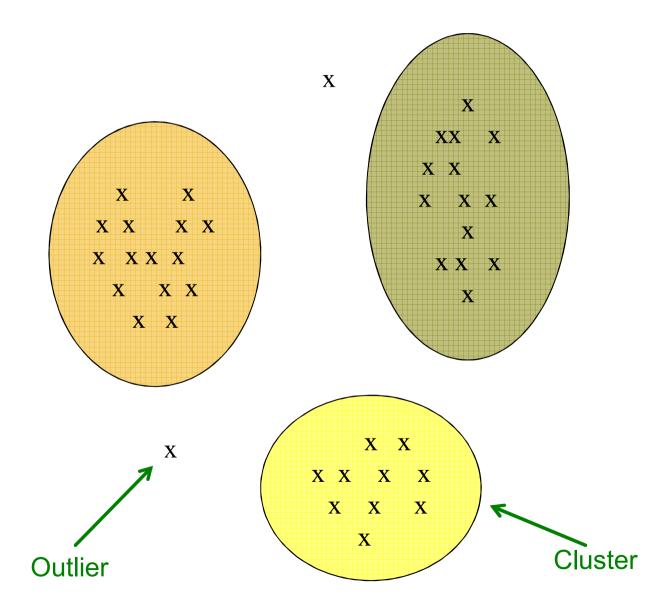
### The Problem of Clustering

- 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, ...

### **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
- 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 at about the same distance

### 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
  - 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 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

- 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

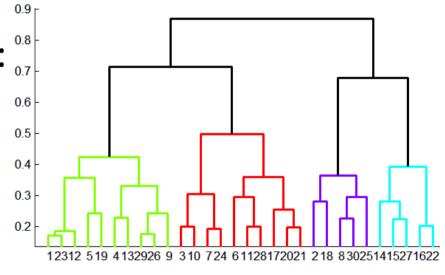
### Overview: Methods of Clustering

### Hierarchical:

- Agglomerative (bottom up): 0.8 0.7
  - 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

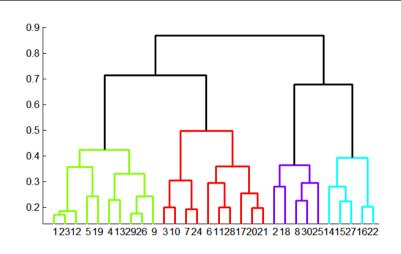
### Point assignment:

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



# **Hierarchical Clustering**

Key operation:Repeatedly combinetwo nearest clusters

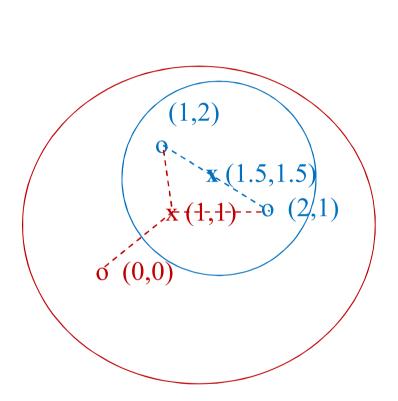


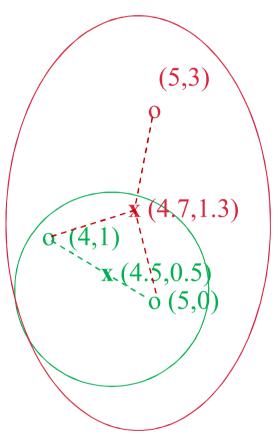
- Three important questions:
  - 1) How do you represent a cluster of more than one point?
  - 2) How do you determine the "nearness" of clusters?
  - 3) When to stop combining clusters?

