CLUSTERING Methods

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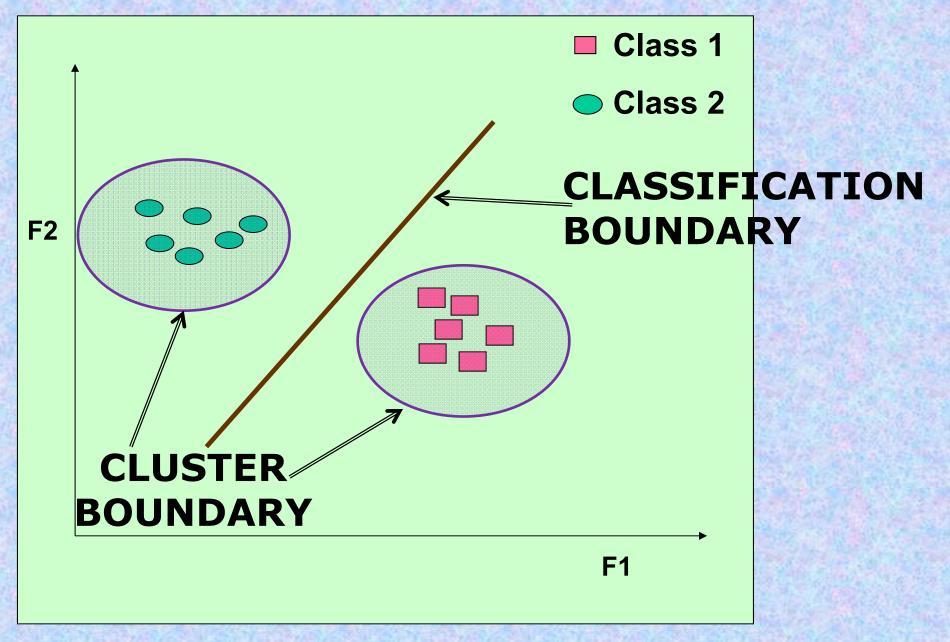
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What is Cluster Analysis?

- 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

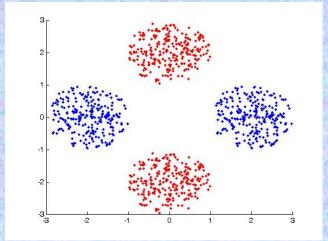
Clustering: Application Examples

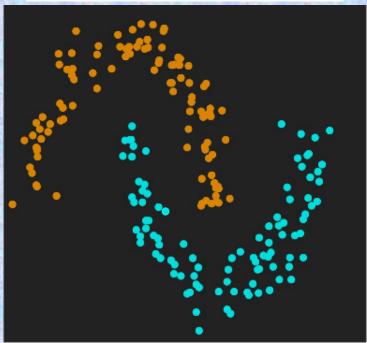
- Biology: taxonomy of living things: kingdom, phylum, class, order, family, genus and species
- Information retrieval: document clustering
- Land use: Identification of areas of similar land use in an earth observation database
- Marketing: Help marketers discover distinct groups in their customer bases, 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
- Economic Science: market research

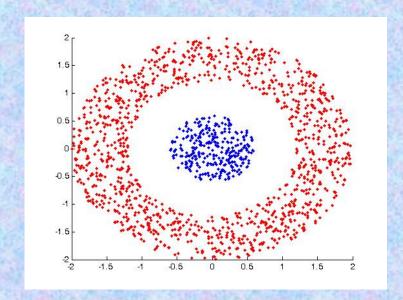


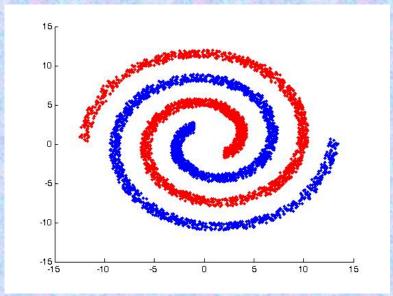
Sample points in a two-dimensional feature space

Complex cases of classification and clustering









CLUSTERING CLASSIFICATION Data Points have no labels Most data points have labels

CLUSTERING

METHODS OF AND CLASSIFICATION

- REPRESENTATIVE POINTS
- Split & MERGE
- LINKAGE
- · SOM
- MODEL-BASED
- VECTOR
 QUANTIZATION

Quality: What Is Good Clustering?

