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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

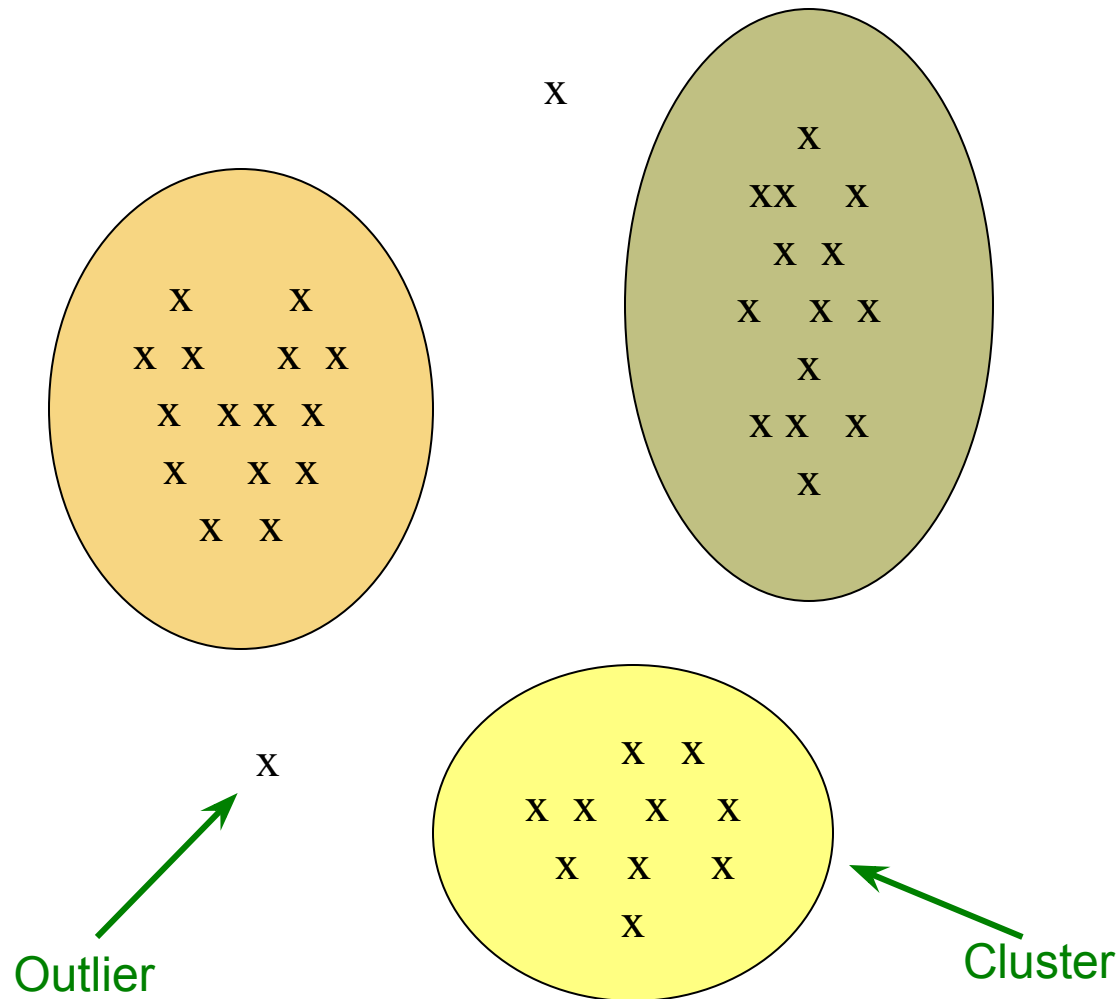
- Given a cloud of data points we want to understand its structure



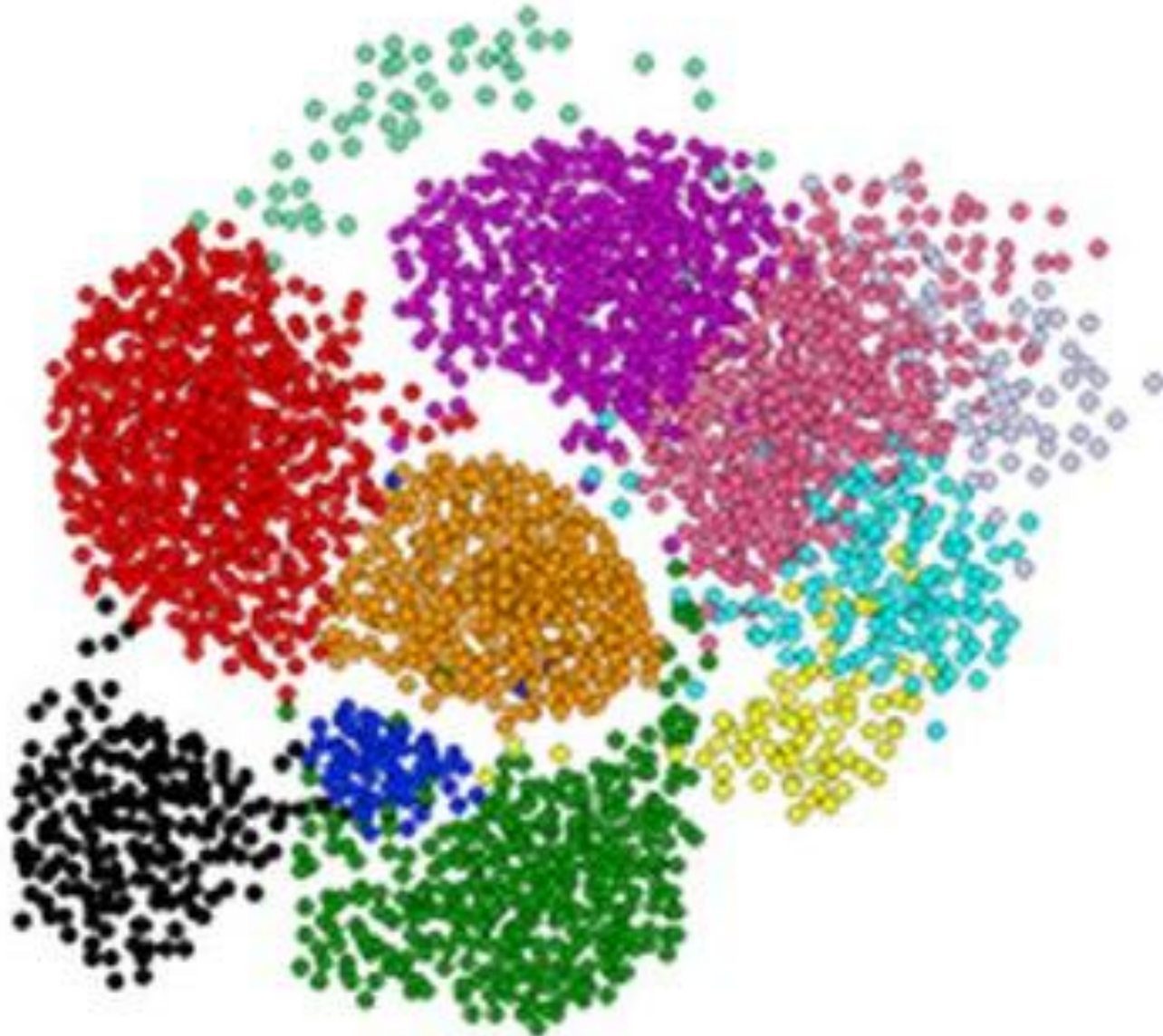
# The Problem of Clustering

- 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, ...

# Example: Clusters & Outliers



# Clustering is a hard problem!

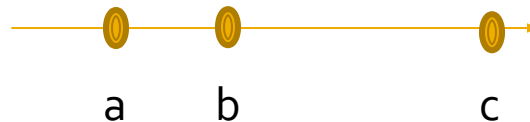


# Why is it hard?

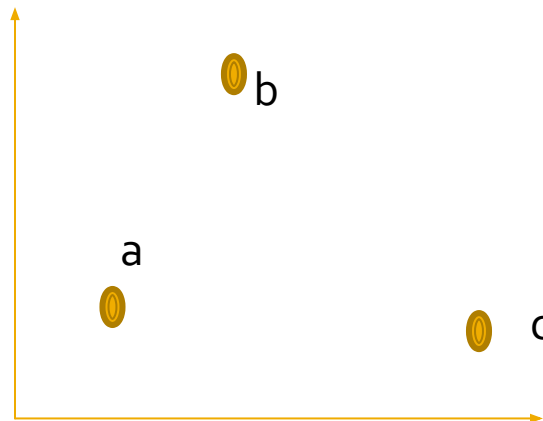
- Clustering in two dimensions looks easy
- Clustering small amounts of data looks easy
- Many applications involve not 2, but 10 or 10,000 dimensions
- **High-dimensional spaces look different:**  
Almost all pairs of points are at about the same distance

# High Dimension: Euclidean

- Consider a set of data points on a line
  - $\text{dist}(a, b) < \text{dist}(a, c)$



- Consider increasing the dimension by 1
  - $\text{dist}(a, b) \sim \text{dist}(a, c)$

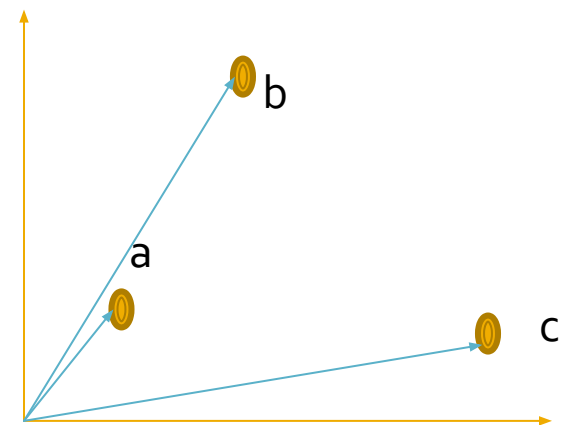


# High Dimension: Cosine

- $\text{Cosine}(a, b) > \text{Cosine}(a, c)$

- Increase d to 3

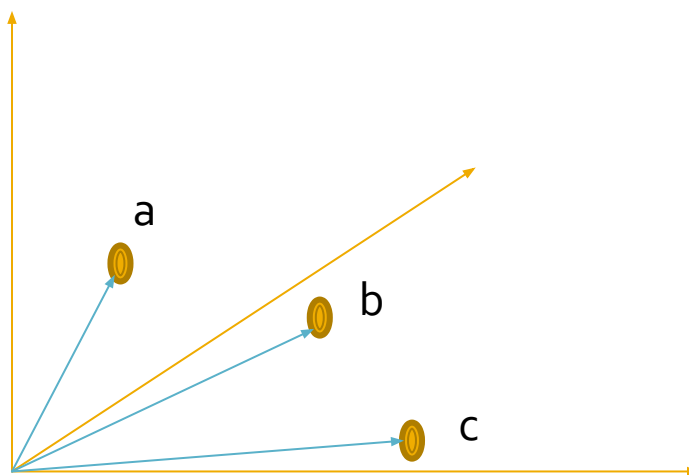
- $\text{Cosine}(a, b) \sim \text{Cosine}(a, c)$



