Cluster Analysis (Part B)

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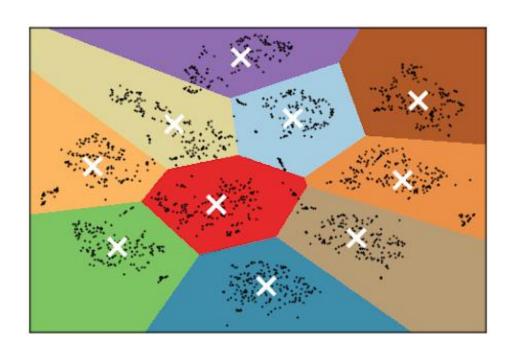
Major Clustering Approaches

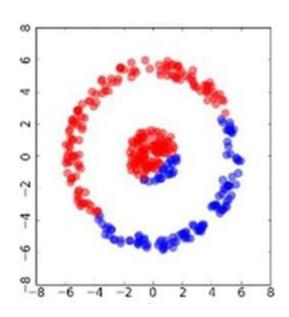
- ✓ Partitioning-based approach
 - Construct various partitions and then evaluate them by some criterion, e.g., minimizing the sum of square errors.
 - o Typical methods: k-means, k-medoids, CLARA, CLARANS
- Density-based approach
 - Based on connectivity and density functions
 - Typical methods: DBSACN, OPTICS, DenClue
- Hierarchical approach
 - Create a hierarchical decomposition of the set of data (or objects) using some criterion
 - o Typical methods: Agnes, Diana, BIRCH, CURE, CAMELEON
- Model-based approach
 - 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

Major Clustering Approaches (cont.)

- Grid-based approach
 - Based on a multiple-level granularity structure
 - Typical methods: STING, CLIQUE, WaveCluster
- Frequent Pattern-based approach
 - Based on the analysis of frequent patterns
 - Typical methods: p-Cluster
- Support Vector approach
 - Based on the idea of mapping data points into higher dimensional feature space via a kernel function.
 - Typical methods: SVC, Kernel K-means
- Graph Theoretic approach
 - Typical methods: Spectral Clustering

When K-means clustering fails





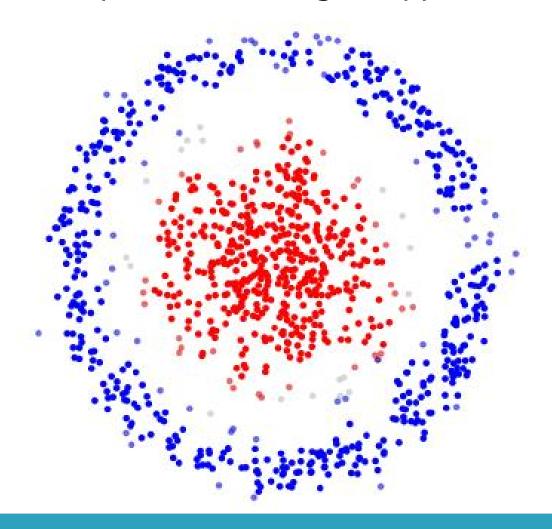
Density-based Approach

It is based on connectivity and density functions

Example: DBSCAN

DBSCAN

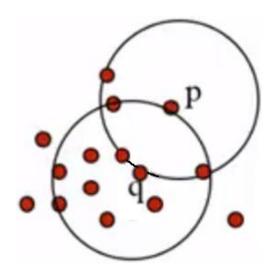
Density-Based Spatial Clustering of Applications with Noise



Density Definition

- Cluster: maximal set of density-connected points
- Parameters:
 - ε: max radius of the neighborhood
 - o minPts: min # of points in a ϵ -neighborhood of a point
 - The ϵ -neighborhood of a point q:

