Data Mining 4. Cluster Analysis

4.4 Hierarchical Methods

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Outline

- Introduction
- BIRCH Algorithm
- References

• A hierarchical clustering method works by grouping data objects into a tree of clusters.

• Types of hierarchical clustering methods:

- Agglomerative: the hierarchical decomposition is formed in a bottom-up (merging) fashion.
- Divisive: the hierarchical decomposition is formed in a topdown (splitting) fashion.

• Agglomerative hierarchical clustering

- This bottom-up strategy starts by placing each object in its own cluster and then merges these atomic clusters into larger and larger clusters, until all of the objects are in a single cluster or until certain termination conditions are satisfied.
- Most hierarchical clustering methods belong to this category.
- They differ only in their definition of intercluster similarity.

• Divisive hierarchical clustering

- This top-down strategy starts with all objects in one cluster.
- It subdivides the cluster into smaller and smaller pieces, until each object forms a cluster on its own or until it satisfies certain termination conditions,
- Termination conditions can be
 - ◆ a desired number of clusters is obtained or
 - the diameter of each cluster is within a certain threshold.

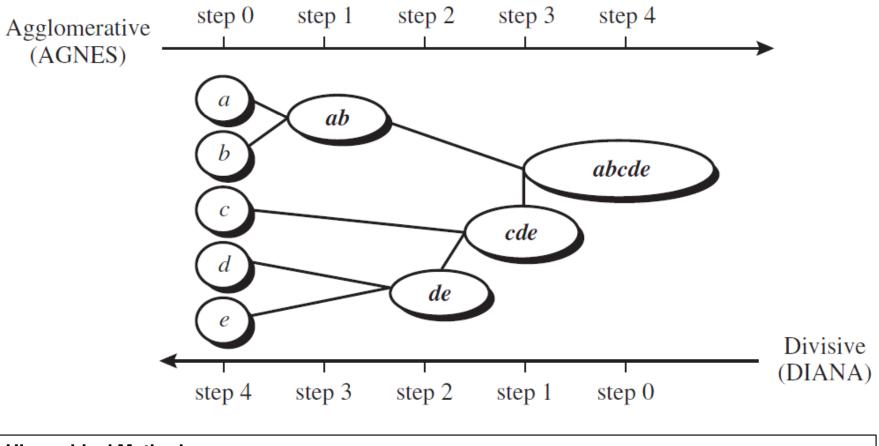
Example

• Example: Agglomerative versus divisive hierarchical clustering

- the application of AGNES (AGglomerative NESting), an agglomerative hierarchical clustering method,
- and DIANA (DIvisive ANAlysis), a divisive hierarchical clustering method, to a data set of five objects, {*a*, *b*, *c*, *d*, *e*}.

Example

• Agglomerative and divisive hierarchical clustering on data objects {*a*, *b*, *c*, *d*, *e*}.



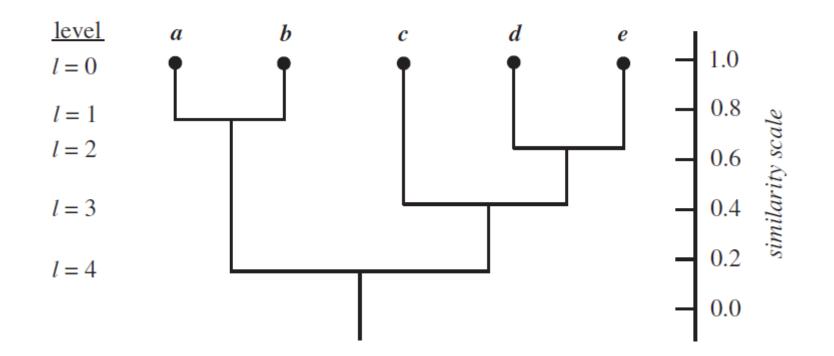
Dendrogram

• Dendrogram

- A tree structure which is commonly used to represent the process of hierarchical clustering.
- It shows how objects are grouped together step by step.

Dendrogram

• Dendrogram representation for hierarchical clustering of data objects {*a*, *b*, *c*, *d*, *e*}.



