Clustering: Similarity-Based Clustering

CS4780/5780 – Machine Learning Fall 2013

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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
 - Novelty detection: find changes in data
 - Outlier detection: find unusual events (e.g. hackers)
 - Only instances required, but no labels

Clustering

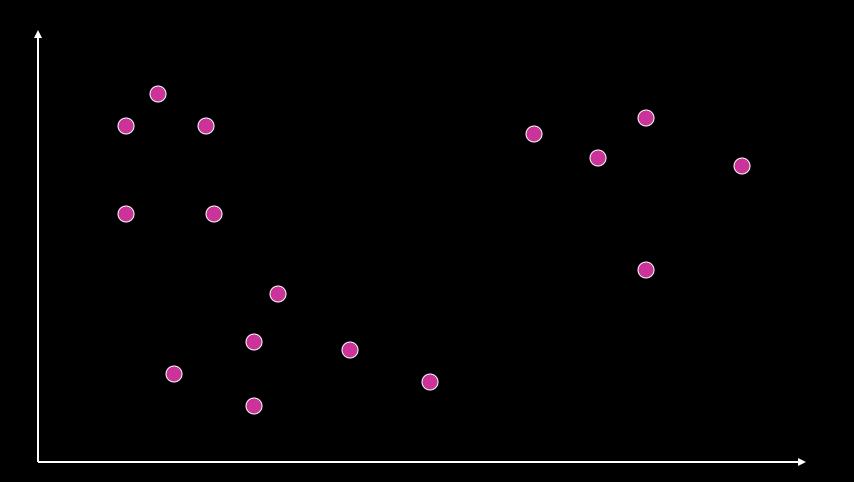
- Partition unlabeled examples into disjoint subsets of clusters, such that:
 - Examples within a cluster are similar
 - Examples in different clusters are different
- Discover new categories in an *unsupervised* manner (no sample category labels provided).

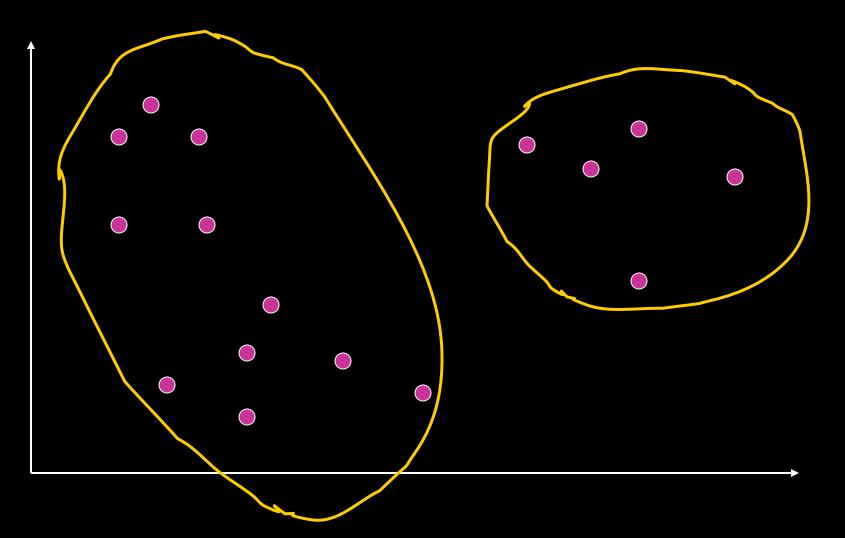
Applications of Clustering

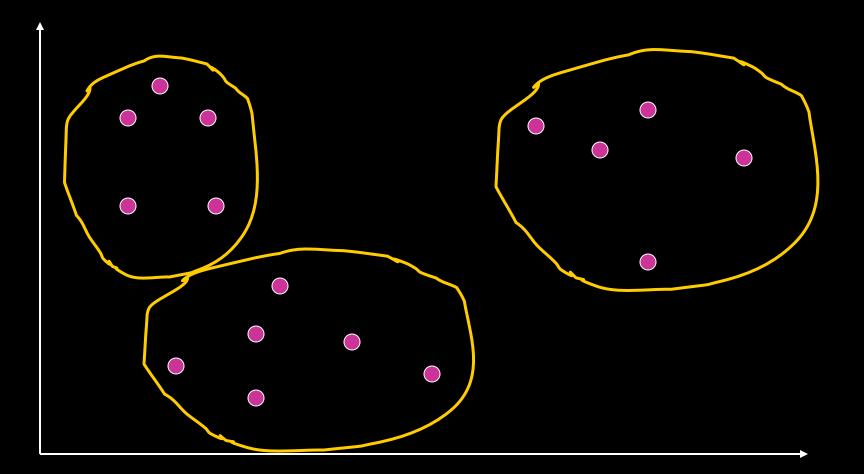
- Cluster retrieved documents
 - to present more organized and understandable results to user → "diversified retrieval"
- Detecting near duplicates
 - Entity resolution
 - E.g. "Thorsten Joachims" == "Thorsten B Joachims"
 - Cheating detection
- Exploratory data analysis
- Automated (or semi-automated)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

