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# Clustering

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# Supervised learning vs. unsupervised learning

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- *Supervised* learning: discover patterns in the data that relate data attributes with target attributes
  - These patterns are then utilized to predict the values of target attributes in future data instances
- *Unsupervised* learning: The data have no target attribute
  - Find some intrinsic structures in data

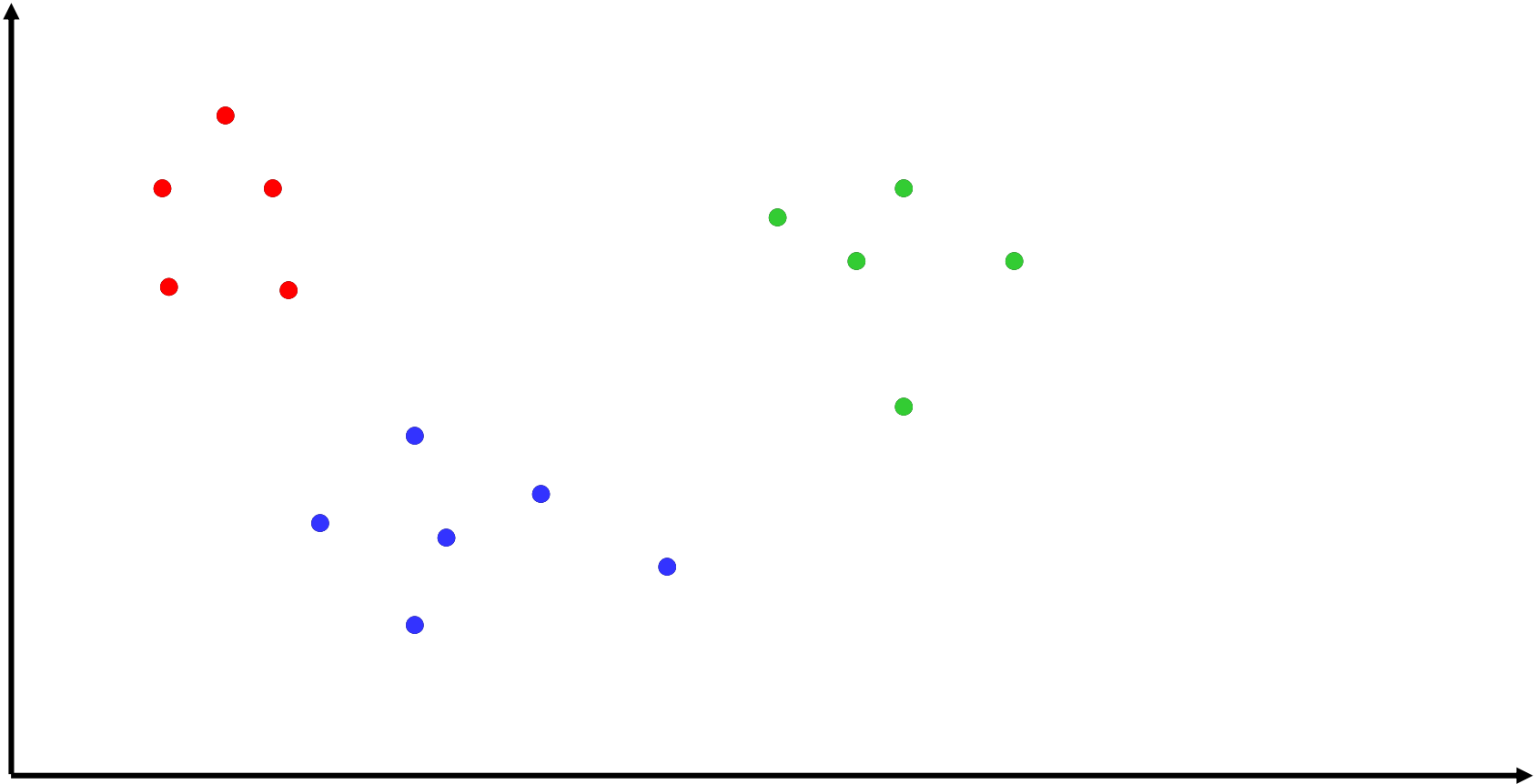
# Clustering

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- Partition unlabeled examples into disjoint subsets of *clusters*, such that:
  - Examples within a cluster are very similar (*infra-cluster* similarity)
  - Examples in different clusters are very different (*inter-cluster* dissimilarity)
- Discover new categories in an *unsupervised* manner
- Due to historical reasons, clustering is often considered synonymous with unsupervised learning.
  - In fact, association rule mining is also unsupervised

# Clustering Example

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# Application examples

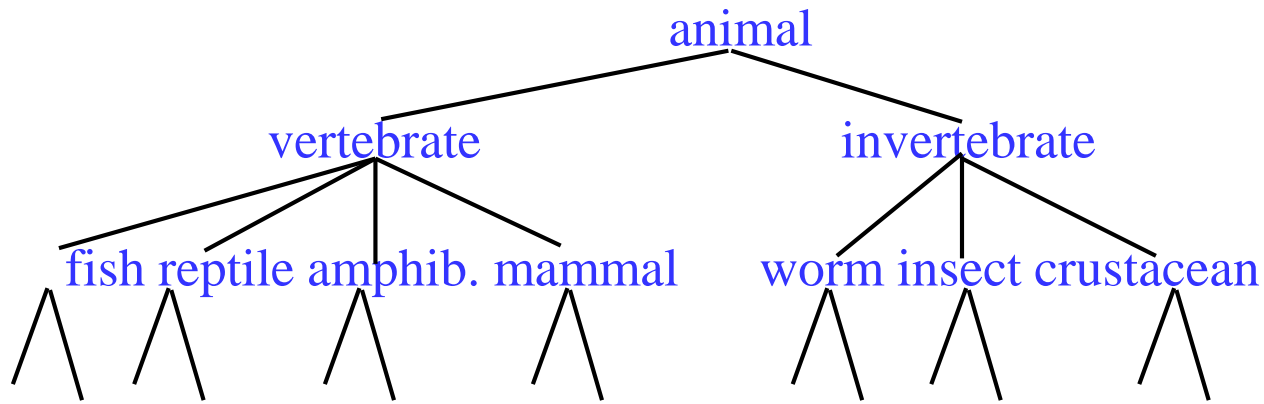
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- **Example 1:** In marketing, segment customers according to their similarities
  - To do targeted marketing
- **Example 2:** Given a collection of text documents, we want to organize them according to their content similarities
  - To produce a topic hierarchy

# Hierarchical Clustering

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- Build a tree-based hierarchical taxonomy (*dendrogram*) from a set of unlabeled examples



- Recursive application of a standard clustering algorithm can produce a hierarchical clustering.

# Direct Clustering

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- *Direct clustering* methods require a specification of the number of desired clusters,  $k$
- A *clustering evaluation function* assigns a real-value quality measure to a clustering.
- The number of clusters can be determined automatically by explicitly generating clusterings for multiple values of  $k$  and choosing the best result according to a clustering evaluation function.

# Aspects of clustering

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- **A clustering algorithm**
  - Single Link Agglomerative Clustering
  - K-Means
  - ...
- **A distance (similarity, or dissimilarity) function**
- **Clustering quality**
  - Inter-clusters distance  $\Rightarrow$  maximized
  - Intra-clusters distance  $\Rightarrow$  minimized
- The **quality** of a clustering result depends on the algorithm, the distance function, and the application



# Hierarchical Clustering: Agglomerative vs. Divisive Clustering

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- *Agglomerative* (*bottom-up*) methods start with each example in its own cluster and iteratively combine them to form larger and larger clusters
- *Divisive* (*partitional, top-down*) : It starts with all data points in one cluster, the root. Splits the root into a set of child clusters. Each child cluster is recursively divided further

# Hierarchical Agglomerative Clustering (HAC)

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- Assumes different *similarity functions* for determining the similarity of two instances
- Starts with all instances in a separate cluster and then repeatedly joins the two clusters that are most similar until there is only one cluster
- The history of merging forms a binary tree or hierarchy

# HAC Algorithm

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Start with all instances in their own cluster.