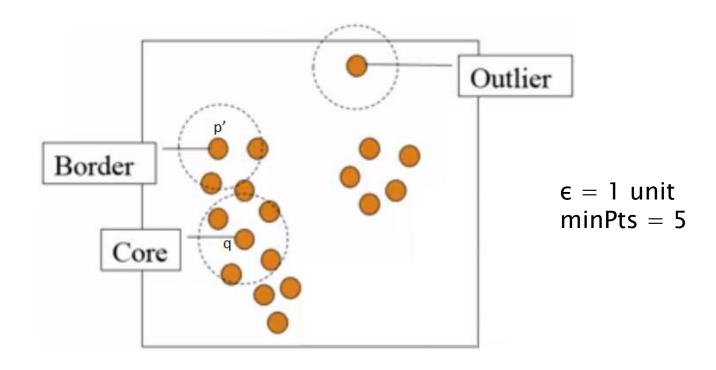
$$N_{\epsilon}(q) = \{p \text{ in } D \mid d(p,q) \leq \epsilon\}$$



$$\epsilon = 1$$
 unit minPts = 5

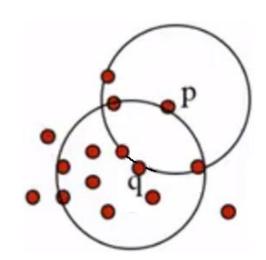
Core, Border, and Outlier

- Core point (q): dense neighborhood
- Border point (p'): in cluster, but neighborhood is not dense (reachable by the cluster)
- Outlier/noise: not in a cluster

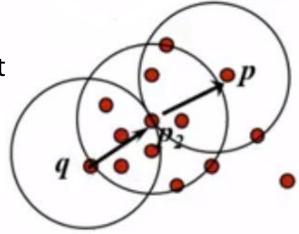


Density Reachability

- Directly density-reachable (ε, minPts):
 - A point p is density directly-reachable from a point q if:
 - p in $N_{\epsilon}(q)$
 - Core point condition: $|N_{\epsilon}(q)| \ge minPts$



- Density-reachable (ε, minPts):
 - A point p is directly-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 .



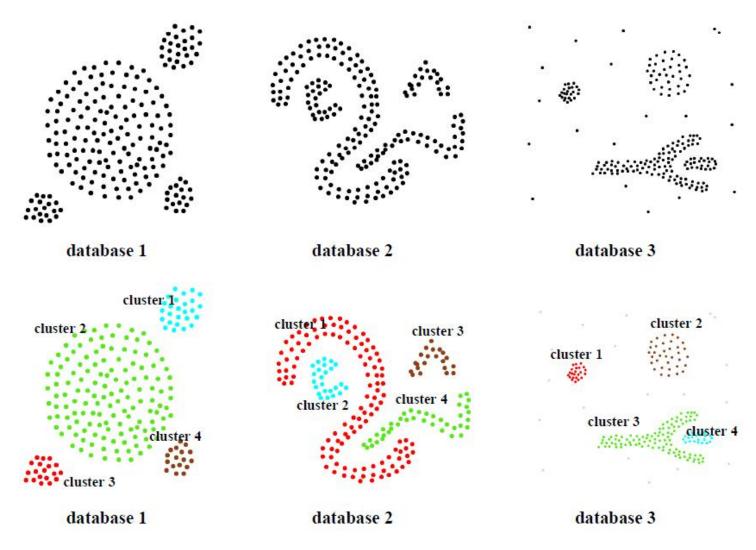
The DBSCAN Algorithm

- □ The DBSCAN algorithm (ϵ , minPts):
 - Start with an arbitrary point p from the database
 - Retrieve all density-reachable points from p
 - If p is a core point, a cluster is formed
 - If p is a border point, no points are density reachable from p.
 Thus DBSCAN visits the next points of the database
 - Continue the process until all of the points have been processed.

