

# Clustering

Advanced Search Techniques for Large Scale Data Analytics

Pavel Zezula and Jan Sedmidubsky

Masaryk University

<http://disa.fi.muni.cz>

# High Dimensional Data

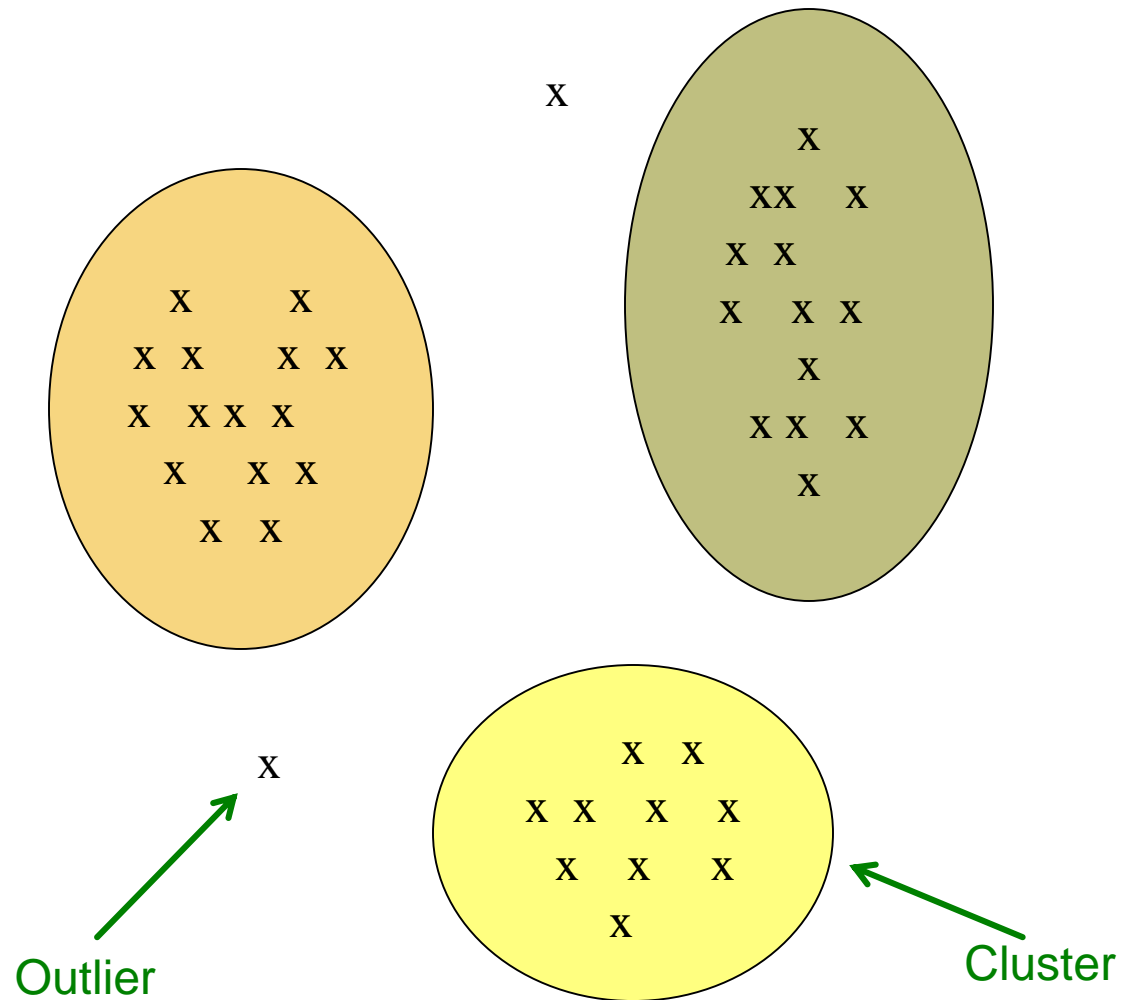
- Given a cloud of data points we want to understand its structure



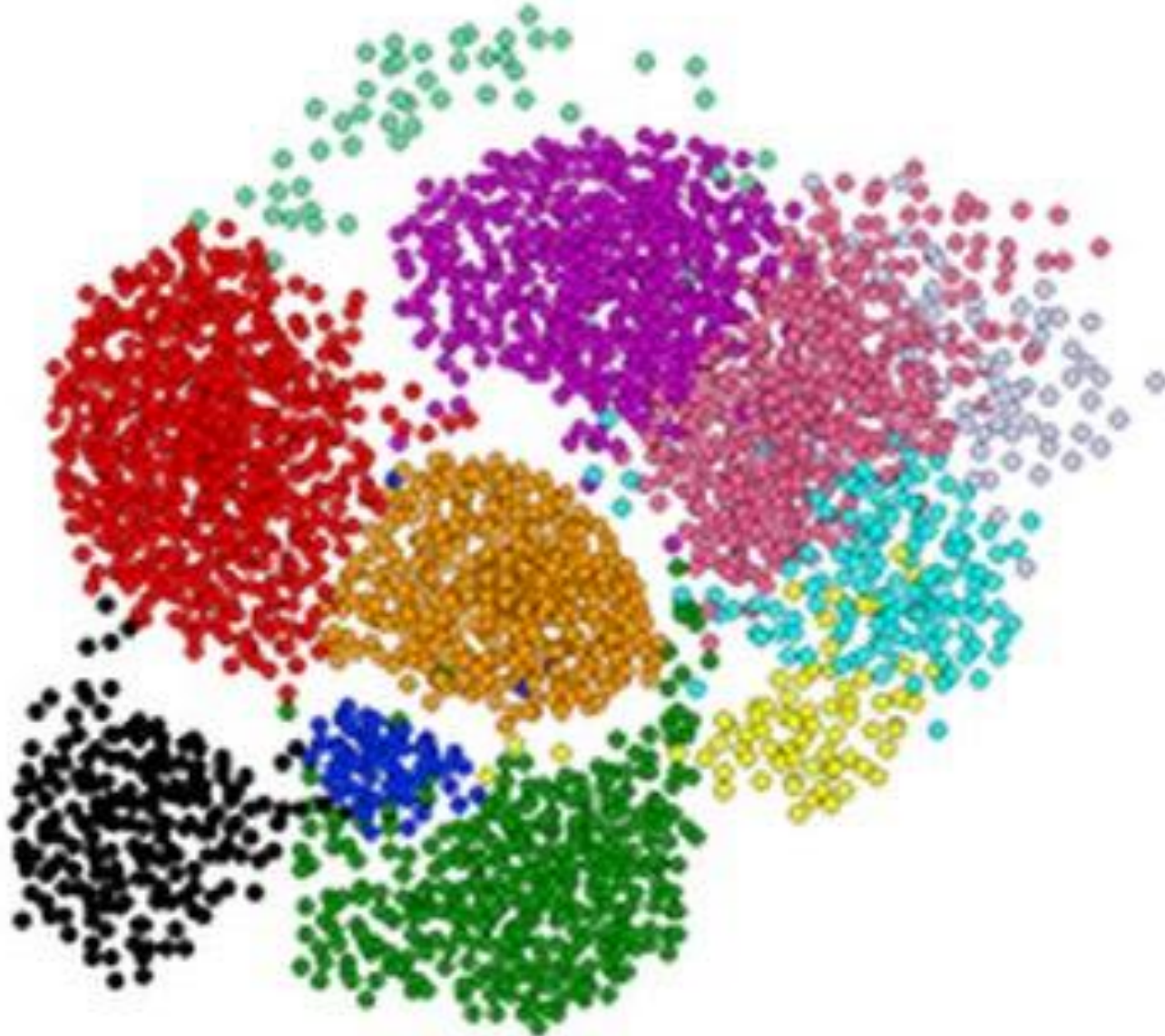
# The Problem of Clustering

- 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  $(x_1, x_2, \dots, x_k)$ , where  $x_i = 1$  iff the  $i^{\text{th}}$  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  $(x_1, x_2, \dots, x_k)$ , where  $x_i = 1$  iff the  $i^{\text{th}}$  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

