

Clustering: Basics & Hierarchical Clustering

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MACS 40500: Computational Methods for American Politics

October 22, 2019

Lecture Outline

- 1 Clustering Basics
- 2 Diagnosing Clusterability
- 3 Conceptualizing and Calculating Distance
- 4 Hierarchical Clustering
- 5 Linkage Methods
- 6 Dendrograms & Tree Cutting
- 7 Divisive Hierarchical Clustering
- 8 Coming Up

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 - ▶ **Clustering**: the group labels are not known a priori
 - ▶ **Classification**: the group labels are known for a trained sample (next week)
- Thus, the typical goal in clustering is to discover the “natural groupings” present in the data

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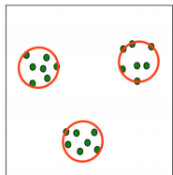
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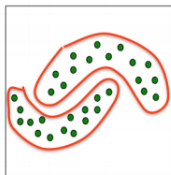
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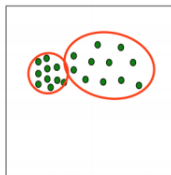
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- *Note:* Next class, also application on presidential vote shares by state in R

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- Importantly, note that with these approaches to assessing clusterability, as with virtually all of clustering, there is great ambiguity
- We can do our best to make sense of a complex feature space, but as our data are unlabeled, we are essentially always “guessing” about what these patterns are revealing
- This is a limitation of UML across the board; yet simultaneously the reason it is so important to combine UML with other data reduction and modeling processes

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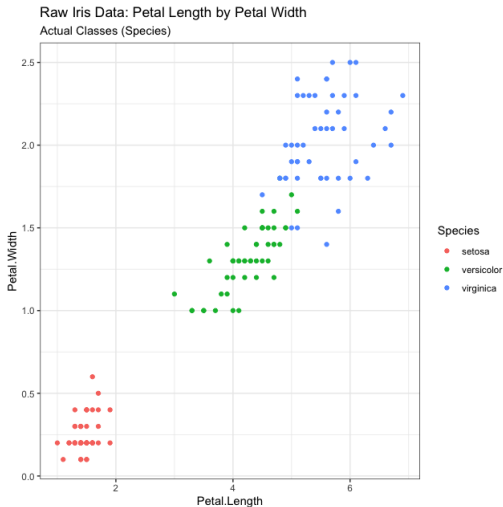
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 - ▶ Sepal Length
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 - ▶ 3 Species: setosa, versicolor, and virginica
 - ▶ 150 observations (50 of each)

Diagnosing Clusterability: Informally (Sepal Length by Sepal Width)



Diagnosing Clusterability: Informally (Petal Length by Petal Width)



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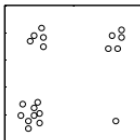
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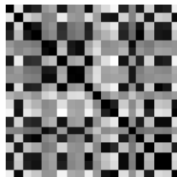
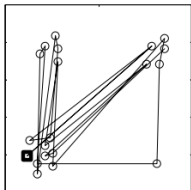
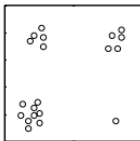
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- The visual result becomes darker blocks along the diagonal reflect greater spatial similarity, compared to lighter shaded blocks, which inversely suggest greater dissimilarity

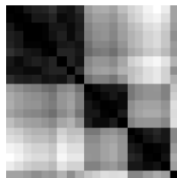
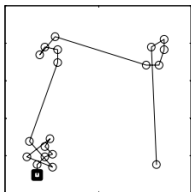
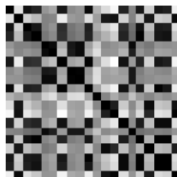
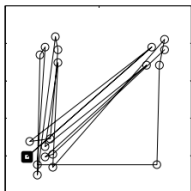
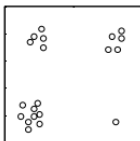
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VAT (ODI): Iris Data

