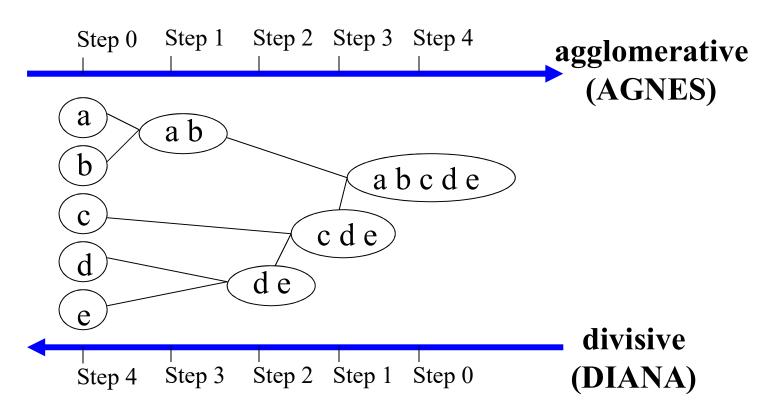
Chapter 7. Cluster Analysis

- 1. What is Cluster Analysis?
- 2. Types of Data in Cluster Analysis
- 3. A Categorization of Major Clustering Methods
- 4. Partitioning Methods
- 5. Hierarchical Methods
- 6. Density-Based Methods
- 7. Clustering High-Dimensional Data
- 8. Constraint-Based Clustering
- 9. Outlier Analysis
- 10.Summary



Hierarchical Clustering

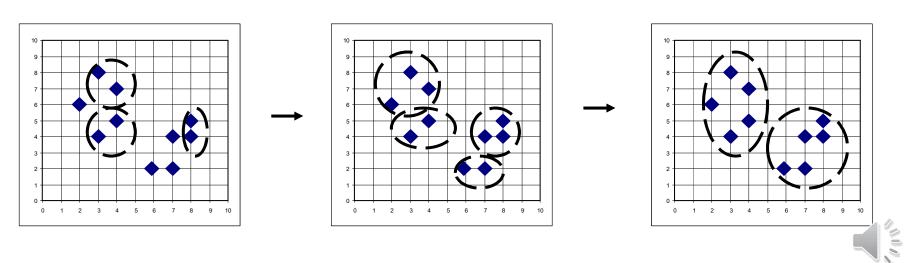
- Use a distance matrix as clustering criteria
- Does not require the number of clusters k as an input, but needs a termination condition



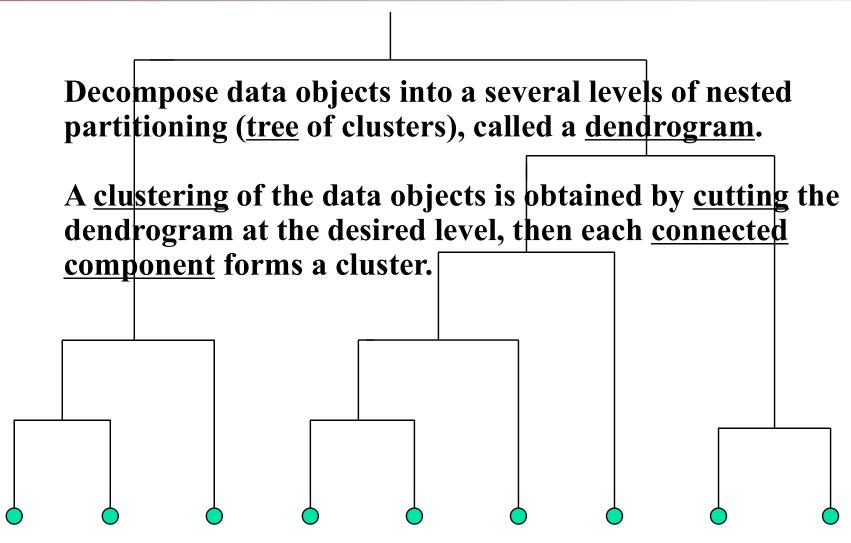


AGNES (Agglomerative Nesting)