### **Hierarchical Clustering**

- Key operation: Repeatedly combine two nearest clusters
- (1) How to represent a cluster of many points?
  - Key problem: As you merge 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 (data)points
- (2) How to determine "nearness" of clusters?
  - Measure cluster distances by distances of centroids

# Example: Hierarchical clustering

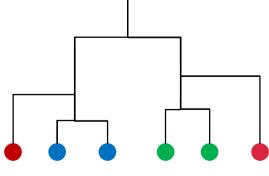




### Data:

o ... data point

x ... centroid



**Dendrogram** 

### And in the Non-Euclidean Case?

### What about the Non-Euclidean case?

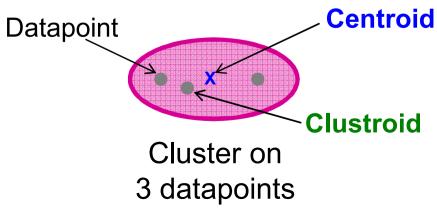
- The only "locations" we can talk about are the points themselves
  - i.e., there is no "average" of two points

### 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? 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

### 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

### Implementation

- Naïve implementation of hierarchical clustering:
  - At each step, compute pairwise distances between all pairs of clusters, then merge
  - O(N³)
- Careful implementation using priority queue can reduce time to  $O(N^2 \log N)$ 
  - Still too expensive for really big datasets that do not fit in memory

# k-means clustering

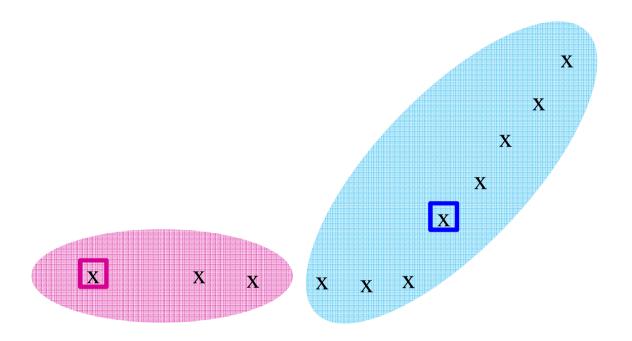
# k-means Algorithm(s)

- Assumes Euclidean space/distance
- 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, 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