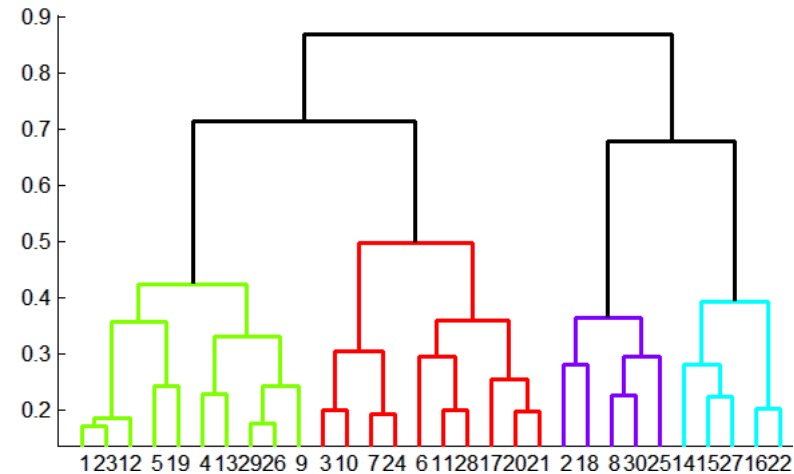
## ■ Hierarchical:

### ■ **Agglomerative** (bottom up):

- Initially, each point is a cluster
- Repeatedly combine the two “nearest” clusters into one

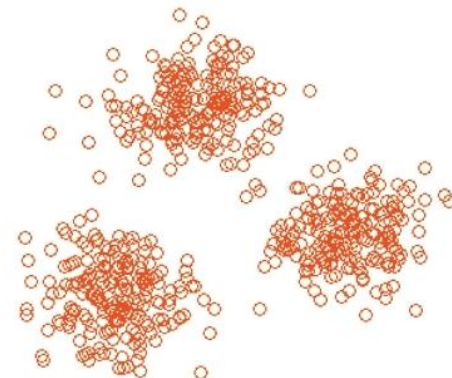
### ■ **Divisive** (top down):

- Start with one cluster and recursively split it



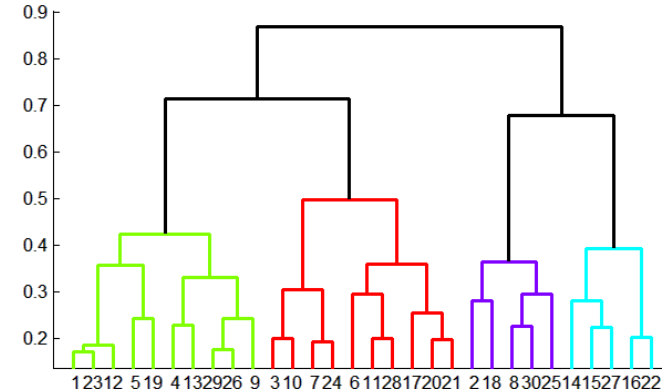
## ■ **Point assignment:**

- Maintain a set of clusters
- Points belong to “nearest” cluster



# Hierarchical Clustering

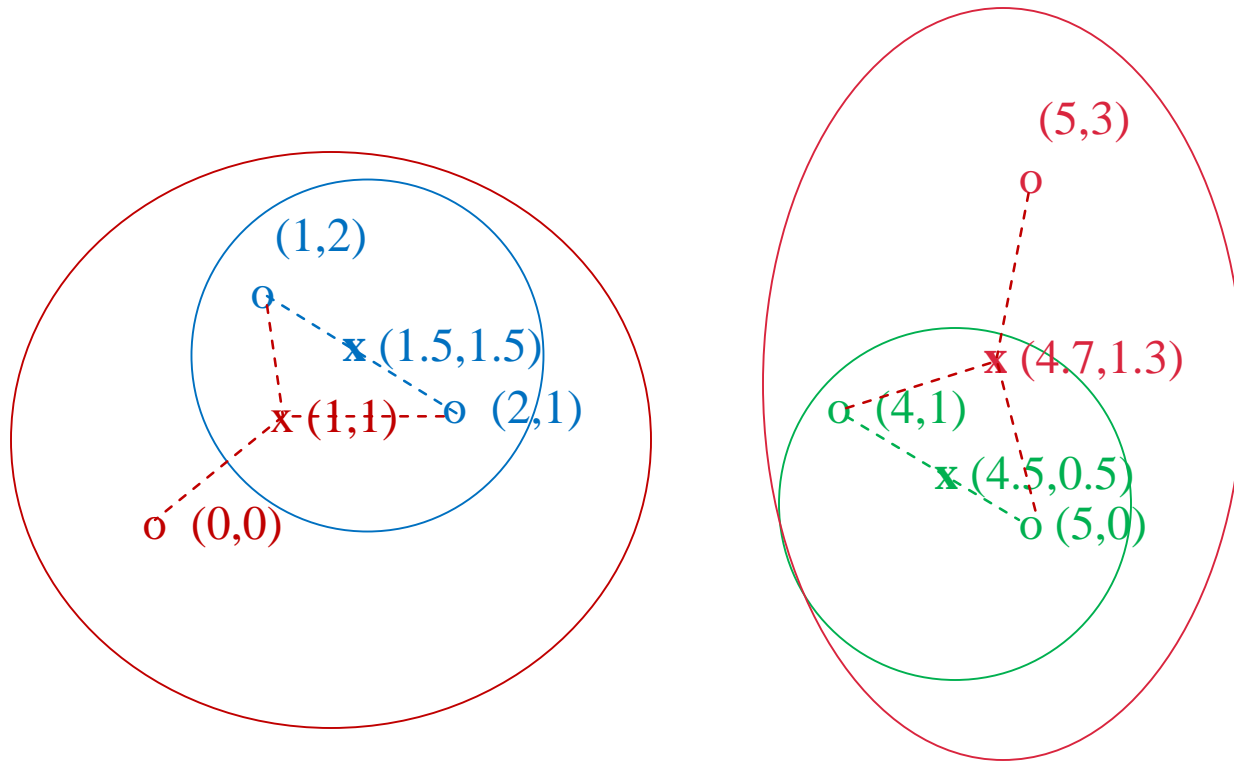
- **Key operation:**  
**Repeatedly combine two nearest clusters**
- **Three important questions:**
  - 1) How do you represent a cluster of more than one point?
  - 2) How do you determine the “nearness” of clusters?
  - 3) When to stop combining clusters?



# 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

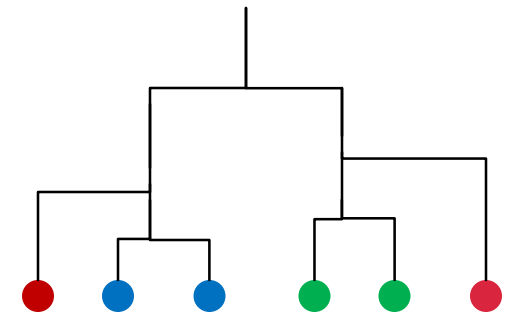
# Example: Hierarchical clustering



## Data:

$\circ$  ... data point

$\mathbf{x}$  ... 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?