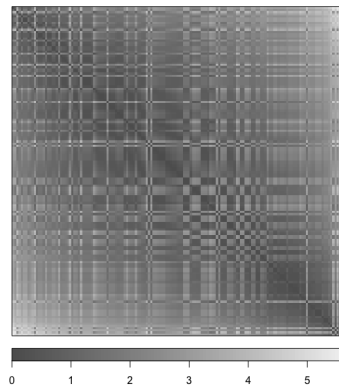


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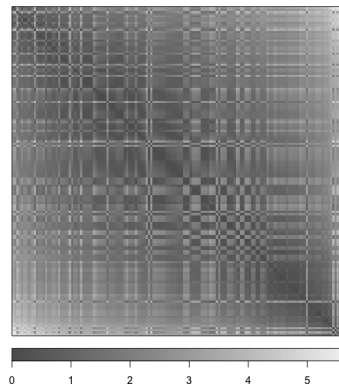


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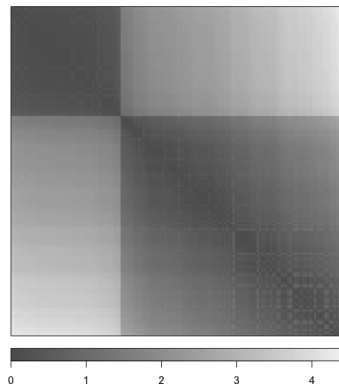


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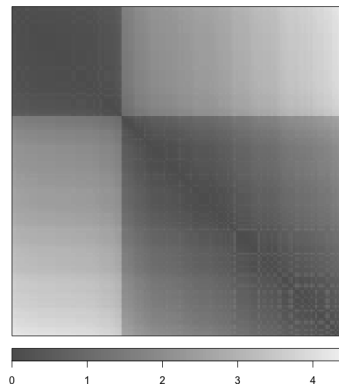


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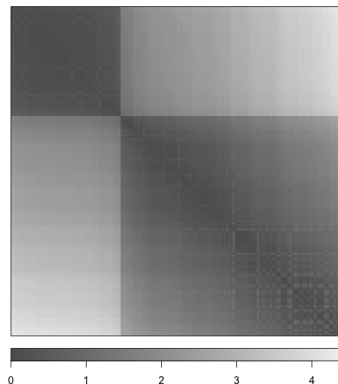


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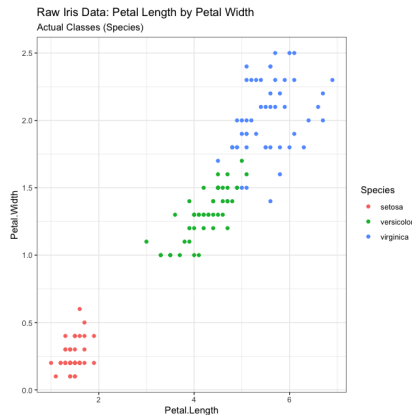


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- The Hopkins (or “H”) statistic tests the the null hypothesis of spatial randomness in the data using a sparse sampling test
- It calculates the probability that a given dataset is generated by a uniform (random noise, with no clusters) distribution or not (non-random, with clustering likely)
- This general procedure is called **sparse sampling**

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- **Goal**: calculate the pairwise dissimilarity across all observations in the **actual** data is compared to a set of **simulated** dataset drawn from some random distribution (usually uniform) with the *same* standard deviation as the original data
- In the form of a question: is the actual data random, compared to the synthetic data set, which we *know* is random?

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- In general, $H > 0.5$ leads to rejection of H_0 , suggesting the data are non-random, and are “clusterable”

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- By contrast, *features* are usually grouped on the basis of correlation coefficients (but so can observations)

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- Suppose you had two features on which you wanted to cluster units: weight (lbs.) and household income (dollars)
- In such a case, different units of measurement *and* distributions (skewed) will always return biased results
- Thus first, always standardize input features to ensure they are “unitless” (commonly setting $\mu = 0$ and $\sigma = 1$)

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- Generally a distance measure $d(p, q)$ between two points p and q satisfies the following properties, where g is any other intermediate point:

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 - ▶ Respondents on large surveys
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 - ▶ Economies
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 - ▶ Perceptual studies
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 - ▶ And, on Friday, partisan voting by state
 - ▶ And, on your HW, state legislative professionalism

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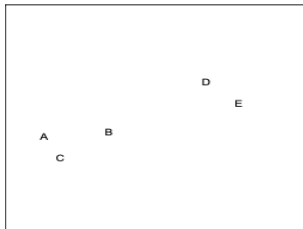
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- Key terms: distance measure, linkage methods, dendrogram, tree-cutting

