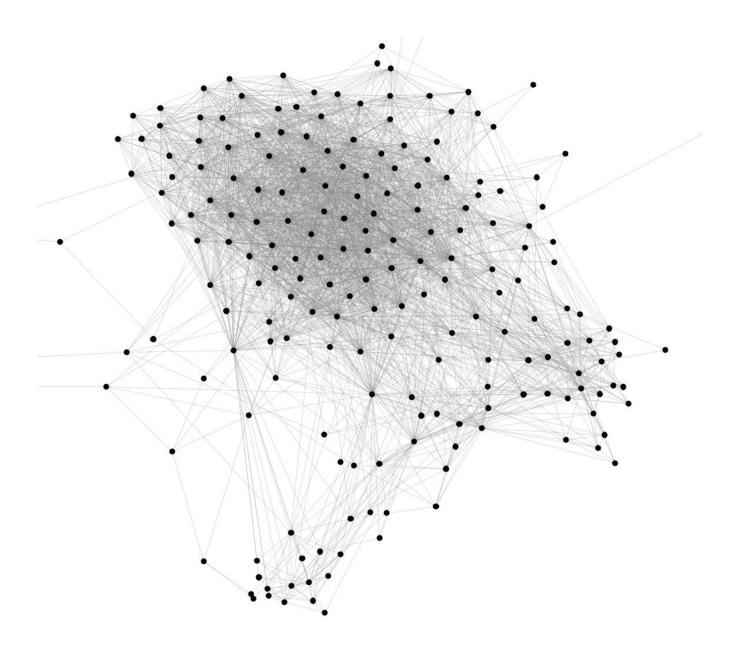
Online Social Networks and Media

Community detection



Introduction

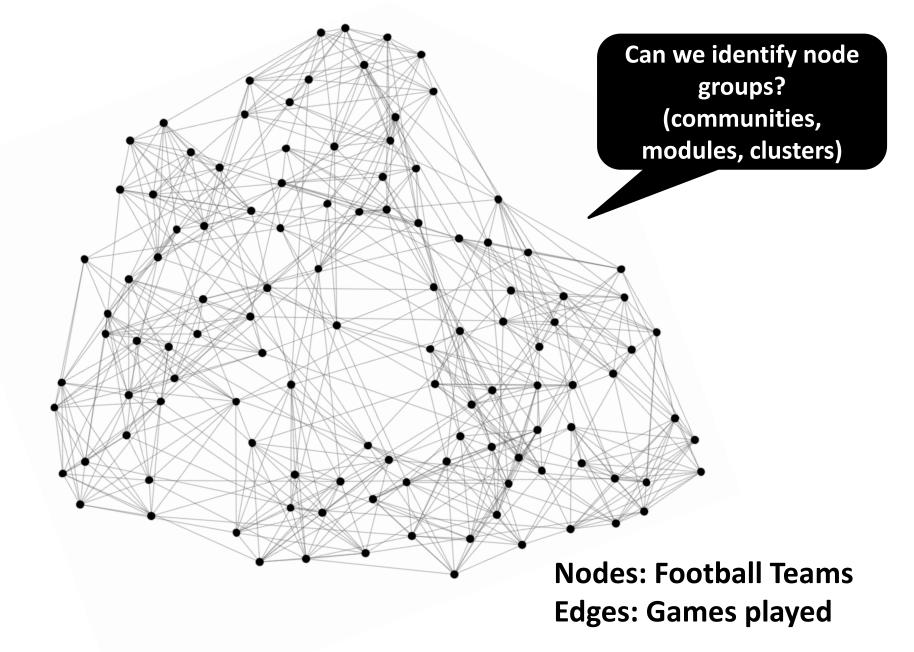
Real networks are *not random graphs*

Communities

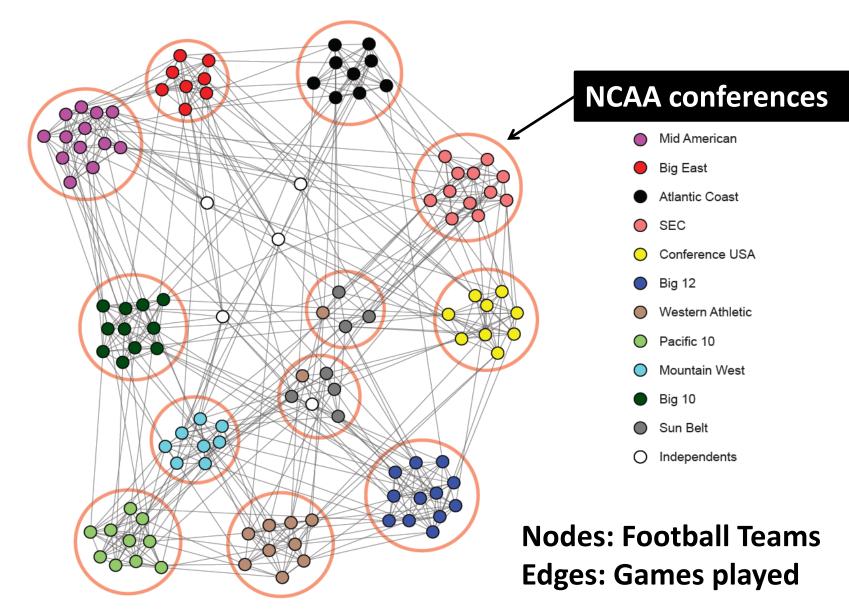
aka: groups, clusters, cohesive subgroups, modules

(informal) Definition: groups of vertices which probably share common properties and/or play similar roles within the graph

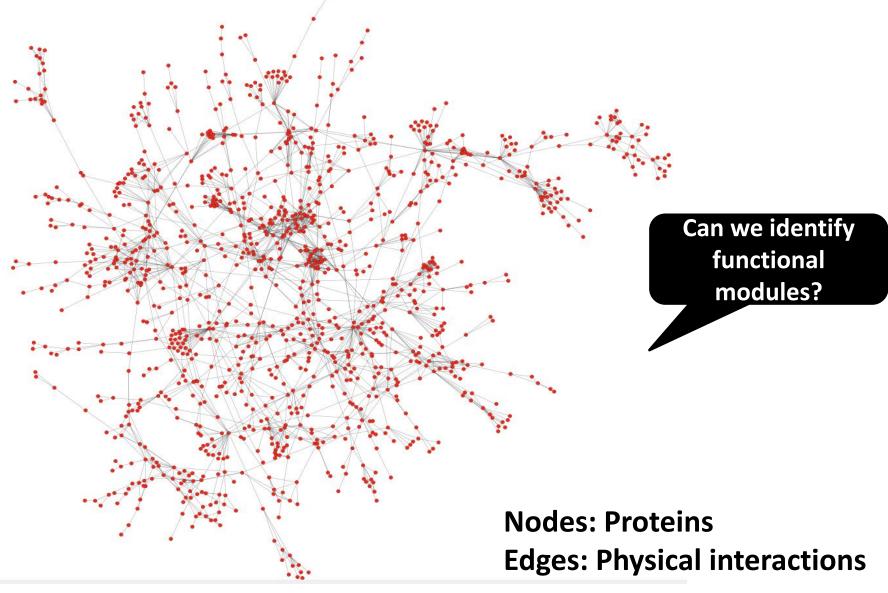
Some are *explicit* (*emic*) (e.g., Facebook (groups), LinkedIn (groups, associations), etc), we are interested in *implicit* (*etic*) ones