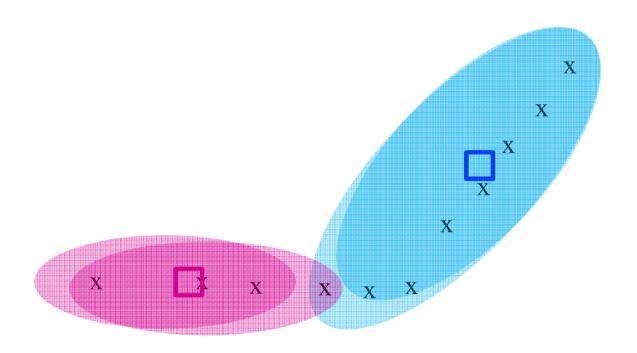
# **Example: Assigning Clusters**



x ... data point ... centroid

**Clusters after round 1** 

# **Example: Assigning Clusters**

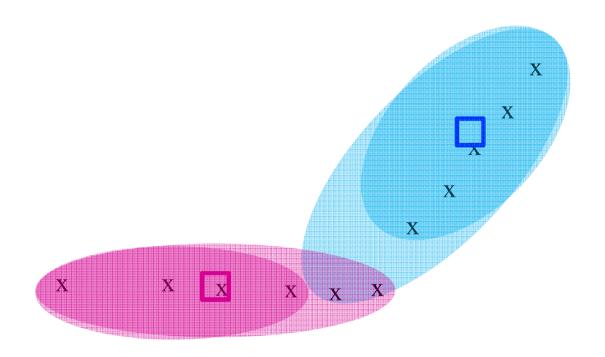


x ... data point

... centroid

### **Clusters after round 2**

# **Example: Assigning Clusters**



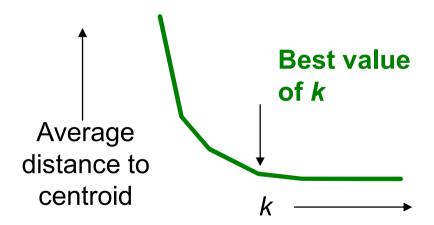
x ... data point ... centroid

Clusters at the end

# Getting the k right

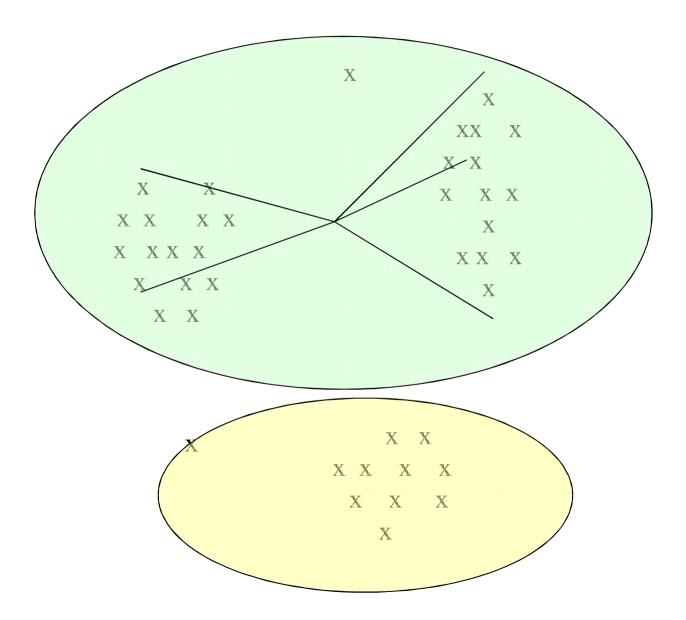
### How to select *k*?

- 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



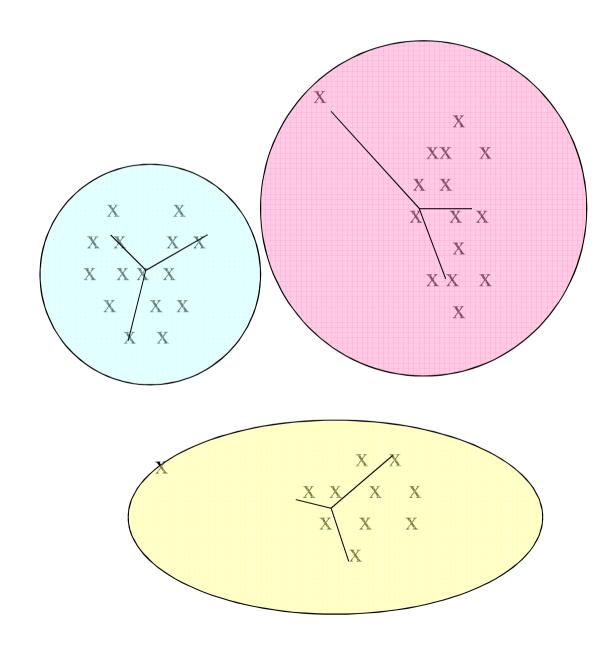
# Example: Picking k

Too few; many long distances to centroid.



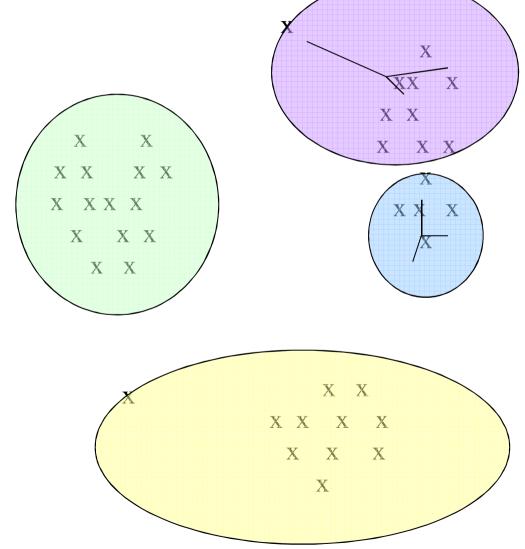
# Example: Picking k

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**

- Gaussian or "normal" distribution  $f_g(x) \\ 0.0214 \\ .00135 \\ .1359 \\ .3413 \\ .3413 \\ .1359 \\ .00135 \\ .00135$
- 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
- Efficient way to summarize clusters
   (want memory required O(clusters) and not O(data))

