

# Hierarchical clustering

Hongning Wang

CS@UVa

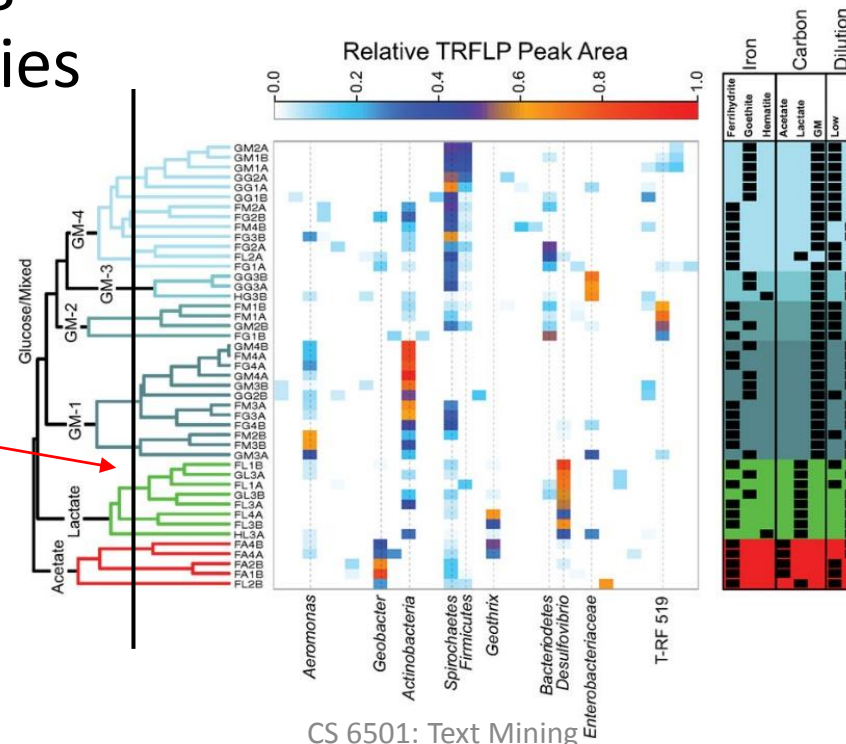
# Today's lecture

- Hierarchical clustering algorithm
  - Bottom-up: agglomerative
  - Distance between clusters
  - Complexity analysis

# Hierarchical clustering















- Build a tree-based hierarchical taxonomy from a set of instances
  - Dendrogram – a useful tool to summarize similarities

*After cutting, each connected component will be a cluster*



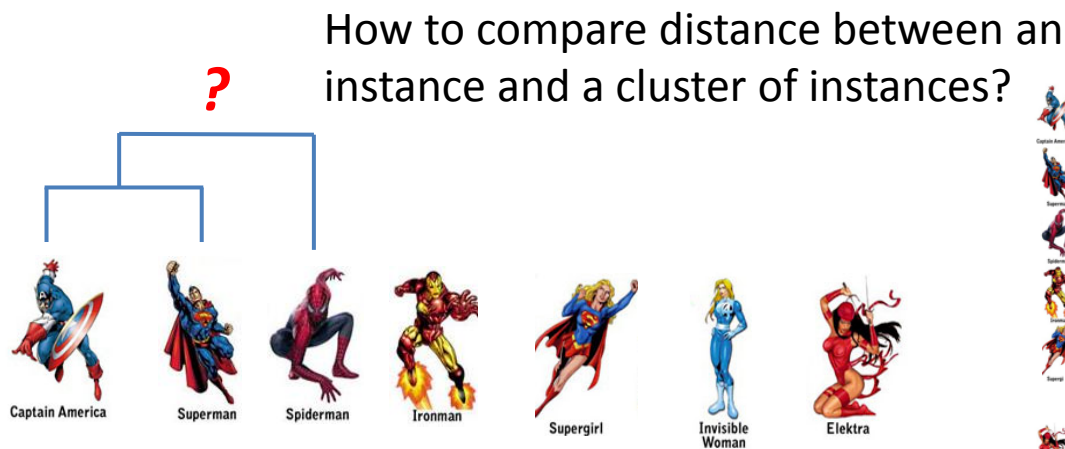
# Agglomerative hierarchical clustering








- Pairwise distance metric between instances

	 Captain America	 Superman	 Spiderman	 Ironman	 Supergirl	 Invisible Woman	 Elektra
 Captain America	0	1	2	2	3	3	4
 Superman		0	2	2	3	3	4
 Spiderman			0	2	3	3	4
 Ironman				0	3	4	4
 Supergirl					0	2	3
 Invisible Woman						0	2
 Elektra							0

