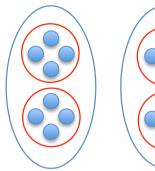
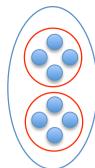
# IAML: Hierarchical Clustering

Victor Lavrenko and Nigel Goddard School of Informatics

Semester 1

## Hierarchical clustering

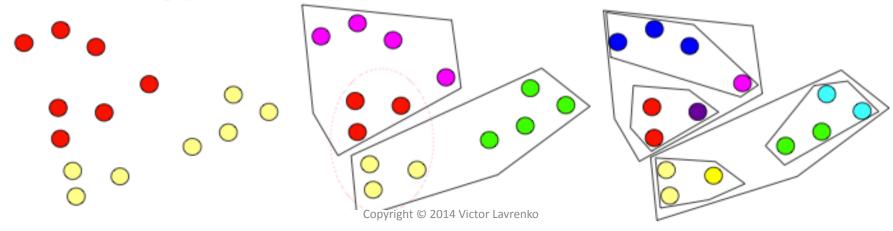




- Selecting K question of granularity
  - how coarse or fine-grained is the structure in your data?
    - analogy: tidal waves or ripples on the surface?
    - real data: both, and probably everything in-between
  - no clustering algorithm able to pick K (some claim to)
- Instead of picking K find a hierarchy of structure
  - top levels coarse effects, low levels fine-grained
    - topmost cluster contains every point in the dataset
    - bottom level set of n singleton clusters, one per data point
  - strategies:
    - top-down: start with all items in one cluster, split recursively
    - bottom-up: start with singletons, merge by some criterion

#### Hierarchical K-means

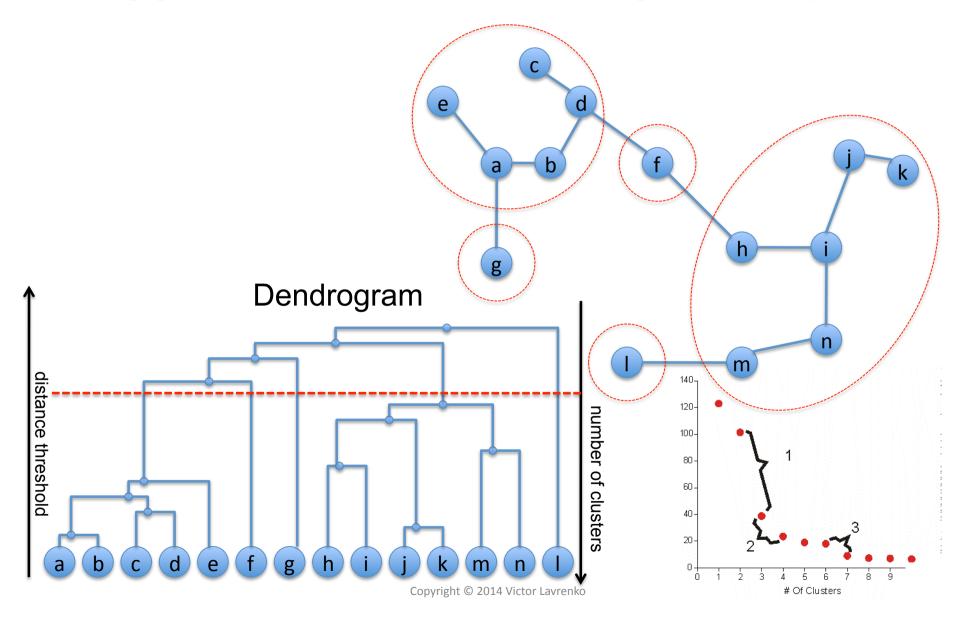
- Top-down approach:
  - run K-means algorithm on the original data x₁...x<sub>n</sub>
  - for each of the resulting clusters c<sub>i</sub>: i = 1 ... K
    - recursively run K-means on points in c<sub>i</sub>
- Fast: recursive calls operate on a slice: O(Knd log<sub>K</sub>n)
- Greedy: can't cross boundaries imposed by top levels
  - nearby points may end up in different clusters



