Clustering: Similarity-Based Clustering

CS4780/5780 – Machine Learning Fall 2014

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Reading: Manning/Raghavan/Schuetze, Chapters 16 (not 16.3) and 17 (http://nlp.stanford.edu/IR-book/)

Outline

- Supervised vs. Unsupervised Learning
- Hierarchical Clustering

 Hierarchical Agglomerative Clustering (HAC)
- Non-Hierarchical Clustering
 - K-means
 - Mixtures of Gaussians and EM-Algorithm

Supervised Learning vs. Unsupervised Learning

- Supervised Learning
 - Classification: partition examples into groups according to pre-defined categories
 - Regression: assign value to feature vectors
 - Requires labeled data for training
- Unsupervised Learning
 - Clustering: partition examples into groups when no pre-defined categories/classes are available
 - Outlier detection: find unusual events (e.g. hackers)
 - Novelty detection: find changes in data
 - Only instances required, but no labels

Clustering

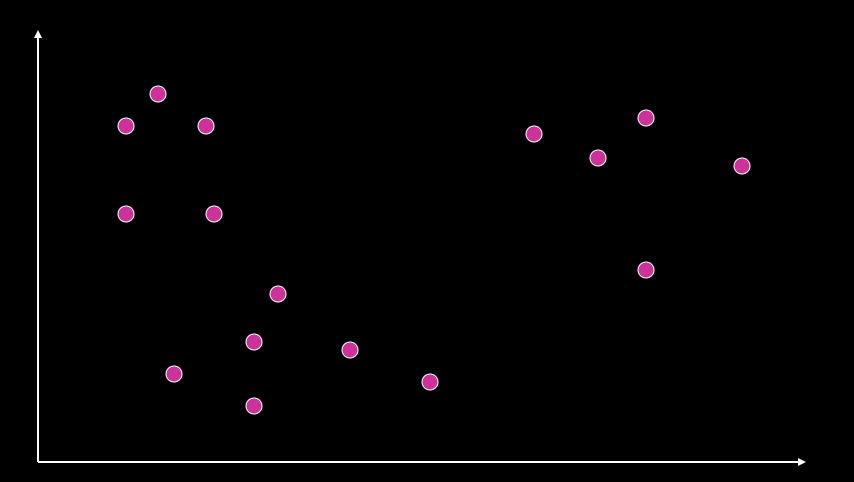
- Partition unlabeled examples into disjoint subsets of clusters, such that:
 - Examples within a cluster are similar
 - Examples in different clusters are different
- Discover new categories in an *unsupervised* manner (no sample category labels provided).

