k-means, hierarchical, and spectral clustering

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k-means clustering

- Partition data into k non-overlapping clusters
- Must specify k
- k-means algorithm creates k clusters with smallest total within-cluster variance
- Thus Euclidean distance measures similarity
- This difficult combinatorial problem, but iterative algorithm does pretty good

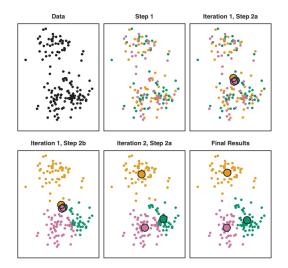
k-means algorithm

Algorithm 10.1 K-Means Clustering

- 1. Randomly assign a number, from 1 to K, to each of the observations. These serve as initial cluster assignments for the observations.
- 2. Iterate until the cluster assignments stop changing:
 - (a) For each of the K clusters, compute the cluster *centroid*. The kth cluster centroid is the vector of the p feature means for the observations in the kth cluster.
 - (b) Assign each observation to the cluster whose centroid is closest (where *closest* is defined using Euclidean distance).

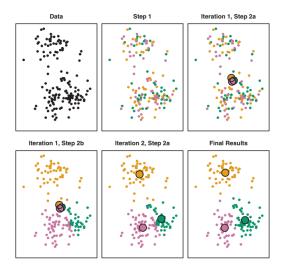
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k-means algorithm



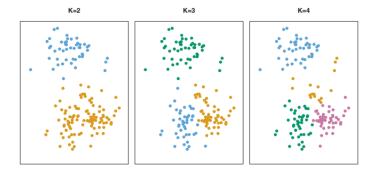
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k-means starting values



Need to do k-means several times, pick best

k-means value of \boldsymbol{k}



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k-means value of \boldsymbol{k}

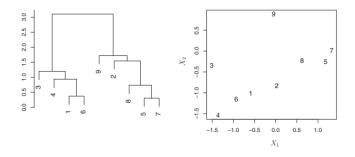
- ► As k increases total within-cluster variance decreases
- Use domain-specific considerations
- Use the elbow rule if no outside info

Hierarchical clustering

- Generates a sequence of clusters
- Split or merging clusters at each step
- Does not require prespecification of k
- ► Has tree-based representation: *dendrogram*
- Top-down and bottom-up (agglomerative) versions
- ► Top-down: start with 1 big cluster, split until N clusters
- ▶ Bottom-up: start with N clusters, merge until 1 cluster

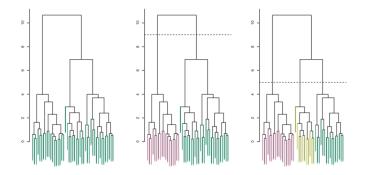
Bottom-up most common

Hierarchical clustering: dendrogram



- Leaf at bottom is single observation
- Most similar observations merged to form cluster
- Most similar clusters merged to form new clusters
- Vertical axis is dissimilarity of merged clusters

Hierarchical clustering: dendrogram clusters



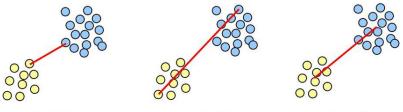
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- Cut the dendrogram to make clusters
- ► Where you cut determines *k*
- Can apply elbow rule to dendrogram

Hierarchical clustering: dissimilarity and linkage

- Must define dissimilarity between observations
- Euclidean distance is common
- Must define dissimilarity or *linkage* between clusters:
- Complete Maximum intercluster dissimilarity
- ► Single Minimal intercluster dissimilarity
- Average Mean intercluster dissimilarity
- Centroid Dissimilarity between centroids
- Different dissimilarity and linkage results in different dendrograms
- Can try a variety and see if patterns consistently emerge

Hierarchical clustering: linkage



single-link

complete-link

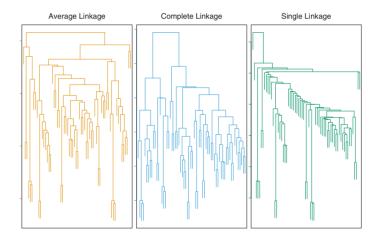
average-link

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Hierarchical clustering: linkage



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Hierarchical clustering: bottom-up algorithm

Algorithm 10.2 Hierarchical Clustering

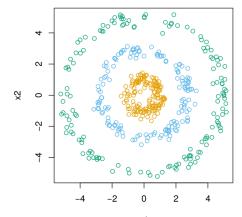
- 1. Begin with n observations and a measure (such as Euclidean distance) of all the $\binom{n}{2} = n(n-1)/2$ pairwise dissimilarities. Treat each observation as its own cluster.
- 2. For $i = n, n 1, \dots, 2$:
 - (a) Examine all pairwise inter-cluster dissimilarities among the *i* clusters and identify the pair of clusters that are least dissimilar (that is, most similar). Fuse these two clusters. The dissimilarity between these two clusters indicates the height in the dendrogram at which the fusion should be placed.
 - (b) Compute the new pairwise inter-cluster dissimilarities among the i-1 remaining clusters.