- Higher d

- Angle  $\rightarrow 90^\circ$

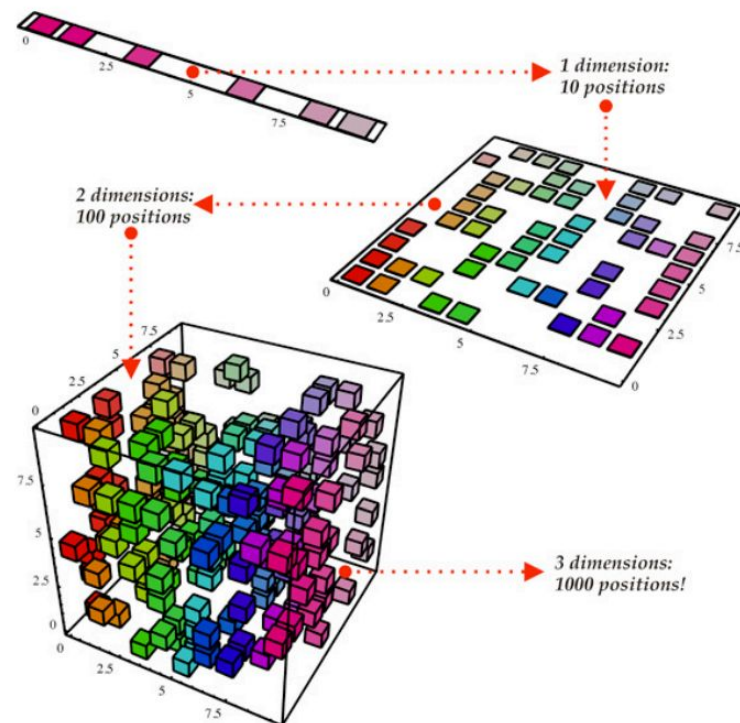
- Cosine  $\rightarrow 0$





# 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

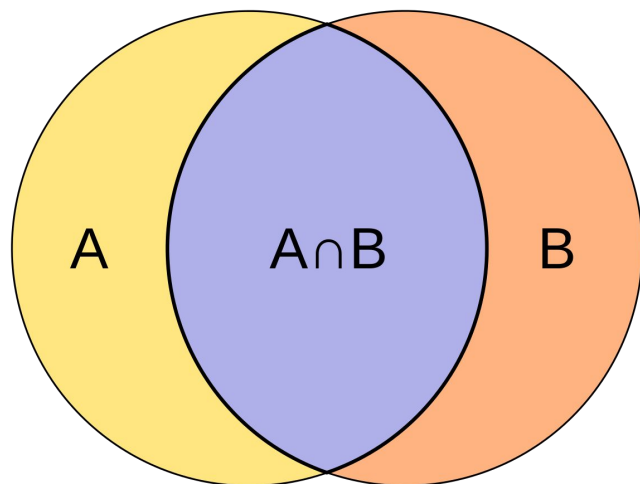
## Space of all CDs:

- 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  $(x_1, x_2, \dots, x_k)$ , 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

# Cosine, Jaccard, and Euclidean

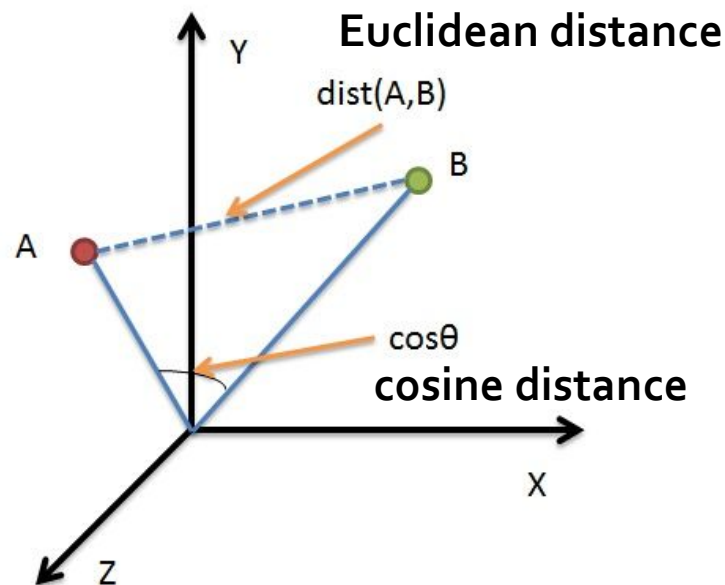
- As with CDs we have a choice when we think of documents as sets of words or shingles:
  - **Sets as vectors:** Measure similarity by the cosine distance
  - **Sets as sets:** Measure similarity by the Jaccard distance
  - **Sets as points:** Measure similarity by Euclidean distance

# Measure similarity

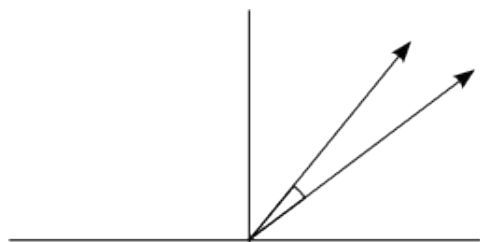


Jaccard distance

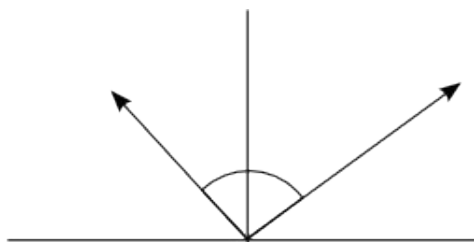
$$J(A, B) = \frac{|A \cap B|}{|A \cup B|}$$



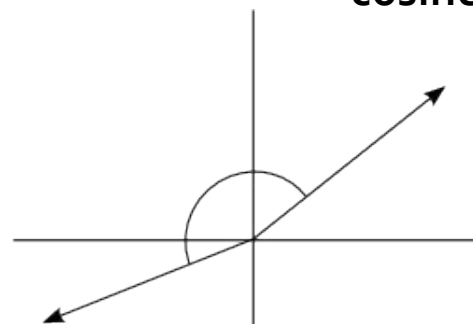
cosine distance



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



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



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

# Overview: Methods of Clustering

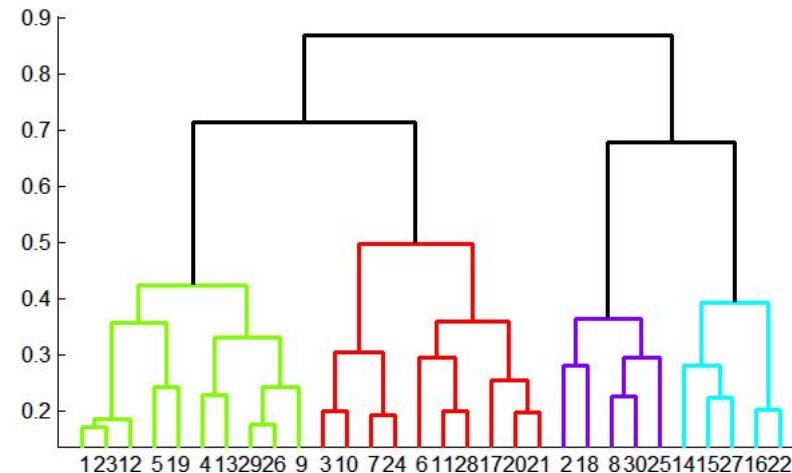
## 1. Hierarchical

- **Agglomerative** (bottom up)

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

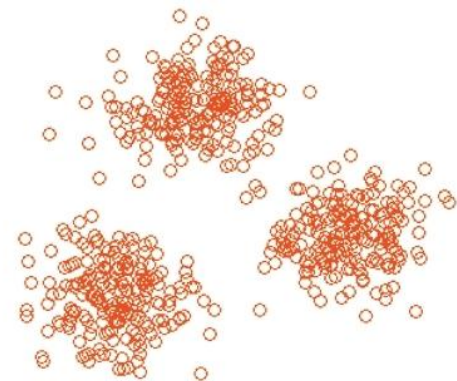
- **Divisive** (top down)