- Common measures for distance between clusters are as follows:
 - Minimum distance
 - Maximum distance
 - Mean distance
 - Average distance

Notation

- |p-p'|: is the distance between two objects or points, *p* and *p'*
- m_i is the mean for cluster, C_i
- n_i is the number of objects in C_i
- m_j is the mean for cluster, C_j

• Minimum distance

$$d_{min}(C_i, C_j) = \min_{\boldsymbol{p} \in C_i, \, \boldsymbol{p}' \in C_j} |\boldsymbol{p} - \boldsymbol{p}'|$$

- When an algorithm uses the minimum distance, it is sometimes called a nearest-neighbor clustering algorithm.
- If the clustering process is terminated when the distance between nearest clusters exceeds an **arbitrary threshold**, it is called **a single-linkage algorithm**.

• Maximum distance

$$d_{max}(C_i, C_j) = \max_{\boldsymbol{p} \in C_i, \, \boldsymbol{p}' \in C_j} |\boldsymbol{p} - \boldsymbol{p}'|$$

- When an algorithm uses the maximum distance, it is sometimes called a farthest-neighbor clustering algorithm.
- If the clustering process is terminated when the maximum distance between nearest clusters exceeds an arbitrary threshold, it is called a complete-linkage algorithm.
- Farthest-neighbor algorithms tend to minimize the increase in diameter of the clusters at each iteration as little as possible.

• Mean distance

$$d_{mean}(C_i, C_j) = |\boldsymbol{m_i} - \boldsymbol{m_j}|$$

- The minimum and maximum measures tend to be overly sensitive to outliers or noisy data.
- The use of mean or average distance is a compromise between the minimum and maximum distances and overcomes the outlier sensitivity problem.

• Average distance

$$d_{avg}(C_i, C_j) = \frac{1}{n_i n_j} \sum_{\boldsymbol{p} \in C_i} \sum_{\boldsymbol{p}' \in C_j} |\boldsymbol{p} - \boldsymbol{p}'|$$

- Whereas the mean distance is the simplest to compute, the average distance is advantageous in that it can handle categoric as well as numeric data.
- The computation of the mean vector for categoric data can be difficult or impossible to define.

The Difficulties with Hierarchical Clustering

- The quality of a pure hierarchical clustering method suffers from its inability to perform adjustment once a merge or split decision has been executed.
- That is, if a particular merge or split decision later turns out to have been a poor choice, the method cannot backtrack and correct it.
- Recent studies have emphasized the integration of hierarchical agglomeration with iterative relocation methods.

The Difficulties with Hierarchical Clustering

- Three such methods are introduced in this chapter, including:
 - BIRCH,
 - begins by partitioning objects hierarchically using tree structures, where the leaf or low-level nonleaf nodes can be viewed as "microclusters" depending on the scale of resolution.
 - It then applies other clustering algorithms to perform macroclustering on the microclusters.
 - ROCK
 - Merges clusters based on their interconnectivity.
 - Chameleon,
 - Explores dynamic modeling in hierarchical clustering.

• BIRCH: Balanced Iterative Reducing and Clustering Using Hierarchies

- BIRCH is designed for clustering a large amount of numerical data
- It integrates the hierarchical clustering (at the initial microclustering stage) and other clustering methods such as iterative partitioning (at the later macroclustering stage).
- It overcomes the two difficulties of agglomerative clustering methods:
 - (1) scalability and
 - ◆ (2) the inability to undo what was done in the previous step.

- BIRCH introduces two concepts:
 - Clustering Feature (CF)
 - Clustering feature tree (CF tree)
- They are used to summarize cluster representations.
- These structures help the clustering method achieve good speed and scalability in large databases and also make it effective for incremental and dynamic clustering of incoming objects.

Given *n d*-dimensional data objects or points in a cluster, we can define the centroid *x₀*, radius *R*, and diameter *D* of the cluster as follows:

$$\boldsymbol{x_0} = \frac{\sum_{i=1}^{n} \boldsymbol{x_i}}{n} \quad R = \sqrt{\frac{\sum_{i=1}^{n} (\boldsymbol{x_i} - \boldsymbol{x_0})^2}{n}} \quad D = \sqrt{\frac{\sum_{i=1}^{n} \sum_{j=1}^{n} (\boldsymbol{x_i} - \boldsymbol{x_j})^2}{n(n-1)}}$$

- Where R is the average distance from member objects to the centroid, and D is the average pairwise distance within a cluster.
- Both R and D reflect the tightness of the cluster around the centroid.