Example - Original Points



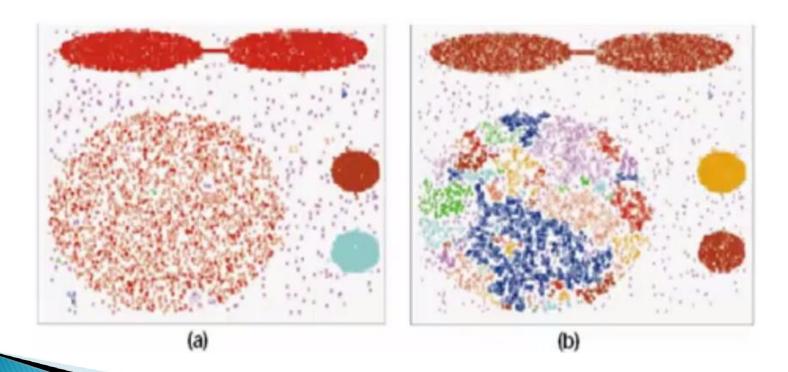
Discovered Clusters



M. Ester et al., A density-based algorithm for discovering clusters in large spatial databases with noise, 1996

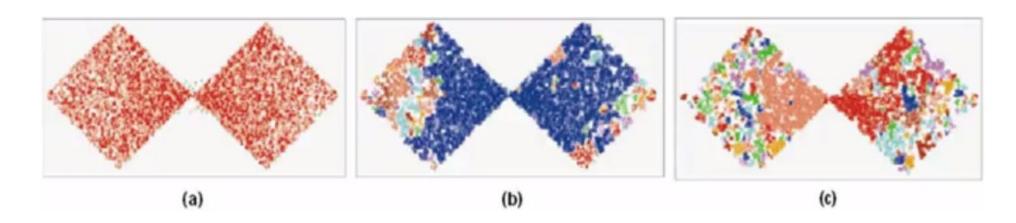
Sensitivity to Parameters

- \square minPts = 4
 - o (a) $\epsilon = 0.5$
 - \circ (b) $\epsilon = 0.4$



Sensitivity to Parameters

- \square minPts = 4
 - o (a) $\epsilon = 5.0$
 - \circ (b) $\epsilon = 3.5$
 - \circ (c) $\epsilon = 3.0$



Further Reading

- The OPTICS Clustering Algorithm
 - M. Ankerst, MM. Breunig, H-P. Kriegel, J. Sander, OPTICS: Ordering Points To Identify the Clustering Structure, ACM SIGMOD international conference on Management of data, pp. 49-60, 1999.

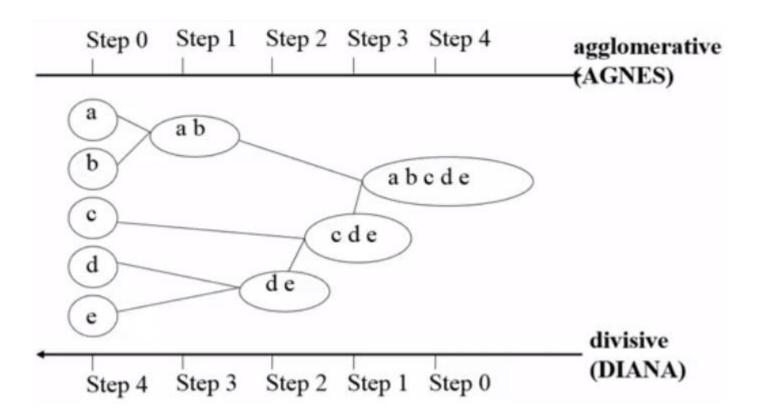
References

- M. Ester, H-P. Kriegel, J. Sander, X. Xu. "<u>A density-based</u> algorithm for discovering clusters in large spatial databases with noise", 1996.
 - [Highest impact paper award, 2014]
- J. Sander, M. Ester, H-P. Kriegel, X. Xu. Density-Based Clustering in Spatial Databases: The Algorithm GDBSCAN and Its Applications, 1998.

