CS145: INTRODUCTION TO DATA MINING

09: Vector Data: Clustering Basics

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Announcement

- About Homework
 - Split HW3 into two: HW3 and HW4

- Optional HW6
 - We will pick the highest 5 homework scores

Methods to Learn

	Vector Data	Set Data	Sequence Data	Text Data
Classification	Logistic Regression; Decision Tree; KNN SVM; NN			Naïve Bayes for Text
Clustering	K-means; hierarchical clustering; DBSCAN; Mixture Models			PLSA
Prediction	Linear Regression GLM*			
Frequent Pattern Mining		Apriori; FP growth	GSP; PrefixSpan	
Similarity Search			DTW	

Vector Data: Clustering Basics

- Clustering Analysis: Basic Concepts
- Partitioning methods
- Hierarchical Methods
- Density-Based Methods
- Summary

What is Cluster Analysis?

- Cluster: A collection of data objects
 - similar (or related) to one another within the same group
 - dissimilar (or unrelated) to the objects in other groups
- Cluster analysis (or clustering, data segmentation, ...)
 - Finding similarities between data according to the characteristics found in the data and grouping similar data objects into clusters
- Unsupervised learning: no predefined classes (i.e., *learning by observations* vs. learning by examples: supervised)
- Typical applications
 - As a stand-alone tool to get insight into data distribution
 - As a preprocessing step for other algorithms

Applications of Cluster Analysis

Data reduction

- Summarization: Preprocessing for regression, PCA, classification, and association analysis
- Compression: Image processing: vector quantization
- Prediction based on groups
 - Cluster & find characteristics/patterns for each group
- Finding K-nearest Neighbors
 - Localizing search to one or a small number of clusters
- Outlier detection: Outliers are often viewed as those "far away" from any cluster

Clustering: Application Examples

- Biology: taxonomy of living things: kingdom, phylum, class, order, family, genus and species
- Information retrieval: document clustering
- Land use: Identification of areas of similar land use in an earth observation database
- Marketing: Help marketers discover distinct groups in their customer bases, and then use this knowledge to develop targeted marketing programs
- City-planning: Identifying groups of houses according to their house type, value, and geographical location
- Earth-quake studies: Observed earth quake epicenters should be clustered along continent faults
- Climate: understanding earth climate, find patterns of atmospheric and ocean

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Partitioning Algorithms: Basic Concept

<u>Partitioning method</u>: Partitioning a dataset *D* of *n* objects into a set of *k* clusters, such that the sum of squared distances is minimized (where c_i is the centroid or medoid of cluster C_i)

$$J = \sum_{j=1}^{k} \sum_{C(i)=j} d(x_i, c_j)^2$$

- Given k, find a partition of k clusters that optimizes the chosen partitioning criterion
 - Global optimal: exhaustively enumerate all partitions
 - Heuristic methods: *k-means* and *k-medoids* algorithms
 - <u>*k-means*</u> (MacQueen'67, Lloyd'57/'82): Each cluster is represented by the center of the cluster
 - <u>k-medoids</u> or PAM (Partition around medoids) (Kaufman & Rousseeuw'87): Each cluster is represented by one of the objects in the cluster

- Given k, the k-means algorithm is implemented in four steps:
 - Step 0: Partition objects into k nonempty subsets
 - Step 1: Compute seed points as the centroids of the clusters of the current partitioning (the centroid is the center, i.e., *mean point*, of the cluster)
 - Step 2: Assign each object to the cluster with the nearest seed point
 - Step 3: Go back to Step 1, stop when the assignment does not change

An Example of K-Means Clustering



Until no change

Theory Behind K-Means

Objective function

•
$$J = \sum_{j=1}^{k} \sum_{C(i)=j} ||x_i - c_j||^2$$

Re-arrange the objective function

•
$$J = \sum_{j=1}^{k} \sum_{i} w_{ij} ||x_i - c_j||^2$$

- $\bullet \, w_{ij} \in \{0,1\}$
- $w_{ij} = 1$, if x_i belongs to cluster j; $w_{ij} = 0$, otherwise
- Looking for:
 - The best assignment w_{ij}
 - The best center c_j