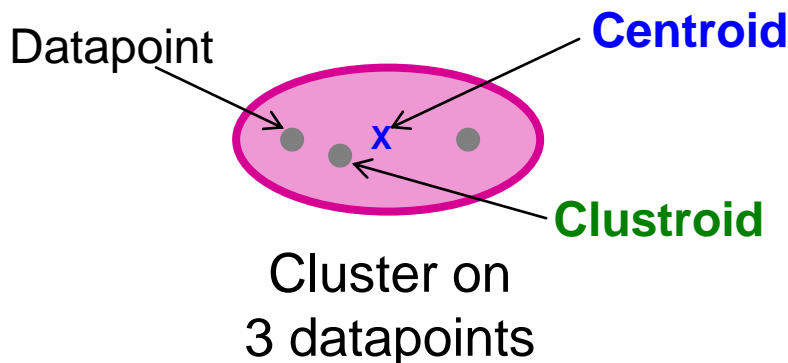
- (1) How to represent a cluster of many points?

**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 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

# Implementation

- **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  $O(N^2 \log N)$ 
  - **Still too expensive for really big datasets that do not fit in memory**

# *k*-means clustering

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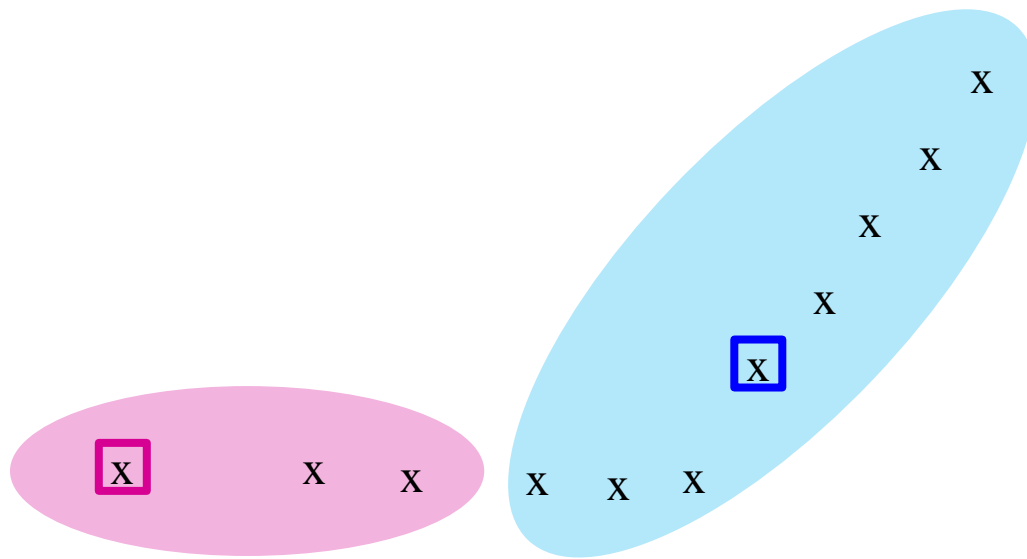
# $k$ -means Algorithm(s)