Hierarchical Approach

- Create a hierarchical decomposition of the set of data (or objects) using some criterion
- Two categories:
 - Agglomerative (bottom-up)
 - Divisive (top-down)

Dendrogram



AGNES (Agglomerative Nesting)
DIANA (Divisive Analysis)

Agglomerative Clustering

- The algorithm:
 - \circ Start with a collection \mathcal{C} on n singleton clusters
 - Each cluster contains one data point: $c_i = \{x_i\}$
 - Repeat
 - Find a pair of clusters that are closest: min $D(c_i, c_j)$
 - Merge the clusters c_i and c_j into a new cluster c_{i+j}
 - Remove c_i and c_j from the collection C, then add c_{i+j}
 - Until only one cluster left

Example

Cluster distance measure: single link

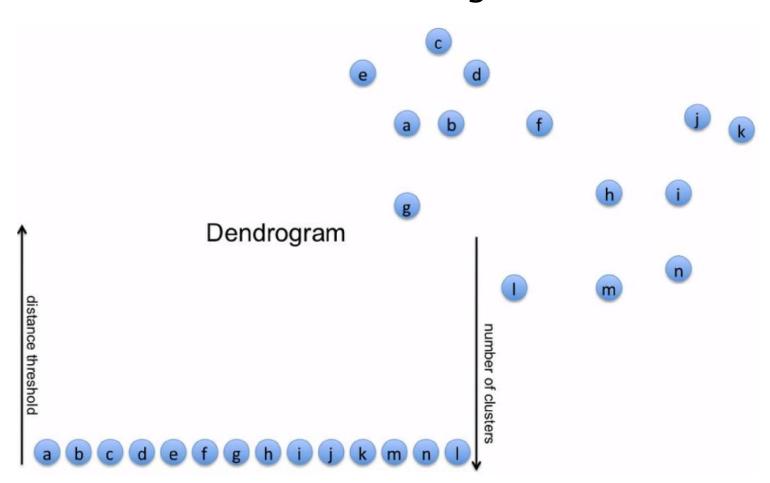


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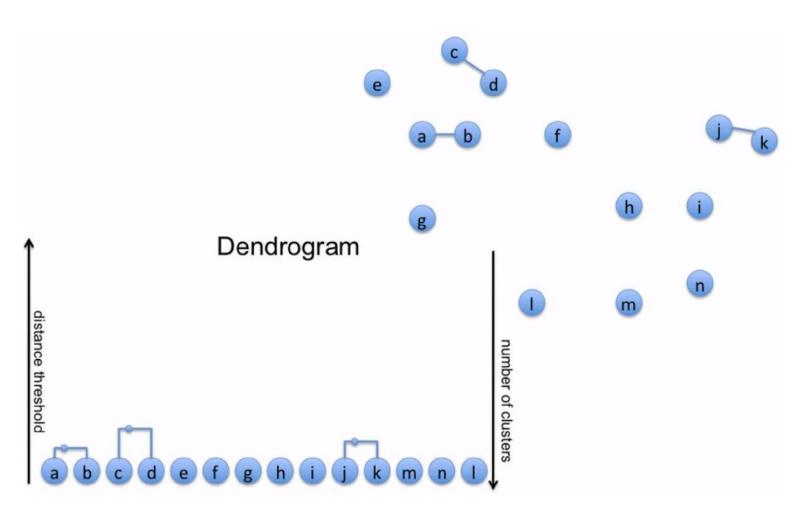


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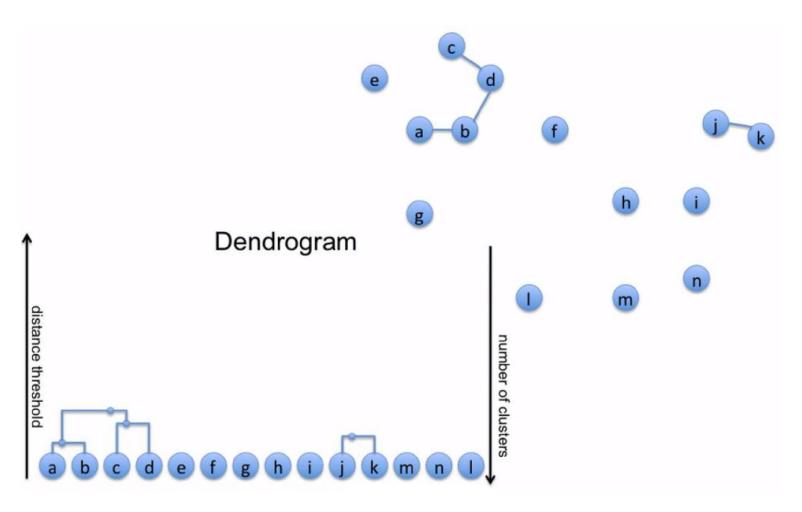