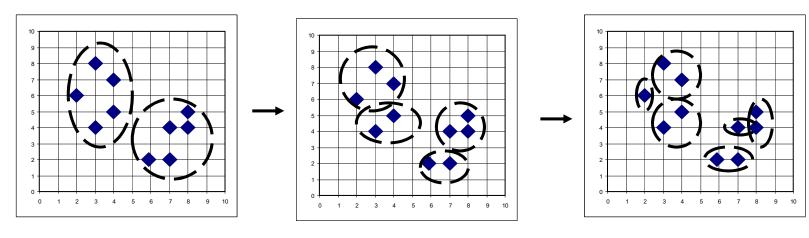
- Introduced in Kaufmann and Rousseeuw (1990)
 - Implemented in statistical analysis packages, Splus
- Use the single-link method and the dissimilarity matrix
- Merge nodes that have the least dissimilarity
- Go on in a non-descending fashion
- Eventually all nodes belong to the same cluster



Dendrogram: How the Clusters are Merged



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



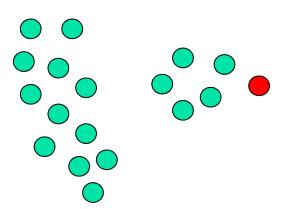


Outline

- Initially, there is one large cluster consisting of all n objects
- At each subsequent step, the largest available cluster is split into two clusters
 - Until finally all clusters comprise of a single object.
 - Thus, the hierarchy is built in n-1 steps.
- Complexity in the first step
 - Agglomerative method: $\frac{n(n-1)}{2}$ possible combinations
 - Divisive method: 2^{n-1} –1 possible combinations
 - Considerably larger than an agglomerative method

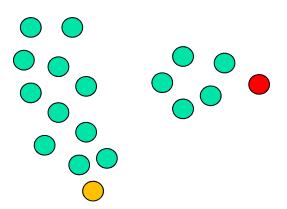


- To avoid considering all possibilities, the algorithm proceeds as follows.
 - 1. Find the object, which has the highest average dissimilarity to all other objects. This object initiates a new cluster— a sort of a *splinter group*.
 - 2. For each object i outside the *splinter group*, compute $D_i = [average \ d(i,j) \ j \notin R_{splinter group}] [average \ d(i,j) \ j \in R_{splinter group}]$
 - 3. Find an object h for which the difference D_h is the largest. If D_h is positive, then h is, on the average close to the splinter group. Put h into the splinter group.





- To avoid considering all possibilities, the algorithm proceeds as follows.
 - 1. Repeat Steps 2 and 3 until all differences $\,^D\!_h$ are negative. The data set is then split into two clusters.
 - 2. Select the cluster with the largest diameter. The diameter of a cluster is the largest dissimilarity between any two of its objects. Then divide this cluster, following steps 1-4.
 - 3. Repeat *Step* 5 until all clusters contain only a single object.





Advacned Hierarchical Clustering Methods

- Major weakness of agglomerative clustering methods
 - do not scale well: time complexity of at least $O(n^2)$, where n is the number of total objects
- Integration of hierarchical with distance-based clustering
 - BIRCH (1996): uses CF-tree and incrementally adjusts the quality of sub-clusters
 - ROCK (1999): clustering categorical data by neighbor and link analysis
 - CHAMELEON (1999): hierarchical clustering using dynamic modeling



BIRCH (1996)

- Birch: Balanced Iterative Reducing and Clustering using Hierarchies (Zhang, Ramakrishnan & Livny, SIGMOD'96)
- Incrementally construct a CF (Clustering Feature) tree (cf. B-tree),
 a hierarchical data structure for multiphase clustering
 - Phase 1: scan DB to build an initial in-memory CF tree (a multi-level compression of the data that tries to preserve the inherent clustering structure of the data)
 - Phase 2: use an arbitrary clustering algorithm to cluster the leaf nodes of the CF-tree
- Scales linearly: finds a good clustering with a single scan and improves the quality with a few additional scans
- Weakness: handles only numeric data, and sensitive to the order of the data records

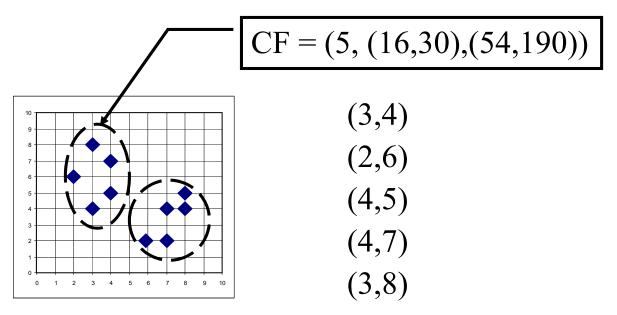
Clustering Feature Vector in BIRCH

Clustering Feature: CF = (N, LS, SS)