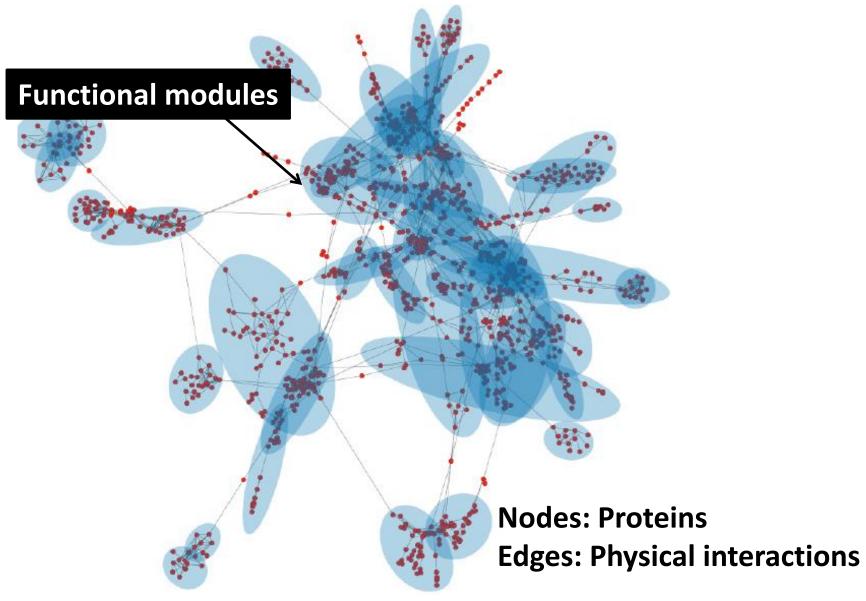
NCAA Football Network



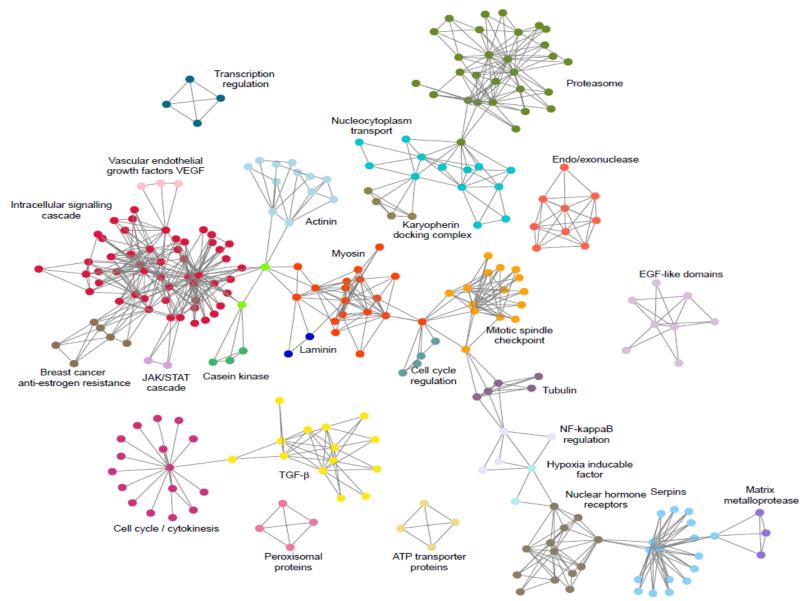
Protein-Protein Interactions



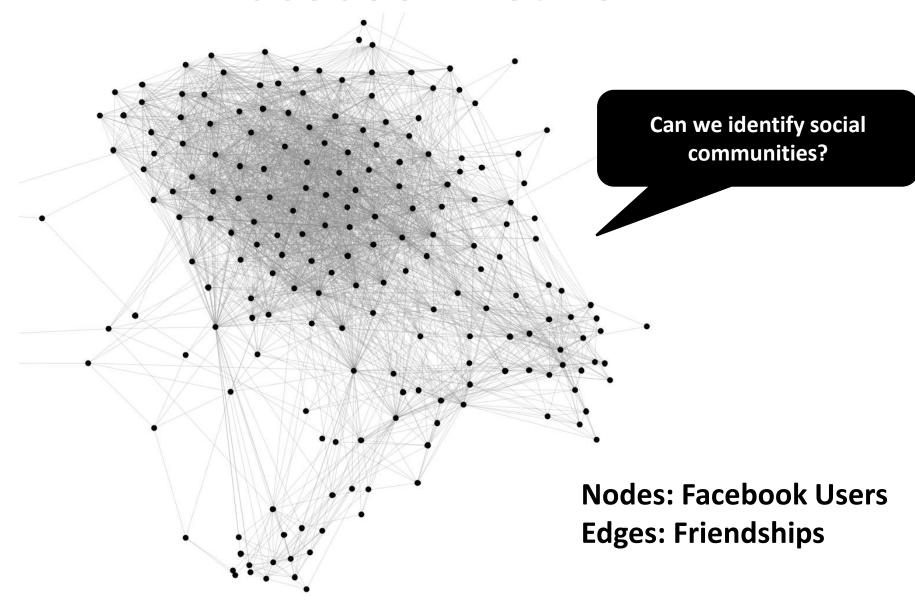
Protein-Protein Interactions



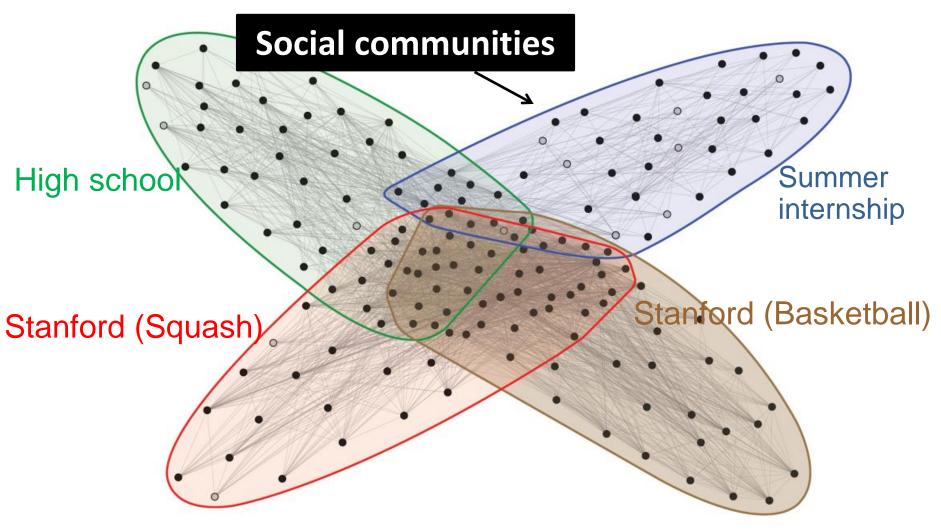
Protein-Protein Interactions



Facebook Network



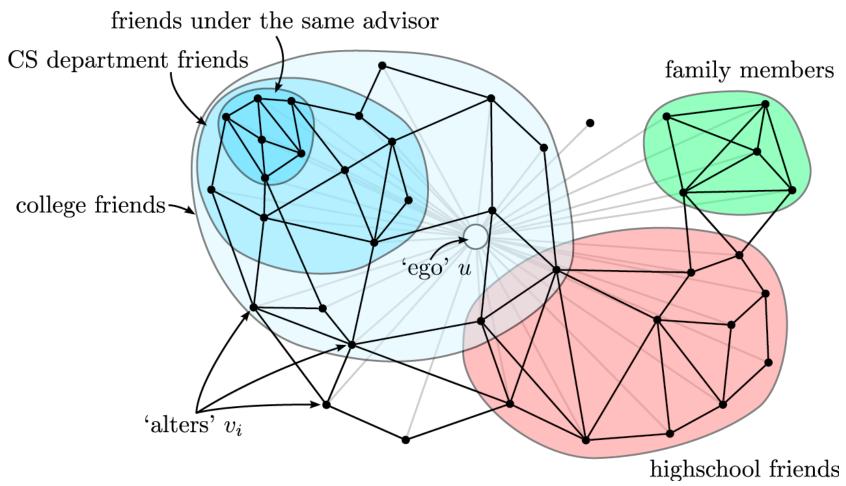
Facebook Network



Nodes: Facebook Users

Edges: Friendships

Twitter & Facebook



social circles, circles of trust

Outline

PART I

- 1. Introduction: what, why, types?
- 2. Cliques
- Background: How it relates to "cluster analysis" (node/edge similarity)
- 4. Betweeness centrality
- 5. Modularity, label propagation
- 6. How to evaluate

PART II (next lecture)