- A good clustering method will produce high quality clusters
 - high intra-class similarity: cohesive within clusters
 - low <u>inter-class</u> similarity: <u>distinctive</u> between clusters
- The <u>quality</u> of a clustering method depends on
 - the similarity measure used by the method
 - its implementation, and
 - Its ability to discover some or all of the <u>hidden</u> patterns

Considerations for Cluster Analysis

Partitioning criteria

- Single level vs. hierarchical partitioning (often, multi-level hierarchical partitioning is desirable)
- Separation of clusters
 - Exclusive (e.g., one customer belongs to only one region)
 vs. non-exclusive (e.g., one document may belong to more than one class)
- Similarity measure
 - Distance-based (e.g., Euclidian, road network, vector) vs. connectivity-based (e.g., density or contiguity)
- Clustering space
 - Full space (often when low dimensional) vs. subspaces (often in high-dimensional clustering)

Major Clustering Approaches (I)

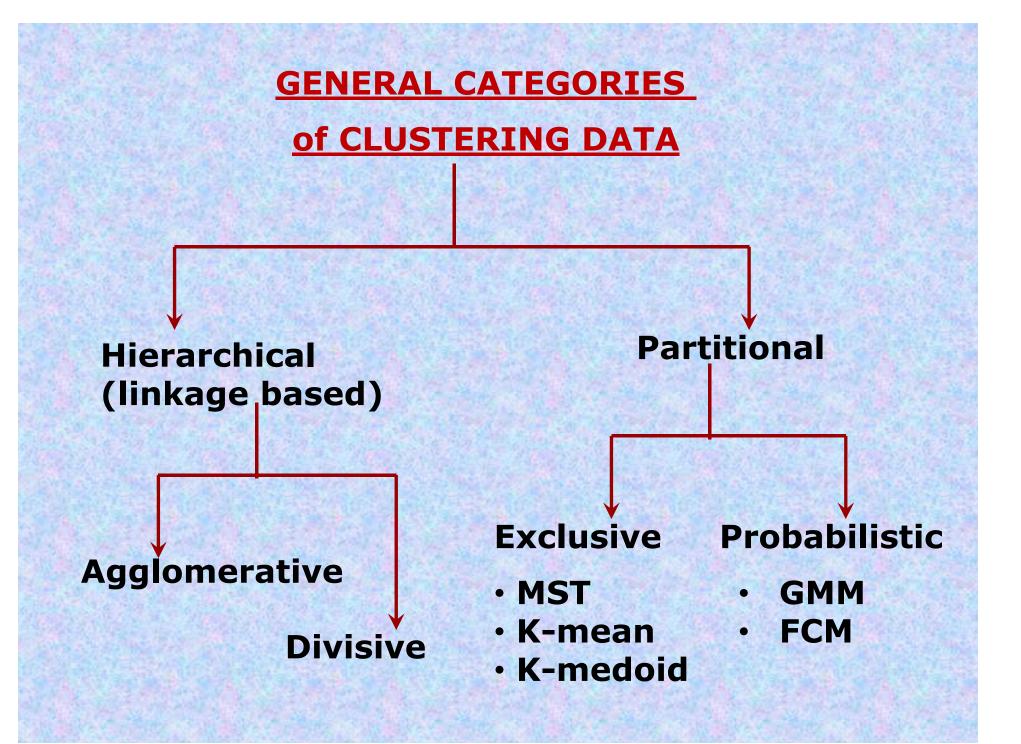
Partitioning approach:

- Construct various partitions and then evaluate them by some criterion, e.g., minimizing the sum of square errors
- Typical methods: k-means, k-medoids, CLARANS
- Hierarchical approach:
 - Create a hierarchical decomposition of the set of data (or objects) using some criterion
 - Typical methods: Diana, Agnes, BIRCH, CAMELEON
- Density-based approach:
 - Based on connectivity and density functions
 - Typical methods: DBSCAN, OPTICS, DenClue
- Grid-based approach:
 - based on a multiple-level granularity structure
 - Typical methods: STING, WaveCluster, CLIQUE

Major Clustering Approaches (II)

Model-based:

- A model is hypothesized for each of the clusters and tries to find the best fit of that model to each other
- Typical methods: EM, SOM, COBWEB
- Frequent pattern-based:
 - Based on the analysis of frequent patterns
 - Typical methods: p-Cluster
- <u>User-guided or constraint-based</u>:
 - Clustering by considering user-specified or applicationspecific constraints
 - Typical methods: COD (obstacles), constrained clustering
- Link-based clustering:
 - Objects are often linked together in various ways
 - Massive links can be used to cluster objects: SimRank,
 LinkClus

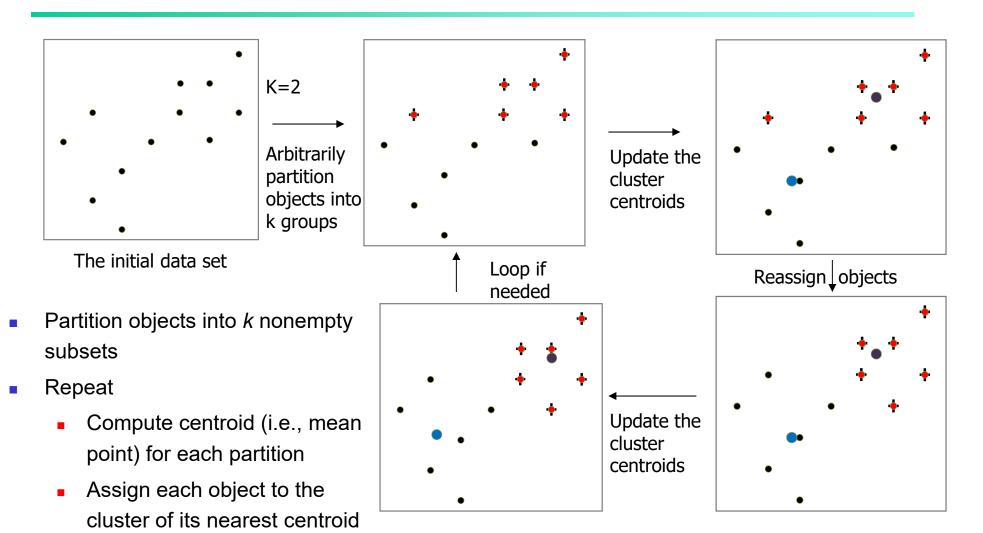


Alternative view of Algorithms for CLUSTERING

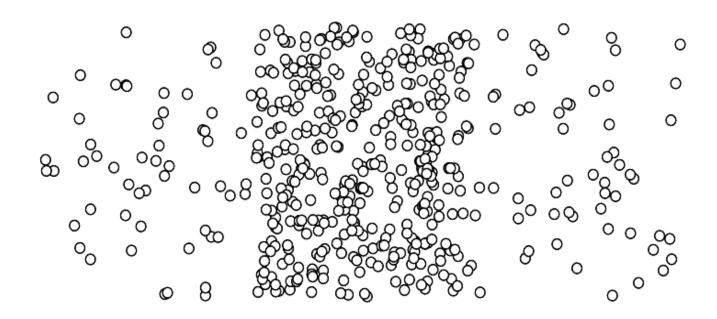
- Unupervised Learning/Classification:
 - K-means; K-medoid
- Density Estimation:
 - (i) Parametric
 - Gaussian
 - MOG (Mixture of Gaussians)
 - Dirichlet, Beta etc.
 - Branch and Bound Procedure
 - Piecewise Quadratic Boundary
 - Nearest Mean Classifier
 - MLE (maximum Likelihood Estimate)