N: Number of data points

$$LS: \sum_{i=1}^{N} = X_i$$

SS:
$$\sum_{i=1}^{N} = X_i^2$$



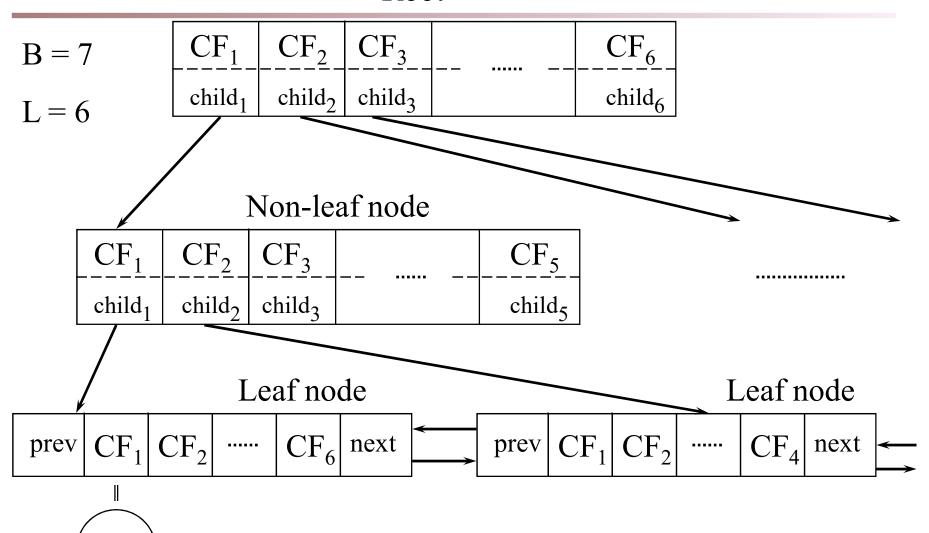
CF-Tree in BIRCH

- Clustering feature:
 - Summary of the statistics for a given cluster: the 0-th, 1st and 2nd moments of the cluster from the statistical point of view
 - Registers crucial measurements for computing cluster and utilizes storage efficiently
- A CF tree is a height-balanced tree that stores the clustering features for a hierarchical clustering
 - A non-leaf node in a tree has descendants or "children"
 - A non-leaf node stores the sum of the CFs of their children
- A CF tree has two parameters
 - Branching factor: specify the maximum number of children
 - threshold: max diameter of a cluster stored at the leaf node



The CF Tree Structure

Root



May 3, 2020

Clustering Categorical Data: The ROCK Algorithm

- ROCK: RObust Clustering using links, ICDE'99
- Major ideas
 - Use the notion of *links* to measure similarity/proximity
 - Not distance-based



Similarity Measure in ROCK

- Traditional measures for categorical data may not work well, e.g., Jaccard coefficient
- Jaccard coefficient-based similarity function:

$$Sim(T_1, T_2) = \frac{\left|T_1 \cap T_2\right|}{\left|T_1 \cup T_2\right|}$$

• Ex. Let $T_1 = \{a, b, c\}, T_2 = \{c, d, e\}$

Sim
$$(T_1, T_2) = \frac{|\{c\}|}{|\{a, b, c, d, e\}|} = \frac{1}{5} = 0.2$$

Similarity Measure in ROCK

- Example: Two groups (clusters) of transactions
 - C₁. <a, b, c, d, e>: {a, b, c}, {a, b, d}, {a, b, e}, {a, c, d}, {a, c, e}, {a, d, e}, {b, c, d}, {b, c, e}, {b, d, e}, {c, d, e}
 - C₂. <a, b, f, g>: {a, b, f}, {a, b, g}, {a, f, g}, {b, f, g}
- Jaccard coefficient may lead to a wrong clustering result
 - C₁: 0.2 ({a, b, c}, {b, d, e}) to 0.5 ({a, b, c}, {a, b, d})
 - $C_1 \& C_2$: could be as high as 0.5 ({a, b, c}, {a, b, f})