Until there is only one cluster:

Among the current clusters, determine the two clusters,  $c_i$  and  $c_j$ , that are most similar.

Replace  $c_i$  and  $c_j$  with a single cluster  $c_i \cup c_j$

# HAC Algorithm: Partition based on Cluster Similarity

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- How to compute similarity of two clusters each possibly containing multiple instances?
  - **Single Link**: Similarity of two most similar members
  - **Complete Link**: Similarity of two least similar members
  - **Group Average**: Average similarity between members

# Single Link Agglomerative Clustering

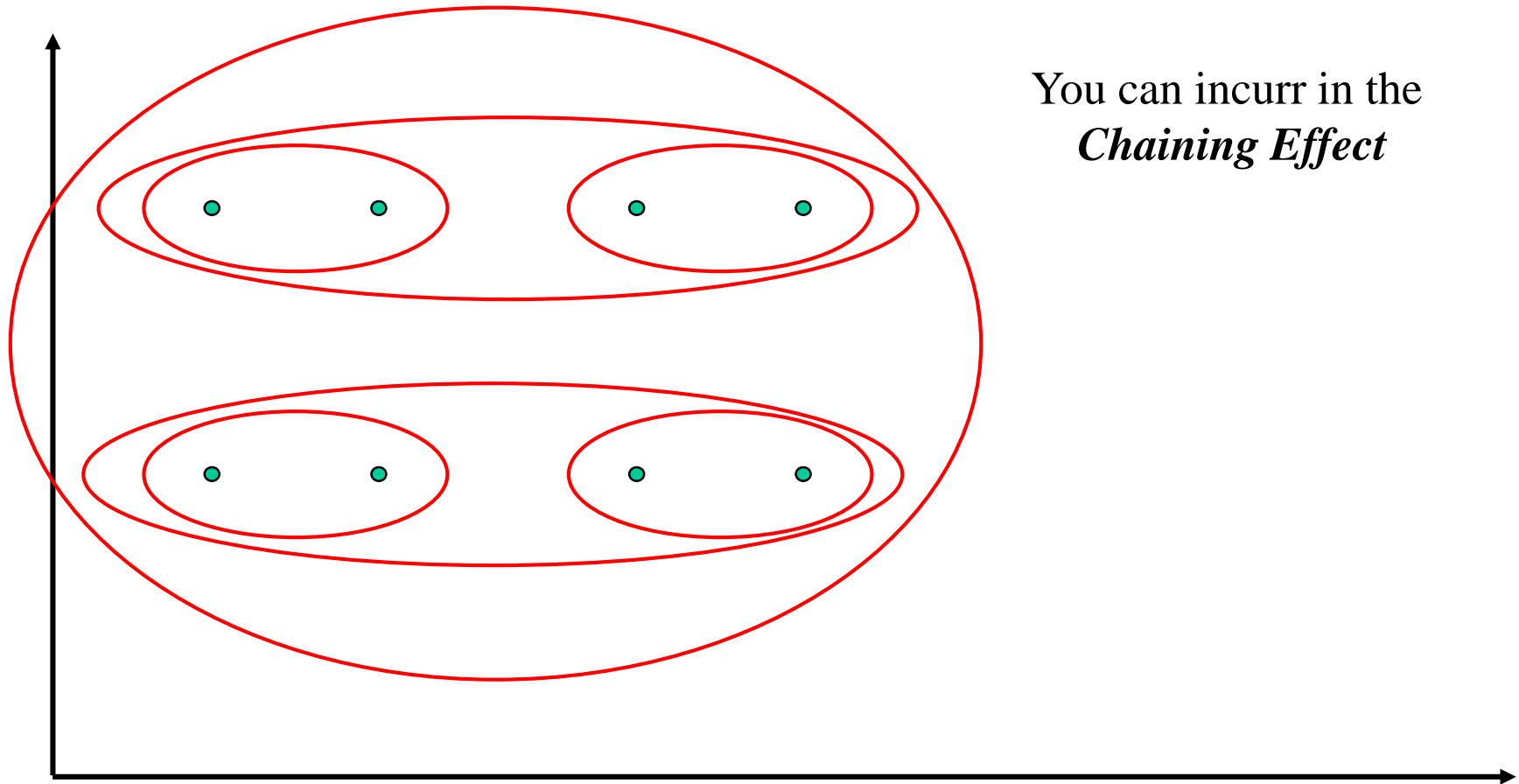
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- The distance between two clusters is the distance between two closest data points in the two clusters
- Use maximum similarity of pairs:

$$\mathit{sim}(C_i, C_j) = \max_{x \in C_i, y \in C_j} \mathit{sim}(x, y)$$

# Single Link Example

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You can incur in the  
*Chaining Effect*

# Complete Link Agglomerative Clustering

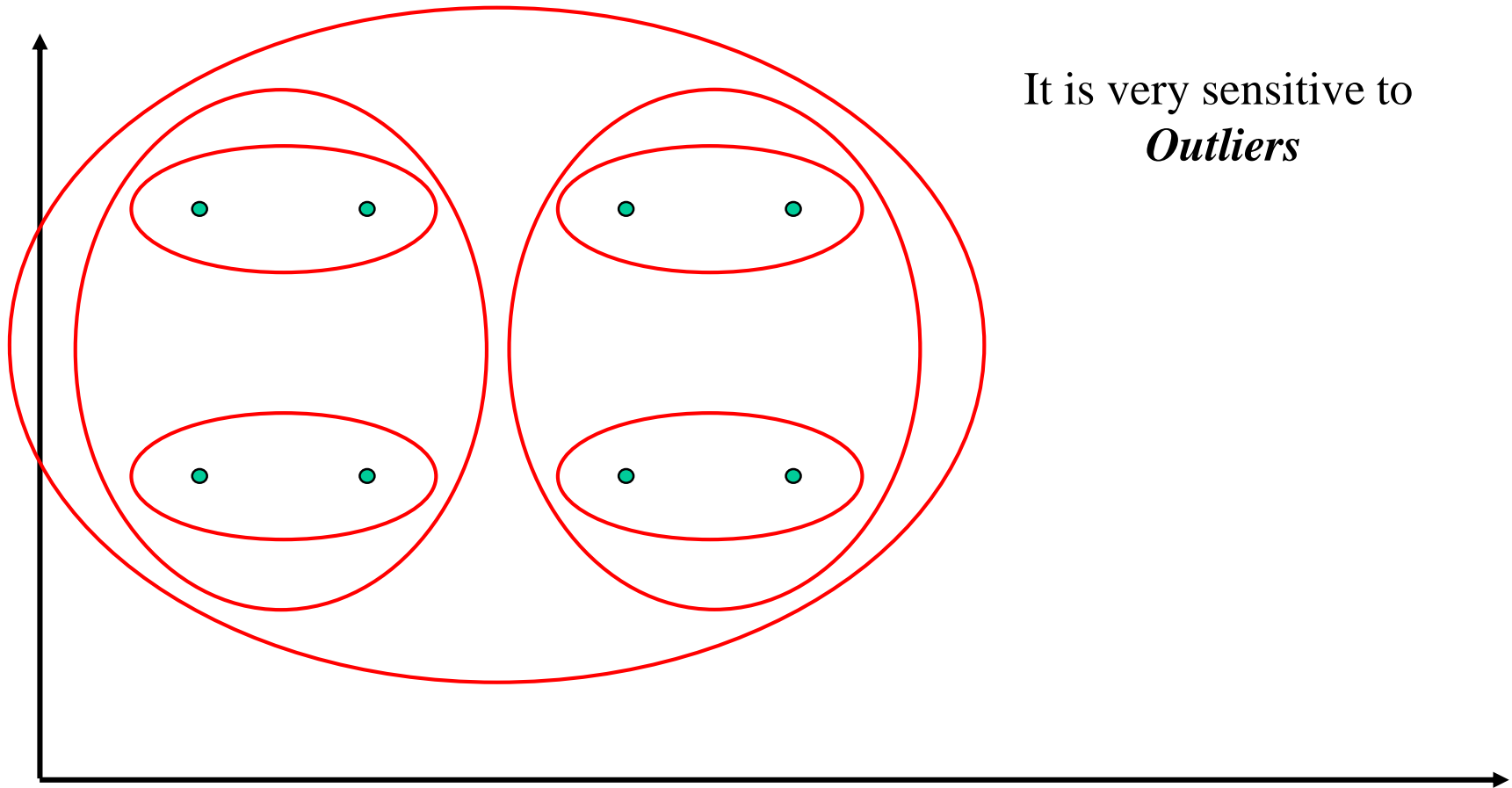
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- The distance between two clusters is the distance of two furthest data points in the two clusters
- Use minimum similarity of pairs:

$$\mathit{sim}(C_i, C_j) = \min_{x \in C_i, y \in C_j} \mathit{sim}(x, y)$$

# Complete Link Example

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It is very sensitive to  
*Outliers*



# Group Average Agglomerative Clustering

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- Use average similarity across all pairs within the merged cluster to measure the similarity of two clusters

$$sim(c_i, c_j) = \frac{1}{|c_i \cup c_j|(|c_i \cup c_j| - 1)} \sum_{\substack{\vec{x} \in (c_i \cup c_j) \\ \vec{y} \in (c_i \cup c_j): \vec{y} \neq \vec{x}}} sim(\vec{x}, \vec{y})$$

- Compromise between single and complete link
- Averaged across all ordered pairs in the merged cluster instead of unordered pairs *between* the two clusters

# Computational Complexity

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- In the first iteration, all HAC methods need to compute similarity of all pairs of  $n$  individual instances which is  $O(n^2)$
- In each of the subsequent  $n-2$  merging iterations, it must compute the distance between all existing clusters
- In order to maintain an overall  $O(n^2)$  performance, computing similarity to each other cluster must be done in constant time.

# Computing Group Average Similarity

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- Assume cosine similarity and normalized vectors with unit length
- Always maintain sum of vectors in each cluster

$$\vec{s}(c_j) = \frac{\sum_{\vec{x} \in c_j} \vec{x}}{|c_j|}$$

- Compute similarity of clusters in constant time:

$$\text{sim}(c_i, c_j) = \frac{(\vec{s}(c_i) + \vec{s}(c_j)) \cdot (\vec{s}(c_i) + \vec{s}(c_j)) - (|c_i| + |c_j|)}{(|c_i| + |c_j|)(|c_i| + |c_j| - 1)}$$

# Non-Hierarchical Clustering

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- Typically must provide the number of desired clusters,  $k$
- Randomly choose  $k$  instances as *seeds*, one per cluster
- Form initial clusters based on these seeds
- Iterate, repeatedly reallocating instances to different clusters to improve the overall clustering
- Stop when clustering converges or after a fixed number of iterations

# K-Means

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- Assumes instances are real-valued vectors
- Clusters based on *centroids*, *center of gravity*, or mean of points in a cluster,  $c$ :

$$\vec{\mu}(c) = \frac{1}{|c|} \sum_{\vec{x} \in c} \vec{x}$$

- Reassignment of instances to clusters is based on distance to the current cluster centroids

# K-Means Algorithm

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Let  $d$  be the distance measure between instances.

Select  $k$  random instances  $\{s_1, s_2, \dots, s_k\}$  as seeds.

Until clustering converges or other stopping criterion:

For each instance  $x_i$ :

Assign  $x_i$  to the cluster  $c_j$  such that  $d(x_i, s_j)$  is minimal.

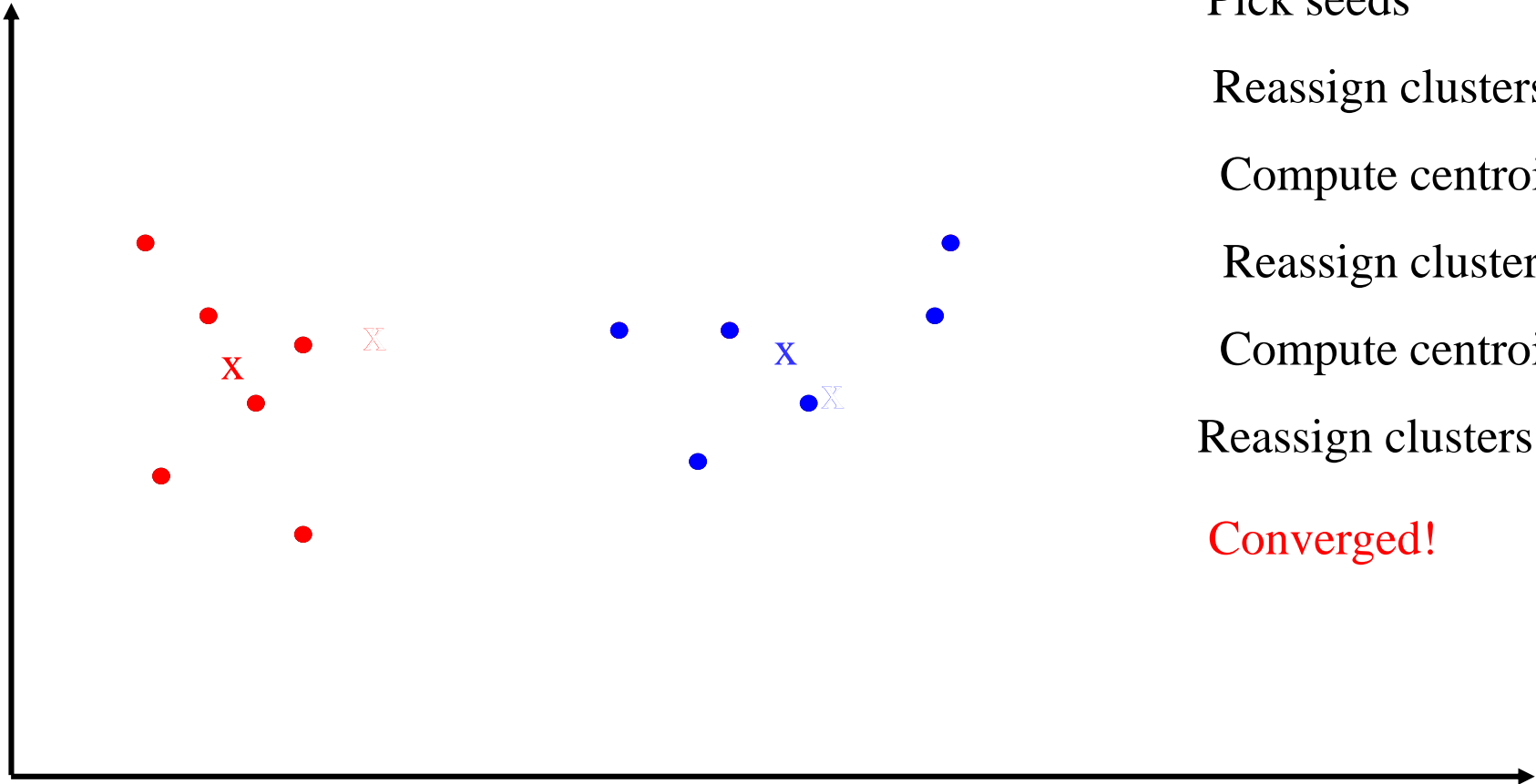
*(Update the seeds to the centroid of each cluster)*

For each cluster  $c_j$

$$s_j = \mu(c_j)$$

# K Means Example (K=2)

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Pick seeds

Reassign clusters

Compute centroids

Reassign clusters

Compute centroids

Reassign clusters

**Converged!**

# K-Means stopping criteria

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- No or minimum reassignment of data in clusters
- No or minimum change in centroids
- Minimum decrease in the sum of squared error (SSE)

$$SSE = \sum_{j=1}^k \sum_{\mathbf{x} \in C_j} \text{dist}(\mathbf{x}, \mathbf{m}_j)^2$$

$C_j$  is the  $j$ -th cluster,  $\mathbf{m}_j$  is the centroid of cluster  $C_j$  (the mean vector of all the data points in  $C_j$ ), and  $\text{dist}(\mathbf{x}, \mathbf{m}_j)$  is the distance between data point  $\mathbf{x}$  and centroid  $\mathbf{m}_j$ .



