Lecture 10: Hierarchical Agglomerative Clustering

Sophie Robert

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Cluster linkage

Agglomerative

Dendograms and

Number of cluster selection

Advantages and limits

Lecture 10: Hierarchical Agglomerative Clustering

Introduction to Machine Learning

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L3 MIASHS — Semestre 2

2022-2023

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Definition: hierarchical clustering

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Hierarchical clustering

Hierarchical clustering* is a clustering method that seeks to build a hierarchy of clusters (clusters within clusters).

Definition: hierarchical clustering

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Hierarchical clustering

Hierarchical clustering* is a clustering method that seeks to **build a hierarchy of clusters** (clusters within clusters).

Two approaches:

Bottom-up approach (agglomerative*): Each observation starts in its own cluster, and pairs of clusters are merged as one moves up the hierarchy. We need to decide when to merge clusters.

Definition: hierarchical clustering

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Hierarchical clustering

Hierarchical clustering* is a clustering method that seeks to build a hierarchy of clusters (clusters within clusters).

Two approaches:

- Bottom-up approach (agglomerative*): Each observation starts in its own cluster, and pairs of clusters are merged as one moves up the hierarchy. We need to decide when to merge clusters.
- Top-down approach (divisive*): All observations start in one cluster, and splits are performed recursively as one moves down the hierarchy. We need to decide when to split clusters.

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To decide when to combine (agglomerative) or when to split (divisive), we need:

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To decide when to combine (agglomerative) or when to split (divisive), we need:

A measure of dissimilarity between observations

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To decide when to combine (agglomerative) or when to split (divisive), we need:

- A measure of dissimilarity between *observations*
- A measure of dissimilarity between *sets* of observations.

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Linkage criterion

A **linkage criterion*** is a function of the pairwise distances of observations to measure dissimilarity of sets.

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Linkage criterion

A **linkage criterion*** is a function of the pairwise distances of observations to measure dissimilarity of sets.

Among the most popular,

- 1 Ward
- 2 Complete
- 3 Average
- 4 Single

Cluster linkage: Ward

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Ward linkage

The **Ward linkage criterion** minimizes the sum of squared differences within all clusters: it measures how much variance is added to the clustering algorithm if the clusters were to be merged.

Cluster linkage: Ward

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Ward linkage

The **Ward linkage criterion** minimizes the sum of squared differences within all clusters: it measures how much variance is added to the clustering algorithm if the clusters were to be merged.

$$W(A, B) = \sum_{x \in A \cup B} ||x - \mu_{A \cup B}||^2 - \sum_{x \in A} ||x - \mu_{A}||^2 - \sum_{x \in B} ||x - \mu_{B}||^2$$
$$= \frac{|A| \times |B|}{|A \cup B|} ||\mu_{A} - \mu_{B}||^2$$

Cluster linkage: Complete

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Complete linkage

The **complete linkage criterion** minimizes the maximum distance between observations of pairs of clusters.

Cluster linkage: Complete

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Complete linkage

The **complete linkage criterion** minimizes the maximum distance between observations of pairs of clusters.

Given a dissimilarity d and two sets A and B,

$$C(A, B) = max_{A \in A, b \in B} d(a, b)$$

Cluster linkage: Single

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Single linkage

The **single linkage criterion** minimizes the distance between the closest observations of pairs of clusters.

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Single linkage

The **single linkage criterion** minimizes the distance between the closest observations of pairs of clusters.

Given a dissimilarity d and two sets A and B,

$$S(A,B) = min_{a \in A, b \in B} d(a,b)$$

Cluster linkage: Average

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Average linkage

The average linkage criterion minimizes the average of the distances between all observations of pairs of clusters.

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Average linkage

The **average linkage criterion** minimizes the average of the distances between all observations of pairs of clusters.

Given a dissimilarity d and two sets A and B,

$$A(A,B) = \frac{1}{|A| \times |B|} \sum_{a \in A} \sum_{b \in B} d(a,b)$$

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Question

Compute the linkage criterion of ward, complete, average, single between the following sets of \mathbb{R}^2 vectors using Manhattan distance:

- A = {[1, 2], [2, 3], [4, 5]}
- B = {[3, 1], [4, 5], [1, 5]}

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For a selected linkage criterion L and dissimilarity d,

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For a selected linkage criterion L and dissimilarity d,

1 Initially put all individuals in their own cluster

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For a selected linkage criterion L and dissimilarity d,

- 1 Initially put all individuals in their own cluster
- 2 Merge together closest elements

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Advantage

For a selected linkage criterion L and dissimilarity d,

- Initially put all individuals in their own cluster
- 2 Merge together closest elements
- 3 Until every elements are in the same cluster: Recursively merge sets of elements that minimize the linkage criterion

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Advantages and limits For a selected linkage criterion L and dissimilarity d,

- 1 Initially put all individuals in their own cluster
- 2 Merge together closest elements
- 3 Until every elements are in the same cluster: Recursively merge sets of elements that minimize the linkage criterion

Clusters are assigned by cutting the algorithm at a selected value for the Linkage criterion.

Example

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Question

Using single linkage criterion and manhattan distance, perform Hierarchical Agglomerative Clustering on the following dataset. Test different cut-off values for final clusters.

Sepal length	Sepal width	Petal length
5	2	1
4	3	1
6	5	4
7	1	5

Dendrograms

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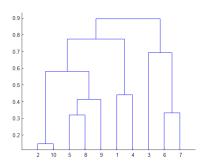
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Dendrograms

A **dendrogram*** is a diagram representing a tree. In the case of hierarchical clustering, it consists in representing graphically the clusters and their linkage criterion.



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Advantages and limits The dendrogram is used to infer information regarding the clustering process:

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The dendrogram is used to infer information regarding the clustering process:

Easily visualize elements grouped together

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The dendrogram is used to infer information regarding the clustering process:

- Easily visualize elements grouped together
- Easily select the number of clusters by setting the cut-off

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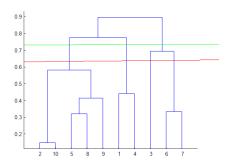
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The dendrogram is used to infer information regarding the clustering process:

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- Easily select the number of clusters by setting the cut-off



Selecting the right number of clusters

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Selecting the right number of clusters is again a problem...

Selecting the right number of clusters

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Selecting the right number of clusters is again a problem...

- The dendogram can show splits that "make sense"
- Plot clustering quality scores vs the number of clusters

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Advantages

- Easy to interprete even for people outside our field
- No need to perform the computations again when changing the number of clusters

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Advantages

- Easy to interprete even for people outside our field
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Advantages

- Easy to interprete even for people outside our field
- No need to perform the computations again when changing the number of clusters

Limits

Number of clusters has to be inferred

Questions

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Questions ?