Cuts, Spectral clustering, Denser subgraphs

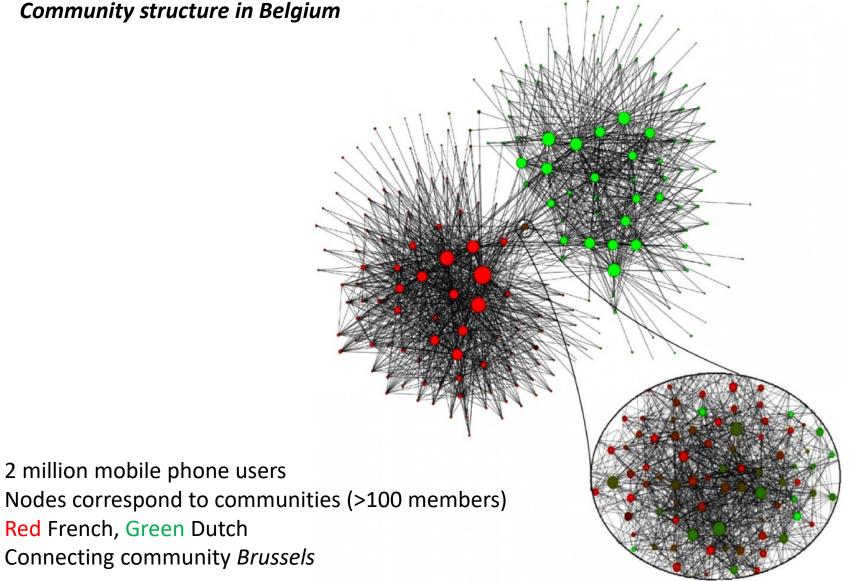
+ signed networks

Why? (some applications)

- Knowledge discovery
- Groups based on common interests, behavior, etc
 (e.g., Canadians who call USA, readings tastes, etc)
 - Recommendations, marketing
- Collective behavior observable at the group, not the individual level, local view is noisy and ad hoc
- Classification of the nodes by identifying modules and their boundaries
- To improve performance: partition a large graph into many machines, assigning web clients to web servers, routing in ad hoc networks, etc
- Summary, visual representation of the graph

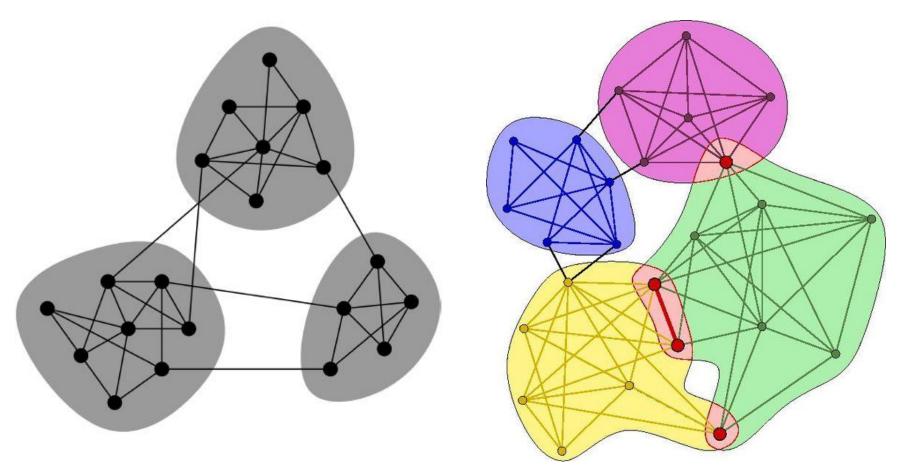
Example: communities in Belgium

59% Flemish, speaking Dutch 40% Walloons speaking French



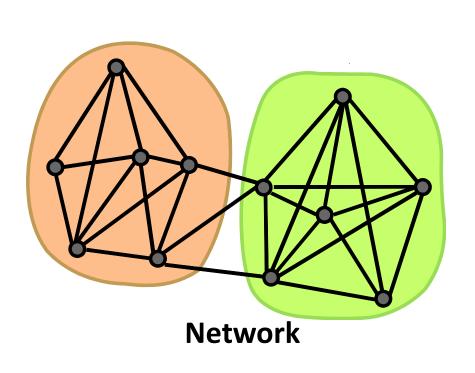
Community Types

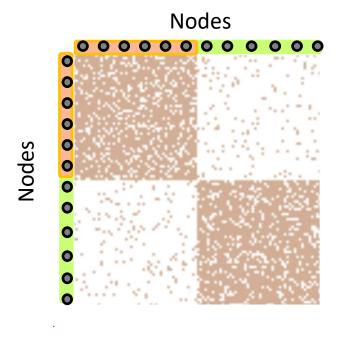
Non-overlapping vs. overlapping communities



Non-overlapping Communities

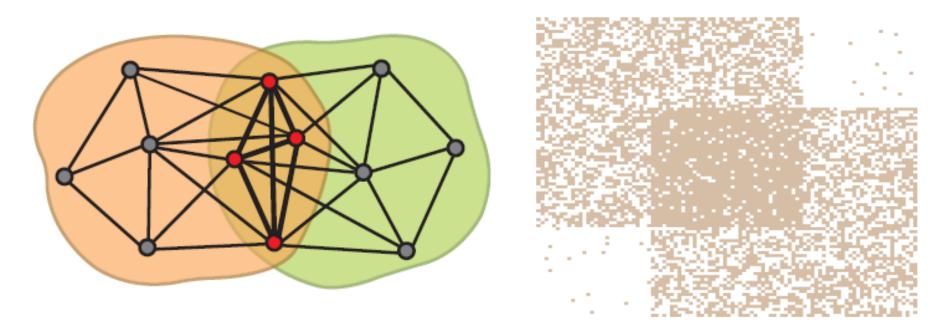
Adjacency matrix





Overlapping Communities

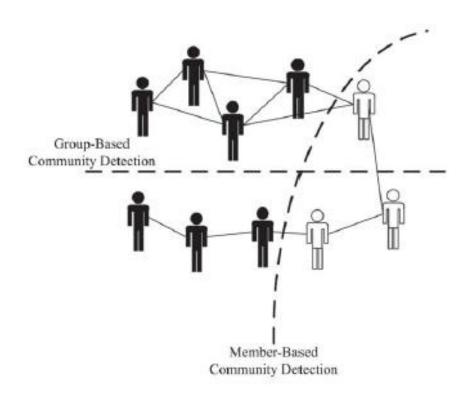
What is the structure of community overlaps: Edge density in the overlaps is higher!



Communities as "tiles"

Community Types

Member-based (local) vs. group-based



Community Detection

Given a graph G(V, E), find subsets C_i of V, such that $\bigcup_i C_i \subseteq V$

Assumptions

- Undirected graphs
- Edges may have
 - weights, (easily extended)
 - labels
 - content or attributes shared by individuals (in the same location, of the same gender, etc)
- Nodes may have labels, attributed, or labeled graphs

Multipartite graphs – e.g., affiliation networks, citation networks, customers-products: reduced to unipartited projections of each vertex class

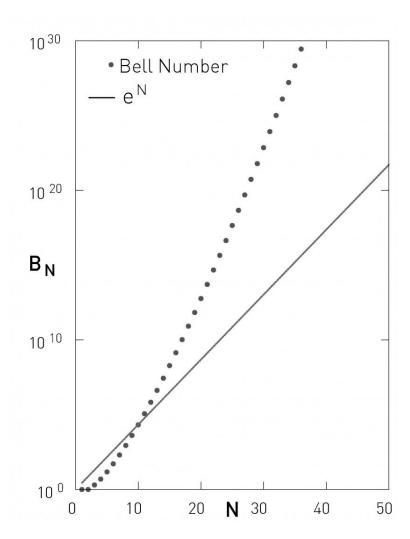
Hardness

Bell Number

Number of all possible partitions of *N* nodes

$$B_N = \frac{1}{e} \sum_{i=0}^{\infty} \frac{j^N}{j!}$$

For N = 50, 1040 partitions



Community Detection

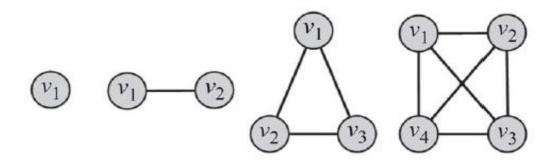
We will see three approaches

- Node degree (familiarity)
 - cliques
- Node similarity
 - cluster
- Node reachability
 - betweeness

Cliques (degree similarity)

Clique: a maximum *complete subgraph* in which all pairs of vertices are connected by an edge.

A *clique of size* k is a subgraph of k vertices where the degree of all vertices in the induced subgraph is k-1.