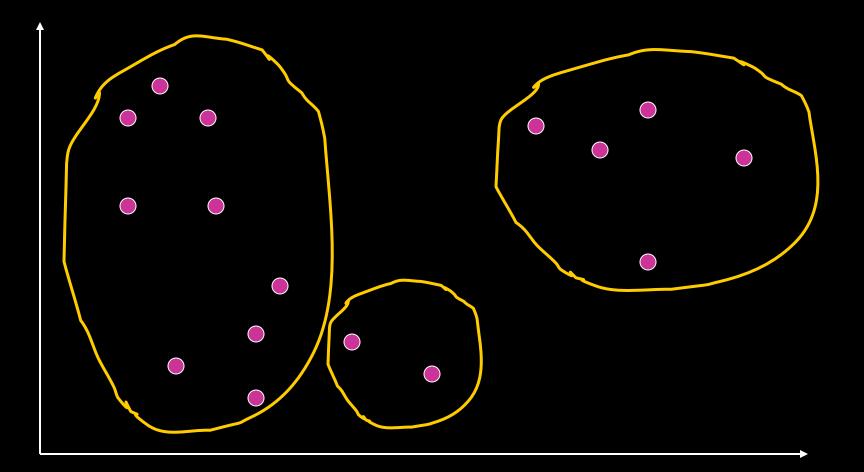
Applications of Clustering

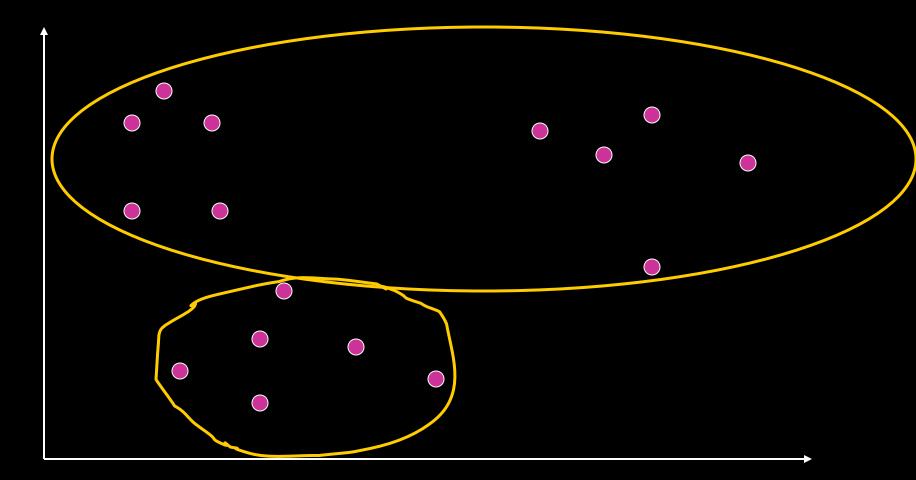
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- Support Vector Machine SVM (11)	In machine learning, support vector machines (SVMs, also support vector networks) are supervised learning models with associated learning algorithms that http://en.wikipedia.org/wiki/Support vector machine [Google, Wikipedia]	=
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A more show all	View the basic SVM stock chart on Yahoo! Finance. Change the date range, chart type and compare Silvercorp Metals Inc Ordinary against other companies.	
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	Up to date analysis of Silvercorp Metals Inc. (SVM) and its stock by hedge fund managers and industry experts. Find out what Silvercorp Metals Inc. is saying	
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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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	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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Query: svm Source: Web (95 results, 1758 ms) Clusterer: Lingo (188 ms) 02/02-2012 Stanislaw Osinski, Dawid Weis		wid Weiss











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:

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

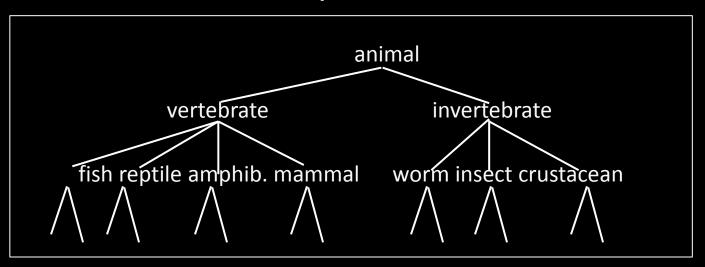
• Cosine similarity:

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

Kernels

Hierarchical Clustering

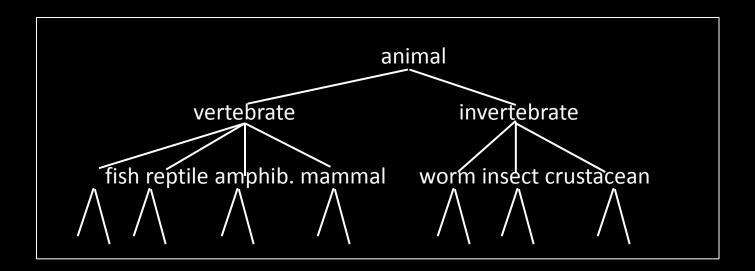
Build a tree-based hierarchical taxonomy from a set of unlabeled examples.