- 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

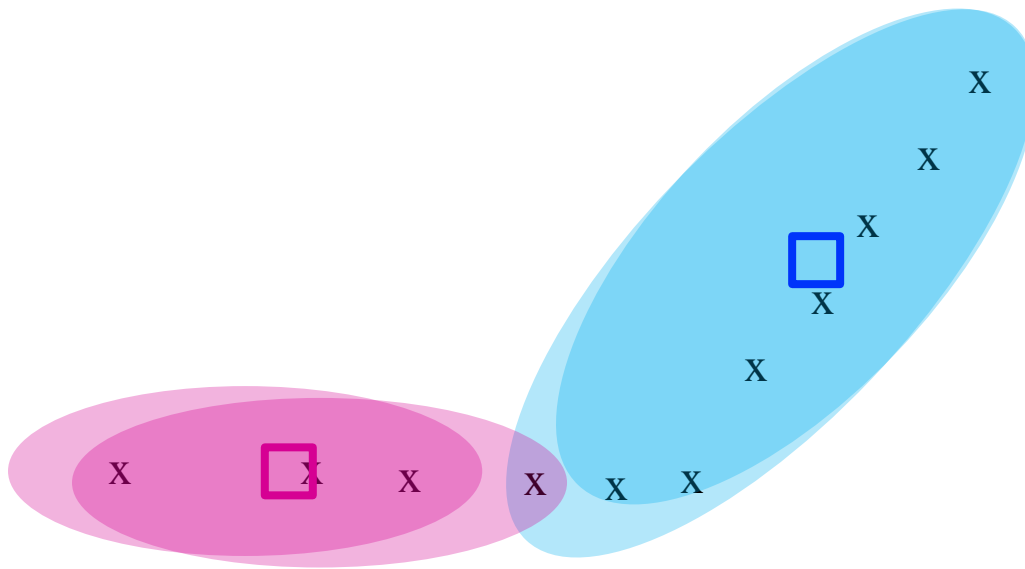
# Example: Assigning Clusters



x ... data point  
□ ... centroid

**Clusters after round 1**

# Example: Assigning Clusters

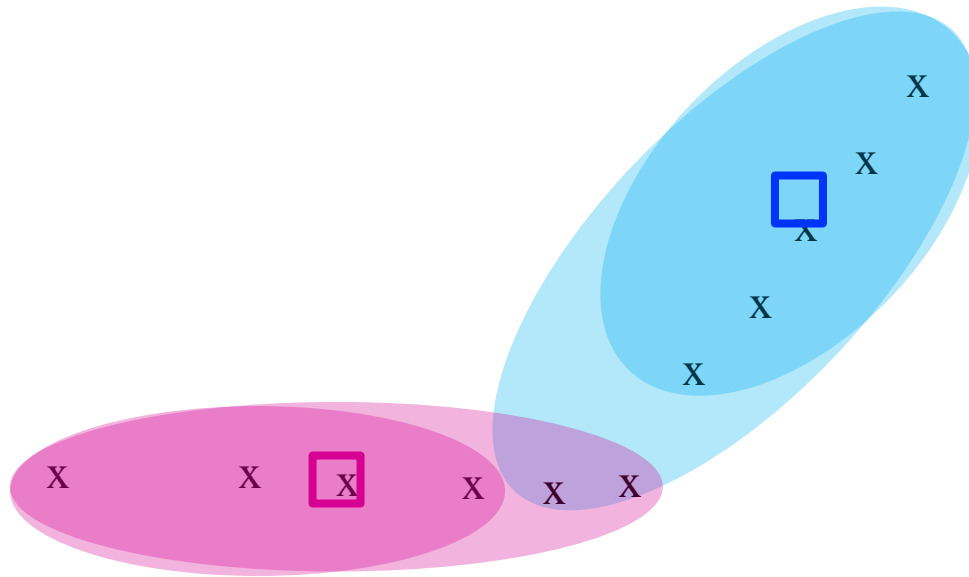


x ... data point  
□ ... centroid

**Clusters after round 2**



# Example: Assigning Clusters



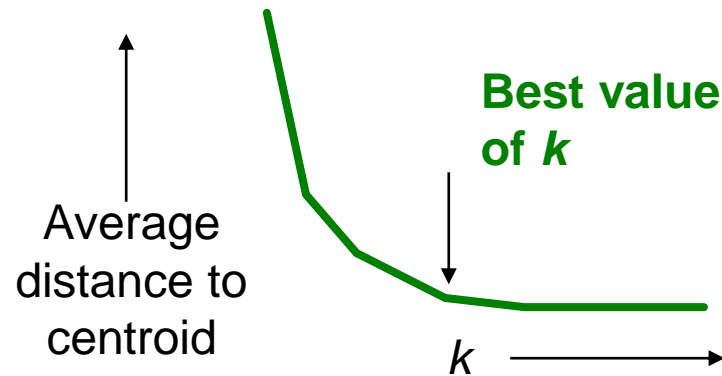
x ... data point  
□ ... centroid

**Clusters at the end**

# Getting the $k$ right

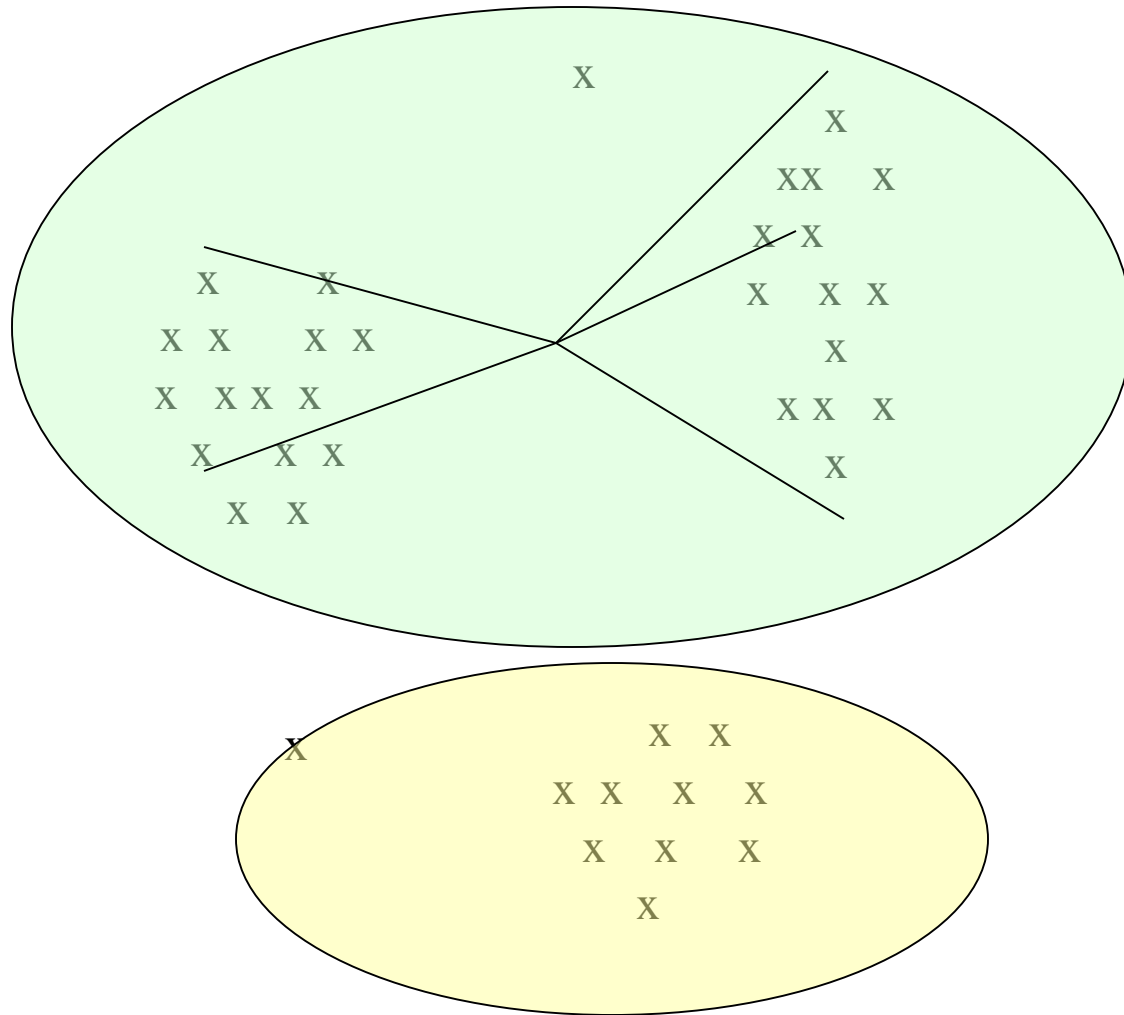
## How to select $k$ ?

- 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



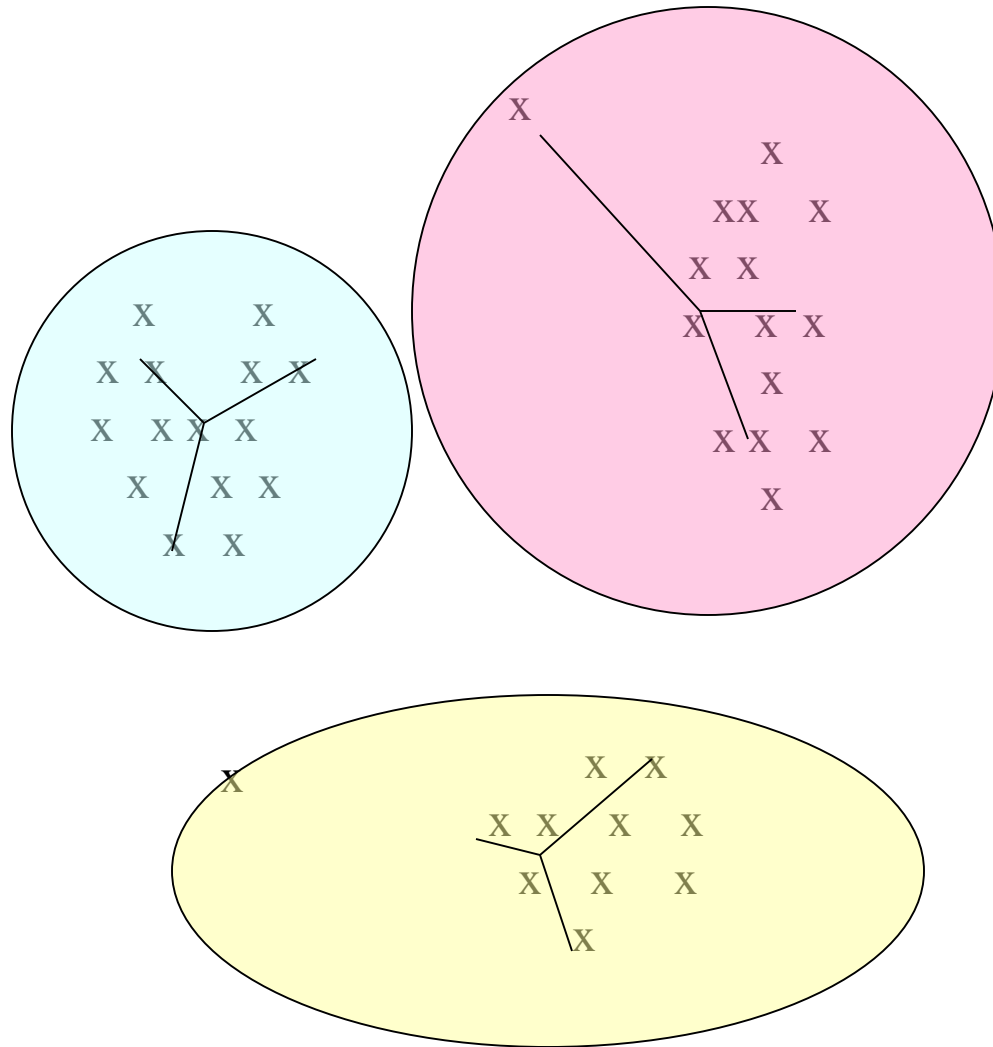
# Example: Picking $k$

Too few;  
many long  
distances  
to centroid.