Solution of K-Means

Iterations

$$J = \sum_{j=1}^{k} \sum_{i=1}^{k} w_{ij} ||x_i - c_j||^2$$

- Step 1: Fix centers c_j , find assignment w_{ij} that minimizes J
 - => $w_{ij} = 1$, if $||x_i c_j||^2$ is the smallest
- Step 2: Fix assignment *w*_{*ij*}, find centers that minimize *J*
 - => first derivative of J = 0

• =>
$$\frac{\partial J}{\partial c_j} = -2\sum_i w_{ij}(x_i - c_j) = 0$$

• => $c_j = \frac{\sum_i w_{ij}x_i}{\sum_i w_{ij}}$

• Note $\sum_i w_{ij}$ is the total number of objects in cluster j

Comments on the K-Means Method

- <u>Strength</u>: *Efficient*: O(tkn), where n is # objects, k is # clusters, and t is # iterations. Normally, k, t << n.
- <u>Comment:</u> Often terminates at a *local optimal*
- Weakness
 - Applicable only to objects in a continuous n-dimensional space
 - Using the k-modes method for categorical data
 - In comparison, k-medoids can be applied to a wide range of data
 - Need to specify *k*, the *number* of clusters, in advance (there are ways to automatically determine the best k (see Hastie et al., 2009)
 - Sensitive to noisy data and *outliers*
 - Not suitable to discover clusters with *non-convex shapes*

Variations of the K-Means Method*

- Most of the variants of the k-means which differ in
 - Selection of the initial k means
 - Dissimilarity calculations
 - Strategies to calculate cluster means
- Handling categorical data: k-modes
 - Replacing means of clusters with modes
 - Using new dissimilarity measures to deal with categorical objects
 - Using a <u>frequency</u>-based method to update modes of clusters
 - A mixture of categorical and numerical data: *k-prototype* method



The K-Medoid Clustering Method*

- K-Medoids Clustering: Find representative objects (medoids) in clusters
 - PAM (Partitioning Around Medoids, Kaufmann & Rousseeuw 1987)
 - Starts from an initial set of medoids and iteratively replaces one of the medoids by one of the non-medoids if it improves the total distance of the resulting clustering
 - PAM works effectively for small data sets, but does not scale well for large data sets (due to the computational complexity)
- Efficiency improvement on PAM
 - CLARA (Kaufmann & Rousseeuw, 1990): PAM on samples
 - CLARANS (Ng & Han, 1994): Randomized re-sampling

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Hierarchical Clustering

 Use distance matrix as clustering criteria. This method does not require the number of clusters k as an input, but needs a termination condition



AGNES (Agglomerative Nesting)

- Introduced in Kaufmann and Rousseeuw (1990)
- Implemented in statistical packages, e.g., Splus
- Use the **single-link** method and the dissimilarity matrix
- Merge nodes that have the least dissimilarity
- Go on in a non-descending fashion
- Eventually all nodes belong to the same cluster



Pseudo Code

- Initialization: Place each data point into its own cluster and compute distance matrix between clusters
- Repeat:
 - Merge the two closest clusters
 - Update the distance matrix for the affected entries
- Until: all the data are merged into a single cluster

Dendrogram: Shows How Clusters are Merged



DIANA (Divisive Analysis)

- Introduced in Kaufmann and Rousseeuw (1990)
- Implemented in statistical analysis packages, e.g., Splus
- Inverse order of AGNES
- Eventually each node forms a cluster on its own



Distance between Clusters



- Single link: smallest distance between an element in one cluster and an element in the other, i.e., dist(K_i, K_j) = min dist(t_{ip}, t_{jq})
- Complete link: largest distance between an element in one cluster and an element in the other, i.e., dist(K_i, K_j) = max dist(t_{ip}, t_{jq})
- Average: avg distance between an element in one cluster and an element in the other, i.e., dist(K_i, K_j) = avg dist(t_{ip}, t_{jq})
- Centroid: distance between the centroids of two clusters, i.e., dist(K_i, K_j) = dist(C_i, C_j)
- Medoid: distance between the medoids of two clusters, i.e., dist(K_i, K_j) = dist(M_i, M_j)
 - Medoid: a chosen, centrally located object in the cluster