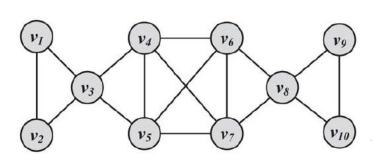
✓ Cliques vs complete graphs

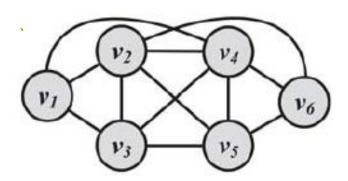
Cliques (degree similarity)

Search for:

- the maximum clique: the one with the largest number of vertices) or
- all maximal cliques: cliques that are not subgraphs of a larger clique; i.e., cannot be expanded further.

Both problems are NP-hard, as is verifying whether a graph contains a clique larger than size k.





Cliques

Algorithm 6.1 Brute-Force Clique Identification

```
Require: Adjacency Matrix A, Vertex v_x
 1: return Maximal Clique C containing v_x
 2: CliqueStack = \{\{v_x\}\}\, Processed = \{\}\;
 while CliqueStack not empty do
      C=pop(CliqueStack); push(Processed,C);
     v_{last} = Last node added to C;
 5:
     N(v_{last}) = \{v_i | A_{v_{last},v_i} = 1\}.
                                       /* Check all neighbors of last node sequentially
 7: for all v_{temp} \in N(v_{last}) do
                                                  if connected with all members in the clique
                                                        new clique -> push */
         if C \cup \{v_{temp}\} is a clique then
            push(CliqueStack, C \cup \{v_{temp}\}\);
 9:
         end if
10:
      end for
11:
12: end while
13: Return the largest clique from Processed
```

Enumerate all cliques (in alphabetical order)
Checks all permutations!
For (complete graph) 100 vertices, 299- 1 different cliques

Cliques

Pruning

- Prune all vertices (and incident edges) with degrees less than k 1.
- Effective due to the power-law distribution of vertex degrees

Exact cliques are *rarely observed* in real networks.

E.g., a clique of 1,000 vertices has (999x1000)/2 = 499,500 edges.

- A single edge removal results in a subgraph that is no longer a clique.
- That represents less than 0.0002% of the edges

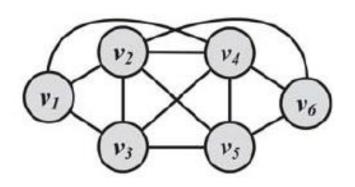
Relaxing Cliques

All vertices have a minimum degree but not necessarily k-1

k-plex

For a set of vertices V_0 , for all $u, d_u \ge |V_0| - k$ where d_u is the degree of v in the induced subgraph

What is k for a clique?



1-plex:
$$\{v_2, v_3, v_4, v_5\}$$

2-plex: $\{v_1, v_2, v_3, v_4, v_5\}, \{v_2, v_3, v_4, v_5, v_6\}$
3-plex: $\{v_1, v_2, v_3, v_4, v_5, v_6\}$

Relaxing Cliques

Clique

$$\forall i \in C, d_i^{int} = |C| - 1$$

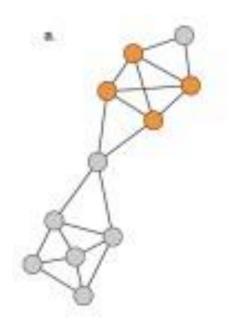
Strong community

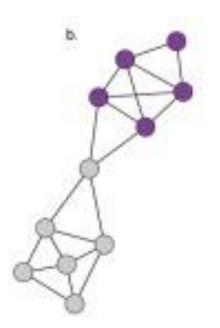
$$\forall i \in C, d_i^{int} > d_i^{ext}$$

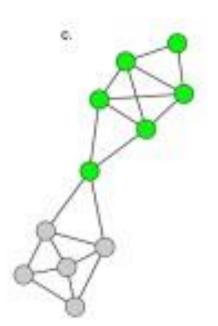
Weak community

$$\sum_{i \in C} d_i^{int} > \sum_{i \in C} d_i^{ext}$$

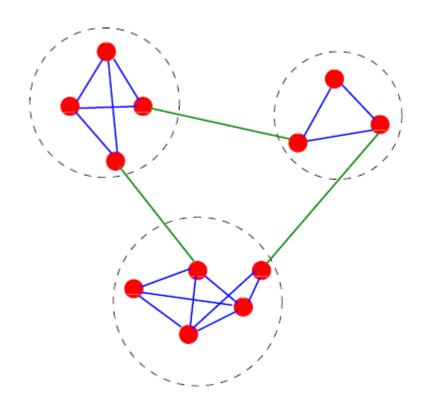
 d_i^{int} degree (#edges) of node i with nodes inside C d_i^{ext} degree (#edges) of node i with nodes outside C

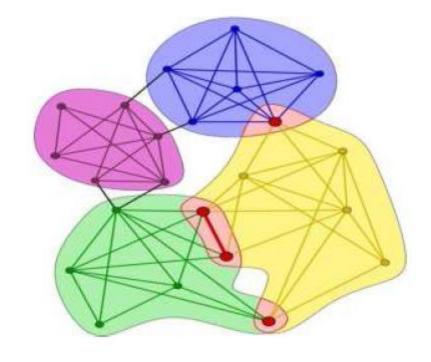






Assumption: communities are formed from a set of cliques and edges that connect these cliques.





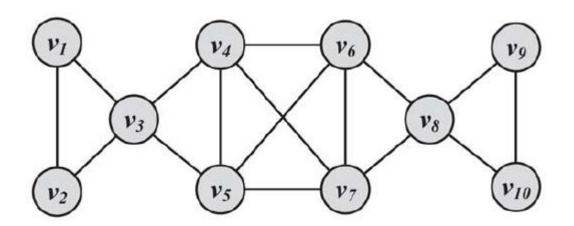
k = 4

Algorithm 6.2 Clique Percolation Method (CPM)

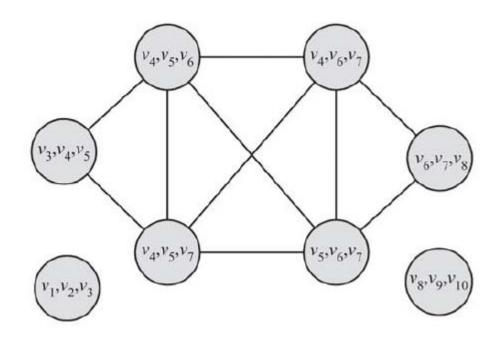
Require: parameter *k*

- 1: return Overlapping Communities
- 2: $Cliques_k = find all cliques of size k$
- 3: Construct clique graph G(V, E), where $|V| = |Cliques_k|$
- 4: $E = \{e_{ij} \mid \text{clique } i \text{ and clique } j \text{ share } k 1 \text{ nodes} \}$
- 5: Return all connected components of G
- 1. Given k, find all cliques of size k.
- Create graph (clique graph) where all cliques are vertices, and two cliques that share k - 1 vertices are connected via an edge.
- 3. Communities are the connected components of this graph.

Input graph, let k = 3

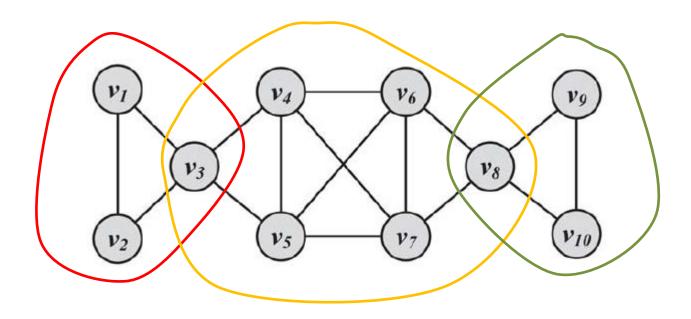


Clique graph for k = 3



(v1, v2, ,v3), (v8, v9, v10), and (v3, v4, v5, v6, v7, v8)

Result



(v1, v2, ,v3), (v8, v9, v10), and (v3, v4, v5, v6, v7, v8)

Clique Percolation Method (CPM)

- By construction, overlapping communities
- Instead of k = 3, maximal cliques
- Theoretical complexity grows exponential with size, but efficient on sparse graphs