# Agglomerative hierarchical clustering

1. Every instance is in its own cluster when initialized
2. Repeat until one cluster left *Enumerate all the possibilities!*
  1. Find the best pair of clusters to merge and break the tie arbitrarily



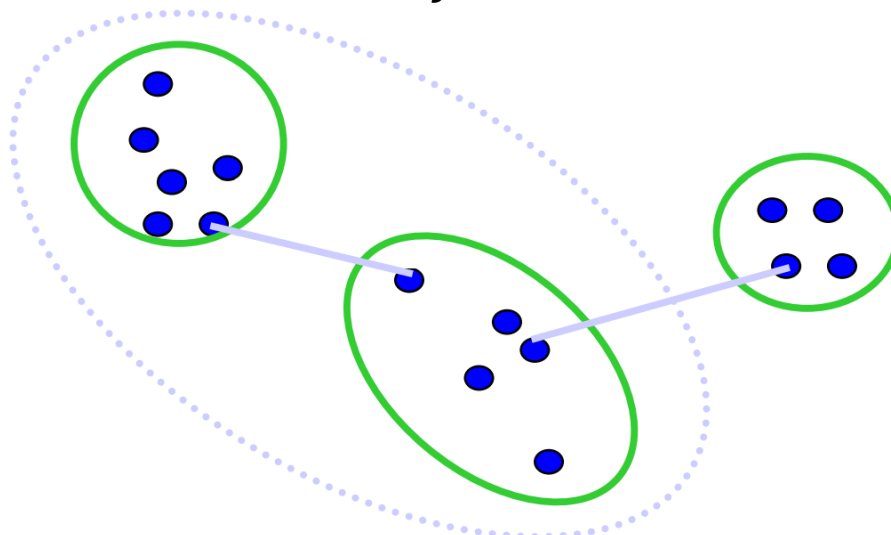
							
	0	1	2	2	3	3	4
		0	2	2	3	3	4
			0	2	3	3	4
				0	3	4	4
					0	2	3
						0	2
							0

# Distance measure between clusters

- Single link
  - Cluster distance = distance of two closest members between the clusters