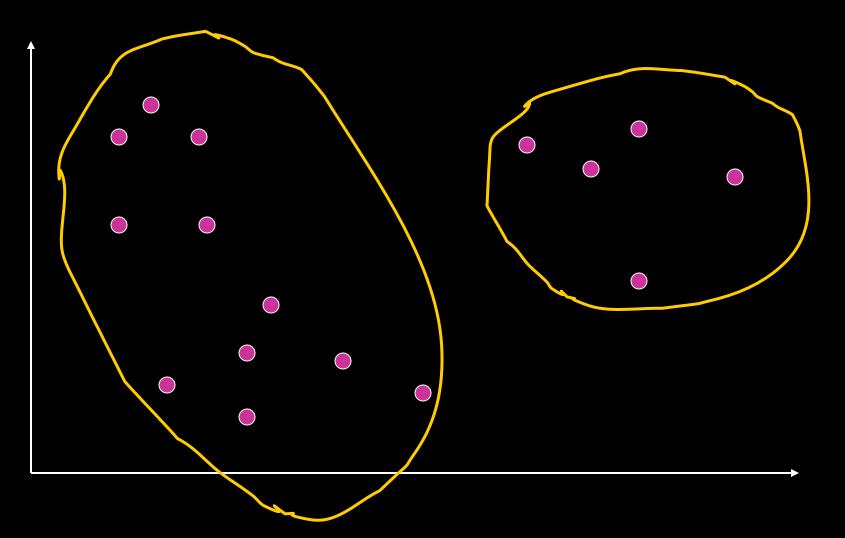
Applications of Clustering

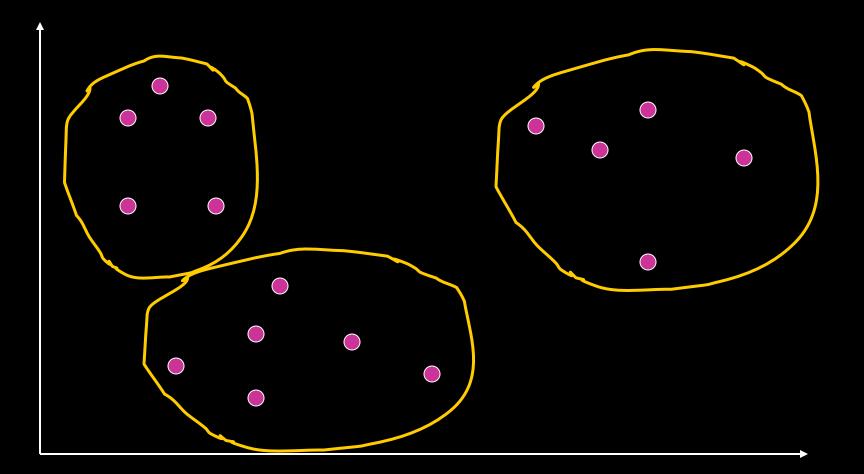
- Exploratory data analysis
- Cluster retrieved documents
 - to present more organized and understandable results to user → "diversified retrieval"
- Detecting near duplicates
 - Entity resolution
 - E.g. "Thorsten Joachims" == "Thorsten B Joachims"
 - Cheating detection
- Automated (or semi-automated) creation of taxonomies
 - e.g. Yahoo, DMOZ
- Compression

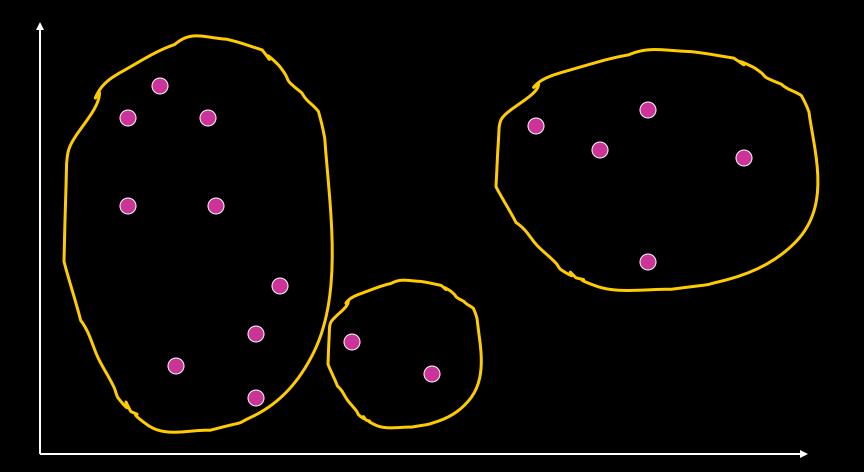
Applications of Clustering

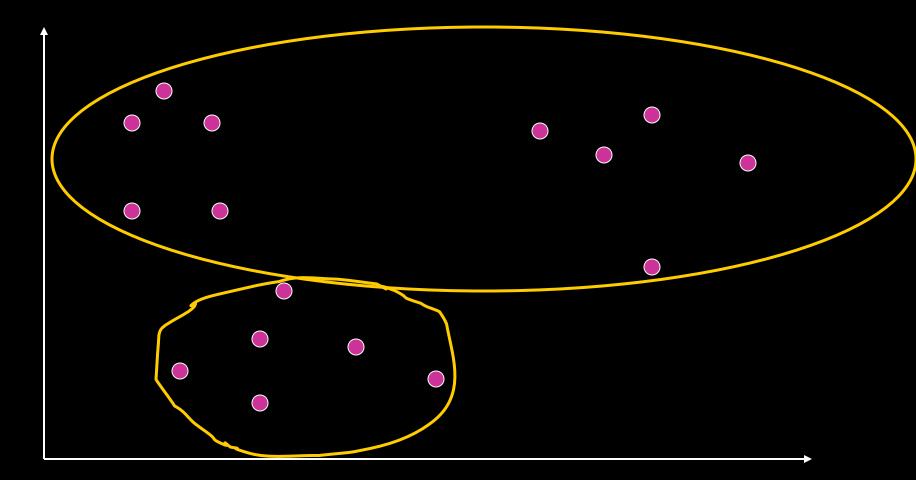
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	6 Society for Vascular Medicine : About SVM : Home 🖏 🕢 🔍	
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	7 SVM - Support Vector Machines 😱 🗗 🔍	
	SVM, support vector machines, SVMC, support vector machines classification, SVMR, support vector machines regression, kernel, machine learning, pattern	
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	8 SVM-Struct Support Vector Machine for Complex Outputs 🞭 🗗 🔍	
	SVMstruct is a Support Vector Machine (SVM) algorithm for predicting multivariate or structured outputs. It performs supervised learning by approximating a	
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Similarity (Distance) Measures