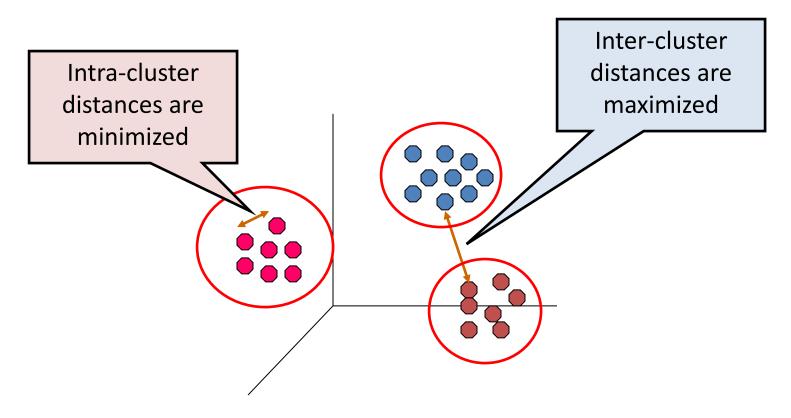
Outline

PART I

- 1. Introduction: what, why, types?
- 2. Cliques
- 3. Background: cluster analysis (node/edge similarity)
- 4. Hierarchical clustering (betweenness)
- 5. Modularity
- 6. How to evaluate

What is Cluster Analysis?

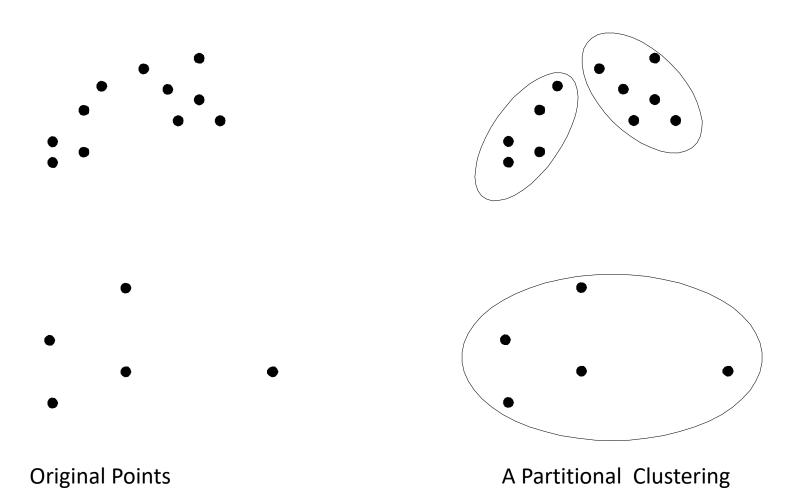
Finding groups of objects such that the objects in a group are similar (or related) to one another and different from (or unrelated to) the objects in other groups



Types of Clustering

- Important distinction between hierarchical and partitional sets of clusters
- Partitional Clustering
 - Division of data objects into subsets (clusters)
 - Assumes that the number of clusters is given
- Hierarchical clustering
 - A set of nested clusters organized as a hierarchical tree

Partitional Clustering

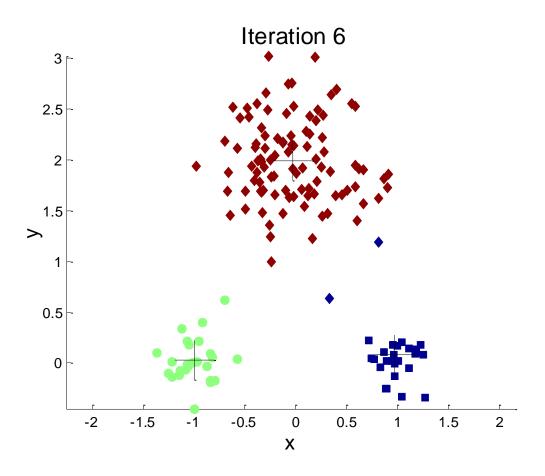


Example Partitioning: K-means Clustering

- 1: Select K points as the initial centroids.
- 2: repeat
- 3: Form K clusters by assigning all points to the closest centroid.
- 4: Recompute the centroid of each cluster.
- 5: **until** The centroids don't change

- Input: Number of clusters, K
- Each cluster is associated with a centroid (center point)
- Each point is assigned to the cluster with the closest centroid

Example



K-means Clustering

- Initial centroids are often chosen randomly.
 - Clusters produced vary from one run to another.
- The centroid is (typically) the mean of the points in the cluster.
- Closeness Similarity is measured by Euclidean distance, cosine similarity, correlation, etc.
- K-means will converge for common similarity measures mentioned above.
 - Most of the convergence happens in the first few iterations.
 - Often the stopping condition is changed to 'Until relatively few points change clusters'
- Complexity is O(n * K * I * d)
 - n = number of points, K = number of clusters,
 I = number of iterations, d = number of attributes

K-means Clusters

- Most common measure is Sum of Squared Error (SSE)
 - For each point, the error is the distance to the nearest cluster
 - To get SSE, we square these errors and sum them.

$$SSE = \sum_{i=1}^{K} \sum_{x \in C_i} dist^2(m_i, x)$$

- x is a data point in cluster C_i and m_i is the representative point for cluster C_i
 - can show that m_i corresponds to the center (mean) of the cluster
- Given two clusters, we can choose the one with the smallest error
- One easy way to reduce SSE is to increase K, the number of clusters
 - A good clustering with smaller K can have a lower SSE than a poor clustering with higher K

Vertex similarity

- Define similarity between two vertices
- Place similar vertices in the same cluster

Use traditional cluster analysis

Vertex similarity

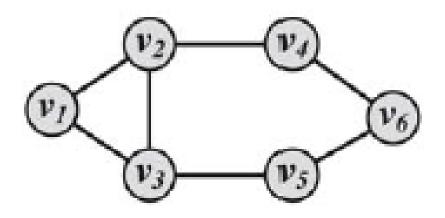
 Structural equivalence: based on the overlap between their neighborhoods

$$\sigma(v_i, v_j) = |N(v_i) \cap N(v_j)|$$

Normalized to [0, 1], e.g.,

$$\sigma_{\text{Jaccard}}(v_i, v_j) = \frac{|N(v_i) \cap N(v_j)|}{|N(v_i) \cup N(v_j)|}$$

Vertex similarity



$$\sigma_{\text{Jaccard}}(v_2,v_5) = \frac{|\{v_1,v_3,v_4\} \cap \{v_3,v_6\}|}{|\{v_1,v_3,v_4,v_6\}|} = 0.25$$

Other definitions of vertex similarity

Use the adjacency matrix A,

$$d_{ij} = \sqrt{\sum_{k \neq i,j} (A_{ik} - A_{jk})^2}$$

Common neighbors (paths of length two)

We can also use A^2

Other definitions of vertex similarity

If we map vertices u, v to n-dimensional points A, B in the Euclidean space,

$$d_{AB}^{E} = \sum_{k=1}^{n} \sqrt{(a_k - b_k)^2}$$

$$d_{AB}^{M} = \sum_{k=1}^{n} |a_k - b_k|$$

$$d_{AB}^{\infty} = \max_{k \in [1,n]} |a_k - b_k|$$

$$\rho_{AB} = \arccos \frac{\mathbf{a} \cdot \mathbf{b}}{\sqrt{\sum_{k=1}^{n} a_k^2} \sqrt{\sum_{k=1}^{n} b_k^2}}$$

Other definitions of vertex similarity

Many more – we shall revisit this issue when we talk about *graph embeddings*

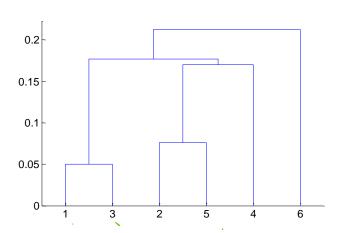
Also useful when there are *attributes* associated with nodes or edges to combine distances

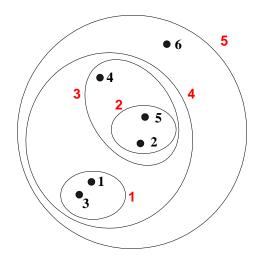
Hierarchical Clustering

- Produces a set of nested clusters organized as a hierarchical tree
- Can be visualized as a dendrogram