- Start with one cluster and recursively split it



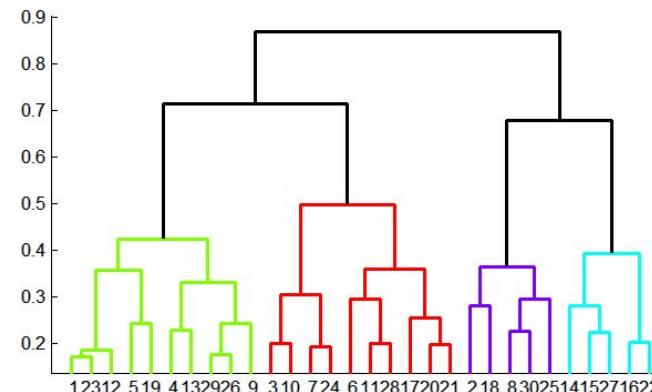
## 2. Point Assignment

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



# Hierarchical Clustering

- **Key operation:**  
**Repeatedly combine**  
**two nearest clusters**



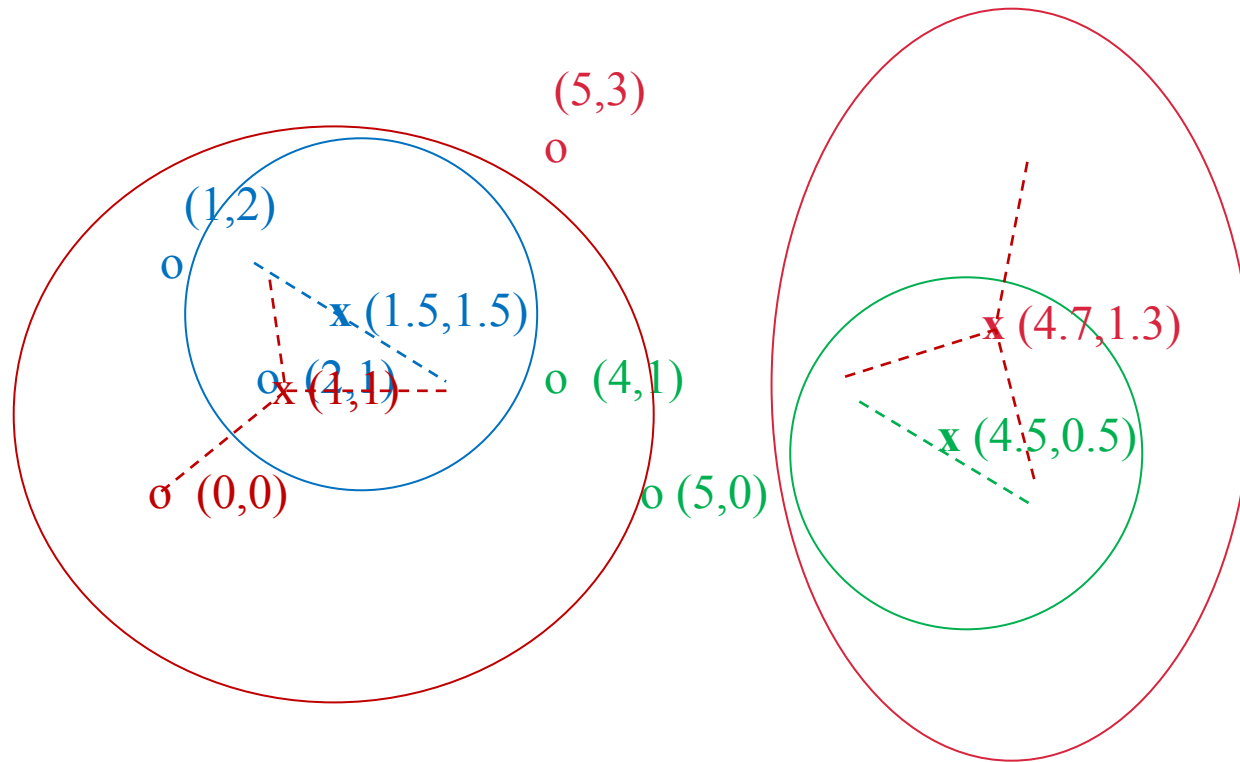
- **Three important questions:**
  - 1) How do you represent a cluster **of more than one point**?
  - 2) How do you determine the “**nearness**” of clusters?
  - 3) When to **stop combining** clusters?

# Hierarchical Clustering

- **Key operation:** Repeatedly combine two nearest clusters
- **(1) How to represent a cluster of many points?**
  - **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



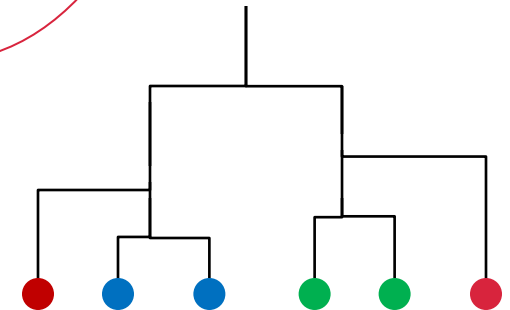
# Example: Hierarchical clustering



**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 (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?  
Treat clustroid as if it were centroid, when computing inter-cluster distances

# “Closest” Point?

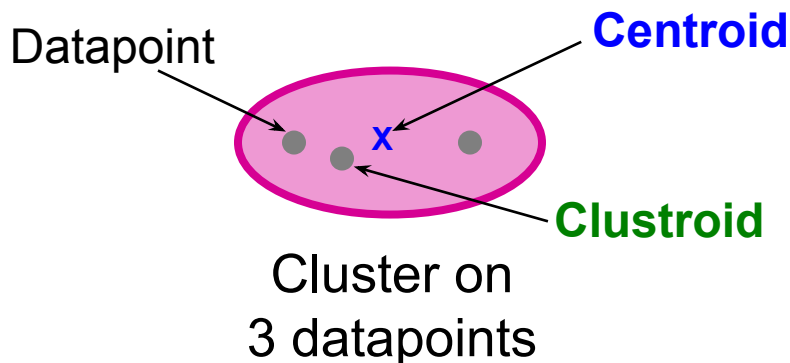
## ■ (1) How to represent a cluster of many points?

**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:

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

# 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	<b>7</b>	<b>17</b>	<b>3</b>
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

	C <sub>1</sub>	C <sub>2</sub>	C <sub>3</sub>	C <sub>4</sub>
C <sub>1</sub>	0	2	3	2
C <sub>2</sub>		0	4	5
C <sub>3</sub>			0	3
C <sub>4</sub>				0



# Complexity of Naive Implementation

- Initially,  $O(n^2)$  for creating matrix and finding pair with minimum distance
- Subsequent merge, assuming matrix:  $k \times k$ 
  - 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:  $O(k^2)$

=> Overall complexity:  $O(n^3)$

# 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**

# Sum Squared Error (SSE)

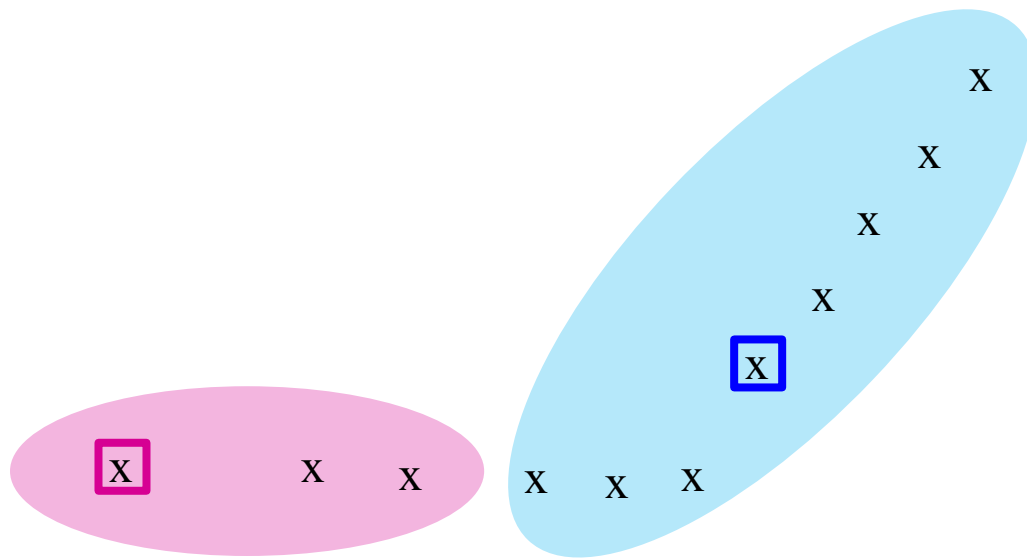
- SSE (sum squared error) is a common measure of the quality of a cluster.
  - It is the sum of the squares of the distances between each of the points of the cluster and the centroid.
- Sometimes, we decide to split a cluster in order to reduce the SSE.
  - Calculate the SSE before split
  - Decide 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
- 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