• Euclidian distance (L₂ norm):

$$L_2(\vec{x}, \vec{x}') = \sqrt{\sum_{i=1}^N (x_i - x_i')^2}$$

• L₁ norm:

$$L_1(\vec{x}, \vec{x}') = \sqrt{\sum_{i=1}^N |x_i - x'_i|}$$

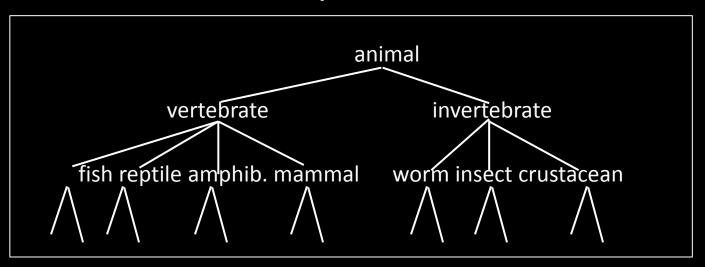
• Cosine similarity:

$$\cos(\vec{x}, \vec{x}') = \frac{\vec{x} * \vec{x}'}{\|\vec{x}\| \|\vec{x}'\|}$$

Kernels

Hierarchical Clustering

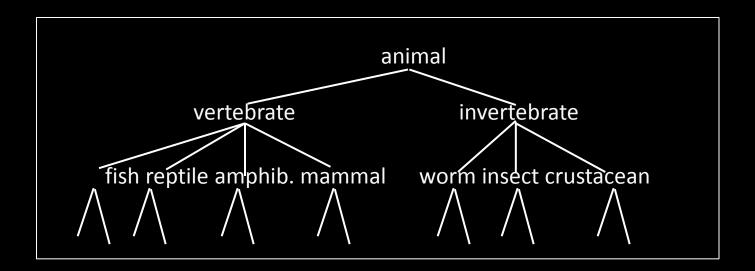
Build a tree-based hierarchical taxonomy from a set of unlabeled examples.



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

Agglomerative vs. Divisive Clustering

- Agglomerative (bottom-up) methods start with each example in its own cluster and iteratively combine them to form larger and larger clusters.
- *Divisive* (*top-down*) separate all examples immediately into clusters.



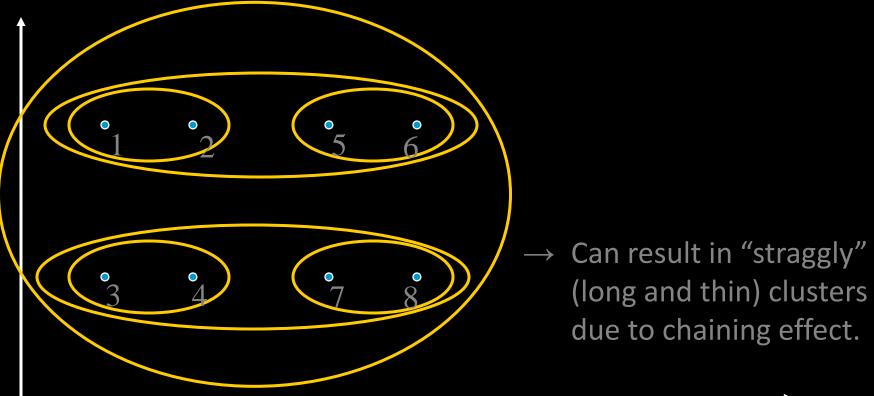
Hierarchical Agglomerative Clustering (HAC)

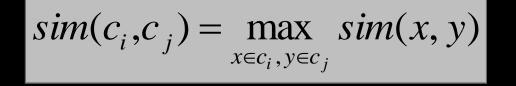
- Assumes a *similarity function* for determining the similarity of two clusters.
- Starts with all instances in a separate cluster and then repeatedly joins the two clusters that are most similar until there is only one cluster.
- The history of merging forms a binary tree or hierarchy.
- Basic algorithm:
 - 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_i, that are most similar.
 - Replace c_i and c_j with a single cluster $c_i \cup c_j$