A tree like diagram that records the sequences of

merges or splits





Hierarchical Clustering

- Two main types of hierarchical clustering
 - Agglomerative:
 - Start with each node as an individual cluster (called singletons)
 - At each step, merge the closest pair of clusters until only one cluster (or k clusters) is left

– Divisive:

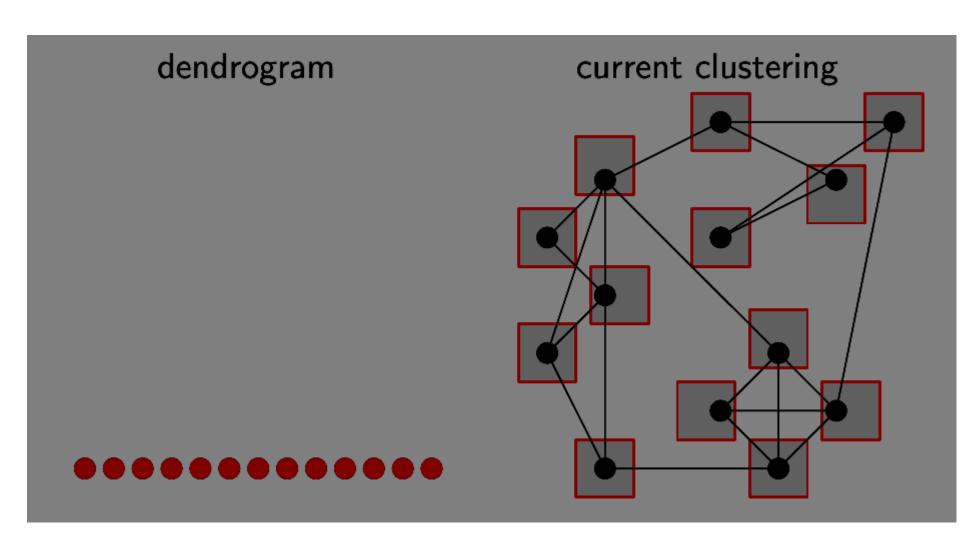
- Start with one, all-inclusive cluster = the whole graph
- At each step, split a cluster until each cluster contains a single node (or there are k clusters)
- Traditional hierarchical algorithms use a similarity or distance matrix
 - Merge or split one cluster at a time

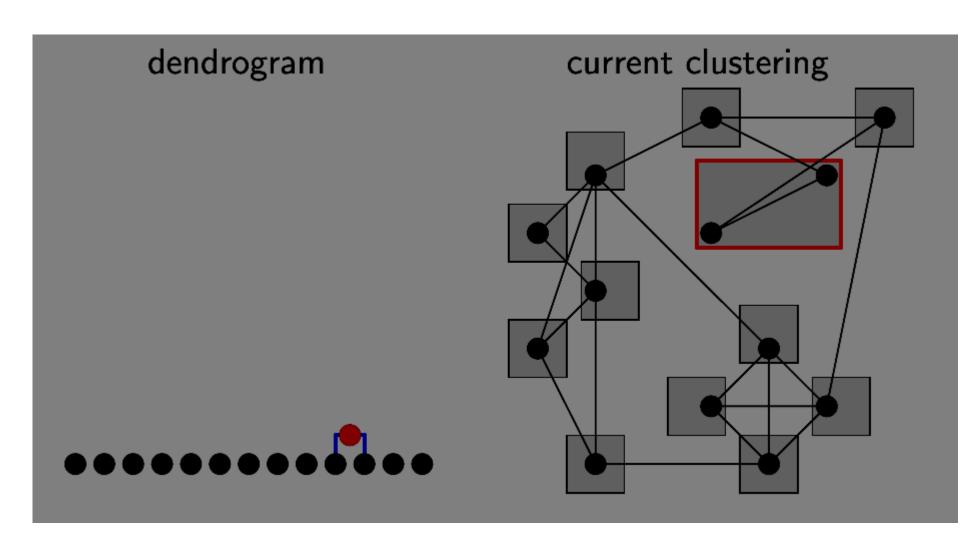
Agglomerative Clustering Algorithm

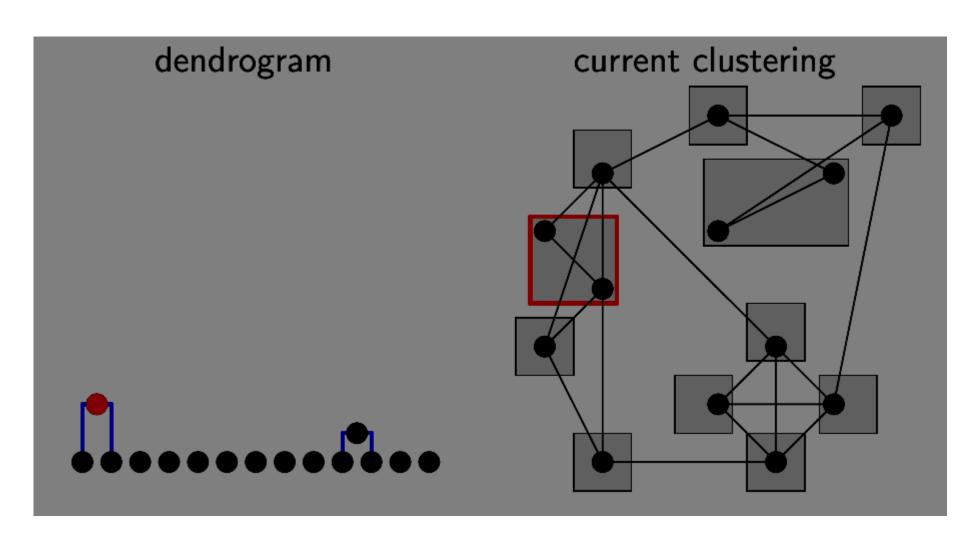
Popular hierarchical clustering technique

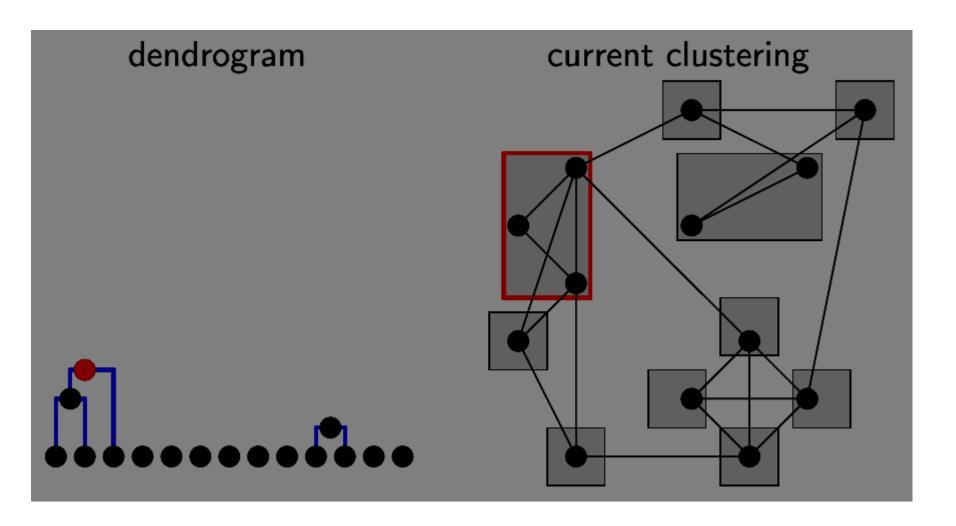
Basic algorithm is straightforward

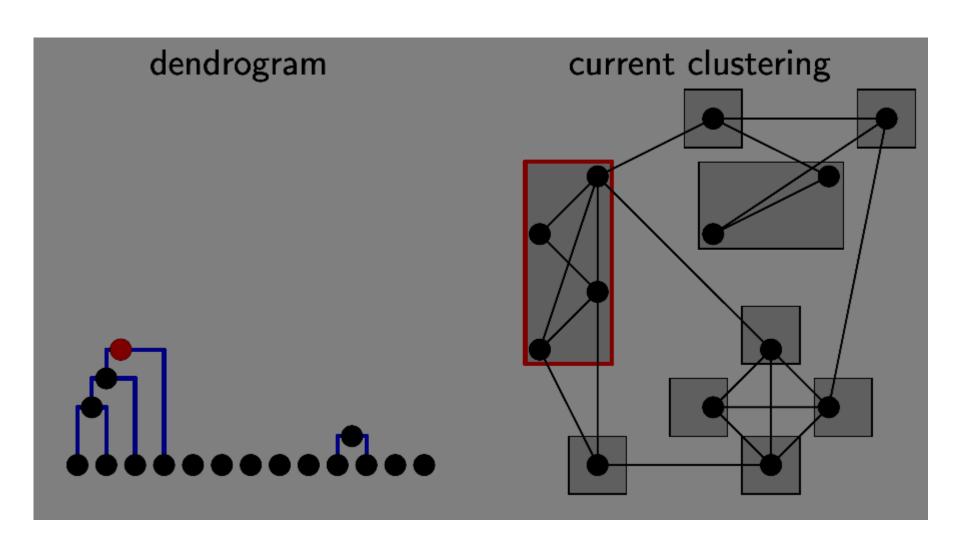
- 1. [Compute the proximity matrix]
- 2. Let each node be a cluster
- 3. Repeat
- 4. Merge the *two closest clusters*
- 5. [Update the proximity matrix]
- **6. Until** only a single cluster remains