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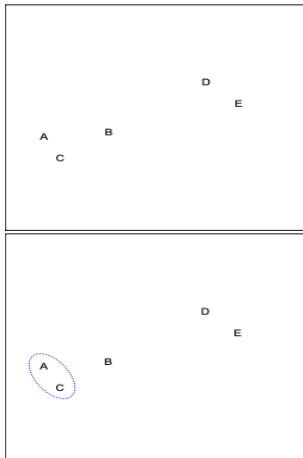
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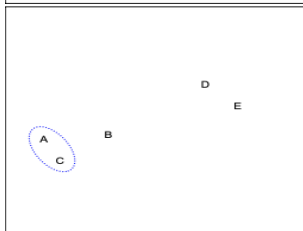
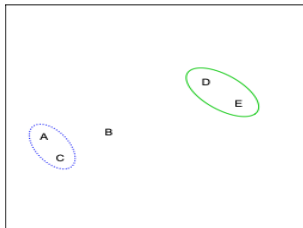
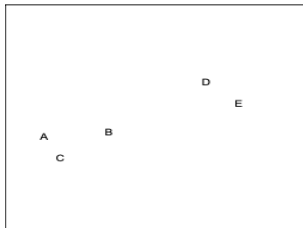
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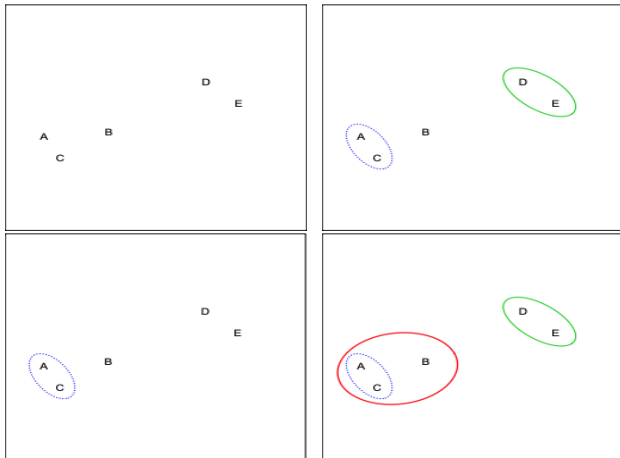
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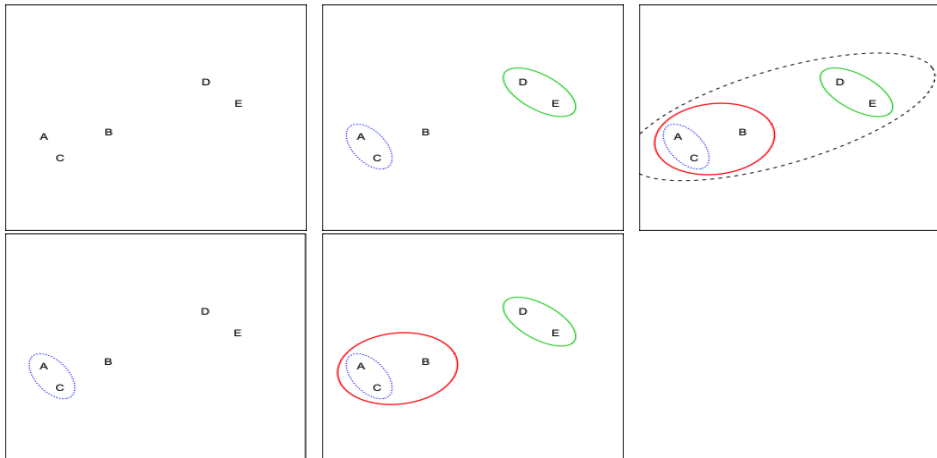
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- Stop when we reach k clusters ($k = 1$ in agglomerative; $k = n$ in divisive)

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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 ,

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

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- **Average** linkage uses the *mean* inter-cluster dissimilarity,

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

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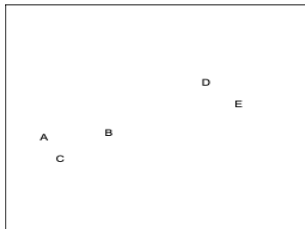
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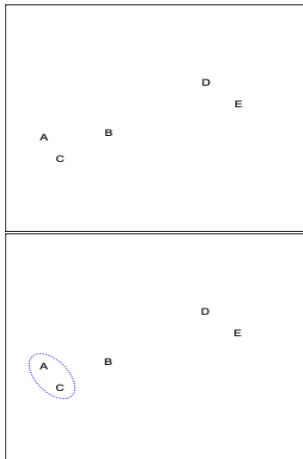
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- So we can get clear clustering when branches along the Y axis are long (suggesting greater distance from other clusters), and less obvious clustering when the branches are shorter