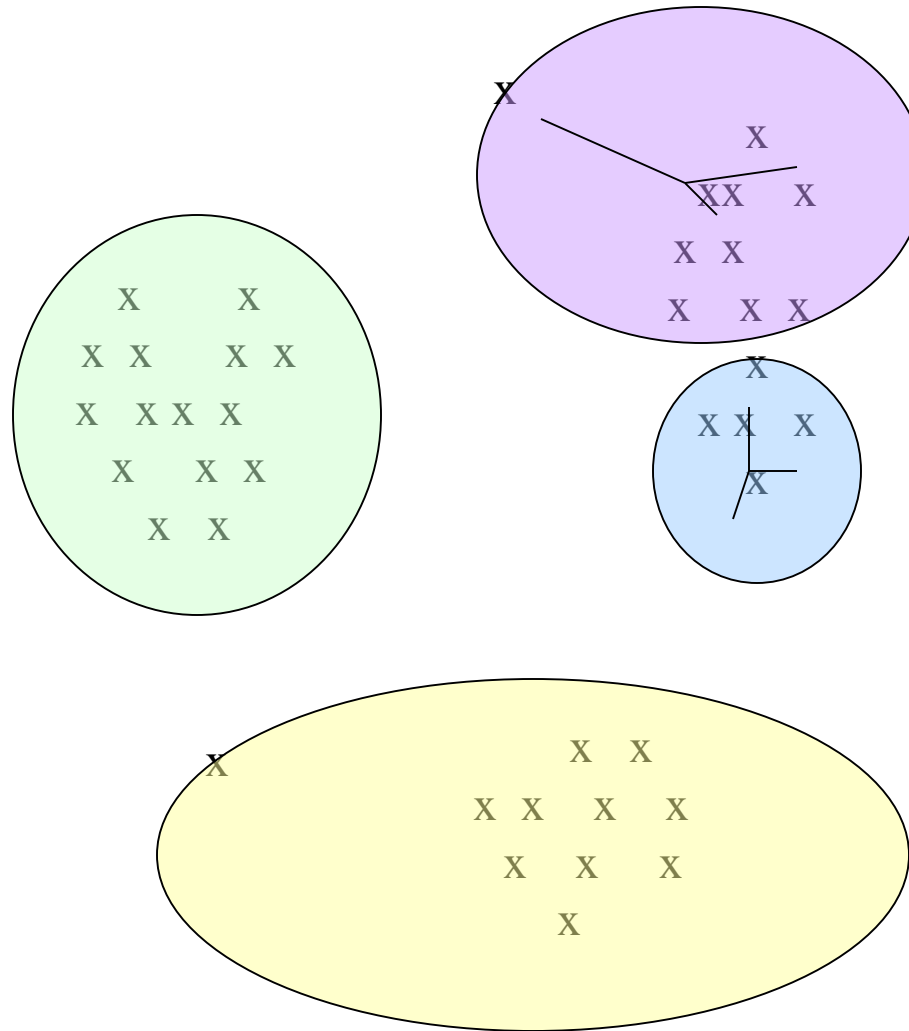
# Example: Picking $k$

Just right;  
distances  
rather short.



# Example: Picking $k$

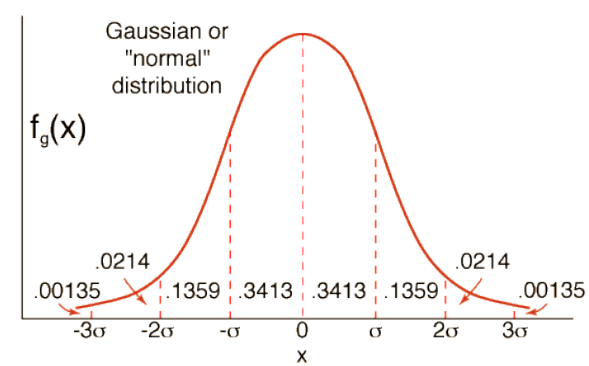
Too many;  
little improvement  
in average  
distance.



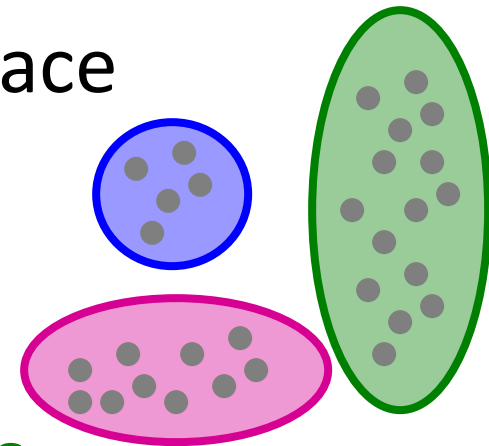
# The BFR Algorithm

Extension of *k*-means to large data

# BFR Algorithm



- **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  $O(\text{clusters})$  and not  $O(\text{data})$ )



# BFR Algorithm

- Points are read from disk 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:
  - 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