# K-Mean - Time Complexity

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- Assume computing distance between two instances is  $O(m)$  where  $m$  is the dimensionality of the vectors
- Reassigning clusters:  $O(kn)$  distance computations, or  $O(knm)$
- Computing centroids: Each instance vector gets added once to some centroid:  $O(nm)$
- Assume these two steps are each done once for  $I$  iterations:  $O(Iknm)$
- Linear in all relevant factors, assuming a fixed number of iterations, more efficient than  $O(n^2)$  HAC

# Distance Metrics

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- Euclidian distance ( $L_2$  norm):

$$L_2(\vec{x}, \vec{y}) = \sqrt{\sum_{i=1}^m (x_i - y_i)^2}$$

- $L_1$  norm:

$$L_1(\vec{x}, \vec{y}) = \sum_{i=1}^m |x_i - y_i|$$

- Cosine Similarity (transform to a distance by subtracting from 1):

$$1 - \frac{\vec{x} \cdot \vec{y}}{|\vec{x}| \times |\vec{y}|}$$

# Distance Metrics

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- Chebychev distance
  - define two data points as "different" if they are different on any one of the attributes

$$\text{dist}(\mathbf{x}_i, \mathbf{x}_j) = \max(|x_{i1} - x_{j1}|, |x_{i2} - x_{j2}|, \dots, |x_{ir} - x_{jr}|)$$

# Data standardization

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- In the Euclidean space, standardization of attributes is recommended so that all attributes can have equal impact on the computation of distances
- Consider the following pair of data points
  - $\mathbf{x}_i$ : (0.1, 20) and  $\mathbf{x}_j$ : (0.9, 720).

$$\text{dist}(\mathbf{x}_i, \mathbf{x}_j) = \sqrt{(0.9 - 0.1)^2 + (720 - 20)^2} = 700.000457,$$

- The distance is almost completely dominated by  $(720-20) = 700$
- **Standardize attributes**: to force the attributes to have a common value range

# Interval-scaled attributes

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- Their values are real numbers following a linear scale
  - The difference in Age between 10 and 20 is the same as that between 40 and 50
  - The key idea is that intervals keep the same importance through out the scale
- Two main approaches to standardize interval scaled attributes, **range** and **z-score**.  $f$  is an attribute

$$\text{range}(x_{if}) = \frac{x_{if} - \min(f)}{\max(f) - \min(f)},$$

# Interval-scaled attributes (cont ...)

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- **Z-score**: transforms the attribute values so that they have a mean of zero and a **mean absolute deviation** of 1. The mean absolute deviation of attribute  $f$ , denoted by  $s_f$ , is computed as follows

$$s_f = \frac{1}{n} (|x_{1f} - m_f| + |x_{2f} - m_f| + \dots + |x_{nf} - m_f|),$$

$$m_f = \frac{1}{n} (x_{1f} + x_{2f} + \dots + x_{nf}),$$

$$z(x_{if}) = \frac{x_{if} - m_f}{s_f}.$$

# Clustering evaluation

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- How can we evaluate produced clusters?
- We can use some *internal criteria* for the quality of a clustering
  - Typical objective functions can formalize the goal of attaining high intra-cluster similarity and low inter-cluster similarity
  - But good scores on an internal criterion do not necessarily translate into good effectiveness in an application
- It is better to adopt some *external criteria*
  - we can use a set of classes in an evaluation benchmark or gold standard
- Or we can use some *indirect evaluation criteria*
  - In some applications, clustering is not the primary task, but used to help perform another task.
  - We can use the performance on the primary task to compare clustering methods.

# Cluster evaluation: External Criteria

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- We use some labeled data (for classification)
- **Assumption:** Each class is a cluster
- After clustering, build a confusion matrix
- From the matrix, compute various measurements: Entropy, Purity, Precision, Recall and F-score
  - Let the classes in the data  $D$  be  $C = (c_1, c_2, \dots, c_k)$ . The clustering method produces  $k$  clusters, which divides  $D$  into  $k$  disjoint subsets,  $D_1, D_2, \dots, D_k$ .
  - We can estimate  $Pr_i(c_j)$ , i.e. the proportion of class  $c_j$  data points in cluster  $i$  or  $D_i$

$$Pr_i(c_j) = |c_j \cap D_i| / |D_i|$$



# Evaluation measures: purity

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**Purity:** This again measures the extent that a cluster contains only one class of data. The purity of each cluster is computed with

$$purity(D_i) = \max_j (\Pr_i(c_j)) \quad (31)$$

The total purity of the whole clustering (considering all clusters) is

$$purity_{total}(D) = \sum_{i=1}^k \frac{|D_i|}{|D|} \times purity(D_i) \quad (32)$$

If all clusters contain one instance only, the purity will be maximim, i.e. equal to 1

# Evaluation measures: Entropy

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**Entropy:** For each cluster, we can measure its entropy as follows:

$$entropy(D_i) = -\sum_{j=1}^k \Pr_i(c_j) \log_2 \Pr_i(c_j), \quad (29)$$

where  $\Pr_i(c_j)$  is the proportion of class  $c_j$  data points in cluster  $i$  or  $D_i$ . The total entropy of the whole clustering (which considers all clusters) is

$$entropy_{total}(D) = \sum_{i=1}^k \frac{|D_i|}{|D|} \times entropy(D_i) \quad (30)$$

# Soft Clustering

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- Clustering typically assumes that each instance is (hard) assigned to exactly one cluster
  - Does not allow uncertainty in class membership or for an instance to belong to more than one cluster
- *Soft clustering* gives probabilities to instances of belonging to each clusters
  - ES: Fuzzy C-mean
- Each instance has a probability distribution across a set of discovered categories (probabilities of all categories must sum to 1)