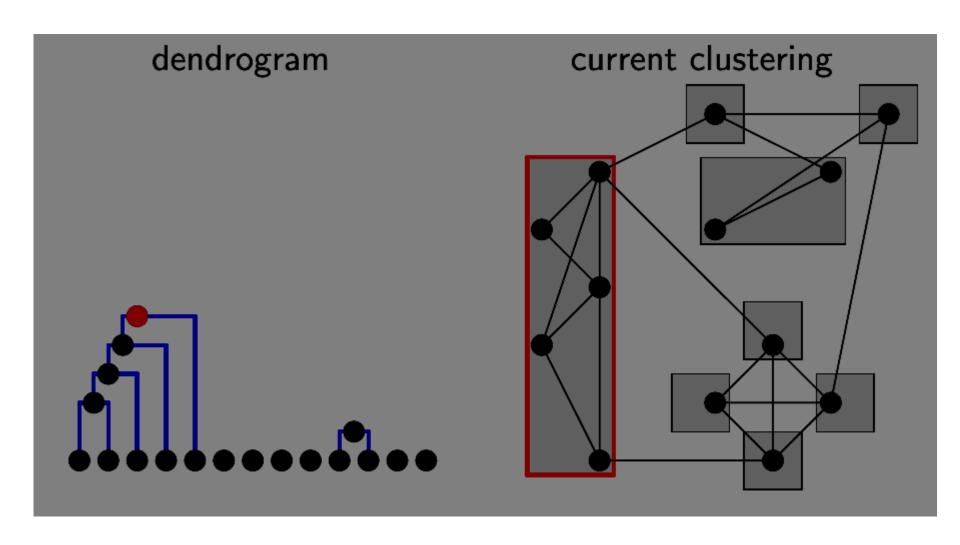


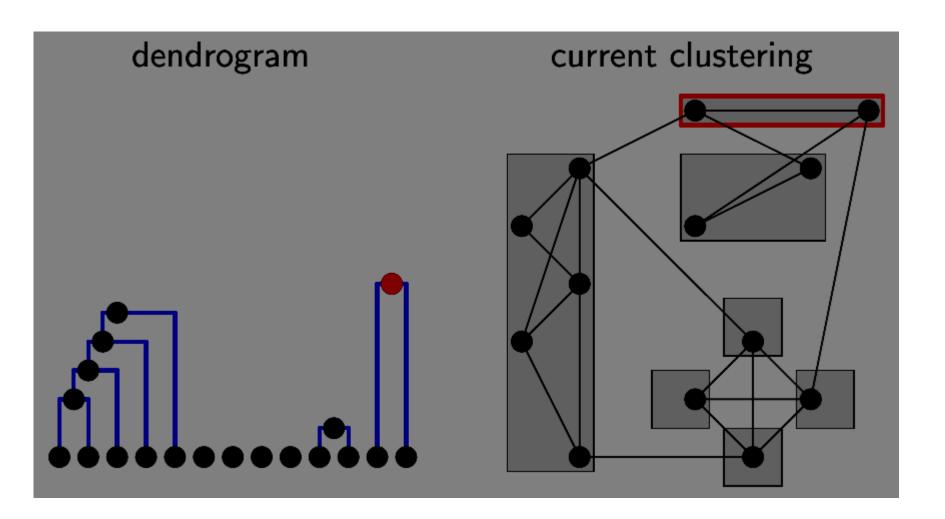


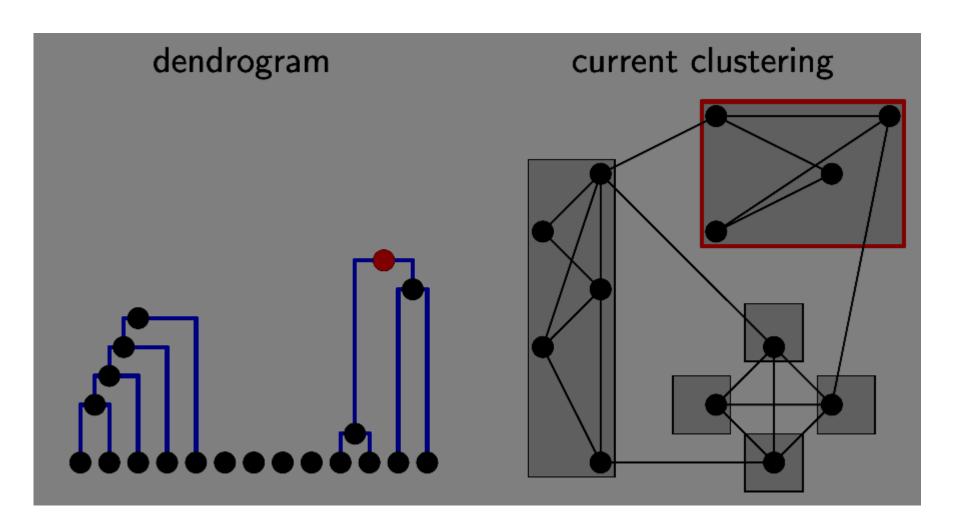


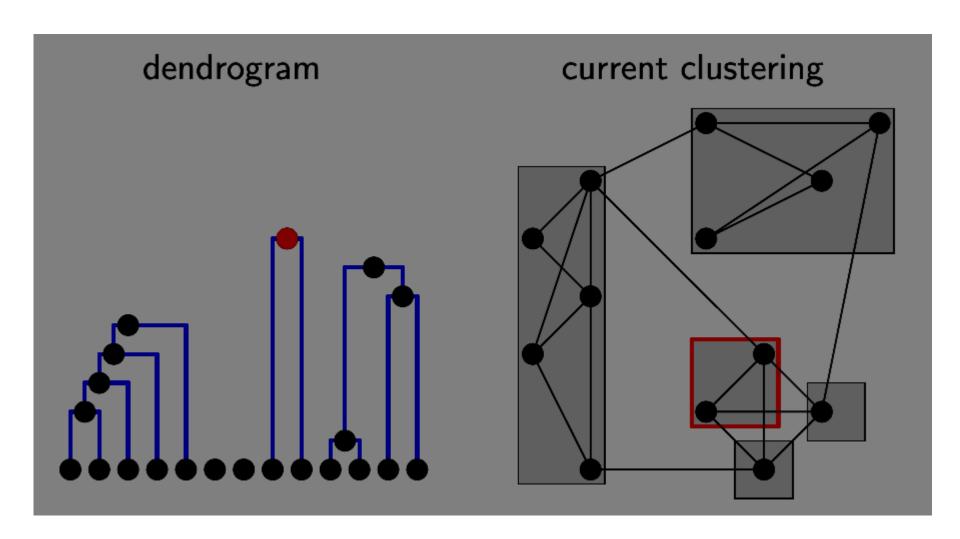


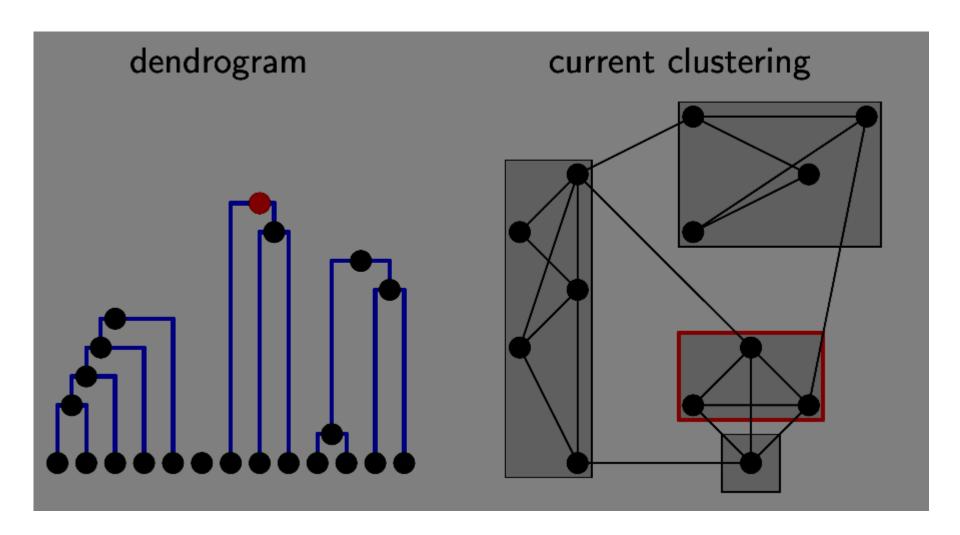


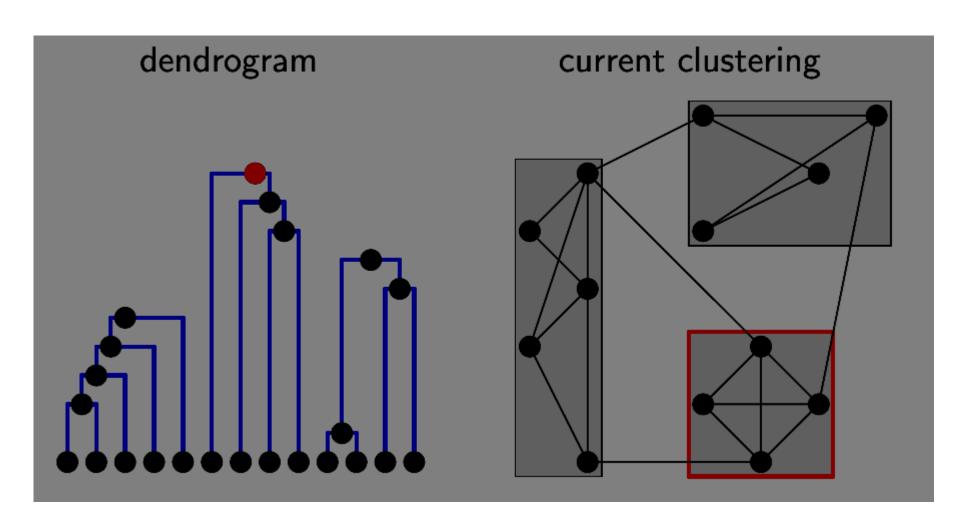


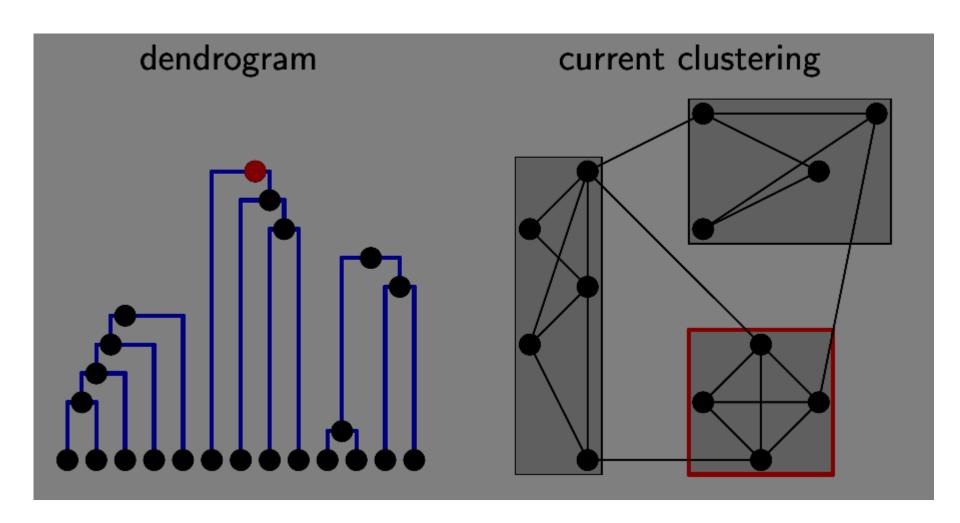


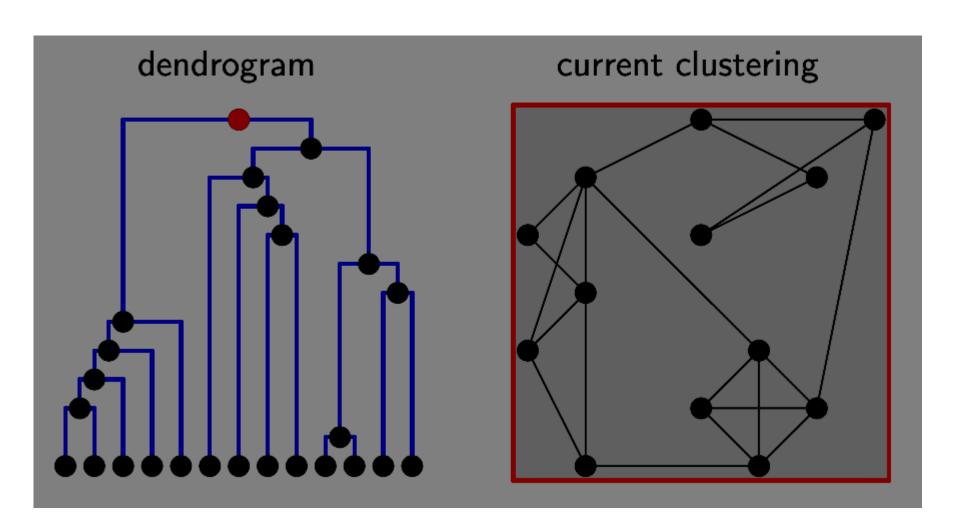