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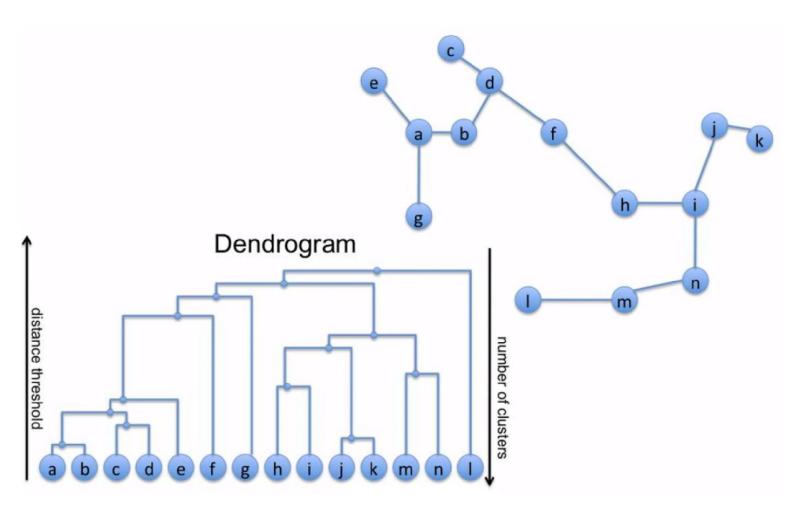


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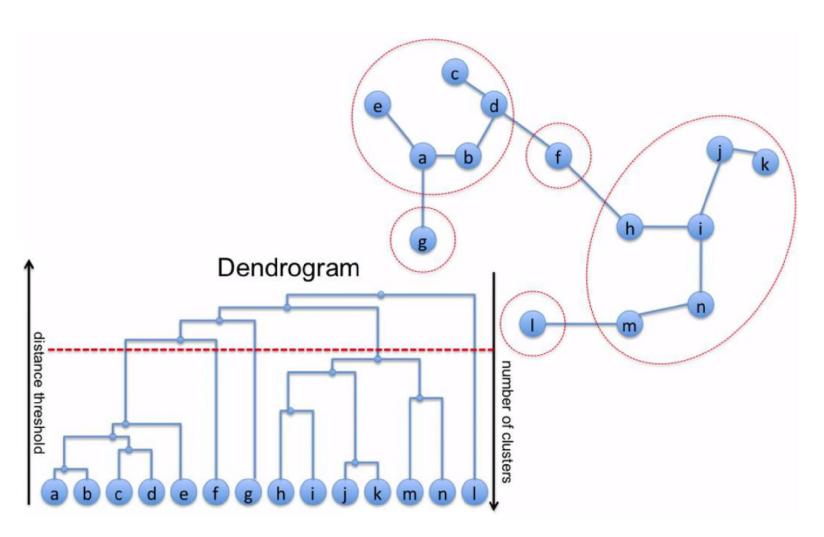


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Cluster Distance Measure

- □ Single link: $D(C_1, C_2) = \min(x_1, x_2)$
 - Distance between the closest elements in clusters
- □ Complete link: $D(C_1, C_2) = \max(x_1, x_2)$
 - Distance between the farthest elements in clusters
- Average link: average all pairwise distances
 - Less affected by outliers
- Centroids: distance between centroids (mean) of clusters
- Ward's method: sum of deviations from the centroid
 - The smaller, the better for merging.

Further Reading

The Birch Clustering Algorithm

 T. Zhang, R. Ramakrishnan, M. Livny, BIRCH: an efficient data clustering method for very large databases, Proceedings of the ACM SIGMOD international conference on Management of data, SIGMOD, pp 103-114, 1996.