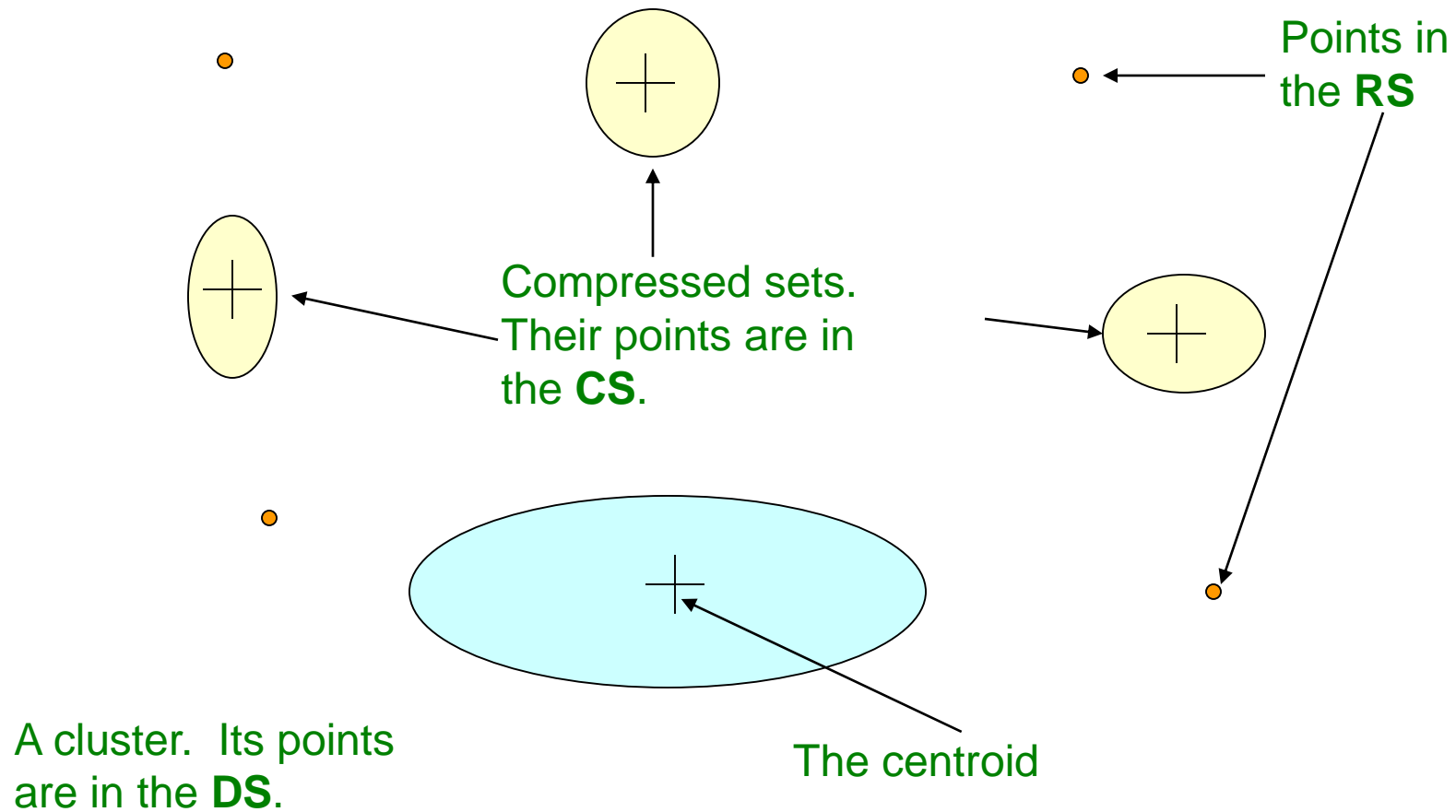


# Three Classes of Points

## 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

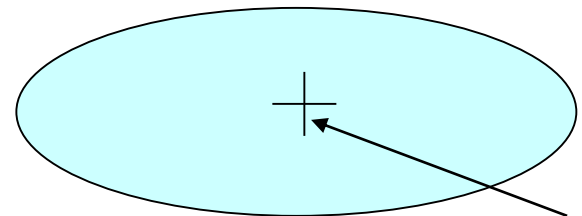


**Discard set (DS):** Close enough to a centroid to be summarized  
**Compression set (CS):** Summarized, but not assigned to a cluster  
**Retained set (RS):** Isolated points

# Summarizing Sets of Points

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

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



A cluster.

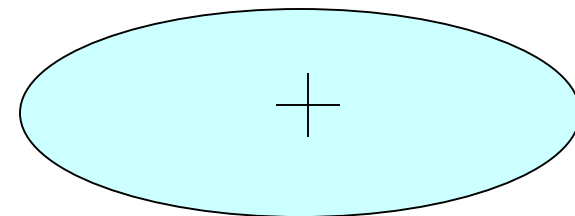
All its points are in the **DS**.

The centroid

# Summarizing Points: Comments

- $2d + 1$  values represent any size cluster
  - $d$  = number of dimensions
- Average in **each dimension** (**the centroid**) can be calculated as  $\text{SUM}_i / N$ 
  - $\text{SUM}_i = i^{\text{th}}$  component of SUM
- Variance of a cluster's discard set in dimension  $i$  is:  $(\text{SUMSQ}_i / N) - (\text{SUM}_i / N)^2$ 
  - 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  $d \times d$  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  $N_s$ ,  $SUM_s$ ,  $SUMSQ_s$
- **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.  $2\sqrt{d}$ 
    - 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  $<$   $\sqrt{d}$
  - For point  $(x_1, \dots, x_d)$  and centroid  $(c_1, \dots, c_d)$ 
    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

$$d(x, c) = \sqrt{\sum_{i=1}^d y_i^2}$$

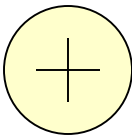
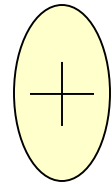
$\sigma_i$  ... standard deviation of points in the cluster in the  $i^{\text{th}}$  dimension