# **BFR Algorithm**

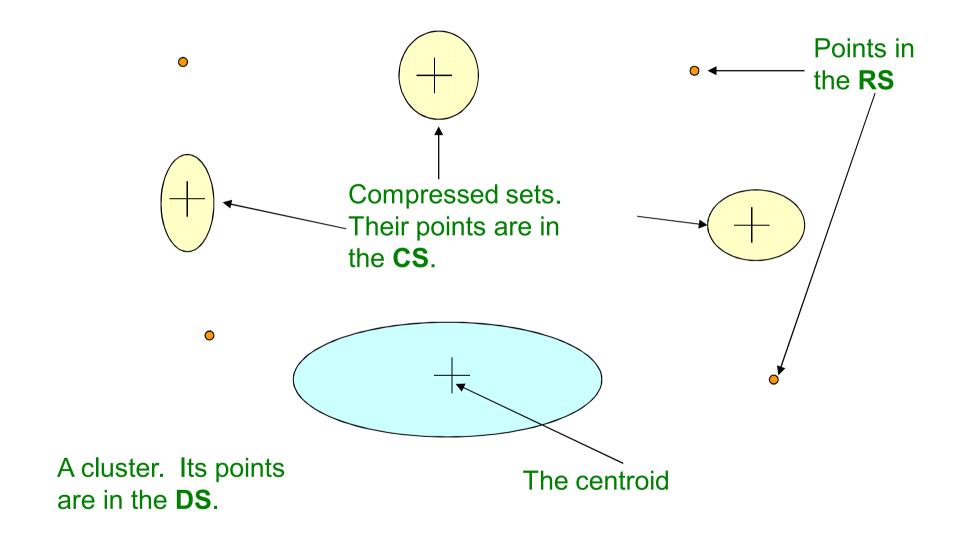
- Points are read from disk one main-memoryfull at a time
- Most points from previous memory loads are summarized by simple statistics
- To begin, from the initial load we select the initial k centroids by some sensible approach:
  - Take k random points
  - Take a small random sample and cluster optimally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

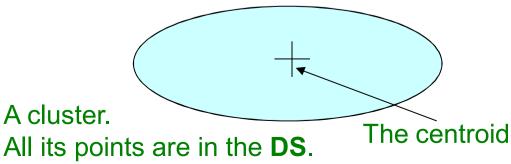


**Discard set (DS):** Close enough to a centroid to be summarized **Compression set (CS):** Summarized, but not assigned to a cluster **Retained set (RS):** Isolated points

### **Summarizing Sets of Points**

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

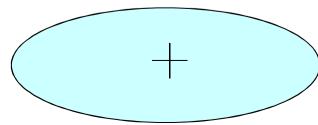
- The number of points, N
- The vector SUM, whose  $i^{th}$  component is the sum of the coordinates of the points in the  $i^{th}$  dimension
- The vector SUMSQ:  $i^{th}$  component = sum of squares of coordinates in  $i^{th}$  dimension



### **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<sub>i</sub> / N
  - $SUM_i = i^{th}$  component of SUM
- Variance of a cluster's discard set in dimension i is: (SUMSQ<sub>i</sub> / N) – (SUM<sub>i</sub> / N)<sup>2</sup>
  - 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
- 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

**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, add all compressed sets in the **CS** and all **RS** points into their nearest cluster

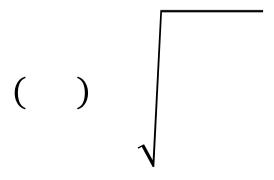
**Discard set (DS):** Close enough to a centroid to be summarized. **Compression set (CS):** Summarized, but not assigned to a cluster **Retained set (RS):** Isolated points

### 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)
  - Using the Mahalanobis distance accept a point for a cluster if its M.D. is < some threshold, e.g.  $2\sqrt{\phantom{a}}$ 
    - If clusters are normally distributed in d dimensions, then after transformation, one standard deviation =  $\sqrt{}$  i.e., 68% of the points of the cluster will have a Mahalanobis distance
  - For point  $(x_1, ..., x_d)$  and 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

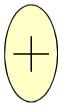


 $\sigma_i$  ... standard deviation of points in the cluster in the  $i^{th}$  dimension

### Should 2 CS clusters be combined?

#### Q2) Should 2 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 combined variance is below some threshold





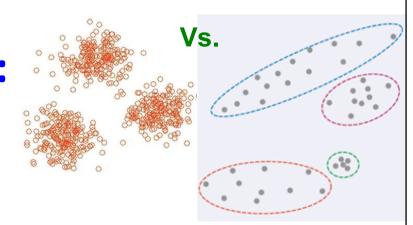
# The CURE Algorithm

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

## The CURE Algorithm

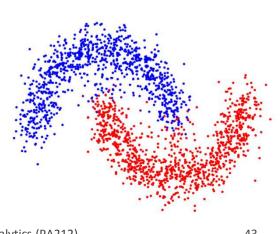
#### Problem with BFR/k-means:

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

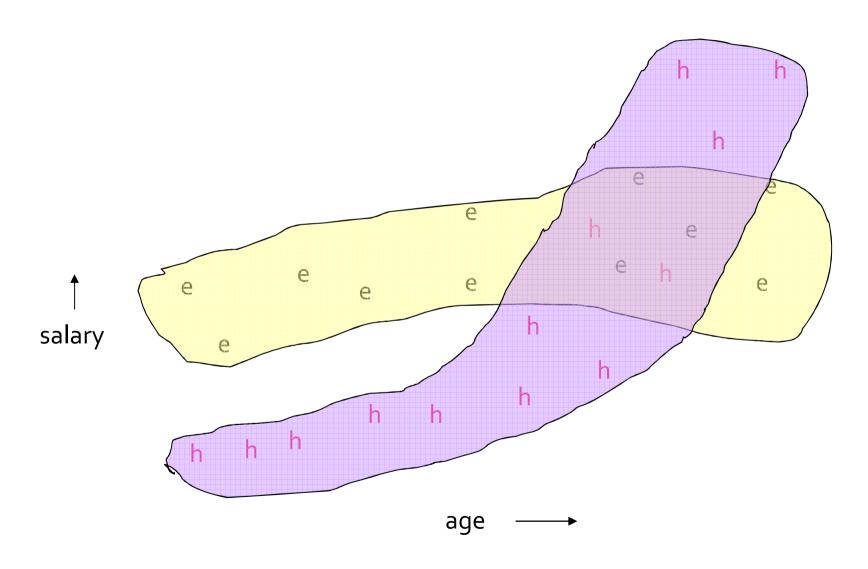


#### CURE (Clustering Using REpresentatives):

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



## **Example: Stanford Salaries**

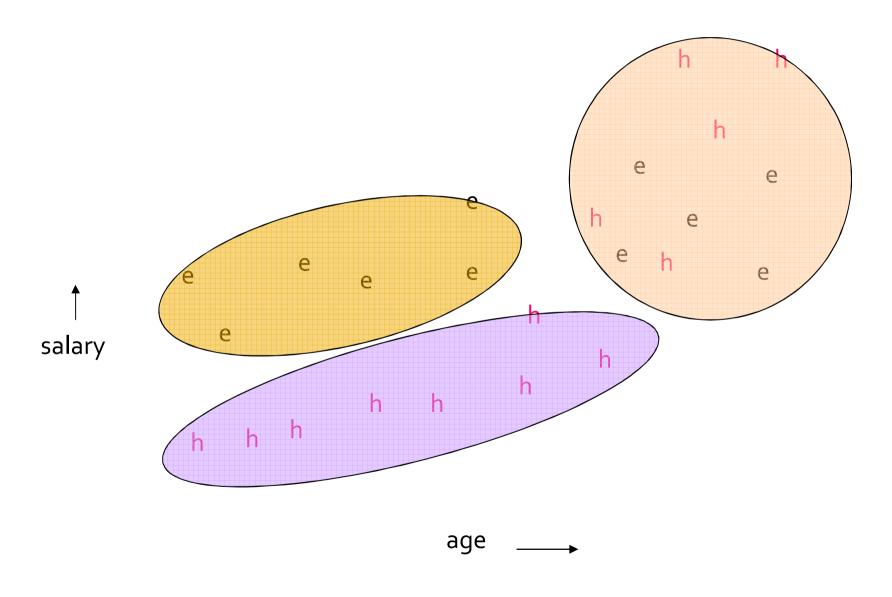


## **Starting CURE**

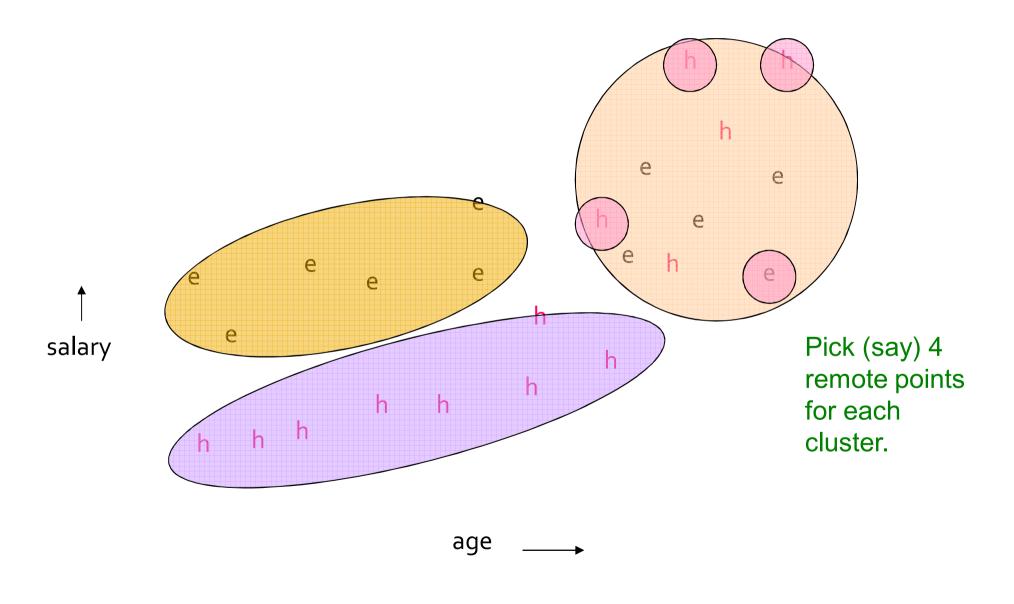
#### 2 Pass algorithm. Pass 1:

- 0) Pick a random sample 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