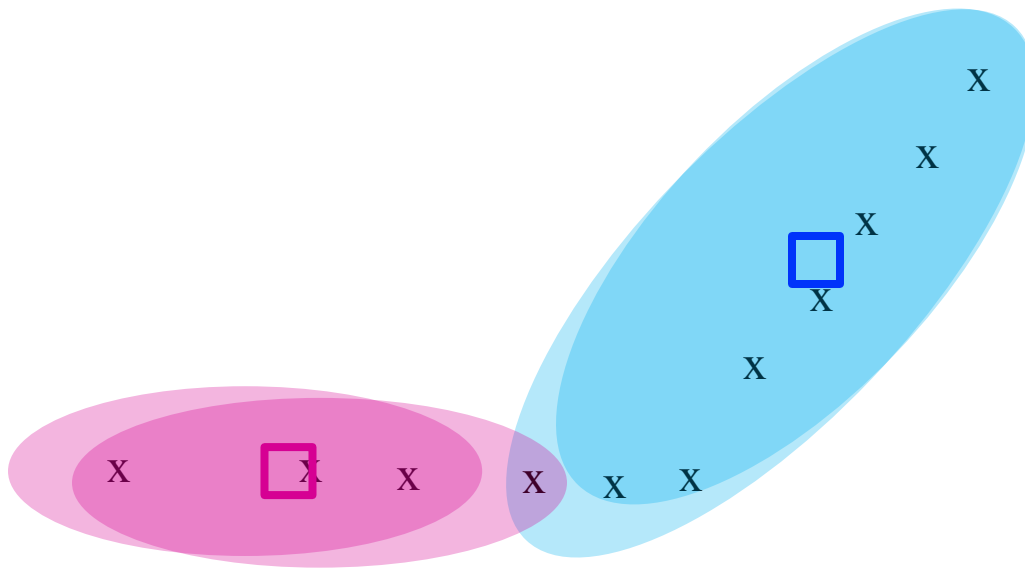


x ... data point

□ ... centroid

**Clusters after round 1**

# Example: Assigning Clusters

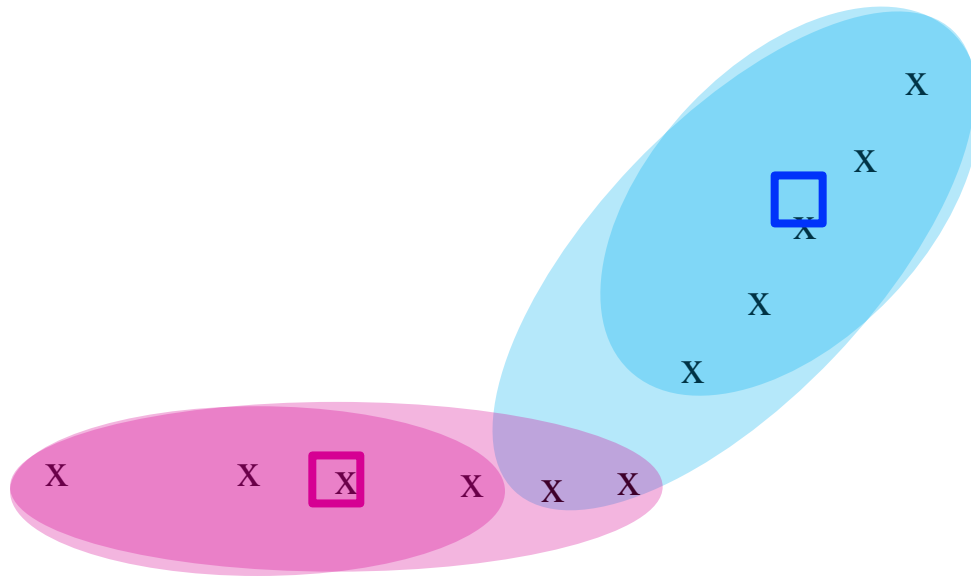


x ... data point

□ ... centroid

**Clusters after round 2**

# Example: Assigning Clusters



x ... data point

□ ... centroid

**Clusters at the end**



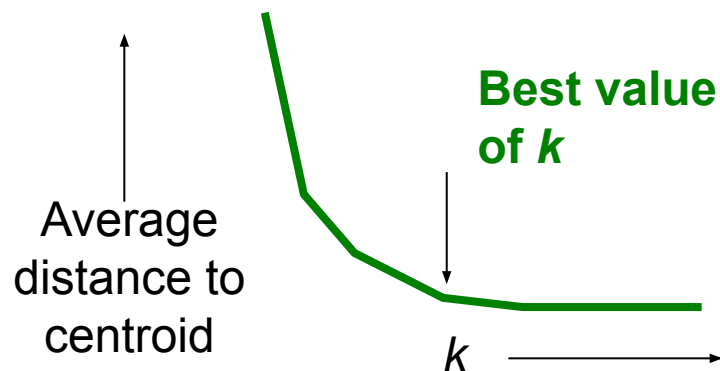
# Populating Clusters

- **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

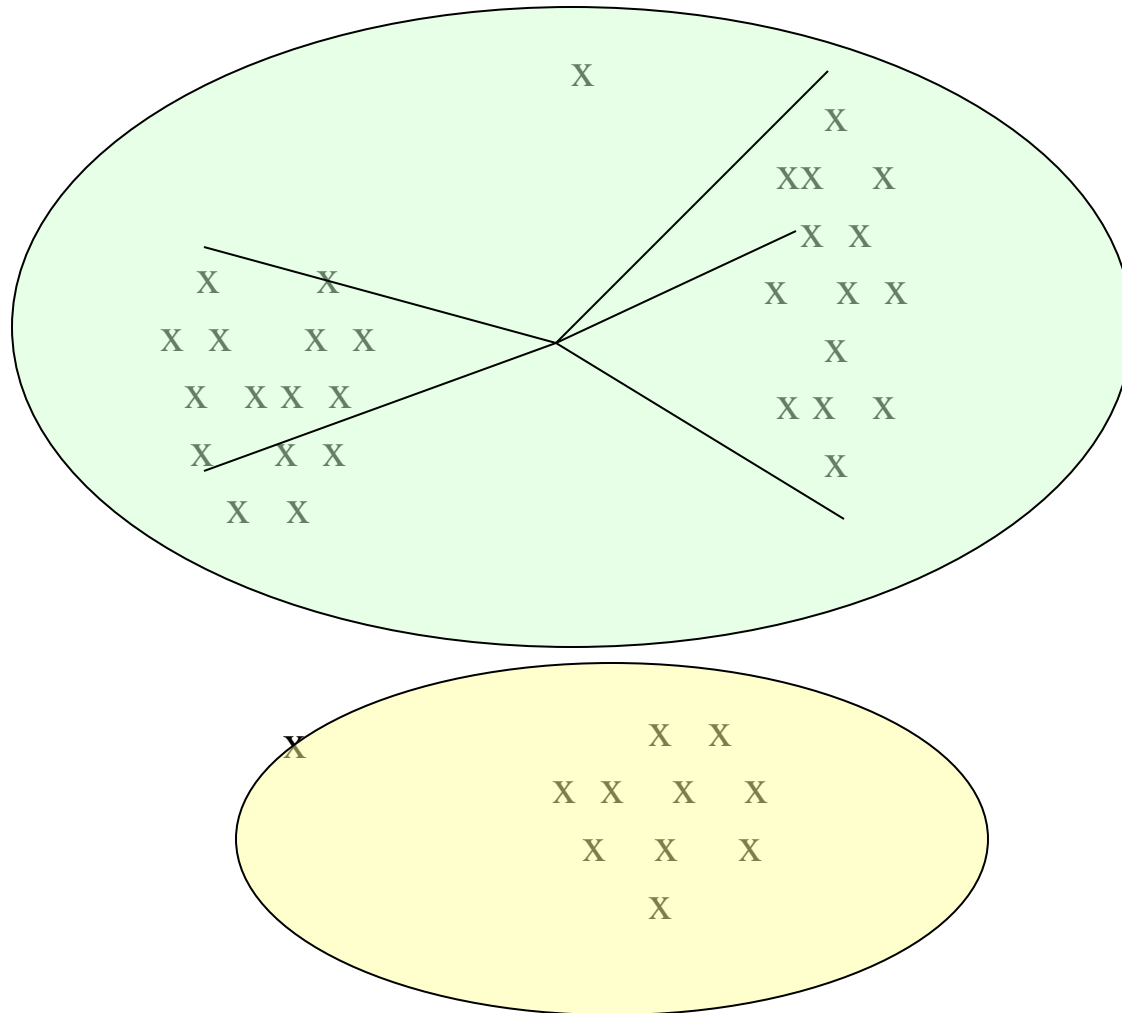
## How to select $k$ ?