Cluster Similarity

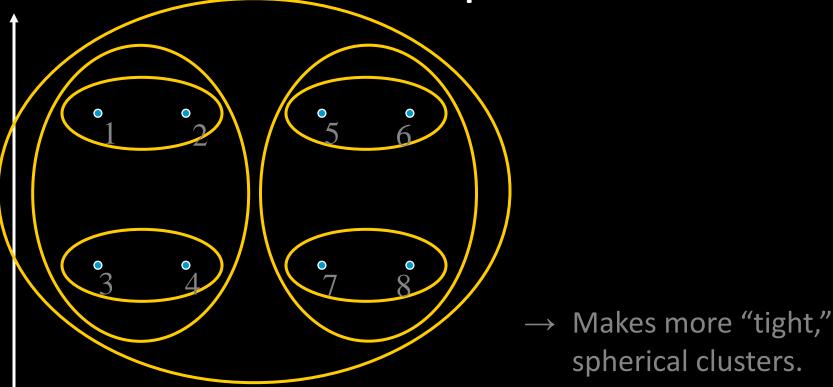
- How to compute similarity of two clusters each possibly containing multiple instances?
 - *Single link*: Similarity of two most similar members.
 - Complete link: Similarity of two least similar members.
 - Group average: Average similarity between members.

Single-Link HAC





Complete-Link HAC



$$sim(c_i,c_j) = \min_{x \in c_i, y \in c_j} sim(x, y)$$

Computational Complexity of HAC

- In the first iteration, all HAC methods need to compute similarity of all pairs of *n* individual instances which is O(n²).
- In each of the subsequent O(n) merging iterations, must find smallest distance pair of clusters → Maintain heap O(n² log n)
- In each of the subsequent O(n) merging iterations, it must compute the distance between the most recently created cluster and all other existing clusters. Can this be done in constant time such that O(n² log n) overall?

Computing Cluster Similarity

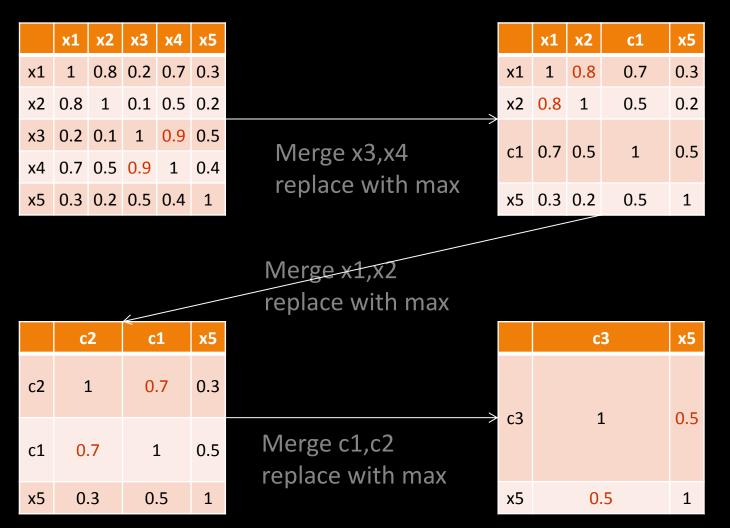
- After merging c_i and c_j, the similarity of the resulting cluster to any other cluster, c_k, can be computed by:
 - Single Link:

$$sim((c_i \cup c_j), c_k) = \max(sim(c_i, c_k), sim(c_j, c_k))$$

- Complete Link:

$$sim((c_i \cup c_j), c_k) = \min(sim(c_i, c_k), sim(c_j, c_k))$$

Single-Link Example



Group Average Agglomerative Clustering

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

$$sim(c_{i}, c_{j}) = \frac{1}{|c_{i} \cup c_{j}| (|c_{i} \cup c_{j}| - 1)} \sum_{\vec{x} \in (c_{i} \cup c_{j})} \sum_{\vec{y} \in (c_{i} \cup c_{j}): \vec{y} \neq \vec{x}} sim(\vec{x}, \vec{y})$$

Compromise between single and complete link.

Computing Group Average Similarity

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

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

 Compute similarity of clusters in constant time:

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

Non-Hierarchical Clustering

- K-means clustering ("hard")
- Mixtures of Gaussians and training via Expectation maximization Algorithm ("soft")