#### Outline

- ▶ What is Clustering?
- ► Types of Data in Cluster Analysis and Similarity Measures
- ▶ Some clustering Methods
  - ▶ K-means
  - ▶ K-medoids
  - ▶ Hierarchical clustering method
  - ▶ DBSCAN: a Density-based Algorithm
- Cluster Validity

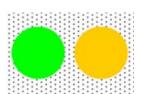
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## **Density-based Clustering**

- A cluster is a dense region of points, which is separated by low-density regions, from other regions of high density.
- ▶ Used when the clusters are irregular or intertwined, and when noise and outliers are present.







## **Density-based Clustering**

- Clustering based on density (local cluster criterion), such as density-connected points
- Major features:
  - Discover clusters of arbitrary shape
  - Handle noise
  - ▶ Do not need to specify k, but need density parameters
- Several interesting studies:
  - ▶ DBSCAN: Ester, et al. (KDD'96)
  - OPTICS: Ankerst, et al. (SIGMOD'99)
  - ▶ DENCLUE: Hinneburg & D. Kein (KDD'98)
  - ► CLIQUE: Agrawal, et al. (SIGMOD'98)

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## DBSCAN: Basic Concepts

 $\triangleright$  *Eps*-neighborhood of point *p* in data set *D*:

$$N_{Eps}(p) = \{q \in D \mid dist(p,q) \leq Eps\}$$

where Eps is called the radius of the neighborhood

• Density of Eps-neighborhood of p:

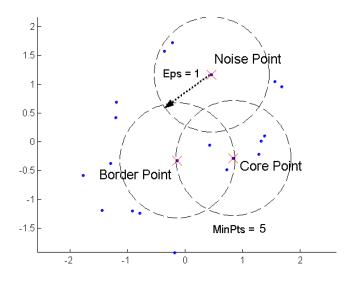
the number of points in  $N_{Eps}(p)$ 

## **DBSCAN: Basic Concepts**

- A point is a **core point** if there are at least *MinPts* number of points in its *Eps*-neighborhood.
  - These are points that are at the interior of a cluster
- A **border point** has fewer than *MinPts* points in its *Eps-neighborhood*, but is in the *Eps-neighborhood* of a core point.
  - These are points that are on or close to the border of a cluster
- A **noise point** is any point that is not a core point or a border point.
  - These are points that are outsider any cluster

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#### DBSCAN: Core, Border and Noise Points



# **DBSCAN**: Input Parameters

 $\triangleright$  *Eps*:

Maximum radius of the neighbourhood

▶ MinPts:

Minimum number of points in an *Eps*-neighbourhood of a core point

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# **DBSCAN: Basic Concepts**

- ▶ Directly density-reachable:
  - ▶ A point *p* is *directly density-reachable* from a point *q* wrt. *Eps* and *MinPts* if
    - 1) p belongs to  $N_{Eps}(q)$
    - 2) q is a core point, that is:

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

Asymmetric in general



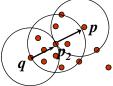
MinPts = 5

Eps = 1 cm

ightharpoonup Symmetric only when both p and q are core points.

## **DBSCAN: Basic Concepts**

- Density-reachable:
  - A point p is density-reachable from a point q wrt. Eps and MinPts if there is a chain of points  $p_1, p_2, ..., p_n, p_1 = q, p_n = p$  such that  $p_{i+1}$  is directly density-reachable from  $p_i$

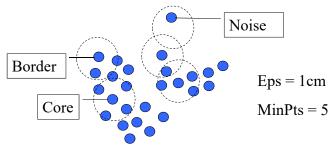


- ► Symmetric only when *p* and *q* are both core points
- ▶ Density-connected:
  - A point *p* is density-connected to a point *q* wrt. *Eps* and *MinPts* if there is a point *o* such that both *p* and *q* are density-reachable from *o* wrt. *Eps* and *MinPts*.
  - Symmetric

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#### **DBSCAN:** Cluster

- ▶ *Density-based* notion of cluster: A *cluster* is defined as a maximal set of density-connected points
  - ► Each pair of points in a cluster are density-connected to each other (Connectivity)
  - ► Core points in other clusters are not density-connected to any core points in this cluster (Maximality)
- ▶ Noise points are not in any cluster