- 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



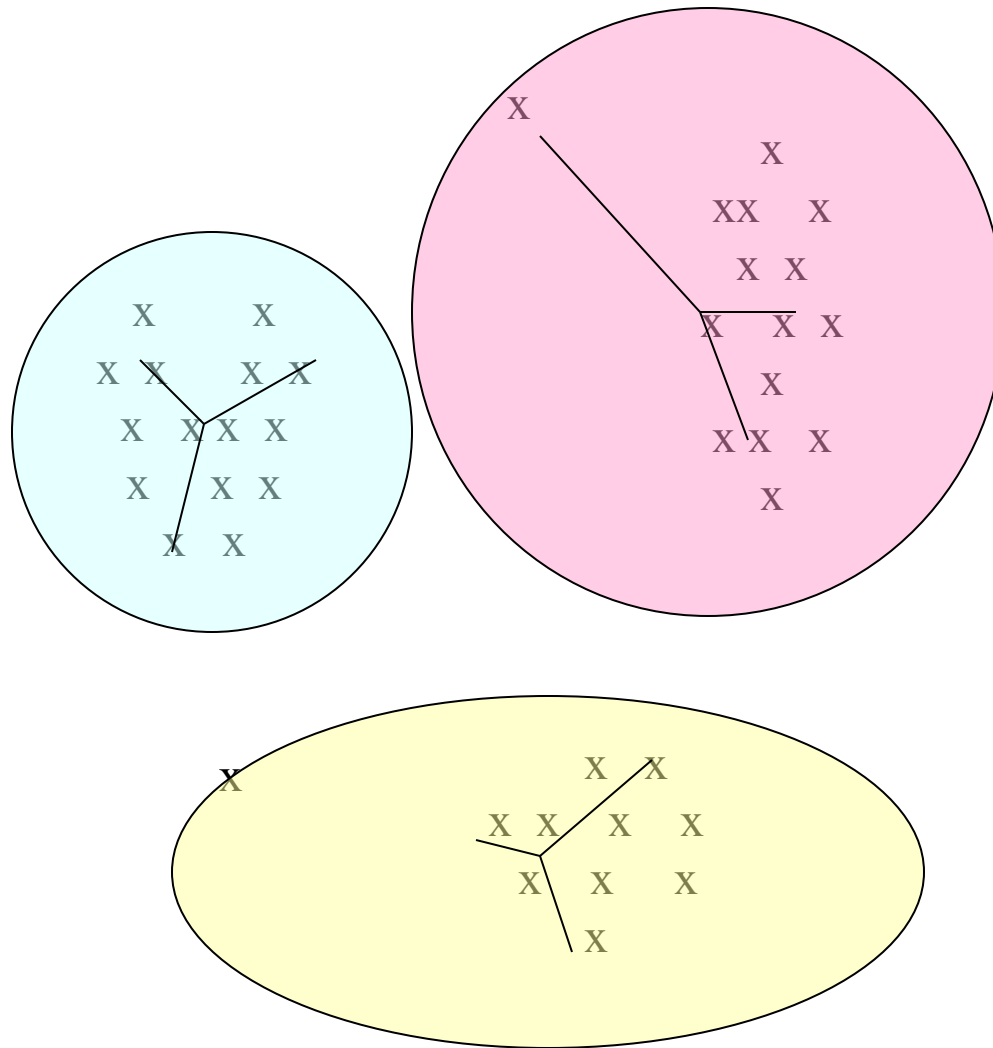
# Example: Picking $k$

**Too few;**  
many long  
distances  
to centroid.



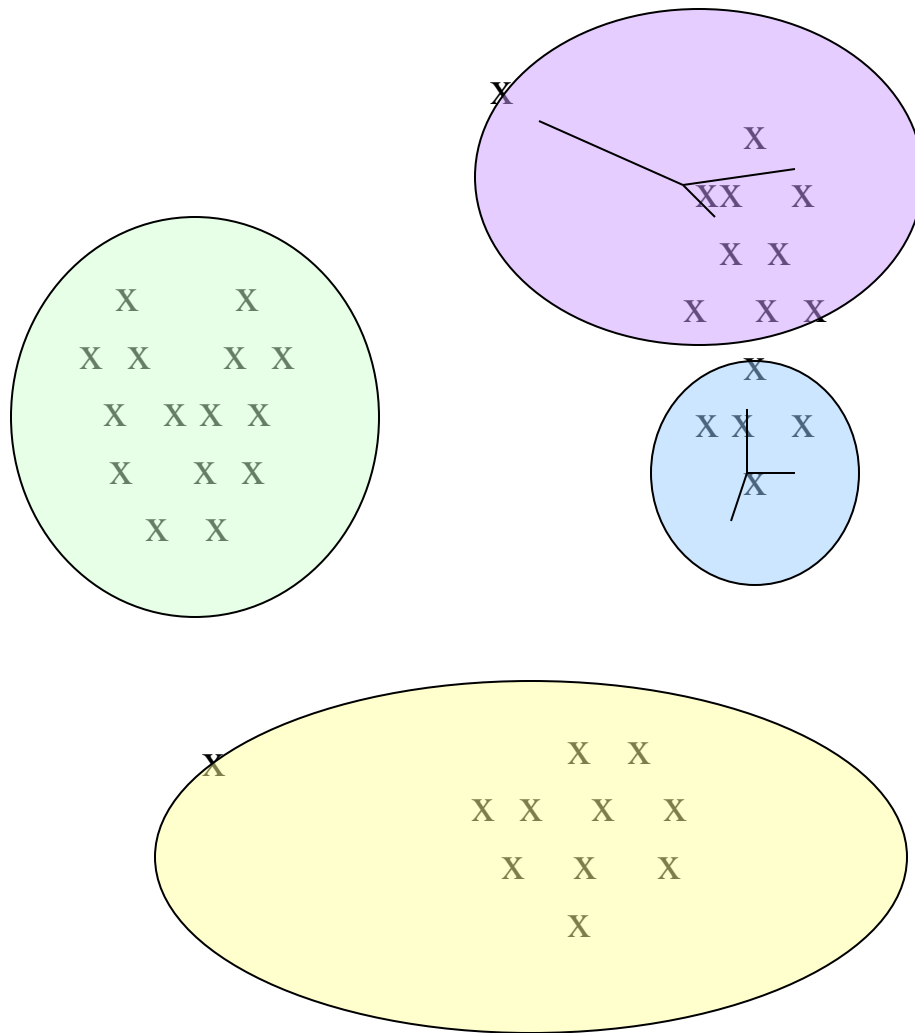
# Example: Picking $k$

**Just right;**  
distances  
rather short.



# Example: Picking $k$

Too many;  
little improvement  
in average  
distance.



# Two Scalable Algorithms

BFR: Bradley-Fayyad-Reina

- for uniform-distributed axis-aligned shapes

CURE: Clustering Using Representitives

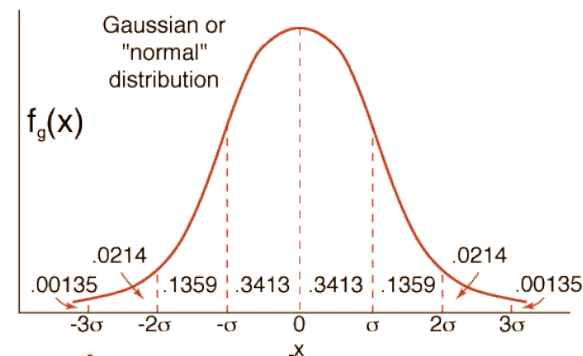
- for arbitrary shapes

# The BFR Algorithm

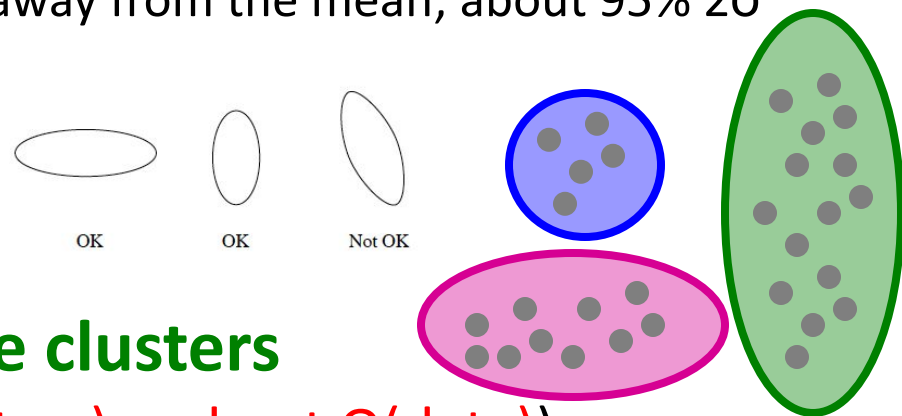
Extension of  $k$ -means to large data

# 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
  - About 68% of values are within  $\sigma$  away from the mean; about 95%  $2\sigma$  away; and about 99.7%  $3\sigma$  away.
  - 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” 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
  - 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

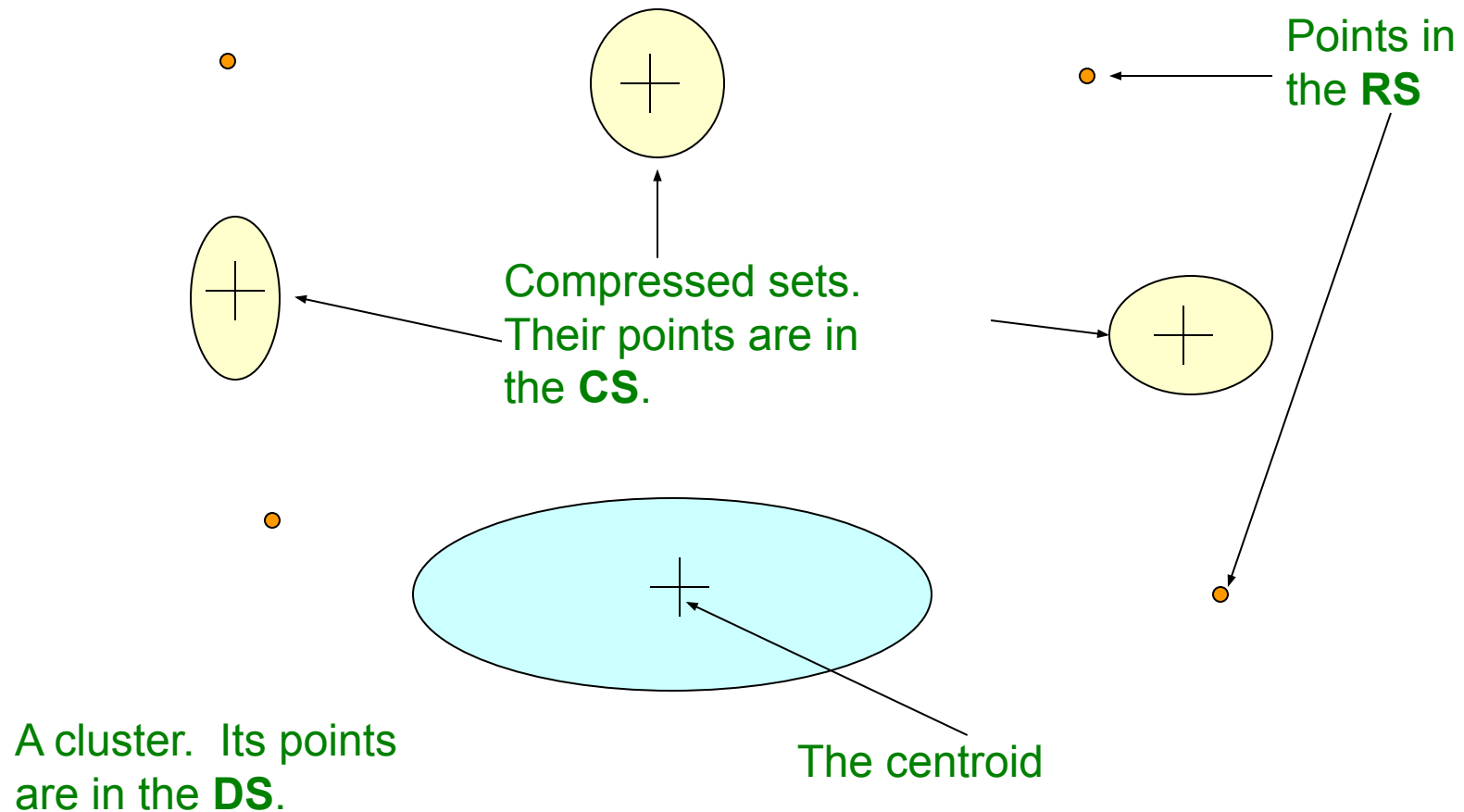
- **Compression set (CS):**

- 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

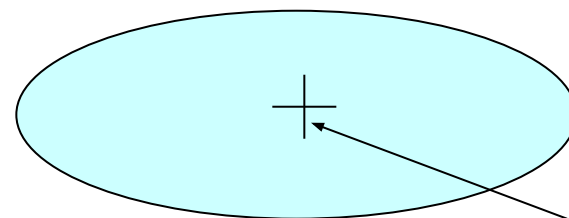


**Discard set (DS):** Close enough to a centroid to be summarized  
**Compression set (CS):** Summarized, but not assigned to a cluster  
**Retained set (RS):** Isolated points

# 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



A cluster.