- Density Estimation: (ii) Non-Parametric
 - Histogram
 - Neighborhood
 - Kernel Methods
 - Graph Theoretic
 - Iterative Valley Seeking

An Example of K-Means Clustering



Until no change



FCM - Fuzzy C-Means Clustering

FCM

- A method of clustering which allows one piece of data to belong to two or more clusters.
- Objective function to be minimized:

$$J_m = \sum_{i=1}^{N} \sum_{j=1}^{C} u_{ij}^m ||x_i - \mu_j||^2, \qquad 1 \le m < \infty$$

Where

- u_{ij} is the degree of membership of x_j in the cluster j.
- x_i is d-dimensional observation
- μ_j is d-dimensional center of cluster j

Updation

- FCM is an iterative optimization approach.
- At each step, the membership u_{ij} and the cluster centers μ_i are updated as follows:

$$u_{ij} = \frac{1}{\sum_{k=1}^{c} \left(\frac{\|x_i - \mu_j\|}{\|x_i - \mu_k\|} \right)^{\frac{2}{m-1}}},$$

$$\mu_{j} = \frac{\sum_{i=1}^{N} u_{ij}^{m}.x_{i}}{\sum_{i=1}^{N} u_{ij}^{m}}$$

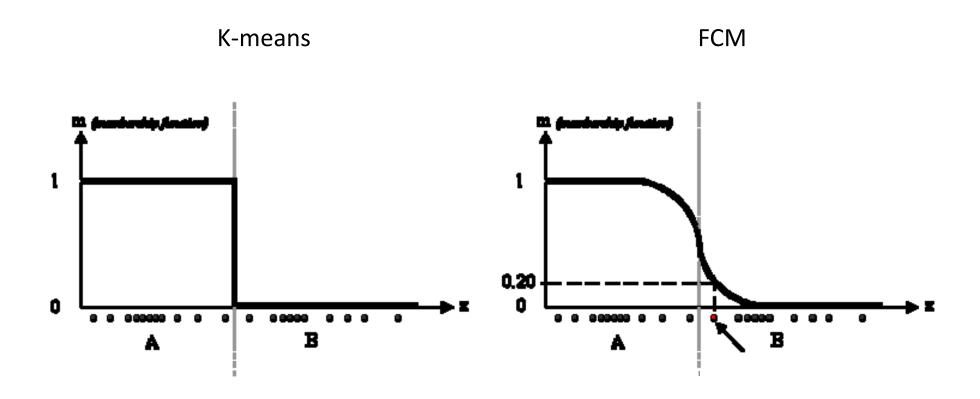
Termination Criterion

• Iteration stops, when $\max_{ij} \left\{ \left| u_{ij}^{(k+1)} - u_{ij}^{(k)} \right| \right\} < \epsilon$

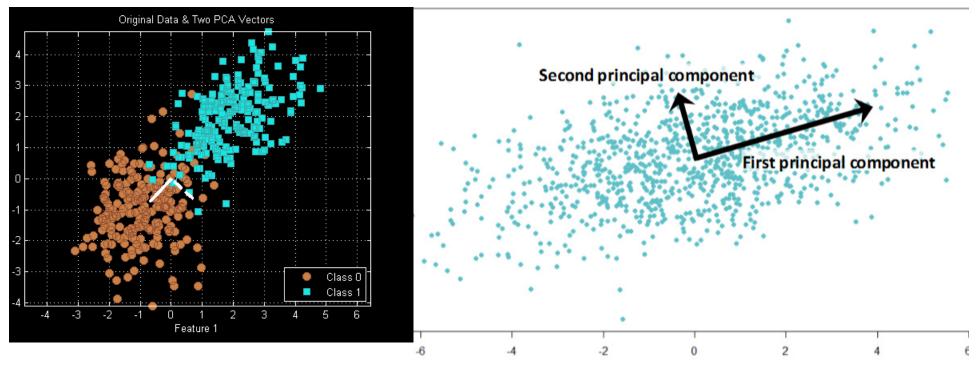
Where k is the iteration number.