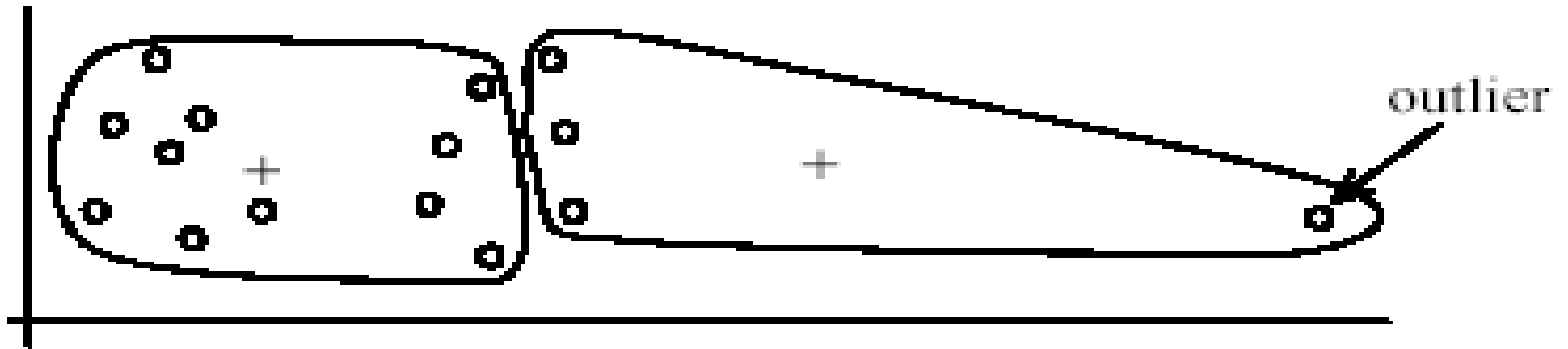
# Weaknesses of k-means

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- The algorithm is only applicable if the **mean** is defined
  - For categorical data, *k*-mode - the centroid is represented by most frequent values
- The algorithm is sensitive to **outliers**
  - Outliers are data points that are very far away from other data points
  - Outliers could be errors in the data recording or some special data points with very different values
- The user needs to specify *k*
- Results can vary based on random seed selection
- Some seeds can result in poor convergence rate, or convergence to sub-optimal clusterings
- Select good seeds using a heuristic or the results of another method

# Weaknesses of k-means: Problems with outliers

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(A): Undesirable clusters



(B): Ideal clusters

# Dealing with outliers

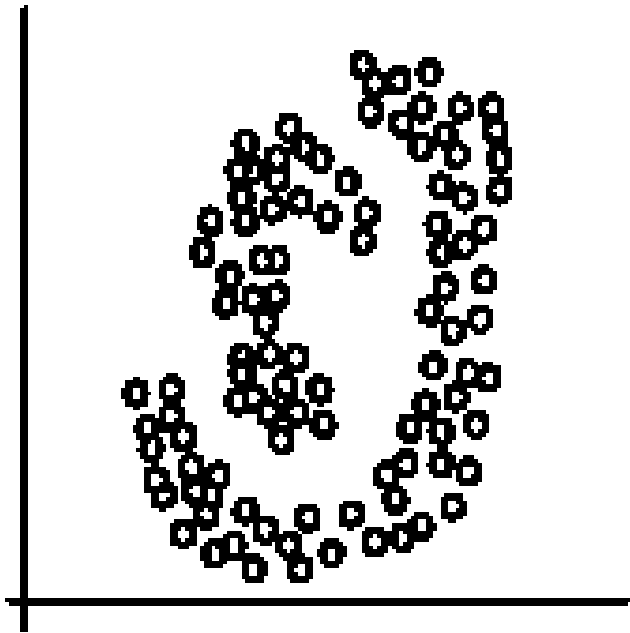
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- How to deal with outliers?
- One method is to remove some data points that are far from centroids
  - However, they can be important data
  - To be safe, monitor these points over multiple iterations before removing them
- Perform random sampling
  - Choose randomly points to partition
  - Choice of outliers is unlikely

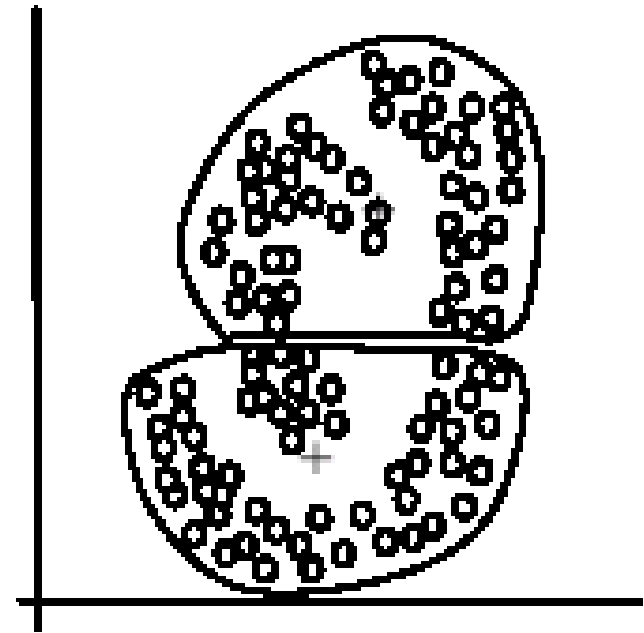
## Weaknesses of $k$ -means (cont ...)

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- The  $k$ -means algorithm is not suitable for discovering clusters that are not hyper-ellipsoids (or hyper-spheres)



(A): Two natural clusters



(B):  $k$ -means clusters

# Advanced Techniques

## QT K-Means Algorithm (1)

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- *Quality Threshold (QT) K-Means* Algorithm is an evolution of basic *K-Means* that dynamically change the number of cluster  $k$
- Use two threshold to consider both *inter-cluster* and *intra-cluster* similarity



# Advanced Techniques

## QT K-Means Algorithm (2)

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Let  $\sigma$  and  $\tau$  be two different thresholds.

Let  $d$  be the distance measure between instances.

Select  $k$  random instances  $\{s_1, s_2, \dots, s_k\}$  as seeds.

Until clustering converges or other stopping criterion:

For each instance  $x_i$ :

Assign  $x_i$  to the cluster  $c_j$  such that  $d(x_i, s_j)$  is minimal but less than  $\sigma$ .

Else create new seed with instance  $x_i$  (the number  $k$  of clusters increase)

*(Update the seeds to the centroid of each cluster)*

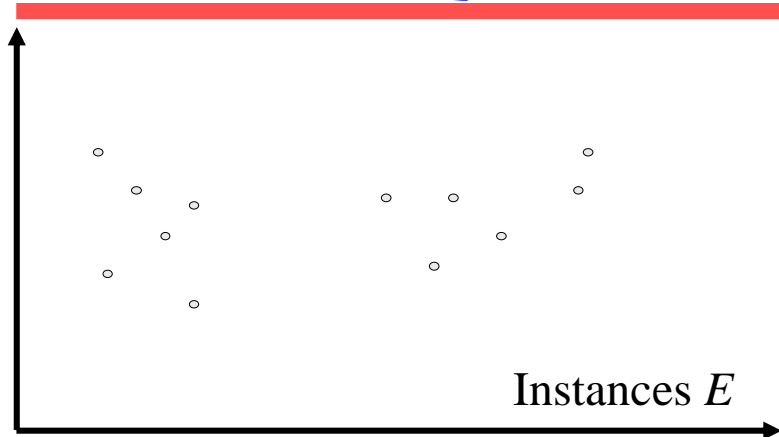
For each cluster pairs  $c_i, c_j$  to  $i \neq j$ :

If  $d(s_i, s_j)$  less than  $\tau$  merge  $c_i$  and  $c_j$  (the number  $k$  of clusters decrease)