## Agglomerative clustering

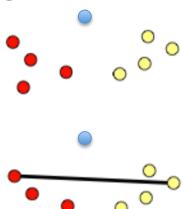
- Idea: ensure nearby points end up in the same cluster
- Start with a collection C of n singleton clusters
  - each cluster contains one data point: c<sub>i</sub>={x<sub>i</sub>}
- Repeat until only one cluster is left:
  - find a pair of clusters that is closest:  $\min_{i,j} D(c_i, c_j)$
  - merge the clusters c<sub>i</sub>, c<sub>j</sub> into a new cluster c<sub>i+j</sub>
  - remove c<sub>i</sub>,c<sub>j</sub> from the collection C, add c<sub>i+j</sub>
- Produces a dendrogram: hierarchical tree of clusters
- Need to define a distance metric over clusters
- Slow: O(n<sup>2</sup>d + n<sup>3</sup>) create, traverse distance matrix

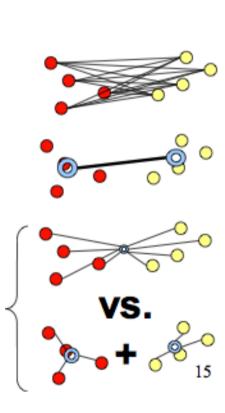
# Agglomerative clustering: example



### Cluster distance measures

- Single link:  $D(c_1, c_2) = \min_{x_1 \in c_1, x_2 \in c_2} D(x_1, x_2)$ 
  - distance between closest elements in clusters
  - produces long chains a→b→c→...→z
- Complete link:  $D(c_1, c_2) = \max_{x_1 \in c_1, x_2 \in c_2} D(x_1, x_2)$ 
  - distance between farthest elements in clusters
  - forces "spherical" clusters with consistent "diameter"
- Average link:  $D(c_1,c_2) = \frac{1}{|c_1|} \frac{1}{|c_2|} \sum_{x_1 \in c_1} \sum_{x_2 \in c_2} D(x_1,x_2)$ 
  - average of all pairwise distances
  - less affected by outliers
- Centroids:  $D(c_1, c_2) = D\left(\left(\frac{1}{|c_1|} \sum_{x \in c_1} \vec{x}\right), \left(\frac{1}{|c_2|} \sum_{x \in c_2} \vec{x}\right)\right)$ 
  - distance between centroids (means) of two clusters
- Ward's method:  $TD_{c_1 \cup c_2} = \sum_{x \in c_1 \cup c_2} D(x, \mu_{c_1 \cup c_2})^2$ 
  - consider joining two clusters, how does it change the total distance (TD) from centroids?

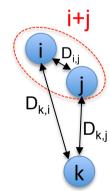




## Lance-Williams Algorithm

- D = {D<sub>i,j</sub>: distance between  $\mathbf{x}_i$  and  $\mathbf{x}_j$  for i,j=1..N}
- for N iterations:

i,j = arg min  $D_{i,j}$  ... pair of closest clusters add cluster: i+j, delete clusters i, j for each remaining cluster k:



$D_{k,i+i} =$	$\alpha_i D_{k,i}$	$+ \alpha_i D_{k,i}$	+ β D <sub>i,i</sub> -	+ γ   D <sub>k,i</sub>	$-D_{k,i}$
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Method	$lpha_i$	$lpha_i$	$oldsymbol{eta}$	$\gamma$
Single linkage	0.5	0.5	0	-0.5
Complete linkage	0.5	0.5	0	0.5
Group average	$\frac{n_i}{n_i + n_j}$	$rac{n_j}{n_i + n_j}$	0	0
Weighted group average	0.5	0.5	0	0
Centroid	$\frac{n_i}{n_i + n_j}$	$\frac{n_j}{n_i + n_j}$	$\frac{-n_i \cdot n_j}{(n_i + n_j)^2}$	0
Ward	$\frac{n_i + n_k}{(n_i + n_j + n_k)}$	$\frac{n_j \!+\! n_k}{(n_i \!+\! n_j \!+\! n_k)}$	$\frac{-n_k}{(n_i+n_j+n_k)}$	0

#### Single link:

$$D_{k,i+j} = \frac{1}{2} (D_{ki} + D_{kj} - |D_{ki} - D_{kj}|)$$
  
= min {D<sub>ki</sub>, D<sub>kj</sub>}

$$\min_{a,b} = \max_{a,b} - |a-b|$$

## Summary

- Clustering: discover underlying sub-populations
- K-means
  - fast, iterative, leads to a local minimum
  - need to pick k: look for unusual reduction in variance
- Mixture models
  - probabilistic version of K-means
  - Expectation Maximization (EM) algorithm
- Hierarchical clustering
  - top-down (K-means) and bottom-up (agglomerative)
  - single / complete / average link variations