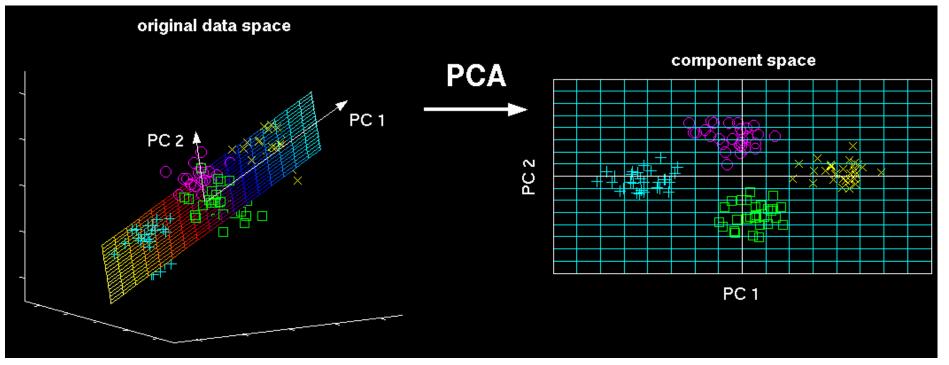
 ϵ is between 0 and 1

K-means Vs FCM



Read about K-medoids





Hierarchical Clustering

Hierarchical Clustering

Builds hierarchy of clusters

- Types:
 - Bottom Up *Agglomerative*
 - Starts by considering each observation as a cluster of it's own
 - Clusters are merged as we move up the hierarchy
 - Top Down *Divisive*
 - Starts by considering all observations in one cluster
 - Clusters are divided as we move down the hierarchy

Distance Functions

Certain mathematical properties are expected of any distance measure, or *metric*:

- 1. $d(x,y) \ge 0$ for all x, y.
- 2. d(x, y) = 0 iff x = y.
- 3. d(x, y) = d(y, x) (symmetry)
- 4. $d(x,y) \le d(x,z) + d(z,y)$ for all x, y, and z. (triangle inequality)

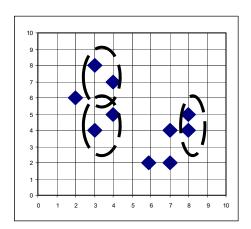
Euclidean distance $d(x,y) = \sqrt{\sum_{i=1}^{d} |x_i - y_i|^2}$ is probably the most commonly used metric. Note that it weights all features/dimensions "equally".

Some commonly used Metrics

- Euclidean distance
- Squared Euclidean distance
- Manhattan distance
- Maximum distance
- Mahalanobis distance

Agglomerative clustering

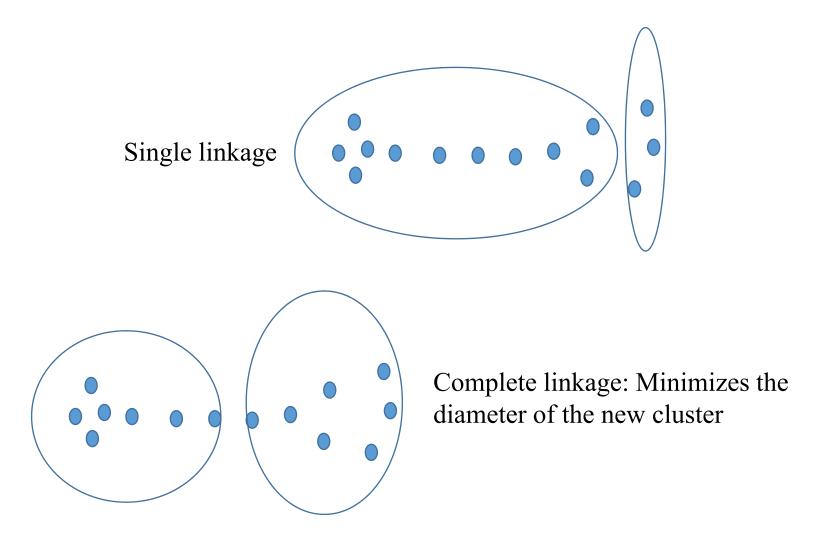
- Each node/object is a cluster initially
- Merge clusters that have the **least** dissimilarity
 - Ex: single-linkage, complete-linkage, etc.
- Go on in a non-descending fashion
- Eventually, all nodes belong to the same cluster