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- unsupervised learning
- useful for unusual clusters

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Toy example



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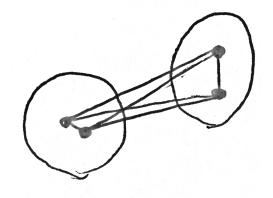
- k-means will fail
- normal mixtures might work
- hierarchical methods might work (linkage?)

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useful for unusual clusters

- main idea is to transform data, then cluster
- ▶ start with $N \times N$ matrix of pairwise similarities $s_{ii'} \ge 0$
- \blacktriangleright organize observations into graph with N nodes connected by edges with length $1/s_{ii'}$
- clustering is then a graph-partition problem; partition graph such that edges that exit clusters have long lengths (small similarity), and edges within clusters have short lengths (large similarity)

Graph partition problem



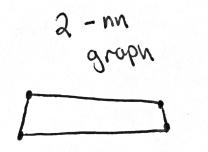
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• consider $x_i \in \mathbb{R}^p$

•
$$s_{ii'} = \exp(-||x_i - x_{i'}||^2/c)$$
 (radial kernel)

- many ways to encode graph based on s_{ii}
- ► e.g., fully connected
- e.g., K-nearest-neighbor graph
- define \mathcal{N}_K to be the set of nearby pairs of points
- obs (i, i') is in \mathcal{N}_K if i is K-nn of i' or vice versa
- only (i, i') in \mathcal{N}_K are connected with weight $w_{ii'} = s_{ii'}$, otherwise no edge $(w_{ii'} = 0)$.
- the K-nn graph is a kind of pre-processing; hard thresholding of similarities)

2-NN Graph



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- adjacency matrix of weights (similarities; $N \times N$): $W = \{w_{ii'}\}$
- degree of node i: $g_i = \sum_{i'} w_{ii'}$
- \blacktriangleright let $G = \operatorname{diag}(q_1, \ldots, q_N)$
- define graph Laplacian:

$$L = G - W$$



- spectral clustering:
 - 1. find m eigenvectors Z $(N \times m)$ corresponding to the m **smallest** eigenvalues of L (ignoring trivial constant eigenvector)
 - 2. use standard clustering method to cluster rows of Z, which serve as the transformed data

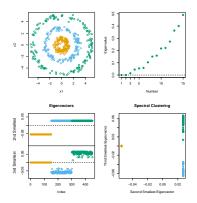


FIGURE 14.29. Toy example illustrating spectral clustering. Data in top left are 450 points falling in three concentric clusters of 150 points each. The points are uniformly distributed in angle, with radius 1,2.8 and 5 in the three groups, and Gaussian noise with standard deviation 0.25 added to each point. Using a k = 10 nearest-neighbor similarity graph, the eigenvector corresponding to the second and third smallest

Spectral clustering: Why it works.

- ► consider tranforming (N × p) data X into (N × 1) vector f for the purpose of clustering
- ▶ for items that are similar (large $s_{ii'}$ and $w_{ii'}$), f_i should have a similar value to $f_{i'}$ and vice versa
- ► if we consider the following quantity

$${}^{T}Lf = \sum_{i} g_{i}f_{i}^{2} - \sum_{i} \sum_{i'} w_{ii'}f_{i}f_{i'}$$
$$= \sum_{i} \sum_{i'} w_{ii'}f_{i}^{2} - \sum_{i} \sum_{i'} w_{ii'}f_{i}f_{i'}$$
$$= \frac{1}{2} \sum_{i} \sum_{i'} w_{ii'}(f_{i} - f_{i'})^{2}$$

- finding f_i similar to $f_{i'}$ for large $w_{ii'}$ implies small $f^T L f$
- now if restrict f to be an eigenvector of L, then f^TLf is the eigenvalue, and we should focus on those vectors with smallest values
- constant vector (1^T) is eigenvector of L with eigenvalue zero