# Advanced Techniques

## QT K-Means Algorithm (3)

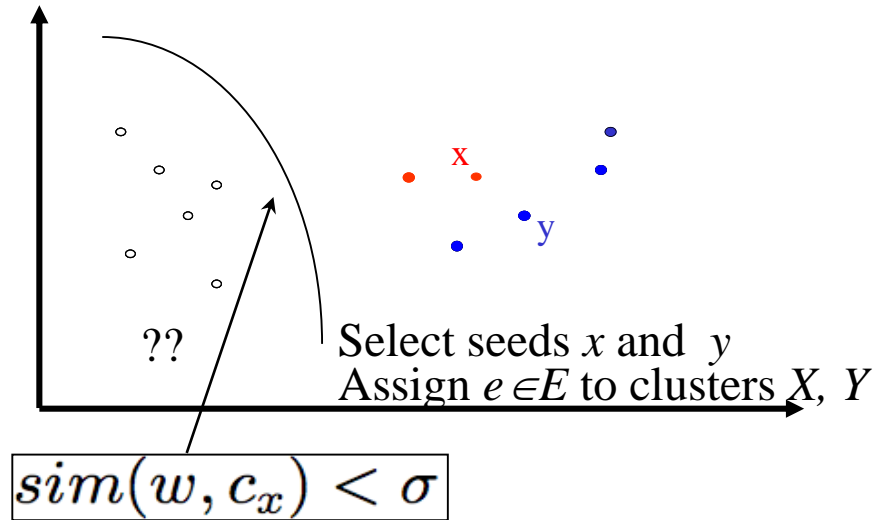
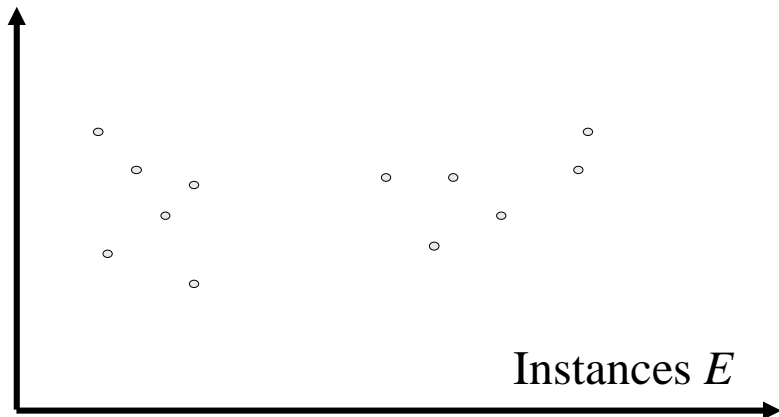
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# Advanced Techniques

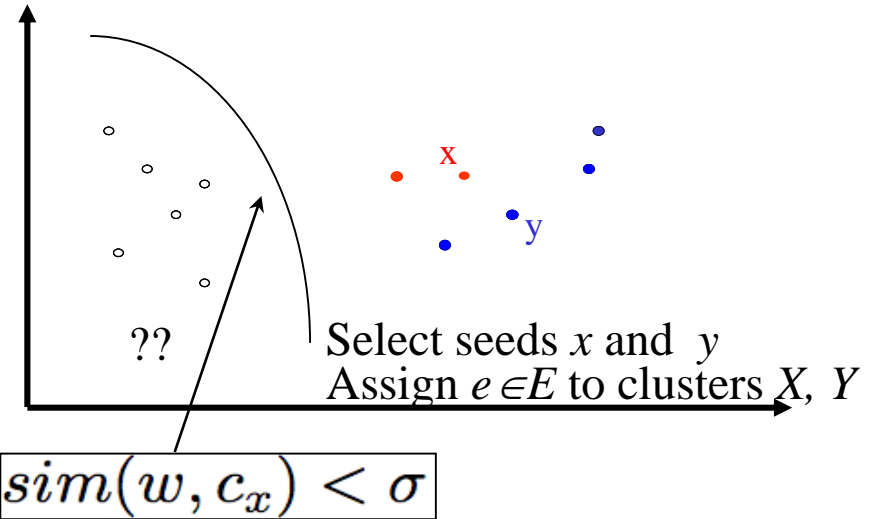
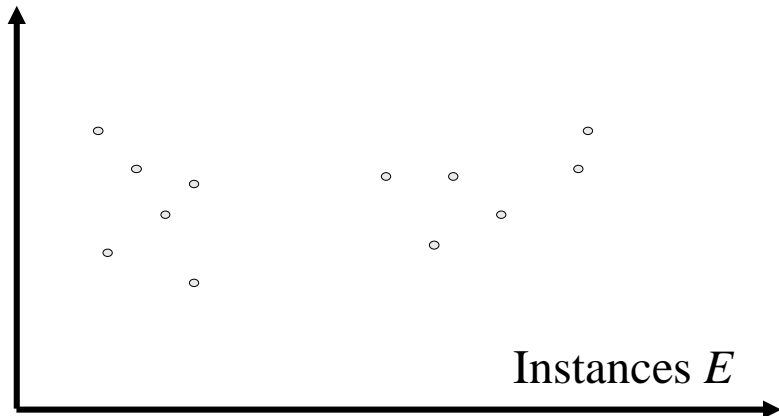
## QT K-Means Algorithm (3)

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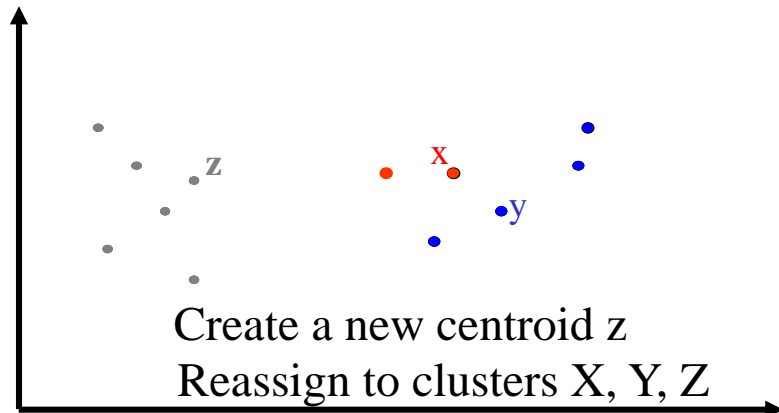


# Advanced Techniques

## QT K-Means Algorithm (3)

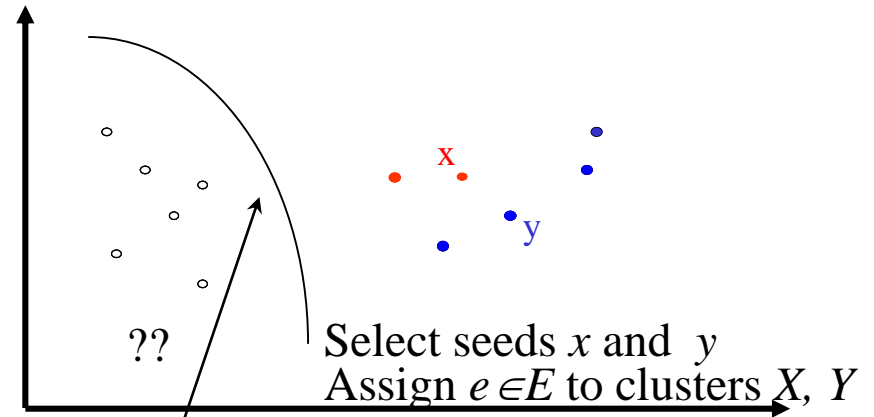
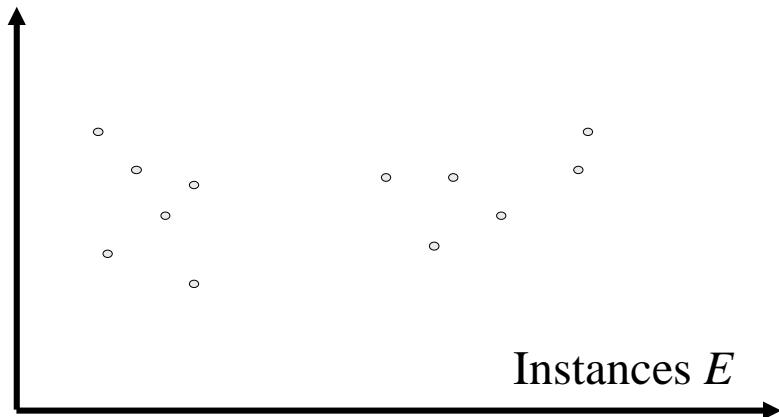


