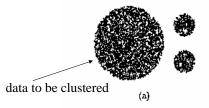
# Cluster Analysis

- What is Cluster Analysis?
- Types of Data in Cluster Analysis
- A Categorization of Major Clustering Methods
- Partitioning Methods
- Hierarchical Methods
- Density-Based Methods
- Grid-Based Methods
- Model-Based Clustering Methods
- Outlier Analysis
- Summary

# CURE (Clustering Using REpresentatives)





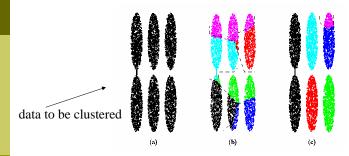




#### ■ CURE: proposed by Guha, Rastogi & Shim, 1998

- Stops the creation of a cluster hierarchy if a level consists of *k* clusters
- Uses multiple representative points to evaluate the distance between clusters, adjusts well to arbitrary shaped clusters and avoids single-link effect

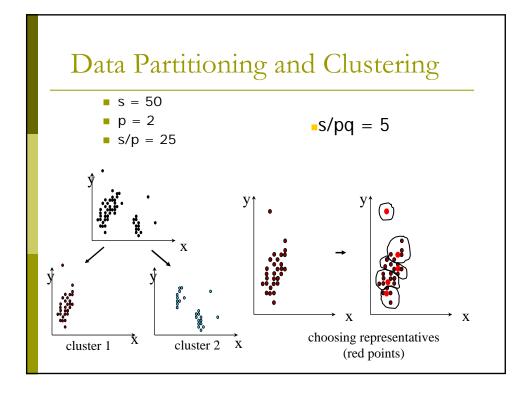
## Drawbacks of Distance-Based Method



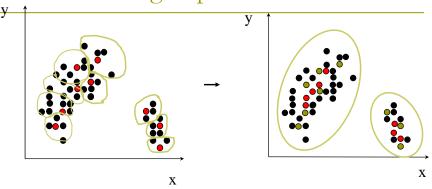
- Drawbacks of single representative methods (b)
  - Consider only one point as representative of a cluster
  - $\blacksquare$  Good only for convex shaped, similar size and density, and if  $\emph{k}$  can be reasonably estimated
- Drawbacks of density-based methods (c)
  - Can merge clusters which are connected by a very narrow dense link

## Cure: The Algorithm

- Draw random sample s.
- Partition sample to p partitions with size s/p
- Partially cluster partitions into s/pq clusters
- Eliminate outliers
  - By random sampling
  - □ If a cluster grows too slow, eliminate it.
- Cluster partial clusters.
- Label data in disk



#### Cure: Shrinking Representative Points



- Shrink the multiple representative points towards the gravity center by a fraction of  $\alpha$ .
- Multiple representatives capture the shape of the cluster

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### 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)
  - OPTICS: Ankerst, et al (SIGMOD'99).
  - DENCLUE: Hinneburg & D. Keim (KDD'98)
  - CLIQUE: Agrawal, et al. (SIGMOD'98)

### Density-Based Clustering: Background

- Neighborhood of point p=all points within distance Eps from p:
- Two parameters:
  - **Eps**: Maximum radius of the neighbourhood
  - MinPts: Minimum number of points in an Eps-neighbourhood of that point
- If the number of points in the Eps-neighborhood of p is at least *MinPts*, then p is called a core object.
- Directly density-reachable: A point **p** is directly density-reachable from a point **q** wrt. **Eps**, **MinPts** if
  - 1) **p** belongs to **N**<sub>Eps</sub>(**q**)
  - 2) core point condition:

 $|N_{Ens}(q)| >= MinPts$ 



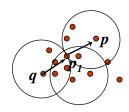
MinPts = 5

Eps = 1 cm

# Density-Based Clustering: Background (II)

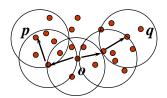
#### Density-reachable:

■ A point p is density-reachable from a point q wrt. 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  $p_i$ 



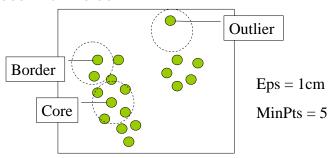
#### Density-connected

A point p is density-connected to a point q wrt. Eps, MinPts if there is a point o such that both, p and q are density-reachable from o wrt. 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 densityconnected points
- Discovers clusters of arbitrary shape in spatial databases with noise



## DBSCAN: The Algorithm

- Arbitrary select a point p
- Retrieve all points density-reachable from p wrt Eps and MinPts.
- If p is a core point, a cluster is formed.
- If p is a border point, no points are density-reachable from p and DBSCAN visits the next point of the database.
- Continue the process until all of the points have been processed.