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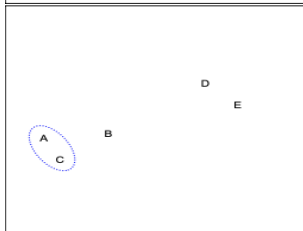
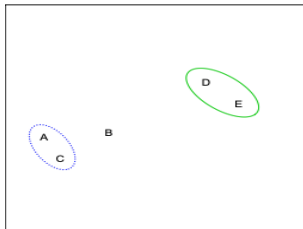
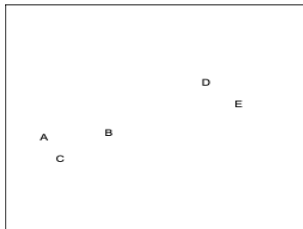
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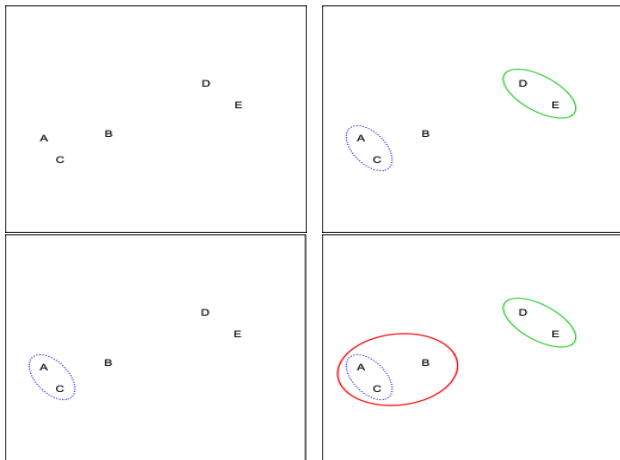
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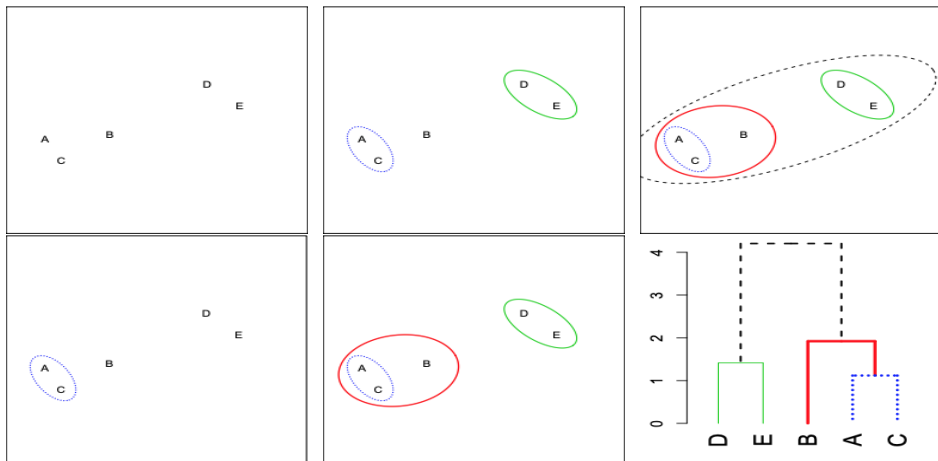
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- Note: HC is computationally inefficient and expensive, especially on large datasets

Lecture Outline

- 1 Clustering Basics
- 2 Diagnosing Clusterability
- 3 Conceptualizing and Calculating Distance
- 4 Hierarchical Clustering
- 5 Linkage Methods
- 6 Dendrograms & Tree Cutting
- 7 Divisive Hierarchical Clustering**
- 8 Coming Up

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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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- Demonstration in R: 2012 state Democratic vote shares

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- Gaussian mixture models
- Demonstration in R: 2012 state Democratic vote shares
- First problem set due **Friday at 5 pm** to our GH repo (quick word on the data and concept)