Example: Single Link vs. Complete Link



Extensions to Hierarchical Clustering

- Major weakness of agglomerative clustering methods
 - Can never undo what was done previously
 - <u>Do not scale</u> well: time complexity of at least $O(n^2)$, where *n* is the number of total objects
- Integration of hierarchical & distance-based clustering
 - <u>*BIRCH (1996)</u>: uses CF-tree and incrementally adjusts the quality of sub-clusters
 - <u>*CHAMELEON (1999)</u>: hierarchical clustering using dynamic modeling

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Summary

Density-Based Clustering Methods

- Clustering based on density (local cluster criterion), such as density-connected points
- Major features:
 - Discover clusters of arbitrary shape
 - Handle noise
 - One scan
 - Need density parameters as termination condition
- Several interesting studies:
 - <u>DBSCAN:</u> Ester, et al. (KDD'96)
 - <u>OPTICS*</u>: Ankerst, et al (SIGMOD'99).
 - <u>DENCLUE*</u>: Hinneburg & D. Keim (KDD'98)
 - <u>CLIQUE*</u>: Agrawal, et al. (SIGMOD'98) (more grid-based)

- Two parameters:
 - *Eps*: Maximum radius of the neighborhood
 - *MinPts*: Minimum number of points in an Epsneighborhood of that point
- $N_{Eps}(q)$: {p belongs to D | dist(p,q) \leq Eps}
- Directly density-reachable: A point p is directly densityreachable from a point q w.r.t. Eps, MinPts if
 - p belongs to $N_{Eps}(q)$
 - q is a core point, core point condition:

 $|N_{Eps}(q)| \ge MinPts$

Density-Reachable and Density-Connected

- Density-reachable:
 - A point *p* is density-reachable from a point *q* w.r.t. *Eps*, *MinPts* if there is a chain of points *p*₁, ..., *p*_n, *p*₁ = *q*, *p*_n = *p* such that *p*_{i+1} is directly density-reachable from *p*_i



- Density-connected
 - A point *p* is density-connected to a point *q* w.r.t. *Eps, MinPts* if there is a point *o* such that both, *p* and *q* are density-reachable from *o* w.r.t. *Eps* and *MinPts*



DBSCAN: Density-Based Spatial Clustering of Applications with Noise

- Relies on a *density-based* notion of cluster: A *cluster* is defined as a maximal set of density-connected points
- Noise: object not contained in any cluster is noise
- Discovers clusters of arbitrary shape in spatial databases with noise



DBSCAN: The Algorithm

(1)	mark all objects as unvisited;
(2)	do
(3)	randomly select an unvisited object p ;
(4)	$\mathrm{mark}\;p\;\mathrm{as}\;\mathtt{visited};$
(5)	if the ϵ -neighborhood of p has at least $MinPts$ objects
(6)	create a new cluster C , and add p to C ;
(7)	let N be the set of objects in the ϵ -neighborhood of p ;
(8)	for each point p' in N
(9)	if p' is unvisited
(10)	$\mathrm{mark}\;p'\;\mathrm{as}\;\mathtt{visited};$
(11)	if the ϵ -neighborhood of p' has at least $MinPts$ points,
	add those points to N ;
(12)	if p' is not yet a member of any cluster, add p' to C ;
(13)	end for
(14)	output C ;
(15)	else mark p as noise;
(16)	until no object is unvisited;

 If a spatial index is used, the computational complexity of DBSCAN is O(nlogn), where n is the number of database objects. Otherwise, the complexity is O(n²)

DBSCAN: Sensitive to Parameters



Figure 9. DBScan results for DS2 with MinPts at 4 and Eps at (a) 5.0, (b) 3.5, and (c) 3.0.



(a)

DBSCAN online Demo:

http://webdocs.cs.ualberta.ca/~yaling/Cluster/Applet/Code/Cluster.html

(b)

(c)

Questions about Parameters

- Fix Eps, increase MinPts, what will happen?
- Fix MinPts, decrease Eps, what will happen?

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Summary

- Cluster analysis groups objects based on their similarity and has wide applications; Measure of similarity can be computed for various types of data
- K-means and K-medoids algorithms are popular partitioningbased clustering algorithms
- AGNES and DIANA are interesting hierarchical clustering algorithms
- DBSCAN, OPTICS*, and DENCLUE* are interesting density-based algorithms

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