# 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



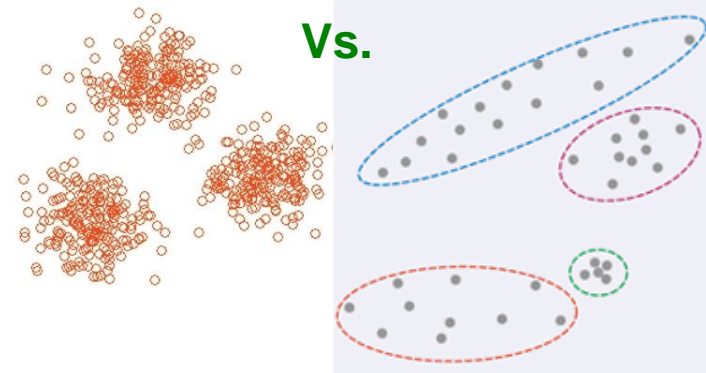
# The CURE Algorithm

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

# The CURE Algorithm

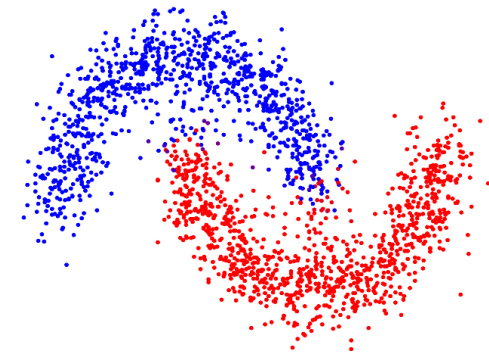
- **Problem with BFR/ $k$ -means:**

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

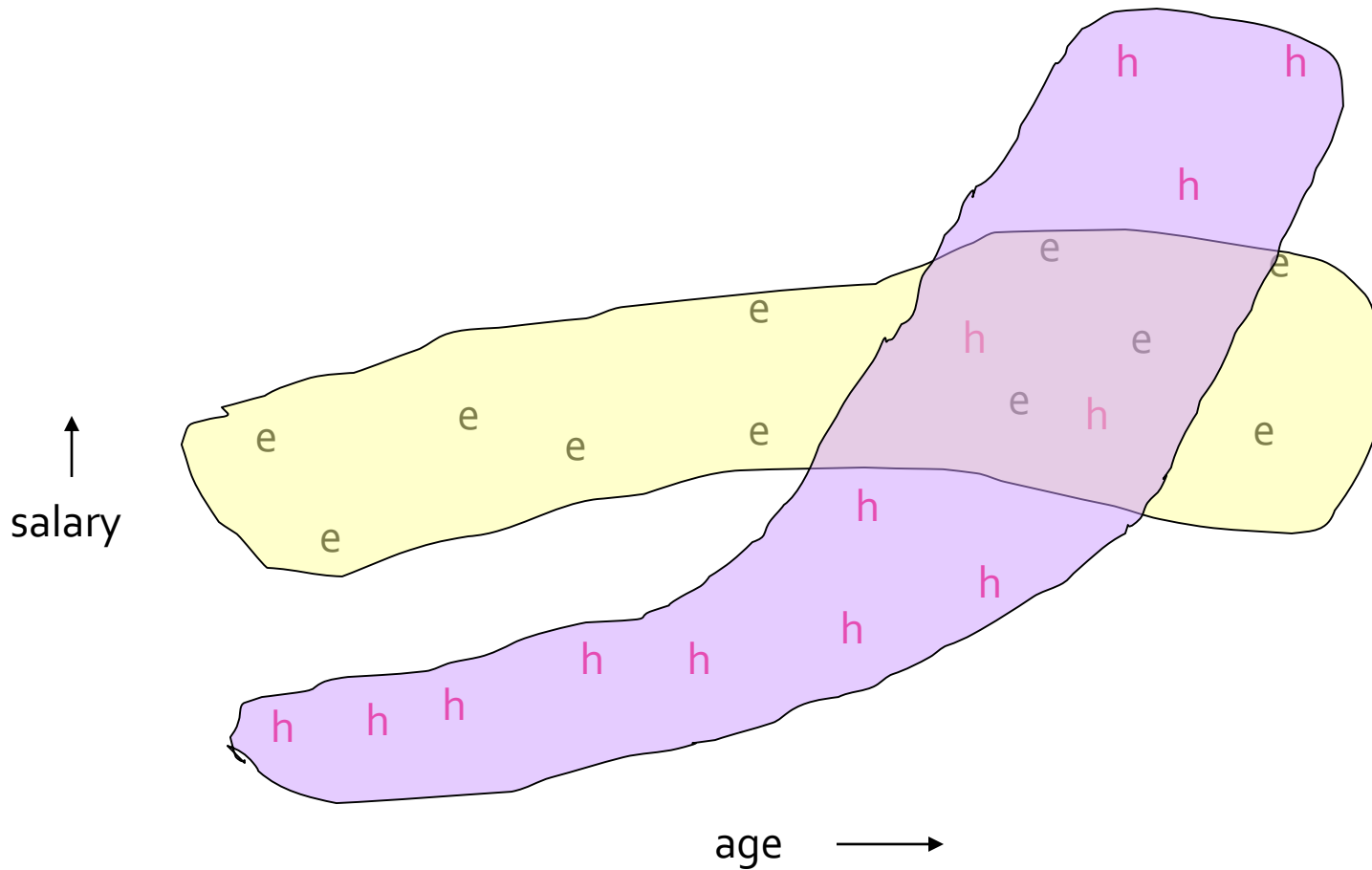


- **CURE (Clustering Using REpresentatives):**

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