$$- d(c_i, c_j) = \min_{x_n \in c_i, x_m \in c_j} d(x_n, x_m)$$

Tend to generate scattered clusters



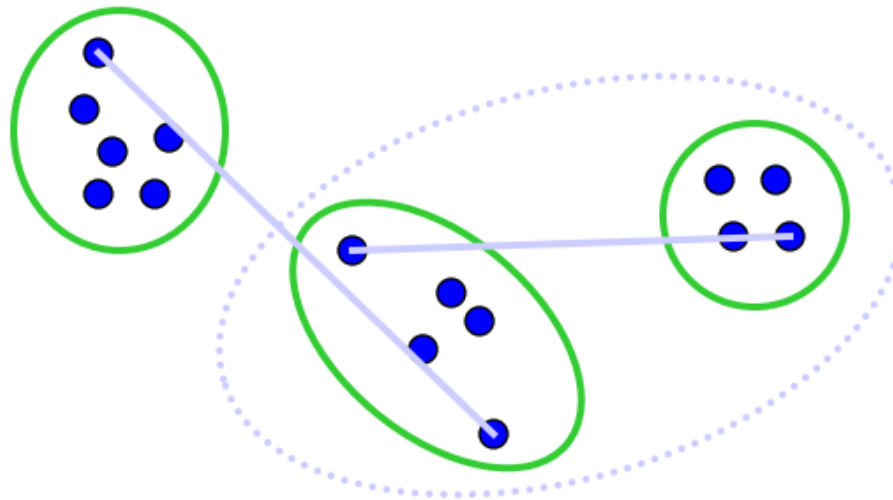
# Distance measure between clusters

- Complete link

- Cluster distance = distance of two farthest members between the clusters

- $d(c_i, c_j) = \max_{x_n \in c_i, x_m \in c_j} d(x_n, x_m)$

Tend to generate  
tight clusters

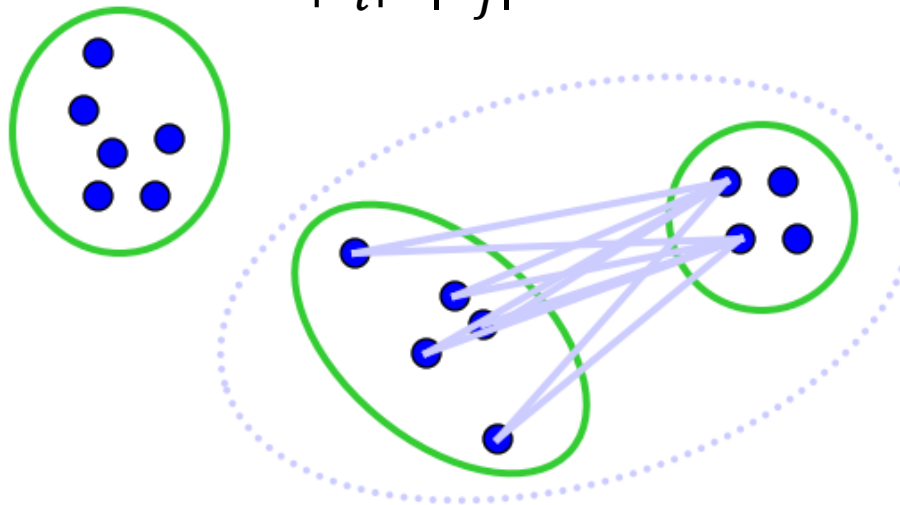


# Distance measure between clusters

- Average link
  - Cluster distance = average distance of all pairs of members between the clusters

$$- d(c_i, c_j) = \frac{\sum_{x_n \in c_i, x_m \in c_j} d(x_n, x_m)}{|c_i| \times |c_j|}$$

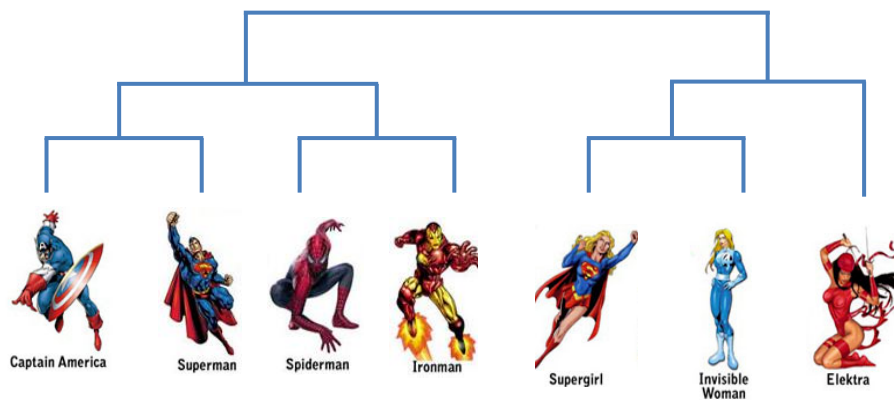
Mostly popularly used  
measure, robust  
against noise








# Agglomerative hierarchical clustering

1. Every instance is in its own cluster when initialized
2. Repeat until one cluster left
  1. Find the best pair of clusters to merge and break the tie arbitrarily



							
	0	1	2	2	3	3	4
		0	2	2	3	3	4
			0	2	3	3	4
				0	3	4	4
					0	2	3
						0	2
							0

# Complexity analysis

- In step one, compute similarity between all pairs of  $n$  individual instances -  $O(n^2)$
- In the following  $n - 2$  steps
  - It could be  $O(n^2 \log n)$  or even  $O(n^3)$  (naïve implementation)

In  $k$ -means, we have  $O(knl)$ ,  
a much faster algorithm

# Comparisons

- Hierarchical clustering
  - Efficiency:  $O(n^3)$ , slow
- Assumptions
  - No assumption
  - Only need distance metric
- Output
  - Dendrogram, a tree
- $k$ -means clustering
  - Efficiency:  $O(knl)$ , fast
- Assumptions
  - Strong assumption – centroid, latent cluster membership
  - Need to specify  $k$
- Output
  - $k$  clusters

# How to get final clusters?

- If  $k$  is specified, find a cut that generates  $k$  clusters
  - Since every time we only merge 2 clusters, such cut must exist
- If  $k$  is not specified, use the same strategy as in  $k$ -means
  - Cross validation with internal or external validation

# What you should know

- Agglomerative hierarchical clustering
  - Three types of linkage function
    - Single link, complete link and average link
  - Comparison with  $k$ -means

# Today's reading

- Introduction to Information Retrieval
  - Chapter 17: Hierarchical clustering
    - 17.1 Hierarchical agglomerative clustering
    - 17.2 Single-link and complete-link clustering
    - 17.3 Group-average agglomerative clustering
    - 17.5 Optimality of HAC