DSC 257R - UNSUPERVISED LEARNING

HIERARCHICAL CLUSTERING

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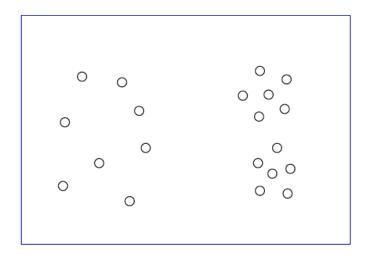


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Hierarchical Clustering

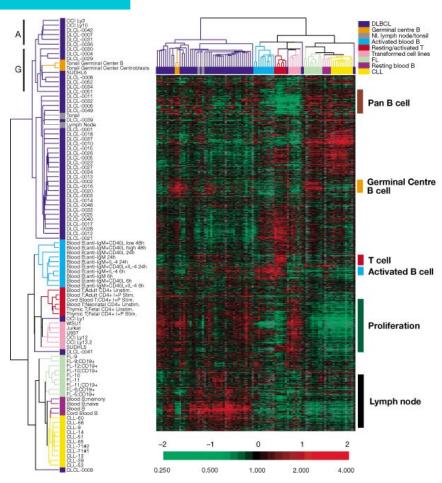
Choosing the number of clusters (k) is difficult.



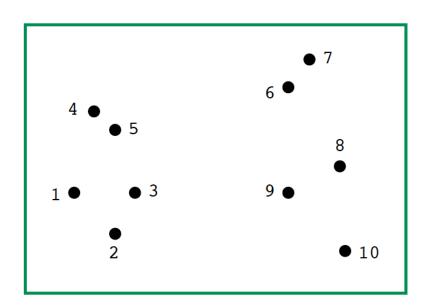
Often: no single right answer, because of multiscale structure.

Hierarchical clustering avoids these problems.

Example: Gene Expression Data



The Single Linkage Algorithm



- Start with each point in its own, singleton, cluster
- Repeat until there is just one cluster:
 - Merge the two clusters with the closest pair of points
- Disregard singleton clusters

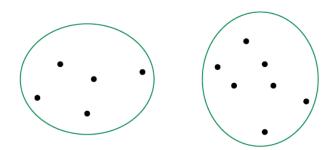
Linkage Methods

- Start with each point in its own, singleton, cluster
- Repeat until there is just one cluster:
 - Merge the two "closest" clusters

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How to measure distance between two clusters C and C'?



Single linkage

$$\operatorname{dist}(C, C') = \min_{x \in C, x' \in C'} ||x - x'||$$

Complete linkage

$$\operatorname{dist}(C, C') = \max_{x \in C, x' \in C'} ||x - x'||$$

Average Linkage

Three commonly-used variants:

1 Average pairwise distance between points in the two clusters

$$dist(C, C') = \frac{1}{|C| \cdot |C'|} \sum_{x \in C} \sum_{x' \in C'} ||x - x'||$$

2 Distance between cluster centers

$$dist(C, C') = ||mean(C) - mean(C')||$$

 $oldsymbol{3}$ Ward's method: the increase in k-means cost occasioned by merging the two clusters

$$dist(C, C') = \frac{|C| \cdot |C'|}{|C| + |C'|} \|mean(C) - mean(C')\|^2$$