The CHAMELEON Clustering Algorithm

 G. Karypis, E-H. Han, V. Kumar, CHAMELEON: A Hierarchical Clustering Algorithm Using Dynamic Modeling, IEEE Computer 32(8): 68-75, 1999.

Model-based Approach

A model is hypothesized for each of the clusters and tries to find the best fit of that model to each other

Example: EM Clustering

EM: Expectation–Maximization

Mixture Models in 1-D

- Case 1
 - Assume the data points come from two Gaussians with <u>unknown parameters</u>
 - We know each point comes from which Gaussian















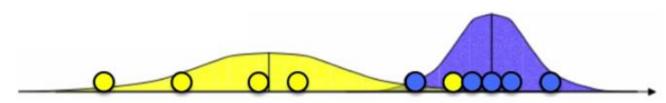
Mixture Models in 1-D

Case 1

- Assume the data points come from two Gaussians with unknown parameters
- We know each point comes from which Gaussian

$$\mu_b = \frac{x_1 + x_2 + \dots + x_{n_b}}{n_b}$$

$$\sigma_b^2 = \frac{(x_1 - \mu_b)^2 + \dots + (x_n - \mu_b)^2}{n_b}$$



Mixture Models of 1-D

- □ Case 2
 - We don't know the sources
 - We know the parameters of the Gaussians

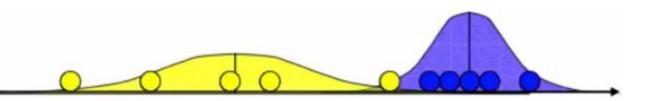


Mixture Models of 1-D

- □ Case 2
 - We don't know the sources
 - We know the parameters of the Gaussians

$$P(b \mid x_i) = \frac{P(x_i \mid b)P(b)}{P(x_i \mid b)P(b) + P(x_i \mid a)P(a)}$$

$$P(x_i | b) = \frac{1}{\sqrt{2\pi\sigma_b^2}} \exp\left\{-\frac{(x_i - \mu_i)^2}{2\sigma_b^2}\right\}$$



Expectation-Maximization

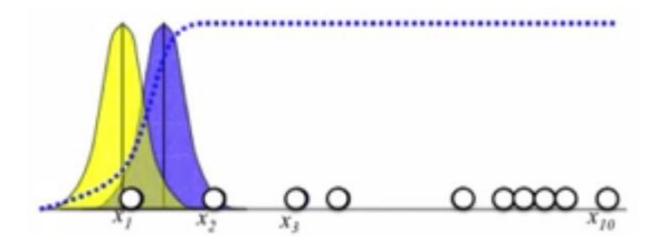
- Chicken and egg problem!
 - We need parameters to guess source of points
 - We need to know source to estimate parameters
- EM algorithm
 - Start with two randomly placed Gaussians
 - REPEAT
 - Use the Gaussians to determine which point comes from which Gaussian (Case 2)
 - Adjust parameters to fit points assigned to them (Case 1)
 - UNTIL Convergence

Expectation-Maximization (cont.)

- Chicken and egg problem!
 - We need parameters to guess source of points
 - We need to know source to estimate parameters
- EM algorithm
 - Start with two randomly placed Gaussians
 - REPEAT
 - Use the Gaussians to determine which point comes from which Gaussian (E-step)
 - Adjust parameters to fit points assigned to them (M-step)
 - UNTIL Convergence

EM Visualization

Initialization



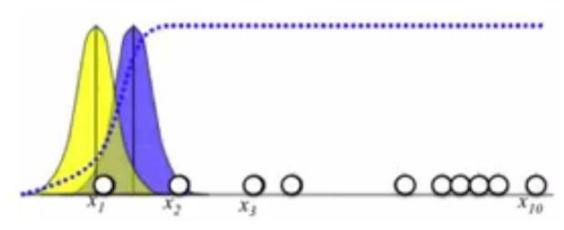
EM Visualization (cont.)