Linkage Criteria

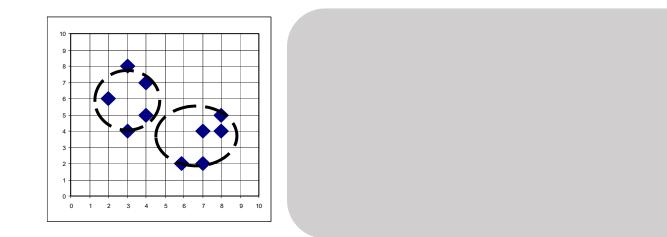
- Determines the distance between sets of observations as a function of the pairwise distances between observations.
- Some commonly used criterias:
 - Single Linkage: Distance between two clusters is the **smallest** pairwise distance between two observations/nodes, each belonging to different clusters.
 - Complete Linkage: Distance between two clusters is the largest pairwise distance between two observations/nodes, each belonging to different clusters.
 - Mean or average linkage clustering: Distance between two clusters is the **average** of all the pairwise distances, each node/observation belonging to different clusters.
 - Centroid linkage clustering: Distance between two clusters is the distance between their centroids.

Single Linkage vs. Complete Linkage



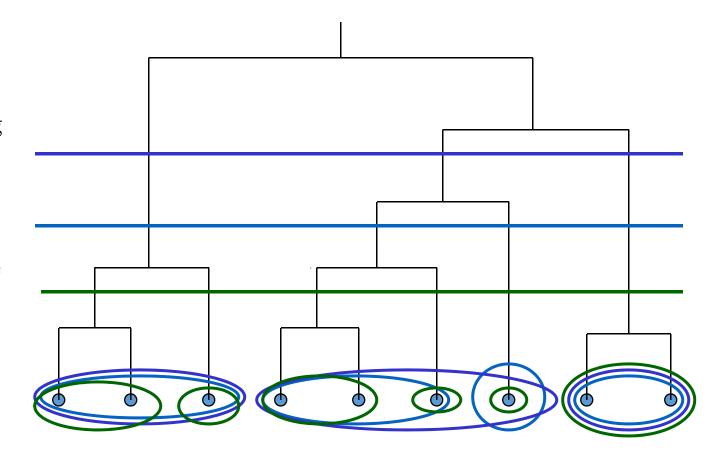
Divisive Clustering

- Initially, all data is in the same cluster
- The largest cluster is split until every object is separate.



What are the true number of clusters?

- Decompose data objects into a several levels of nested partitioning (tree of clusters), called a dendrogram.
- A <u>clustering</u> of the data objects is obtained by <u>cutting</u> the dendrogram at the desired level, then each <u>connected</u> <u>component</u> forms a cluster.