• 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.



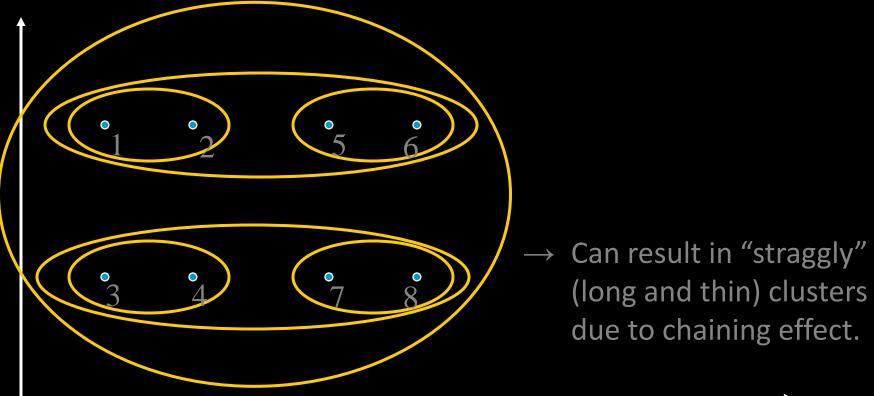
Hierarchical Agglomerative Clustering (HAC)

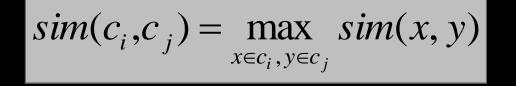
- Assumes a *similarity function* for determining the similarity of two clusters.
- Starts with all instances in a separate cluster and then repeatedly joins the two clusters that are most similar until there is only one cluster.
- The history of merging forms a binary tree or hierarchy.
- Basic algorithm: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

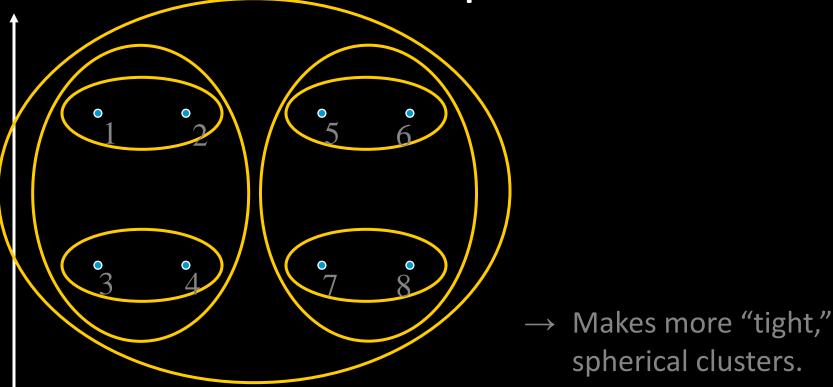
- How to compute similarity of two clusters each possibly containing multiple instances?
 - *Single link*: Similarity of two most similar members.
 - Complete link: Similarity of two least similar members.
 - Group average: Average similarity between members.

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

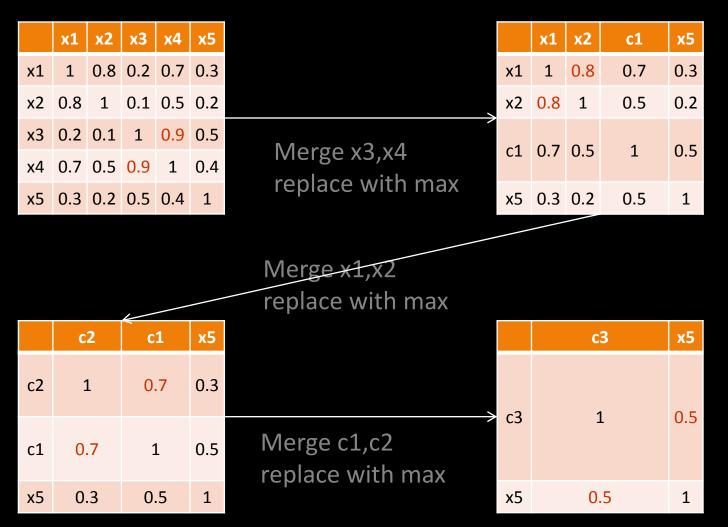
- After merging c_i and c_j, the similarity of the resulting cluster to any other cluster, c_k, can be computed by:
 - Single Link:

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

- 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")

Clustering Criterion

- Evaluation function that assigns a (usually real-valued) value to a clustering
 - Clustering criterion typically function of
 - within-cluster similarity and
 - between-cluster dissimilarity
- Optimization
 - Find clustering that maximizes the criterion
 - Global optimization (often intractable)
 - Greedy search
 - Approximation algorithms

Centroid-Based Clustering

- Assumes instances are real-valued vectors.
- Clusters represented via *centroids* (i.e. average of points in a cluster) *c*:

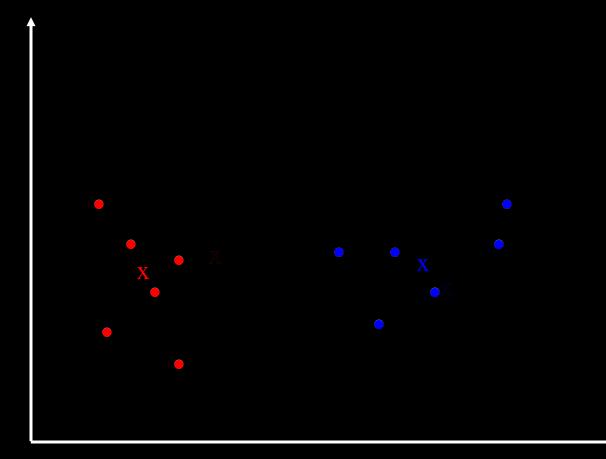
$$\vec{\mu}(c) = \frac{1}{|c|} \sum_{\vec{x} \in c} \vec{x}$$

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

K-Means Algorithm

- Input: k = number of clusters, distance measure d
- 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_i such that $d(x_i, s_i)$ is min.
 - For each cluster c_i //update the centroid of each cluster
 - $s_j = \mu(c_j)$

K-means Example (k=2)



Pick seeds Reassign clusters Compute centroids Reassign clusters Compute centroids Reassign clusters

Converged!

Time Complexity

- Assume computing distance between two instances is O(N) where N is the dimensionality of the vectors.
- Reassigning clusters for n points: O(kn) distance computations, or O(knN).
- Computing centroids: Each instance gets added once to some centroid: O(nN).
- Assume these two steps are each done once for *i* iterations: O(*iknN*).
- Linear in all relevant factors, assuming a fixed number of iterations, more efficient than HAC.

Buckshot Algorithm

Problem

- Results can vary based on random seed selection, especially for high-dimensional data.
- Some seeds can result in poor convergence rate, or convergence to sub-optimal clusterings.
- Idea: Combine HAC and K-means clustering.
- First randomly take a sample of instances of size
- Run group-average HAC on this sample n^{1/2}
- Use the results of HAC as initial seeds for K-means.
- Overall algorithm is efficient and avoids problems of bad seed selection.

Clustering as Prediction

Setup

- Learning Task: P(X)
- Training Sample: $S = (\vec{x}_1, ..., \vec{x}_n)$
- Hypothesis Space: $H = \{h_1, \dots, h_{|H|}\}$ each describes $P(X|h_i)$ where h_i are parameters
- Goal: learn which $P(X|h_i)$ produces the data
- What to predict?

Predict where new points are going to fall

Gaussian Mixtures and EM

- Gaussian Mixture Models
 - Assume

 $P(X = \vec{x} | h_i) = \sum_{j=1}^k P(X = \vec{x} | Y = j, h_i) P(Y = j)$

where $P(X = \vec{x} | Y = j, h) = N(X = \vec{x} | \vec{\mu}_j, \Sigma_j)$ and $h = (\vec{\mu}_1, ..., \vec{\mu}_k, \Sigma_1, ..., \Sigma_k)$.

- EM Algorithm
 - Assume P(Y) and k known and $\Sigma_i = 1$.

– REPEAT

•
$$\vec{\mu}_j = \frac{\sum_{i=1}^n P(Y=j|X=\vec{x}_i,\vec{\mu}_j)\vec{x}_i}{\sum_{i=1}^n P(Y=j|X=\vec{x}_i,\vec{\mu}_j)}$$

•
$$P(Y = j | X = \vec{x}_i, \vec{\mu}_j) = \frac{P(X = \vec{x}_i | Y = j, \vec{\mu}_j) P(Y = j)}{\sum_{l=1}^k P(X = \vec{x}_i | Y = l, \vec{\mu}_j) P(Y = l)} = \frac{e^{-0.5(\vec{x}_i - \vec{\mu}_j)^2} P(Y = j)}{\sum_{l=1}^k e^{-0.5(\vec{x}_i - \vec{\mu}_l)^2} P(Y = l)}$$