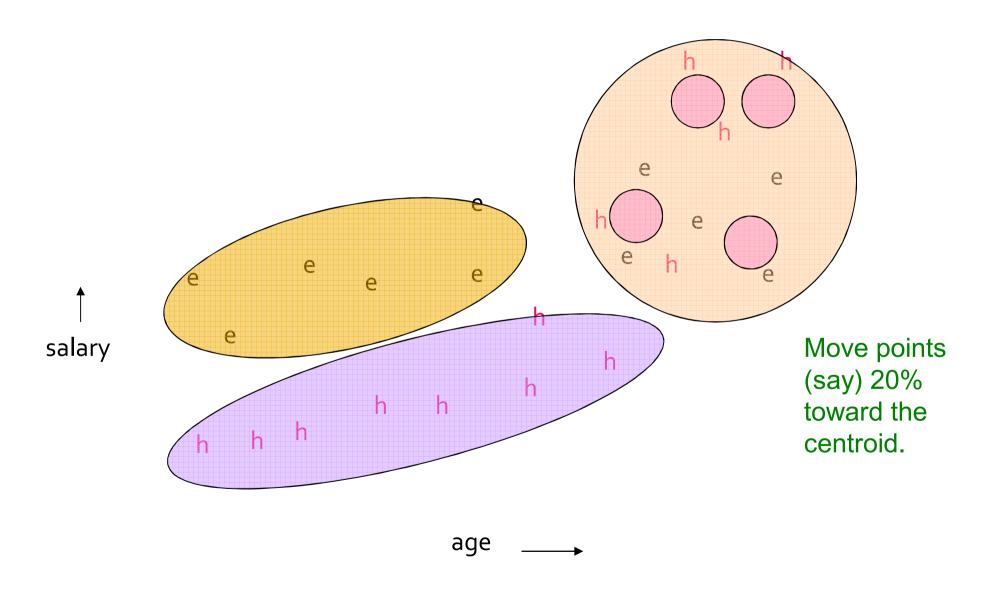
# **Example: Initial Clusters**



# **Example: Pick Dispersed Points**



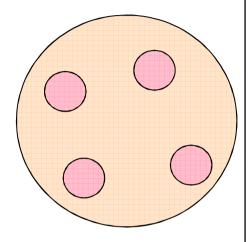
# Example: Pick Dispersed Points



## 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":
     Find the closest representative to p and assign it to representative's cluster

a

### 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