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

The centroid

# Variance

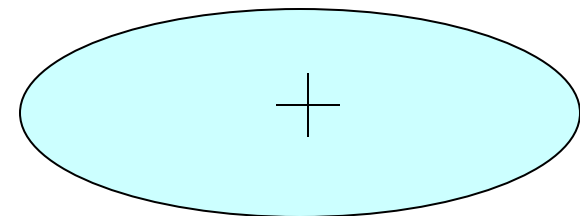
$$\begin{aligned}\text{Var}(X) &= E \left[ (X - E(X))^2 \right] \\ &= E[X^2 - 2XE[X] + (E[X])^2] \\ &= E[X^2] - 2E[X]E[X] + (E[X])^2 \\ &= E[X^2] - (E[X])^2\end{aligned}$$

$$\text{Var}(X) = \frac{1}{n} \sum_{i=1}^n (x_i - \mu)^2, \quad \mu = \frac{1}{n} \sum_{i=1}^n x_i.$$

# Summarizing Points: Comments

- **$2d + 1$  values** represent any size cluster
  - $d$  = number of dimensions
- Average in **each dimension** (**the centroid**) can be calculated as  $\text{SUM}_i / N$ 
  - $\text{SUM}_i = i^{\text{th}}$  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, **merge** all compressed sets in the **CS** and all **RS** points into their **nearest** cluster

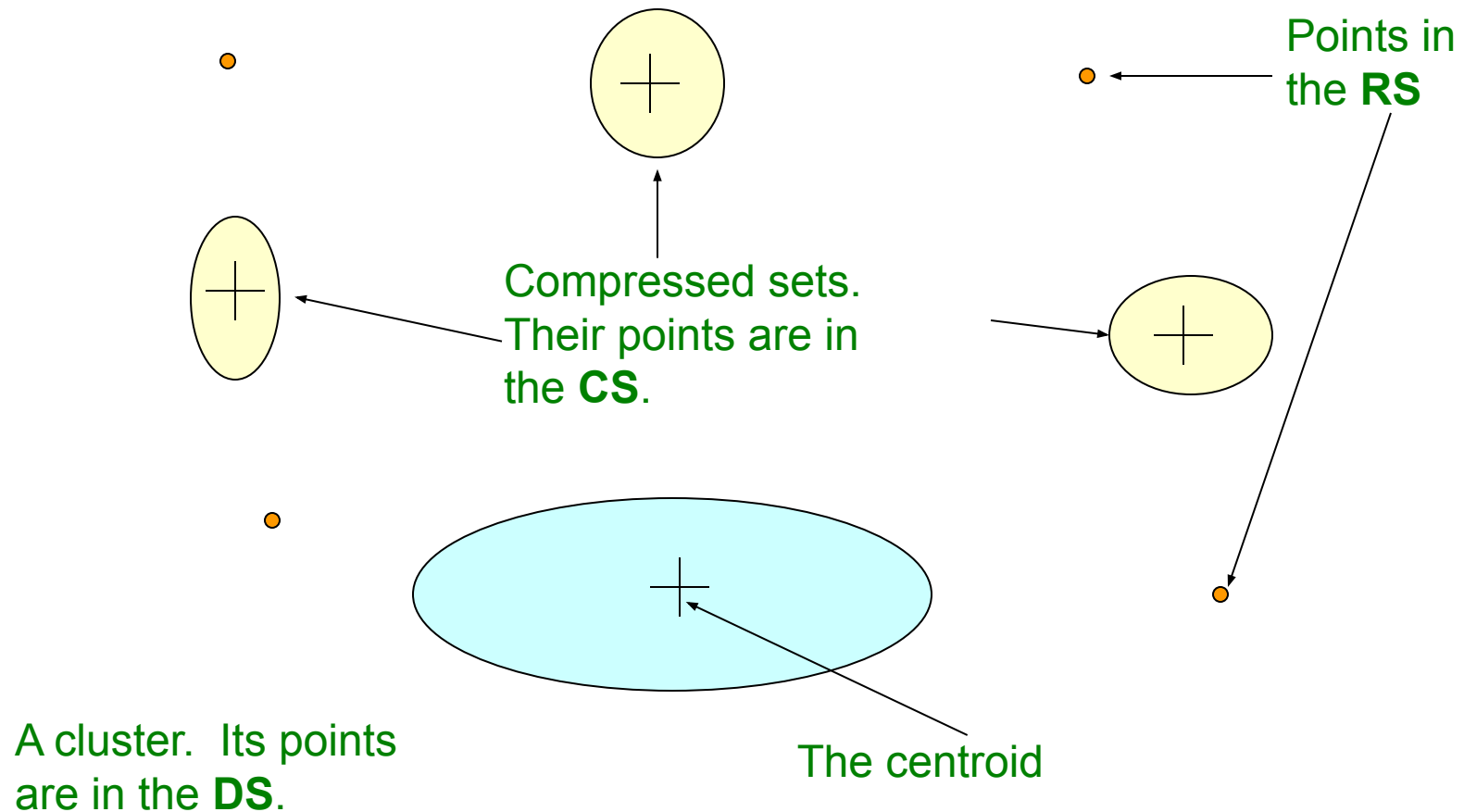
**Discard set (DS):** Close enough to a centroid to be summarized.

**Compression set (CS):** Summarized, but not assigned to a cluster

**Retained set (RS):** Isolated points



# BFR: “Galaxies” Picture



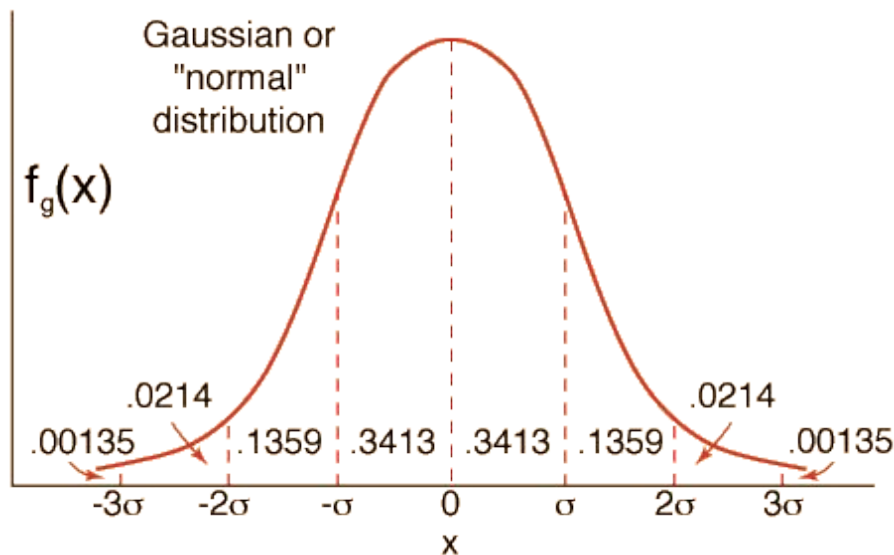
**Discard set (DS):** Close enough to a centroid to be summarized  
**Compression set (CS):** Summarized, but not assigned to a cluster  
**Retained set (RS):** Isolated points

# A Few Details...

- 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?**

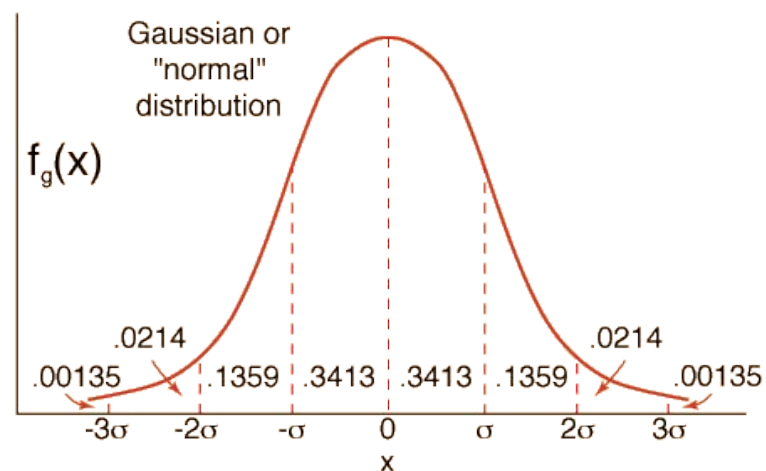
# Define “Sufficiently Close”

- ~68% of points:  $1\sigma$  away from mean
- ~95% of points:  $2\sigma$  away
- ~99% of points:  $3\sigma$  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 likelihood of the point belonging to currently nearest centroid



# Mahalanobis Distance

- **Normalized Euclidean distance from centroid**
- For point  $(x_1, \dots, x_d)$  and centroid  $(c_1, \dots, c_d)$ 
  1. Normalize by  $\sigma_i$  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

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

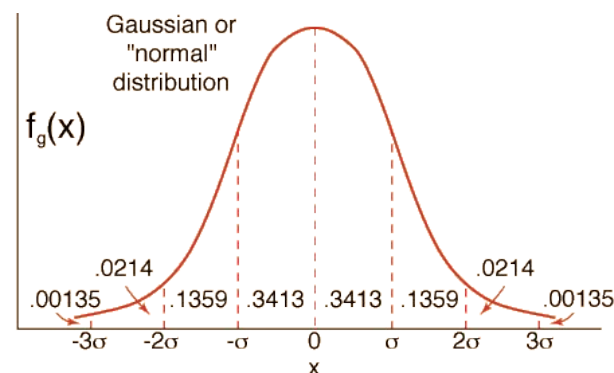
$\sigma_i$  ... standard deviation of points in the cluster in the  $i^{\text{th}}$  dimension

# 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**

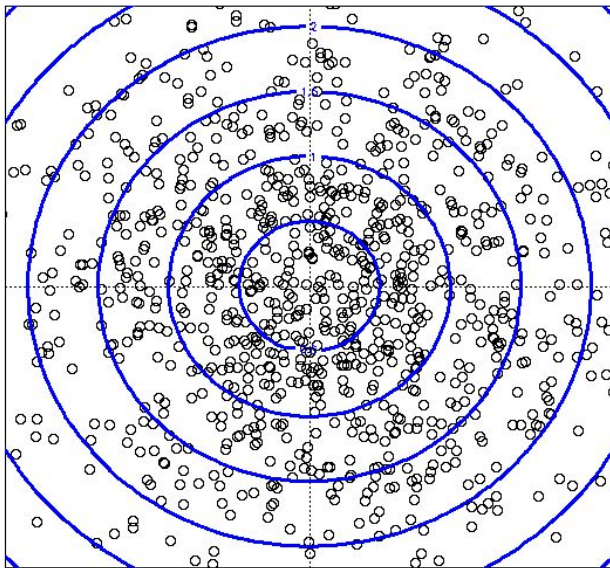
- About 68% of values are within  $\sigma$  away from the mean; about 95%  $2\sigma$  away; and about 99.7%  $3\sigma$  away.