# DBSCAN: The Algorithm

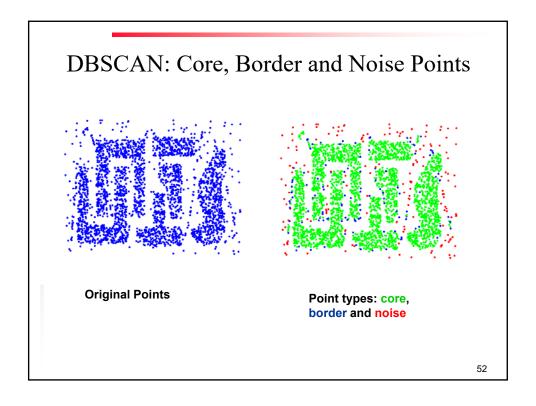
- Arbitrarily select an unprocessed point p
- These points and p are in a cluster because they are density-connected through p.
  - ▶ Retrieve all points density-reachable from *p* wrt/*Eps* and *MinPts*.
  - $\blacktriangleright$  A cluster is formed which includes all the points density-reachable from p
  - Mark all the points in the cluster as "processed"
- ▶ If *p* is not a core point, no points are density-reachable from *p* and DBSCAN visits the next unprocessed point of the database.
- ► Continue the process until all of the core points have been processed.

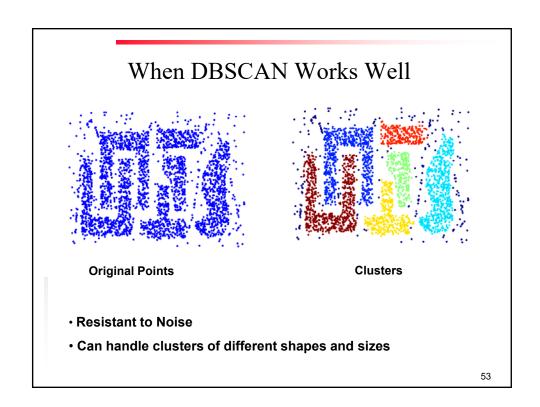
Note: Membership of a border point depends on the order of the points being processed if it is density-connected to core points in two or more clusters.

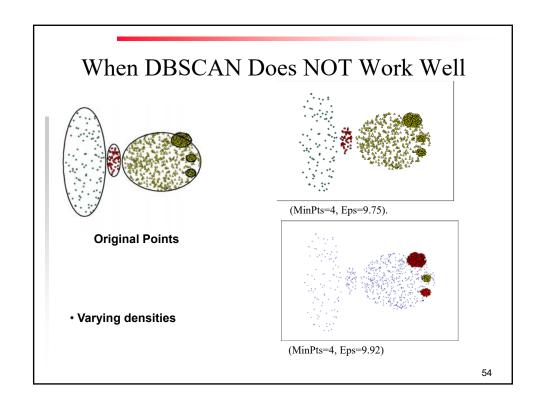
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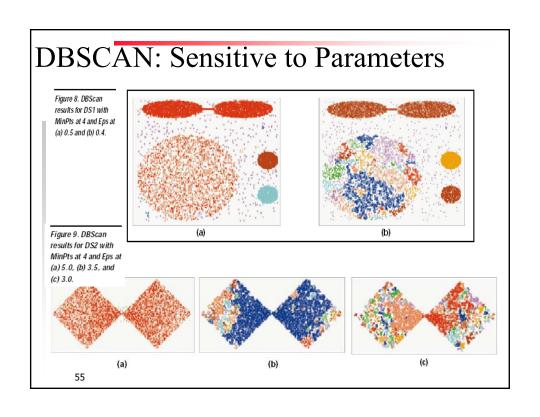
# Another Version of DBSCAN (for simplicity, not efficiency)

- Label all points as core, border or noise points
- ▶ Eliminate noise points
- ▶ Put an edge between all core points that are within *Eps* of each other
- ► Make each group of connected core points into a separate cluster
- ► Assign each border point to one of the clusters of its associated core points.









## Comparing DBSCAN and K-means

- Cluster shapes
  - K-means: spheres
  - DBSCAN: arbitrary shape





- Cluster sizes
  - K-means may have a problem when clusters are of different sizes
  - DBSCAN can handle clusters of different sizes
- Noise and outliers
  - K-means is sensitive to noise or outliers
  - DBSCAN is not strongly affected by noise or outliers 56

# Comparing DBSCAN and K-means (Cont'd)

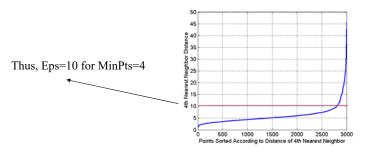
Not well-separated clusters



- K-means can find clusters that are not well separated, even if they overlap
- DBSCAN merges clusters that overlap
- - K-means' result depends on the random initialization of centroids
  - DBSCAN produces the same set of clusters from one run to another (except that the membership of some border points depends on the order of the points being processed)
- Number of clusters
  - For k-means, the number of clusters needs to be specified as a parameter
  - DBSCAN automatically determines the number of clusters. However, it has two other parameters: Eps and MinPts