$$Sim(T_1, T_2) = \frac{\left|T_1 \cap T_2\right|}{\left|T_1 \cup T_2\right|}$$

Link Measure in ROCK

- Links: # of common neighbors (threshold = 0.5 in jC)
 - C₁ <a, b, c, d, e>: {a, b, c}, {a, b, d}, {a, b, e}, {a, c, d}, {a, c, e}, {a, d, e}, {b, c, d}, {b, c, e}, {b, d, e}, {c, d, e}
 - C₂ <a, b, f, g>: {a, b, f}, {a, b, g}, {a, f, g}, {b, f, g}
- Let $T_1 = \{a, b, c\}, T_2 = \{c, d, e\}, T_3 = \{a, b, f\}$
 - $link(T_1, T_2) = 4$, since they have 4 common neighbors
 - {a, c, d}, {a, c, e}, {b, c, d}, {b, c, e}
 - link(T_1, T_3) = 3, since they have 3 common neighbors
 - {a, b, d}, {a, b, e}, {a, b, g}
- Thus, link is a better measure than Jaccard coefficient

CHAMELEON: Hierarchical Clustering Using Dynamic Modeling (1999)

- CHAMELEON: by G. Karypis, E.H. Han, and V. Kumar'99
- Measures the similarity based on a dynamic model
 - Two clusters are merged only if the interconnectivity and closeness (proximity) between two clusters are high
 - Relative to the internal interconnectivity of the clusters and internal closeness
 of items within the clusters

CHAMELEON: Hierarchical Clustering Using Dynamic Modeling (1999)

- Draw a k-nearest neighbor graph first
 - Node: object, edge: k-nearest neighbor's link, weight: similarity
- A two-phase algorithm
 - Use a graph partitioning algorithm:
 - Cluster objects into a large number of relatively small sub-clusters
 - Use an agglomerative hierarchical clustering algorithm:
 - Find the genuine clusters by repeatedly combining these sub-clusters

CHAMELEON: Hierarchical Clustering Using Dynamic Modeling (1999)

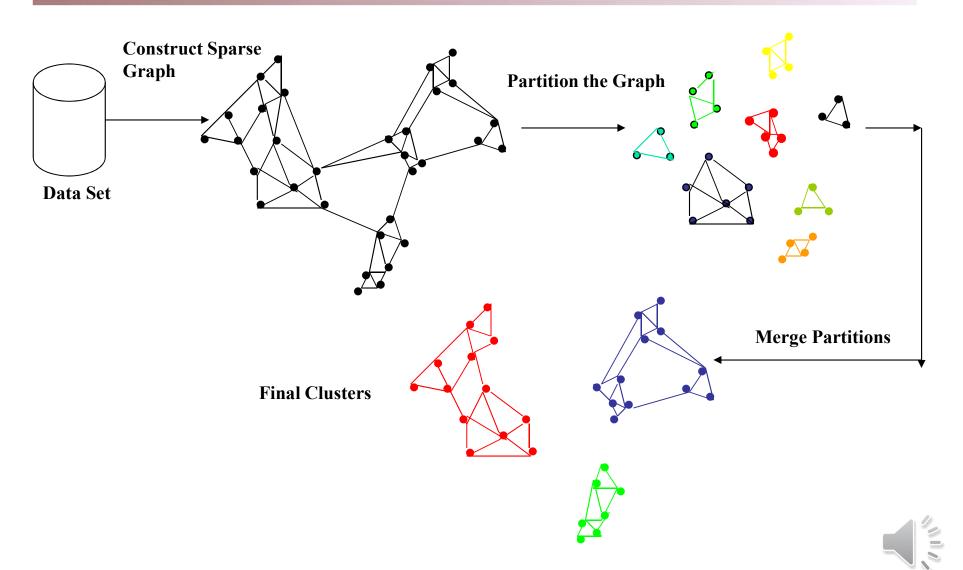
- Partitioning
 - To minimize the edge cut (METIS)
 - Tries to split a graph into two subgraphs of nearly equal sizes
- Relative interconnectivity

$$RI(C_i, C_j) = \frac{|EC_{\{C_i, C_j\}}|}{\frac{1}{2}(|EC_{C_i}| + |EC_{C_j}|)},$$

Relative closeness

$$RC(C_i, C_j) = \frac{S_{EC_{\{C_i, C_j\}}}}{\frac{|C_i|}{|C_i| + |C_j|} \overline{S}_{EC_{C_i}} + \frac{|C_j|}{|C_i| + |C_j|} \overline{S}_{EC_{C_j}}},$$

Overall Framework of CHAMELEON



CHAMELEON (Clustering Complex Objects)