Initialization

$$P(x_i | b) = \frac{1}{\sqrt{2\pi\sigma_b^2}} \exp\left\{-\frac{(x_i - \mu_b)^2}{2\sigma_b^2}\right\}$$

$$b_i = P(b | x_i) = \frac{P(x_i | b)P(b)}{P(x_i | b)P(b) + P(x_i | a)P(a)}$$

$$a_i = P(a | x_i) = 1 - b_i$$

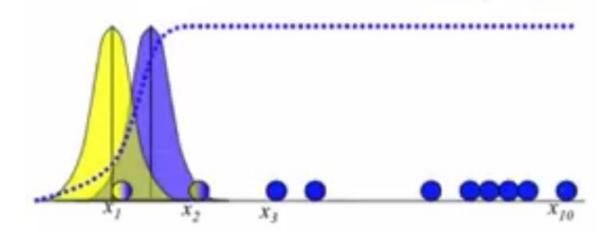


EM Visualization (E-Step)

$$P(x_{i}|b) = \frac{1}{\sqrt{2\pi\sigma_{b}^{2}}} \exp\left\{-\frac{(x_{i} - \mu_{b})^{2}}{2\sigma_{b}^{2}}\right\}$$

$$b_{i} = P(b|x_{i}) = \frac{P(x_{i}|b)P(b)}{P(x_{i}|b)P(b) + P(x_{i}|a)P(a)}$$

$$a_{i} = P(a|x_{i}) = 1 - b_{i}$$



EM Visualization (M-Step)

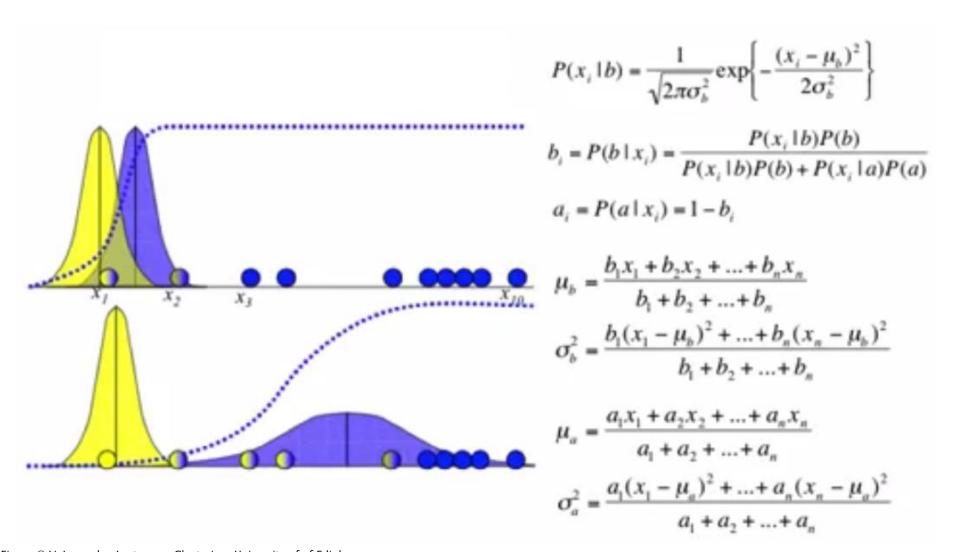


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EM Visualization (E-Step)