## DBSCAN: Determining Eps and MinPts

- Idea is that for points in a cluster, their k<sup>th</sup> nearest neighbors are at roughly the same distance
- Noise points have the k<sup>th</sup> nearest neighbor at farther distance
- So, plot sorted distance of every point to its k<sup>th</sup> nearest neighbor (e.g., k=4)



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- ▶ Cluster Validity

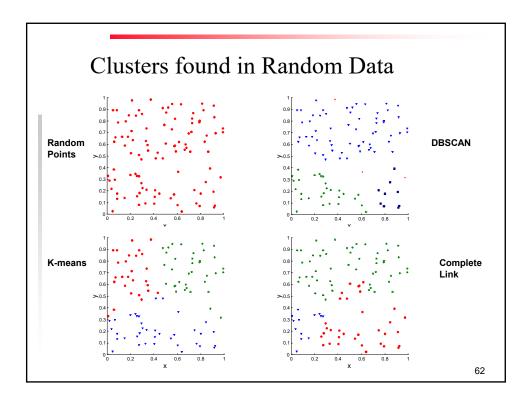
# Cluster Validity

- ► For supervised classification we have a variety of measures to evaluate how good our model is
  - Accuracy, error rate, confusion matrix, misclassification cost
- ► For cluster analysis, the analogous question is how to evaluate the "goodness" of the resulting clusters?
- ▶ But "clusters are in the eye of the beholder"!
- ▶ Then why do we want to evaluate them?
  - ▶ To compare clustering algorithms
  - ▶ To compare two sets of clusters
  - To determine the 'correct' number of clusters.

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#### Different Aspects of Cluster Validation

- Determining whether an algorithm can identify the clustering tendency of a set of data, i.e., whether it can distinguish whether non-random structure actually exists in the data.
- ▶ Determining whether a clustering result is good:
  - Comparing the results of a cluster analysis to externally known results, e.g., to externally given class labels.
  - Evaluating how well the results of a cluster analysis fit the data *without* reference to external information.
    - Use only the data



# Measures of Cluster Validity

- Numerical measures that are applied to judge various aspects of cluster validity, are classified into the following two types.
  - ► External Index: Used to measure the extent to which cluster labels match externally supplied class labels.
    - ▶ E.g., HA index, entropy
  - ▶ Internal Index: Used to measure the goodness of a clustering structure *without* external information.
    - ▶ E.g., Sum of Squared Error (SSE)
- Sometimes these are referred to as criteria instead of indices
  - ▶ However, sometimes criterion is the general strategy and index is the numerical measure that implements the criterion.

#### HA Index: an External Index

$$HA index = \frac{a+d}{a+b+c+d}$$

- ▶ *U* is the true partition in the data set.
- ▶ *V* is the clustering result by some algorithm.
- ullet a is the number of pairs of objects that are placed in the same class in U and in the same cluster in V
- b is the number of pairs of objects in the same class in U but not in the same cluster in V,
- c is the number of pairs of objects in the same cluster in V but not in the same class in U,
- d is the number of pairs of objects in different classes and different clusters in both partitions.

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#### SSE: an Internal Index

▶ Sum of Squared Error (SSE):

$$SSE = \sum_{i=1}^{k} \sum_{x \in C_i} dist(x, v_i)^2$$

 $v_i$  is the center of the cluster  $C_i$  k is the number of clusters

Considers only the compactness (i.e., the intracluster distances) of the clusters

#### DB: an Internal Index

Davies-Bouldin index (DB) :

$$DB = \frac{1}{k} \sum_{j=1..k, j \neq i} \max_{j=1..k, j \neq i} (d_{ij}) \quad \text{where} \quad d_{ij} = \frac{\sigma_i + \sigma_j}{d(v_i, v_j)}$$

- $\triangleright$  k is the number of clusters,
- $\sigma_i$  is the average distance between cluster points and the center in the *i*th cluster
- $d(v_i, v_j)$  is the distance between the *i*th and *j*th cluster centers.
- $d_{ij}$  decreases when the clusters are more compact and when the distance between them is larger.
- For each cluster  $C_i$ , the  $\max(d_{ij})$  identifies its "worst" neighbour. The DB index is the average of such value for all clusters.
- The DB index varies on the interval  $[0,\infty)$  and is small when the clusters are *compact and well separated*.

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# Final Comment on Cluster Validity

"The validation of clustering structures is the most difficult and frustrating part of cluster analysis.

Without a strong effort in this direction, cluster analysis will remain a black art accessible only to those true believers who have experience and great courage."

Algorithms for Clustering Data, Jain and Dubes

# Summary

- ► Cluster analysis groups objects based on their similarity and has wide applications
- ▶ We have looked at three clustering algorithms:
  - K-means
  - ▶ K-medoids
  - ▶ Hierarchical clustering
  - ▶ Density-based clustering
- ▶ Cluster Validity