$$\text{sim}(w, c_x) < \sigma$$

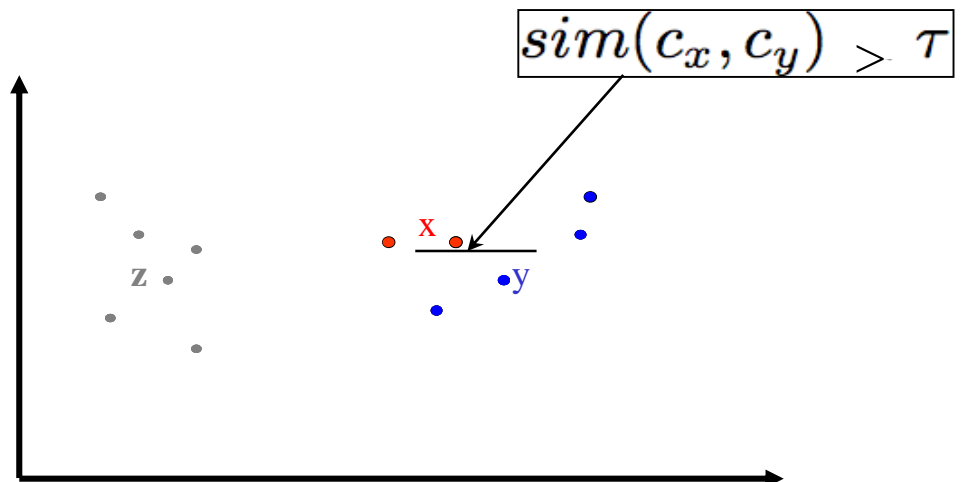
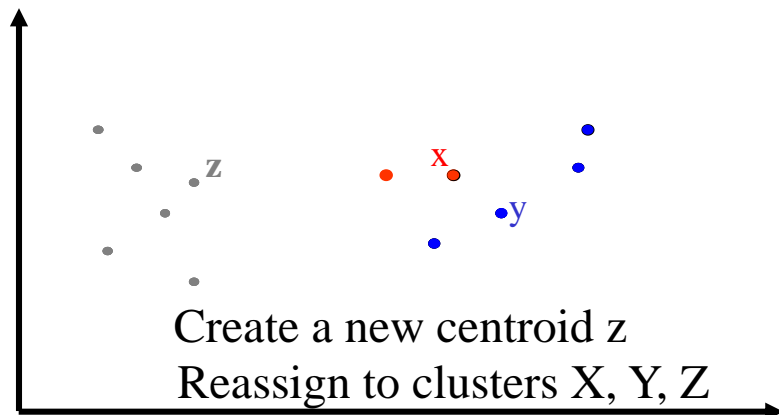


# Advanced Techniques

## QT K-Means Algorithm (3)



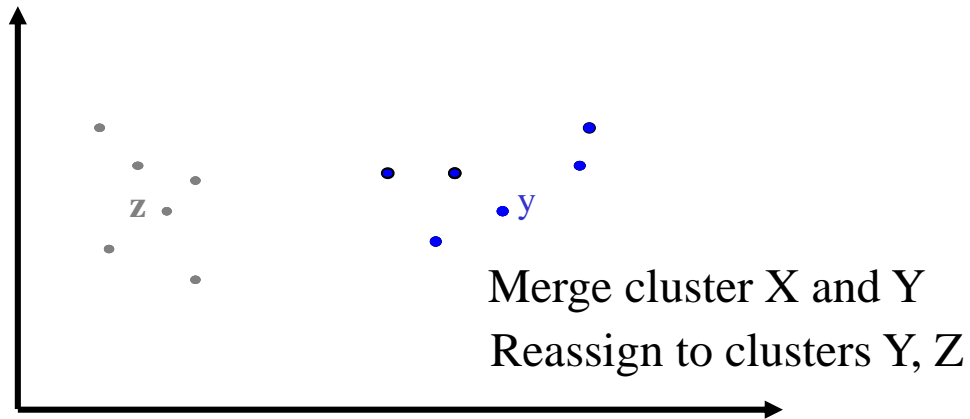
$$\text{sim}(w, c_x) < \sigma$$



# Advanced Techniques

## QT K-Means Algorithm (3)

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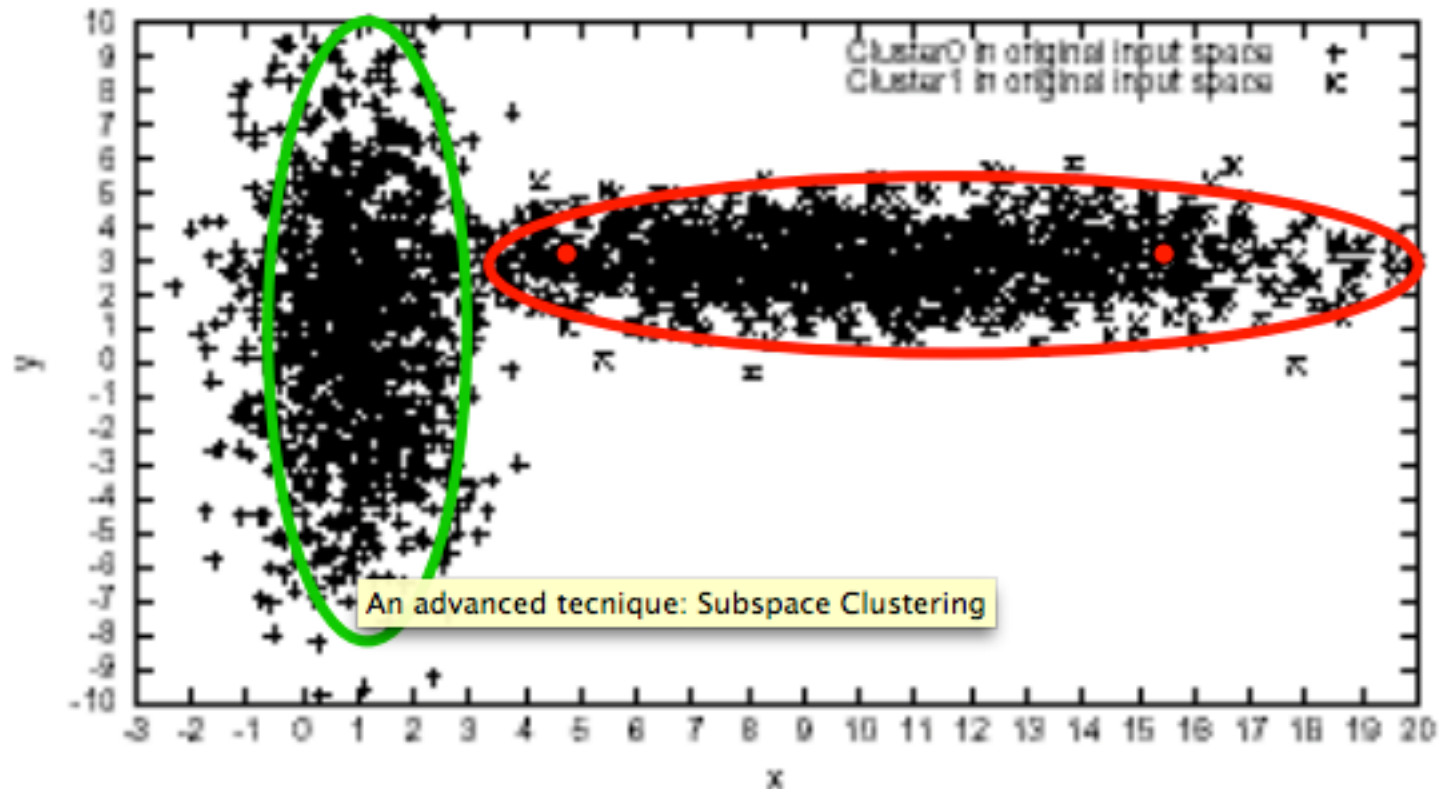
Convergence

# Advanced Techniques

## Subspace Clustering (1)

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- In high dimensional spaces, few dimensions can exist on which the points are far apart from each other