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#### Grid-Based Clustering Method

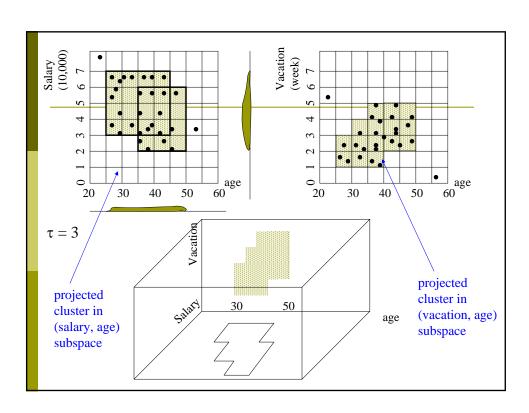
- Using multi-resolution grid data structure
- Several interesting methods
  - STING (a STatistical Information Grid approach) by Wang, Yang and Muntz (1997)
  - WaveCluster by Sheikholeslami, Chatterjee, and Zhang (VLDB'98)
    - A multi-resolution clustering approach using wavelet method
  - CLIQUE: Agrawal, et al. (SIGMOD'98)

#### CLIQUE (Clustering In QUEst)

- Agrawal, Gehrke, Gunopulos, Raghavan (SIGMOD'98).
- Automatically identifying subspaces of a high dimensional data space that allow better clustering than original space
- CLIQUE can be considered as both density-based and grid-based
  - It partitions each dimension into the same number of equal length interval
  - It partitions an m-dimensional data space into nonoverlapping rectangular units
  - A unit is dense if the fraction of total data points contained in the unit exceeds the input model parameter
  - A cluster is a maximal set of connected dense units within a subspace

#### CLIQUE: The Major Steps

- Partition the data space and find the number of points that lie inside each cell of the partition.
- Identify the subspaces that contain clusters using the Apriori principle
- Identify clusters:
  - Determine dense units in all subspaces of interests
  - Determine connected dense units in all subspaces of interests.
- Generate minimal description for the clusters
  - Determine maximal regions that cover a cluster of connected dense units for each cluster
  - Determination of minimal cover for each cluster
- CLIQUE can find projected clusters in subspaces of the dimensional space



# Strength and Weakness of CLIQUE

#### Strength

- It <u>automatically</u> finds subspaces of the <u>highest</u> <u>dimensionality</u> such that high density clusters exist in those subspaces
- It is insensitive to the order of records in input and does not presume some canonical data distribution
- It scales *linearly* with the size of input and has good scalability as the number of dimensions in the data increases

#### Weakness

 The accuracy of the clustering result may be degraded at the expense of simplicity of the method

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#### Model based clustering

- Assume data generated from K probability distributions
- Typically Gaussian distribution Soft or probabilistic version of K-means clustering
- □ Need to find distribution parameters.
- EM Algorithm

### EM Algorithm

- Initialize K cluster centers
- Iterate between two steps
  - Expectation step: assign points to clusters

$$P(d_i \in c_k) = w_k \Pr(d_i \mid c_k) / \sum_j w_j \Pr(d_i \mid c_j)$$

$$w_k = \frac{\sum_i \Pr(d_i \in c_k)}{N}$$

■ Maximation step: estimate model parameters

$$\mu_{k} = \frac{1}{m} \sum_{i=1}^{m} \frac{d_{i} P(d_{i} \in c_{k})}{\sum_{k} P(d_{i} \in c_{j})}$$

### Cluster Analysis

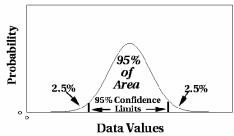
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## What Is Outlier Discovery?

- What are outliers?
  - The set of objects are considerably dissimilar from the remainder of the data (exceptions or noise)
- Problem
  - Find top n outlier points
- Applications:
  - Credit card fraud detection
  - Telecom fraud detection
  - Customer segmentation
  - Medical analysis

# Outlier Discovery: Statistical

# Approaches



- Assume a model underlying distribution that generates data set (e.g. normal distribution)
- Use discordancy tests depending on
  - data distribution
  - distribution parameter (e.g., mean, variance)
  - number of expected outliers
- Drawbacks
  - most tests are for single attribute (not applicable for multidimensional data)
  - In many cases, data distribution may not be known

# Outlier Discovery: Distance-Based Approach

- Introduced to counter the main limitations imposed by statistical methods
  - We need multi-dimensional analysis without knowing data distribution.
- Distance-based outlier: A DB(p, D)-outlier is an object O in a dataset T such that at least a fraction p of the objects in T lies at a distance greater than D from O
- Algorithms for mining distance-based outliers
  - Index-based algorithm
  - Nested-loop algorithm
  - Cell-based algorithm