DBSCAN: Density Based Spatial Clustering of Applications with Noise

Density-Based Clustering Methods

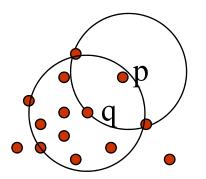
 Clustering based on density (local cluster criterion), such as density-connected points

- Major features:
 - Discover clusters of arbitrary shape
 - Handle noise
 - Need density parameters as termination condition
- Several interesting studies:
 - DBSCAN: Ester, et al. (KDD'96)
 - OPTICS: Ankerst, et al (SIGMOD'99).
 - DENCLUE: Hinneburg & D. Keim (KDD'98)
 - CLIQUE: Agrawal, et al. (SIGMOD'98) (more grid-based)

Density-Based Clustering: Basic Concepts

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

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



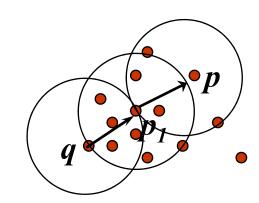
MinPts = 5

Eps = 1 cm

Density-reachable & Density-connected

• Density-reachable:

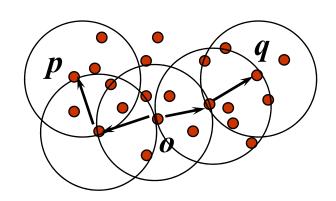
• A point p is density-reachable from a point q if there is a chain of points $p_1, ..., p_n, p_1 = q, p_n =$ p such that p_{i+1} is directly density-reachable from p_i



• This is not symmetric

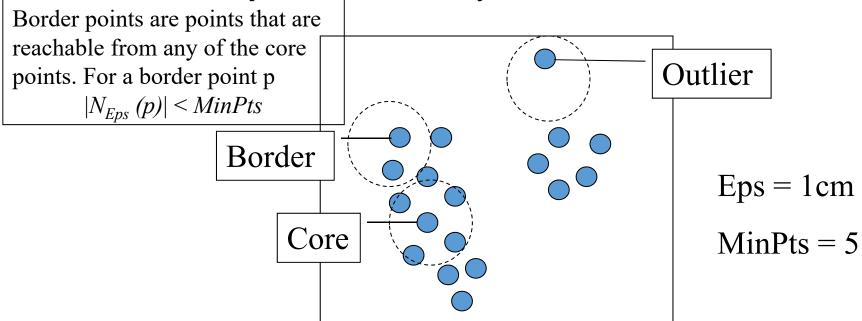
Density-connected

• 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

- A set of points C is a cluster, if
 - For any two points $p, q \in C$, p and q are density-connected
 - There does not exist any pair of points, $p \in C$ and $s \notin C$ such that p and s are density-connected.



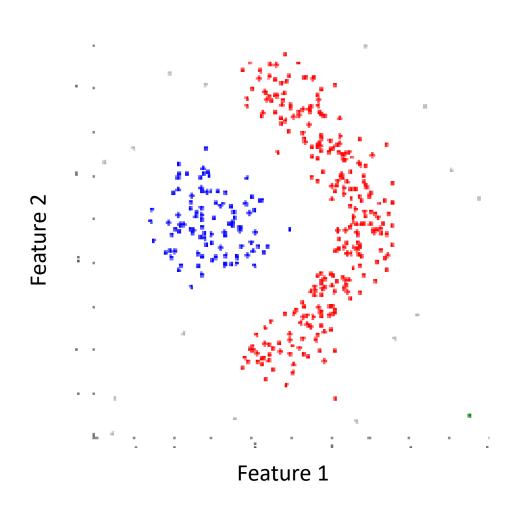
Algorithm

- Select a point p
- Retrieve all points directly density-reachable from p wrt. Eps and MinPts.
- If p is a not a core point, p is marked as noise
- Else a cluster is initiated.
 - p is marked as classified with a cluster ID
 - *seedSet* = all directly reachable points from *p*.
 - For each point p_i in seedSet till it is empty
 - If p_i is a noise point, assign p_i to the current cluster ID
 - If p_i is unclassified, identify if it is a core point. If yes, then add all directly reachable point to seed set and add p_i to cluster ID
 - Delete p_i from seedSet

DBSCAN: Properties

- Can discover clusters of arbitrary shapes
- Complexity
 - Time
 - $O(n^2)$
 - O(nlog^{d-1}n) with range tree. But requires more storage
 - d dimensions
- Weakness:
 - Parameter sensitive

DBSCAN - non-linearly separable clusters



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