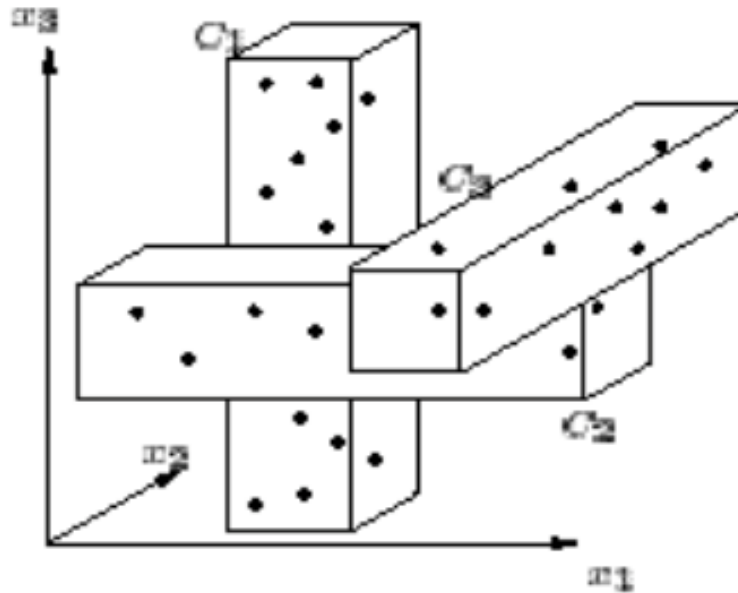


# An advanced technique

## Subspace Clustering (2)

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- **Subspace Clustering:** seek to find clusters in a dataset by selecting the most relevant dimensions for each cluster separately



Each dimension is relevant to at least one cluster



# A Subspace Clustering algorithm

## Locally Adaptive Clustering

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- We cannot prune off dimensions without incurring a loss of crucial information
- The data presents local structure:
  - To capture the local correlations of data a proper feature selection procedure should operate locally
  - A local operation would allow to embed different distance measures in different regions
- **IDEA:** apply a *Co-clustering* approach
  - simultaneous clustering of *both* data and dimensions

*Locally adaptive metrics for clustering high dimensional data*

Domeniconi et al, 2007

# A Subspace Clustering algorithm

## Locally Adaptive Clustering

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- LAC is a variant of K-Means where cluster are weighted
  - Each centroid is weighted so that only few dimensions are considered when associating data point to clusters
  - At each step the centroid weighting schema is update
  - In each cluster the weights determine the informative dimensions

*Locally adaptive metrics for clustering high dimensional data*

Domeniconi et al, 2007

# Some applications: Text Clustering

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- HAC and K-Means have been applied to text in a straightforward way
- Typically use *normalized*, TF/IDF-weighted vectors and cosine similarity
- Optimize computations for sparse vectors
- Applications:
  - During retrieval, add other documents in the same cluster as the initial retrieved documents to improve recall
  - Clustering of results of retrieval to present more organized results to the user (à la Northernlight folders)
  - Automated production of hierarchical taxonomies of documents for browsing purposes (à la Yahoo & DMOZ)


# Some applications: Clustering and search (1)

Vivísimo - Clustered search results - Microsoft Internet Explorer fornito da PC Professionale

File Modifica Visualizza Preferiti Strumenti ?

Indietro Cerca Preferiti Multimedia

Indirizzo <http://vivísimo.com/search?query=Roberto+Basili&y%3Asources=Web&x=63&y=9>

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### Clustered Results

- [Roberto Basili \(166\)](#)
- [Tor Vergata \(34\)](#)
- [Roberto Basili, Alessandro Moschitti \(15\)](#)
- [Linguistic \(15\)](#)
- [Fabio Massimo Zanzotto \(16\)](#)
- [TANLPS Workshop \(8\)](#)
- [Processing, Natural Language \(12\)](#)
- [Query, Meo-Evoli \(9\)](#)
- [ACL, Anthology \(8\)](#)
- [ROMAND, CfP \(8\)](#)
- [Authors \(6\)](#)
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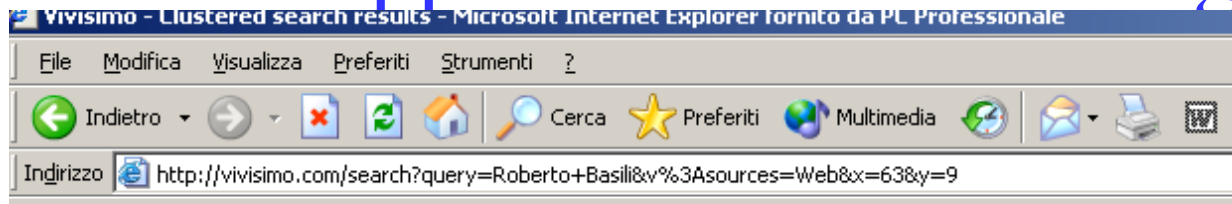
Find in clusters:

### Top 166 results retrieved for the query **Roberto Basili** ([Details](#))

[Roberto on eBay](#) [new window] [preview]  
Find **Roberto** items at low prices. With over 5 million items for sale every day, you'll find all kinds of items on the Online Marketplace.  
[www.ebay.com](http://www.ebay.com)

- [DBLP: Roberto Basili](#) [new window] [frame] [preview]  
dblp.uni-trier.de **Roberto Basili** . ... 2004. 27, EE, Alessandro Moschitti, **Roberto Basili** : Classification: A Comprehensive Study. ...  
URL: [www.informatik.uni-trier.de/~a-tree/b/Basili:Roberto.html](http://www.informatik.uni-trier.de/~a-tree/b/Basili:Roberto.html) - [show in clusters](#)  
Sources: [Netscape 1](#), [Lycos 2](#)
- [Mail archive: CFP: ECML'98 TANLPS Workshop: First Call for Paper](#) [new window] [frame]  
CFP: ECML'98 TANLPS Workshop: First Call for Paper. **Roberto Basili** ( [basili@info.utovr.it](mailto:basili@info.utovr.it) )  
URL: [www.comp.lancs.ac.uk/~research/ucrel/public/0893.html](http://www.comp.lancs.ac.uk/~research/ucrel/public/0893.html) - [show in clusters](#)  
Sources: [Netscape 3](#), [Wisenuit 4](#)
- [3rd Summer Convention on Information Extraction - SCIE 2002](#) [new window] [frame] [p

# Some applications: Clustering and search (2)



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  - Tor Vergata (34)**
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    - [Maria Teresa Pazienza \(5\)](#)
    - [Roberto Basili University Of Roma \(5\)](#)
    - [Nathalie Aussenac-Gilles \(6\)](#)
    - [Programme Committee Roberto Basili \(4\)](#)
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# Current Challenges in Clustering

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Many traditional clustering techniques do not perform satisfactorily in data mining scenarios due to a variety of reasons

- **Data Distribution**

- **Large number of samples**

- The number of samples to be processed is very high. Clustering in general is NP-hard, and practical and successful data mining algorithms usually scale linear or log-linear. Quadratic and cubic scaling may also be allowable but a linear behavior is highly desirable.

- **High dimensionality**

- The number of features is very high and may even exceed the number of samples. So one has to face the curse of dimensionality

- **Sparsity**

- Most features are zero for most samples, i.e. the object-feature matrix is sparse. This property strongly affects the measurements of similarity and the computational complexity.

- **Significant outliers**

- Outliers may have significant importance. Finding these outliers is highly non-trivial, and removing them is not necessarily desirable.

# Current Challenges in Clustering (cont.)

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- **Application context**
  - **Legacy clusterings**
    - Previous cluster analysis results are often available. This knowledge should be reused instead of starting each analysis from scratch.
  - **Distributed data**
    - Large systems often have heterogeneous distributed data sources. Local cluster analysis results have to be integrated into global models.