• Clustering Feature (CF)

- CF is a three-dimensional vector summarizing information about clusters of objects.
- Given *n d*-dimensional objects or points in a cluster, $\{x_i\}$, then the CF of the cluster is defined as:

 $CF = \langle n, LS, SS \rangle$

- where *n* is the number of points in the cluster,
- LS is the linear sum of the *n* points, i.e.,

$$\sum_{i=1}^{n} \mathbf{x}_i$$

- SS is the square sum of the data points, i.e.,

$$\sum_{i=1}^{n} x_i^2$$

- Clustering features are **additive**.
- For example, suppose that we have two disjoint clusters, C_1 and C_2 , having the clustering features, CF_1 and CF_2 , respectively.
- The clustering feature for the cluster that is formed by merging C_1 and C_2 is simply $CF_1 + CF_2$.
- Clustering features are sufficient for calculating all of the measurements that are needed for making clustering decisions in BIRCH.

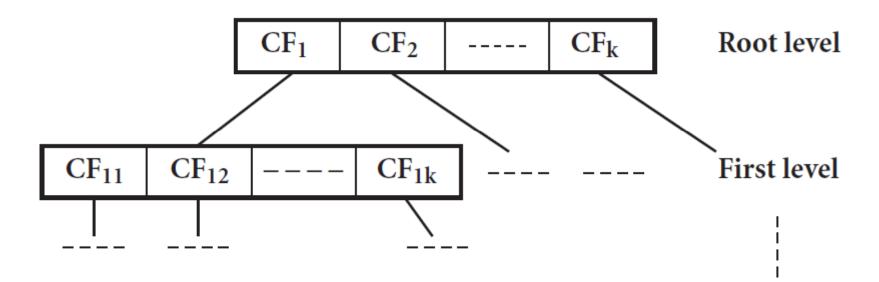
• Example: Clustering feature.

- Suppose that there are three points, (2, 5), (3, 2), and (4, 3), in a cluster, C_I . The clustering feature of C_I is: $CF_1 = \langle 3, (2+3+4,5+2+3), (2^2+3^2+4^2,5^2+2^2+3^2) \rangle$ $= \langle 3, (9,10), (29,38) \rangle.$
- Suppose that C_1 is joint to a second cluster, C_2 , where $CF_2 = \langle 3, (35, 36), (417, 440) \rangle$.
- The clustering feature of a new cluster, C_3 , that is formed by merging C_1 and C_2 , is derived by adding CF_1 and CF_2 . That is:

$$CF_3 = \langle 3+3, (9+35, 10+36), (29+417, 38+440) \rangle$$

= $\langle 6, (44, 46), (446, 478) \rangle.$

• A **CF tree** is a height-balanced tree that stores the clustering features for a hierarchical clustering.



- By definition, a nonleaf node in a tree has **children**.
- The nonleaf nodes store sums of the CFs of their children, and thus summarize clustering information about their children.
- A CF tree has two parameters:
 - Branching factor, B
 - specifies the maximum number of children per nonleaf node.
 - Threshold, T
 - specifies the maximum diameter of subclusters stored at the leaf nodes of the tree.
- These two parameters influence the size of the resulting tree.

BIRCH Algorithm Phases

- The primary phases of BIRCH are:
- Phase 1:
 - BIRCH scans the database to build an initial in-memory CF tree
- **Phase 2**:
 - BIRCH applies a (selected) clustering algorithm to cluster the leaf nodes of the CF tree, which removes sparse clusters as outliers and groups dense clusters into larger ones.

BIRCH Algorithm Phases

• Phase 1:

- the CF tree is built dynamically as objects are inserted.
- Thus, the method is **incremental**.
- An object is inserted into the closest leaf entry (subcluster).
- If the diameterc of the subcluster stored in the leaf node after insertion is larger than the threshold value, then the leaf node and possibly other nodes are split.
- After the insertion of the new object, information about it is passed toward the root of the tree.
- The size of the CF tree can be changed by modifying the threshold.

BIRCH Algorithm Phases

• Phase 2:

 Once the CF tree is built, any clustering algorithm, such as a typical partitioning algorithm, can be used with the CF tree in Phase 2.

Computation Complexity of the Algorithm

- The computation complexity of the algorithm is O(n),
 - were n is the number of objects to be clustered.
- Experiments have shown the linear scalability of the algorithm with respect to the number of objects and good quality of clustering of the data.

Weakness of BIRCH

- However, since each node in a CF tree can hold only a limited number of entries due to its size, a CF tree node does not always correspond to what a user may consider a natural cluster.
- Moreover, if the clusters are not spherical in shape, BIRCH does not perform well, because it uses the notion of radius or diameter to control the boundary of a cluster.

References

References

• J. Han, M. Kamber, **Data Mining: Concepts and Techniques**, Elsevier Inc. (2006). (Chapter 7)

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