# Example: Stanford Salaries

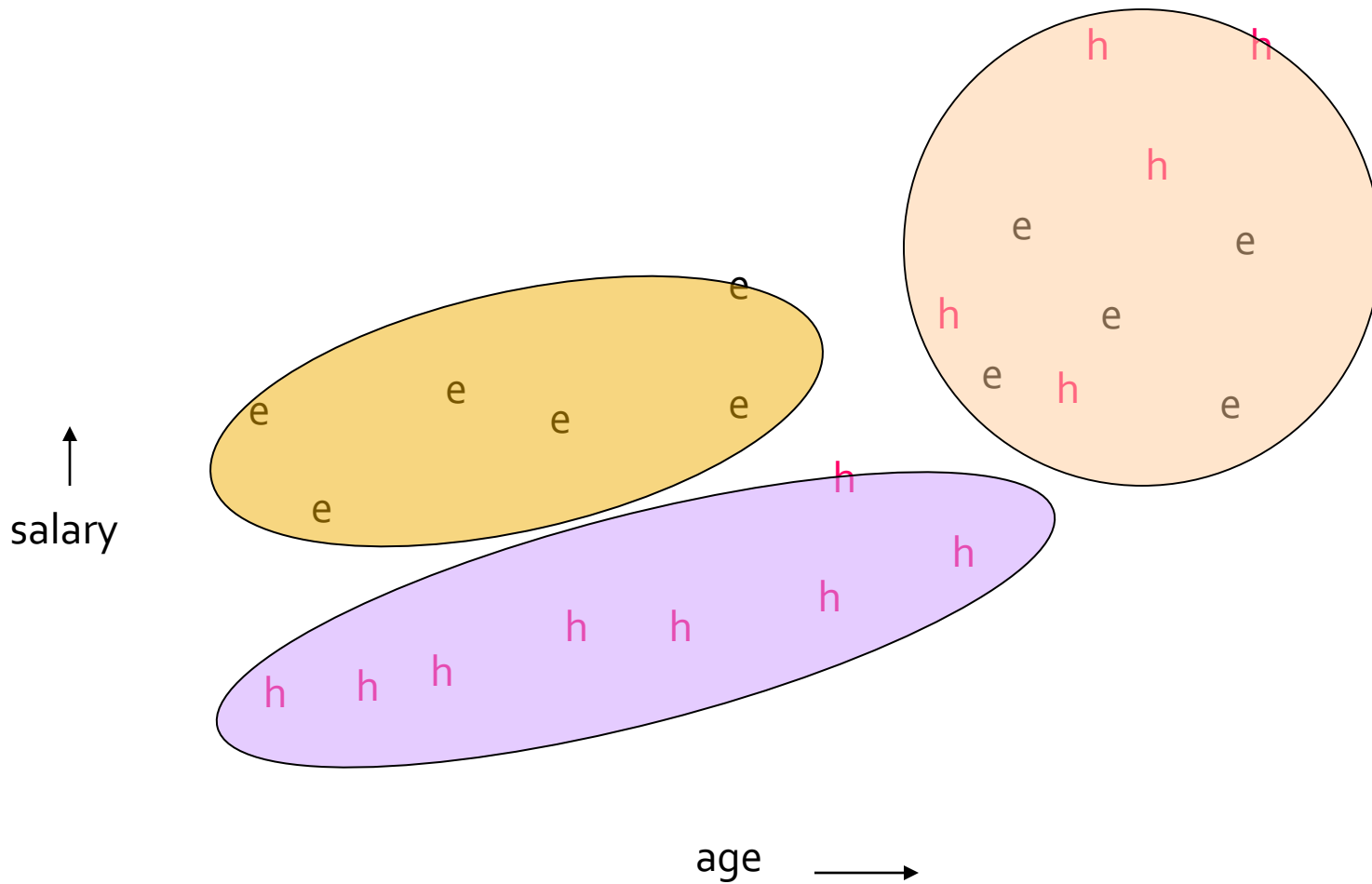


# Starting CURE

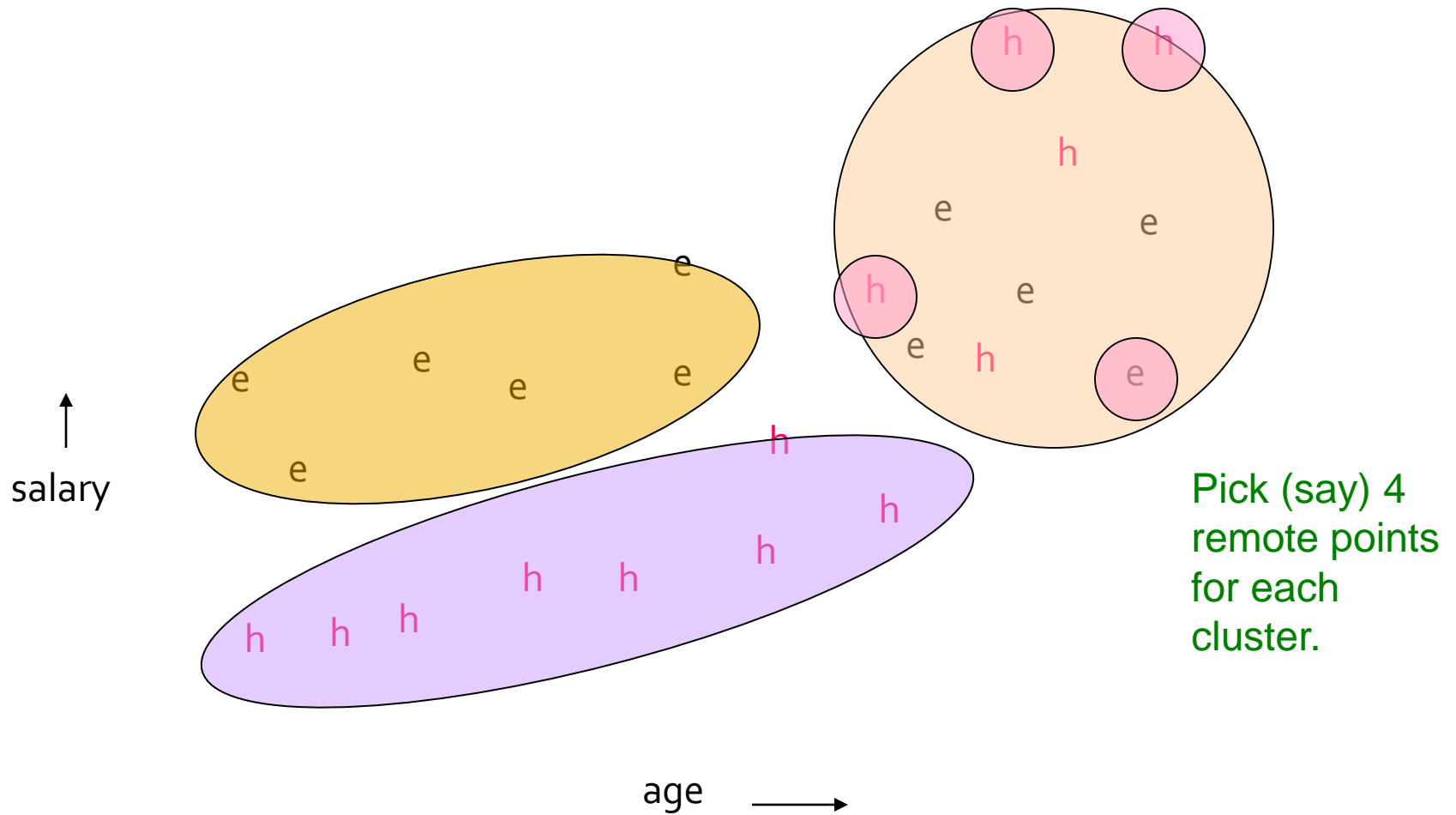
## 2 Pass algorithm. Pass 1:

- **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

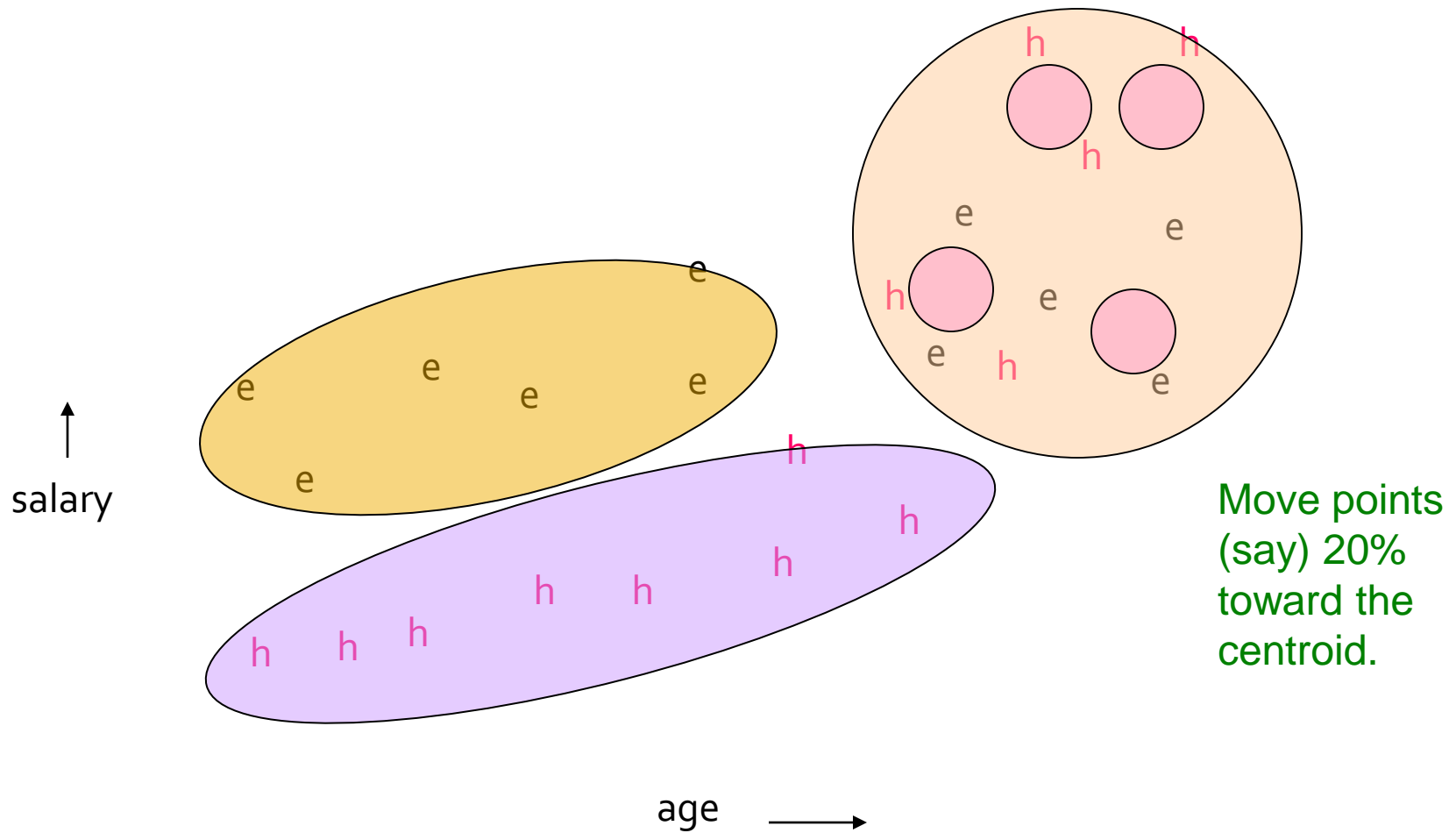
# Example: Initial Clusters



# Example: Pick Dispersed Points



# Example: Pick Dispersed Points

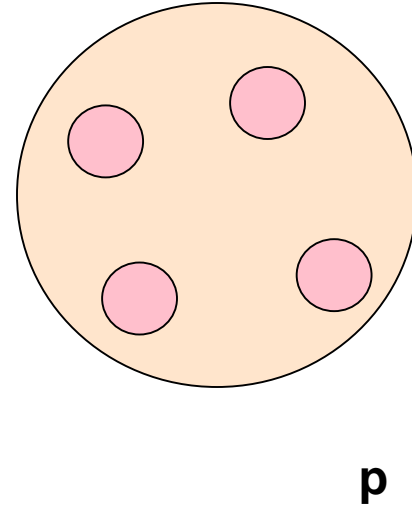




# Finishing CURE

## 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



# 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**