Hierarchical Clustering

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MACS 40800: Unsupervised Machine Learning

October 17, 2019

Lecture Outline

- Quick Review
- 2 Hierarchical Clustering
- 3 Linkage Methods
- 4 Dendrograms & Tree Cutting
- 5 Divisive Hierarchical Clustering
- 6 Demonstration in R
- Due Dates

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- Key difference between clustering approaches: subdividing the data
 - ▶ Hierarchical ~→ No
 - ▶ Partitioning ~> Yes

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 - Check validation, especially if comparing across clustering algorithms (more next week)

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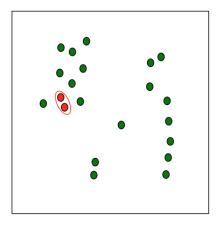
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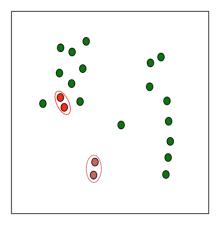
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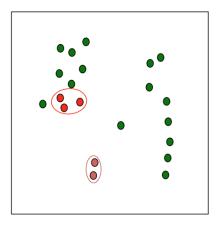
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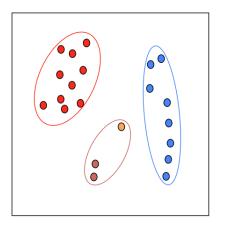
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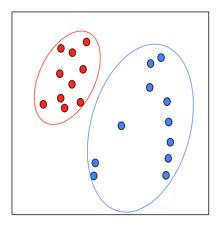
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- Key terms and necessary ingredients: distance measure, linkage methods, dendrogram, tree-cutting

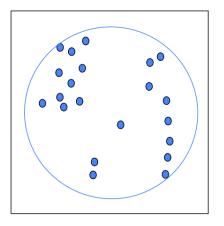












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- Stop when we reach k clusters (k = 1 in agglomerative; k = n in divisive)

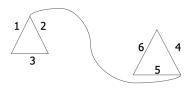
Inter-Cluster Dissimilarity

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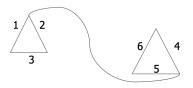
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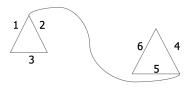


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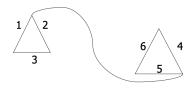
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- So how do we determine what constrains cluster fusion...?

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- There are five common types of linkage: complete, single, Ward's method, average, and centroid

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 Ward's linkage method joins the two clusters whose fusion is constrained by the smallest increase in SSE calculated per cluster, C,

$$\sum_{i=1}^n (x_i - \bar{x})^2$$

Average linkage uses the mean inter-cluster dissimilarity,

$$d_{average}(C_x, C_y) = \frac{\sum_i \sum_j d_{ij}}{N_{C_x} N_{C_y}}$$

where, d_{ij} is the pairwise distance between observations i and j, and N_* is the total number of observations in the computed cluster, C

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 pairwise calculations; centroid
 → fuses across centroids, which are
 intra-cluster averages

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 - Ward's method is based on minimizing the "loss of information" from joining two groups

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- Therefore, the input for a hierarchical clustering algorithm is an N × N distance matrix, from which inter-cluster distances are calculated via the selected linkage method

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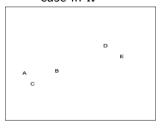
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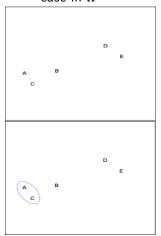
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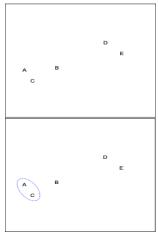
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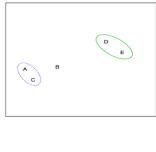
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- There is only one value-axis: the Y axis is the measured distance
- So we can get clear clustering when branches along the Y axis are long (suggesting greater distance from other clusters), and less obvious clustering when the branches are shorter

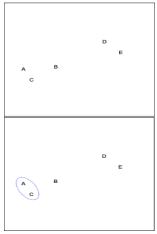
 Let's consider another simple toy example, and then move to a real case in R

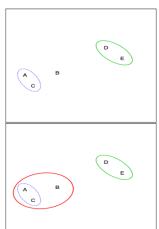


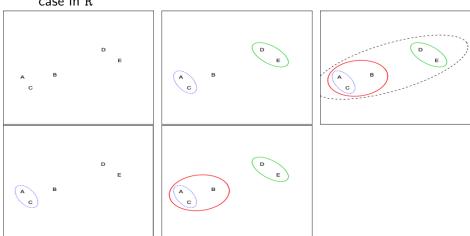


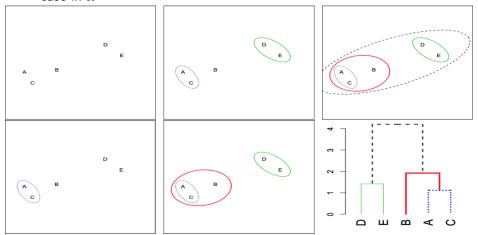












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- Good to note: HC is pretty computationally inefficient and expensive, especially on large datasets

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- We then partition the cluster into two least similar clusters, of all possible split values
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- This is significantly more expensive even than agglomerative, given the many split calculations required at each split (hence its less popular)

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 ${\tt HCA_demo.R}$

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- Take off a bit early and use the time while you're together to polish your proposals