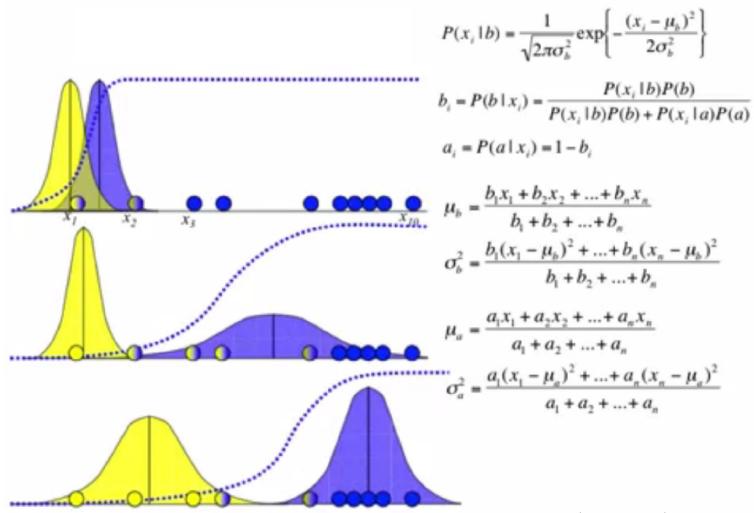


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EM Clustering

- Soft computing clustering
 - Clusters overlap
- K-means clustering is a special case of (hard) EM.
- Sensitive to the initialization
- How to pick K?

Further Reading

- The SOM and Spectral Clustering Algorithms
- Grid-based Clustering
- Subspace Clustering
- Collaborative Clustering
- **U** . . .

Measuring Clustering Quality

- Two methods: extrinsic vs. intrinsic
- Extrinsic: supervised, i.e., the ground truth is available
 - Compare clustering results against the ground truth labels
- Intrinsic: unsupervised, i.e., the ground truth is unavailable
 - Evaluate the goodness of a clustering by considering how well the clusters are separated
 - e.g. Entropy (purity)

Measuring Clustering Quality (cont.)

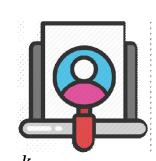
- □ Clustering quality measure: $Q(C, C_g)$, for a clustering C given the ground truth C_g .
- \square Q is good if it satisfies the following 4 essential criteria
 - Cluster homogeneity: the purer, the better
 - Cluster completeness: should assign objects belong to the same category in the ground truth to the same cluster
 - Rag bag: putting a heterogeneous object into a pure cluster should be penalized more than putting it into a rag bag (i.e., miscellaneous or the "other" category)
 - Small cluster preservation: splitting a small category into pieces is more harmful than splitting a large category into pieces

Measuring Clustering Quality (cont.)

Clustering quality:

User inspection





$$\rightarrow$$

$$SSE = \sum_{j=1}^{\kappa} \sum_{x \in C_j} \left| \left| x - \mu_j \right| \right|^2$$

$$\rightarrow$$

$$E_{total}(\Omega) = \sum_{w \in \Omega} \frac{m_w}{m} \times E(w)$$

Silhouette Index

 \rightarrow

...

O ...

Measuring Clustering Quality (cont.)

Total clustering entropy

$$E_{total}(\Omega) = \sum_{w \in \Omega} \frac{m_w}{m} \times E(w)$$
,

where $\Omega = \{w_1, w_2, ..., w_k\}$, m_w is the number of points in cluster w, m is the total number of points

Entropy of a single cluster w

$$E(w) = -\sum_{c \in C} \frac{|w_c|}{m_w} \times \log_2 \frac{|w_c|}{m_w},$$

where c is a class in the set C of all classes, $|w_c|$ is the number of points classified as c in cluster w

References

- Jiawei Han, Micheline Kamber and Jian Pei, Data Mining: Concepts and Techniques, 3rd edition, 2006.
- M. Ankerst, MM. Breunig, H-P. Kriegel, J. Sander, OPTICS: Ordering Points To Identify the Clustering Structure, ACM SIGMOD international conference on Management of data, pp. 49-60, 1999.
- M. Ester, H-P. Kriegel , J. Sander, X. Xu. "A density-based algorithm for discovering clusters in large spatial databases with noise", 1996.
- J. Sander, M. Ester, H-P. Kriegel, X. Xu. Density-Based Clustering in Spatial Databases: The Algorithm GDBSCAN and Its Applications, 1998.
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- □ G. Karypis, E-H. Han, V. Kumar, CHAMELEON: A Hierarchical Clustering Algorithm Using Dynamic Modeling, IEEE Computer 32(8): 68-75, 1999.