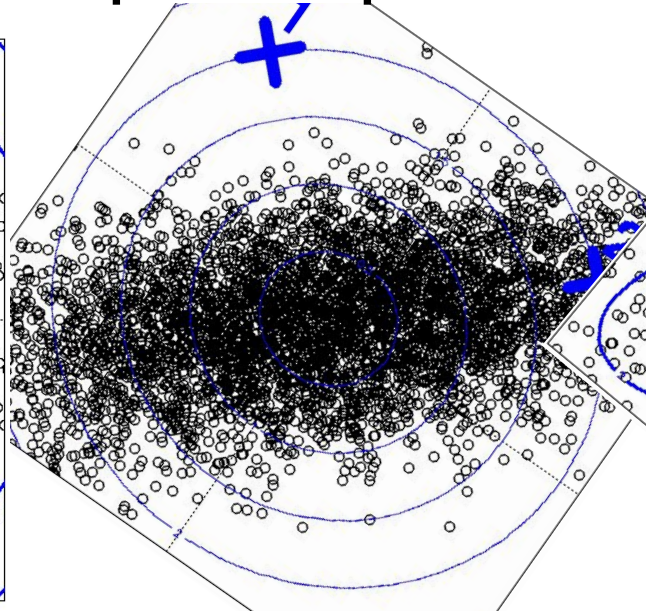
# Picture: Equal M.D. Regions

## ■ Euclidean vs. Mahalanobis distance

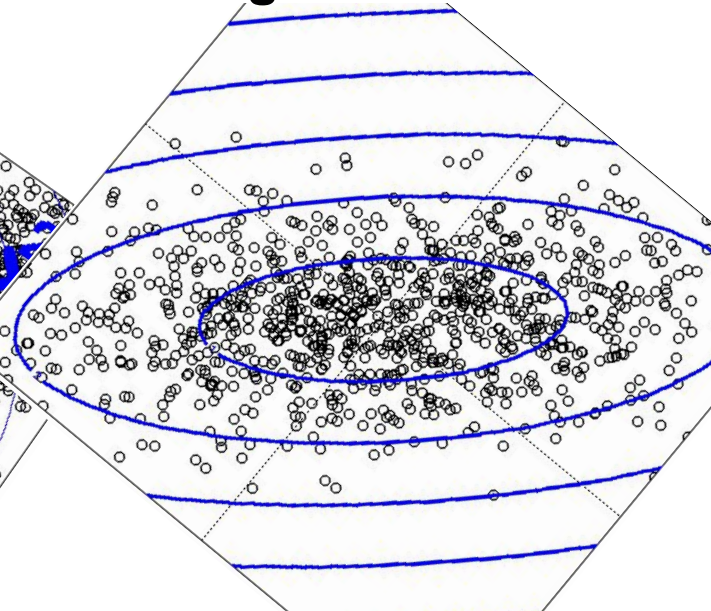
Contours of equidistant points from the origin



Uniformly distributed points,  
Euclidean distance



Normally distributed points,  
Euclidean distance

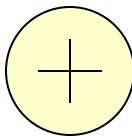
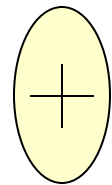


Normally distributed points,  
Mahalanobis distance

# 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





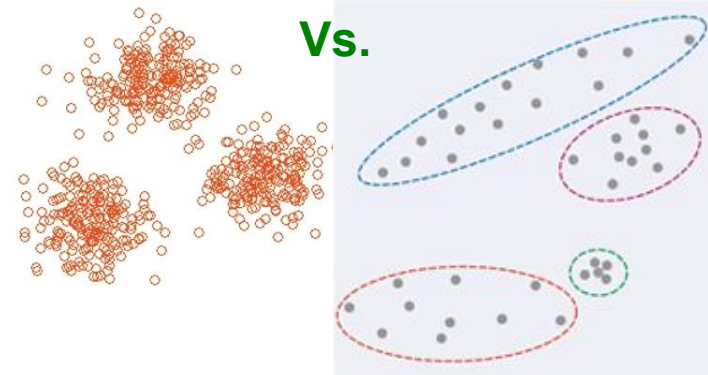
# The CURE Algorithm

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

# The CURE Algorithm

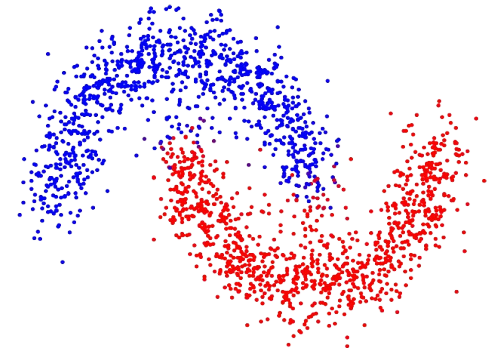
## ■ Problem with BFR/ $k$ -means:

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

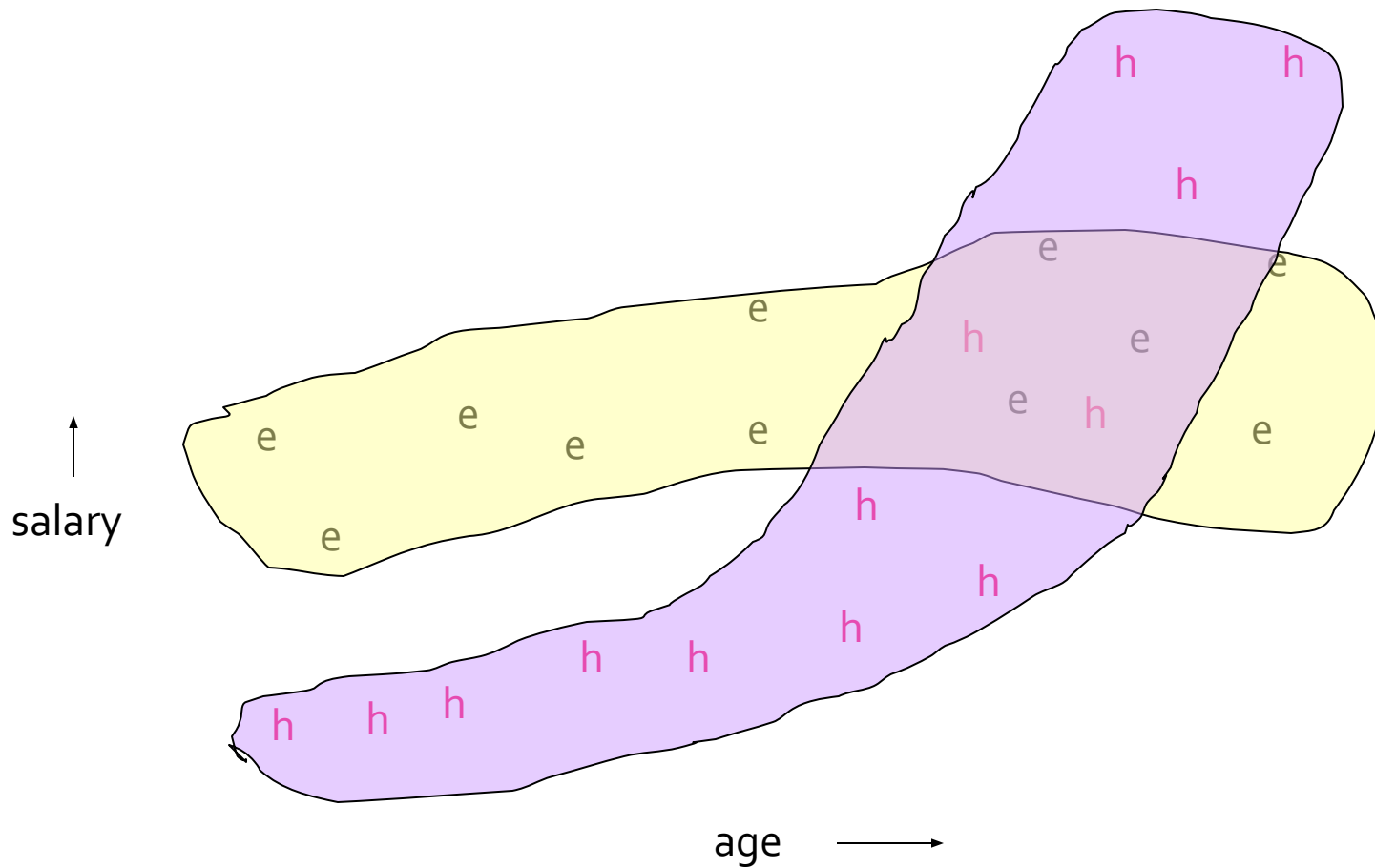


## ■ CURE (Clustering Using REpresentatives):

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



# Example: Stanford Salaries

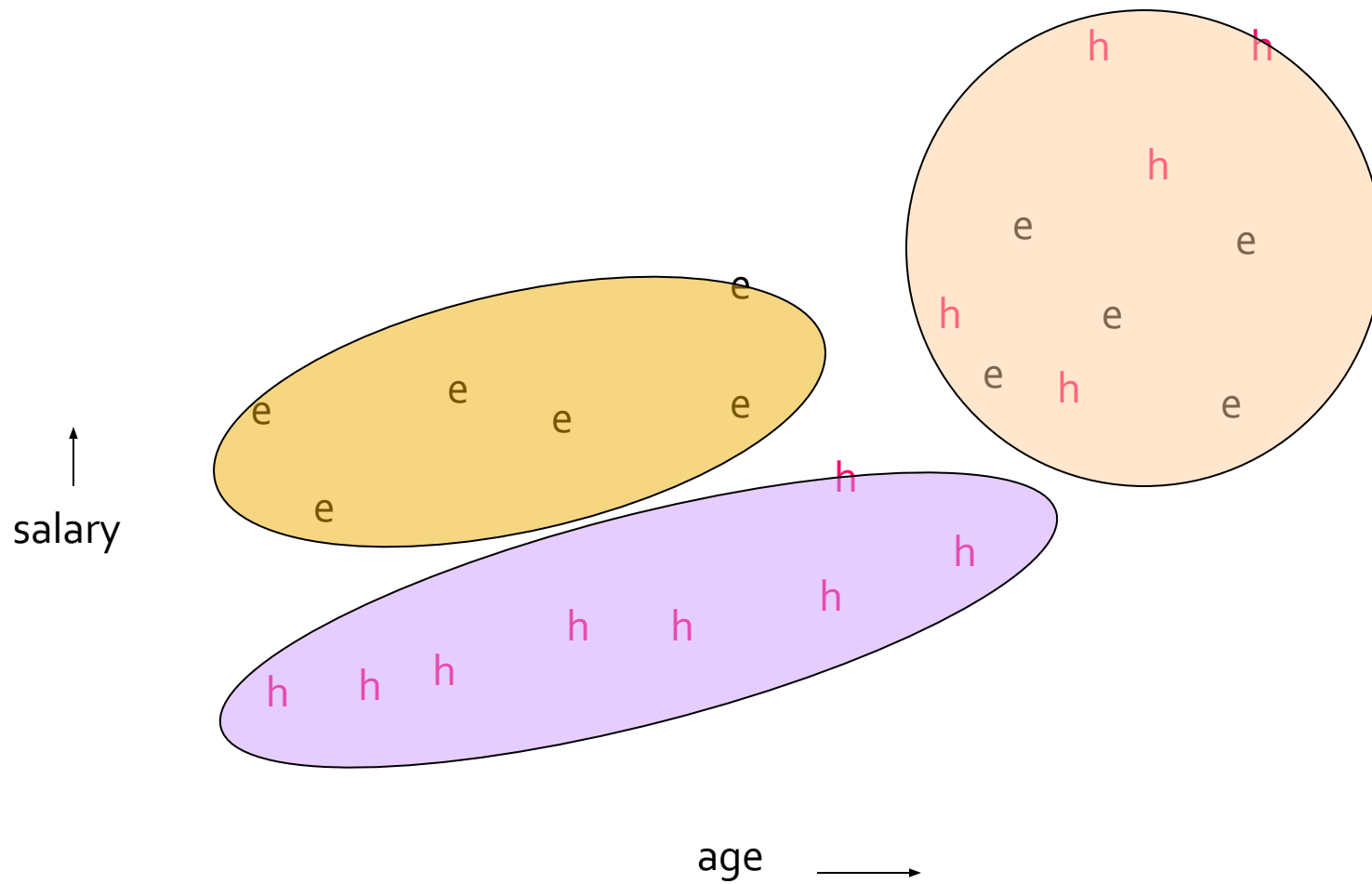


# Starting CURE

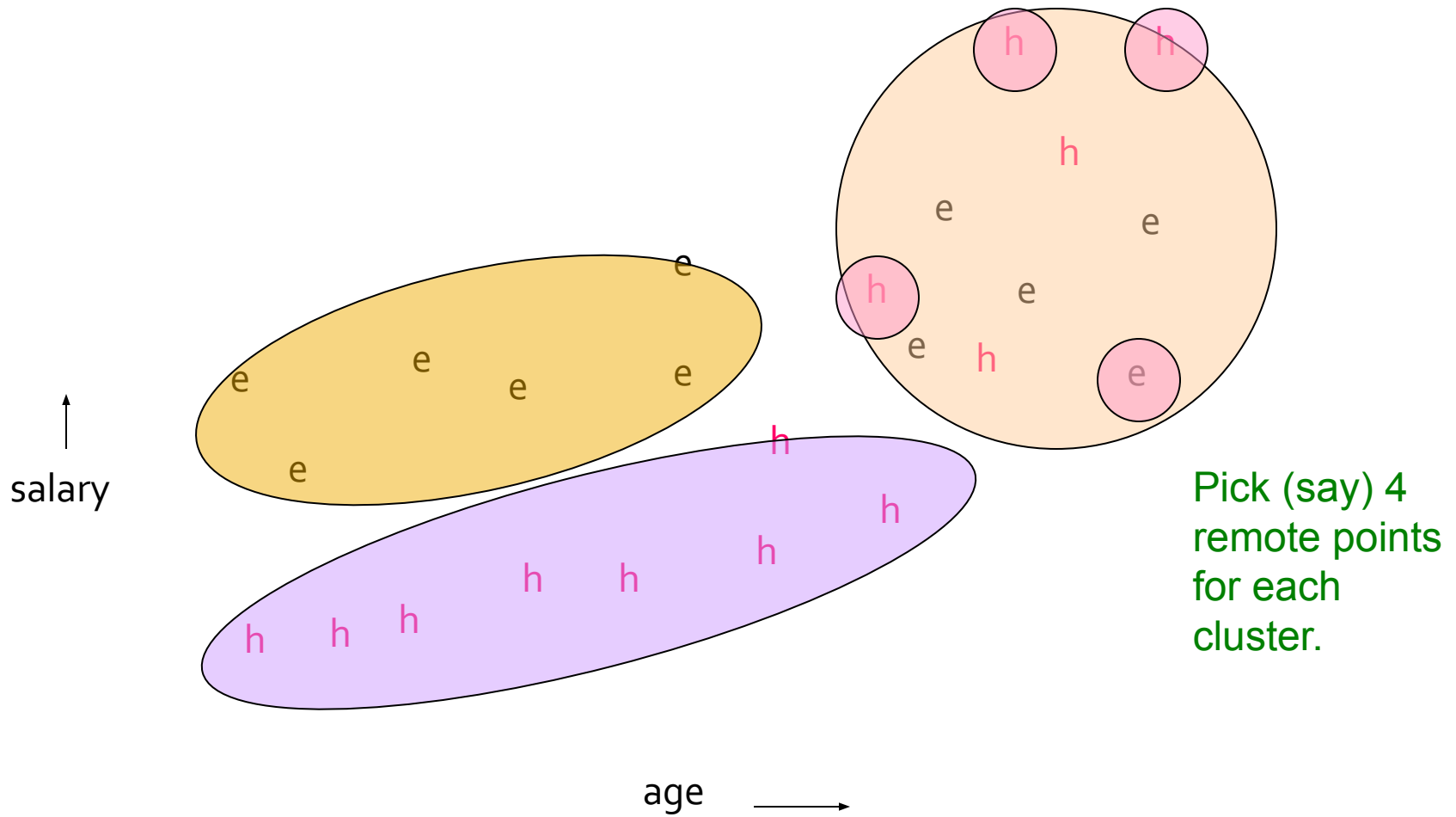
## Two-Pass Algorithm: Pass 1

- 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

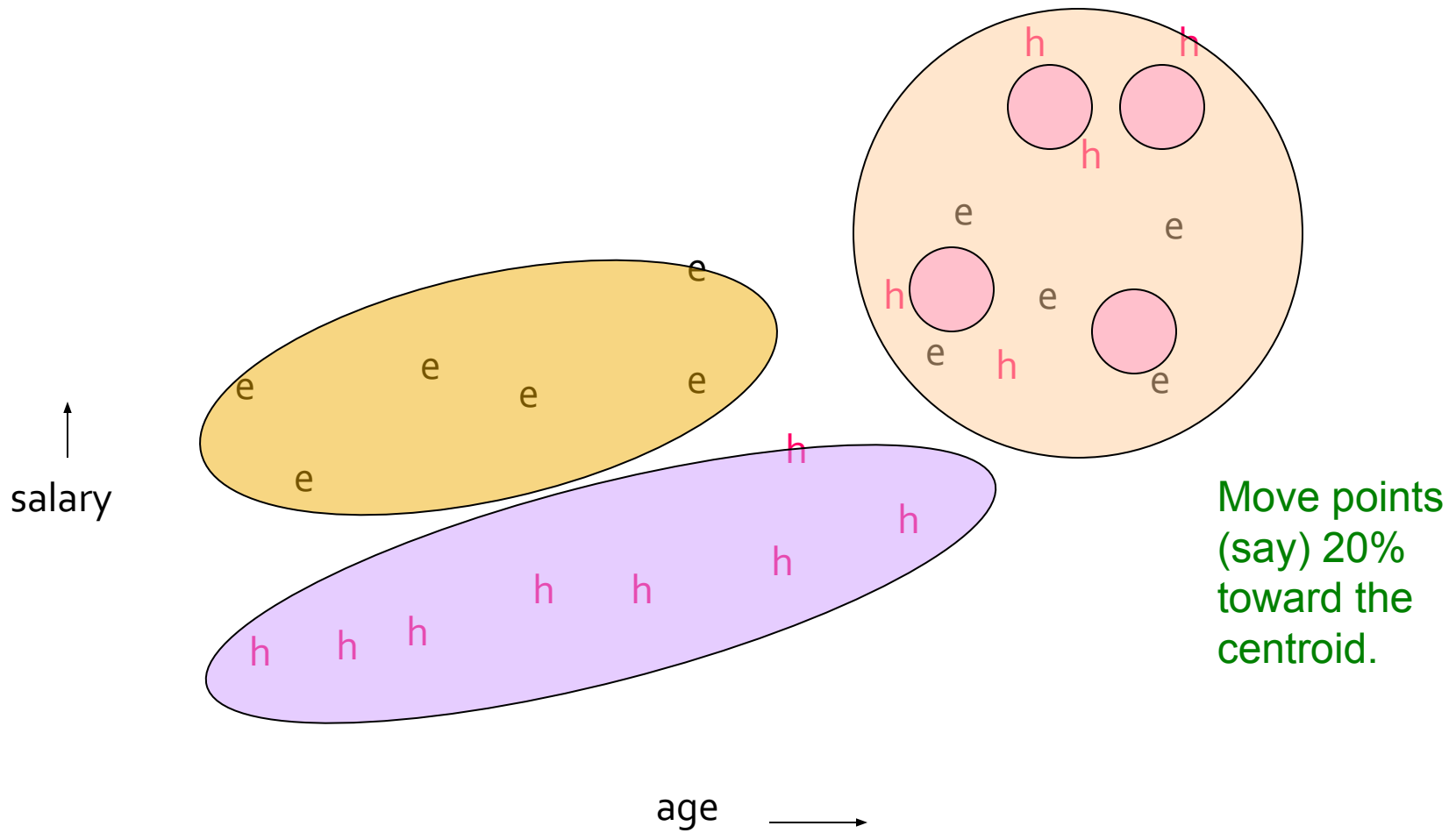
# Example: Initial Clusters



# Example: Pick Dispersed Points



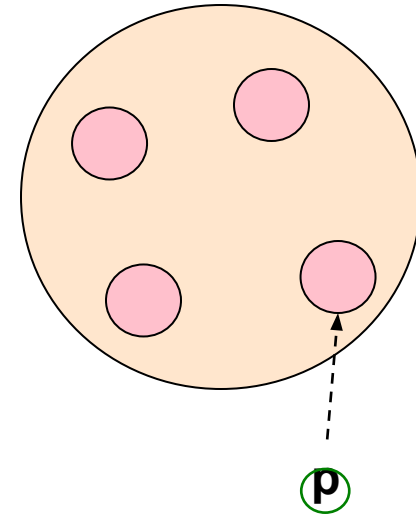
# Example: Pick Dispersed Points



# 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





# 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