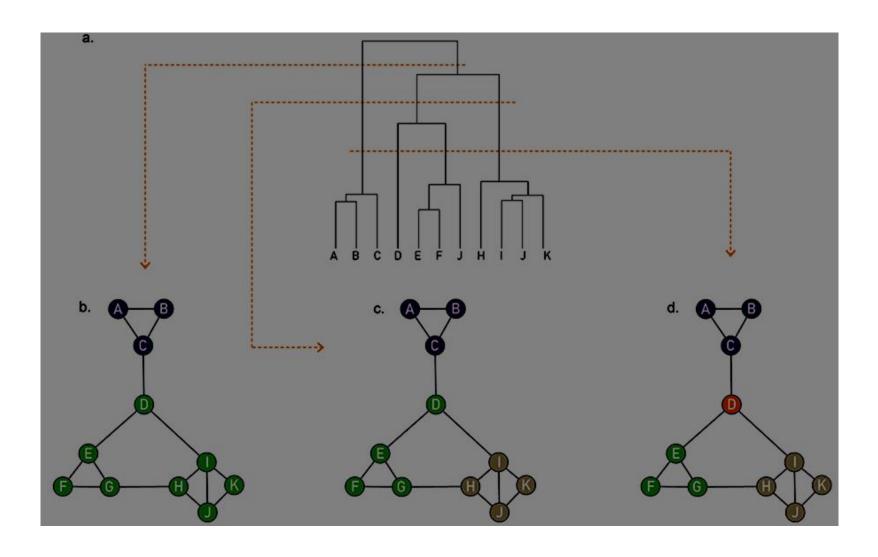


Strengths of Hierarchical Clustering

- Do not have to assume any particular number of clusters
 - Any desired number of clusters can be obtained by 'cutting' the dendogram at the proper level

- They may correspond to meaningful taxonomies
 - Example in biological sciences (e.g., animal kingdom, phylogeny reconstruction, ...)

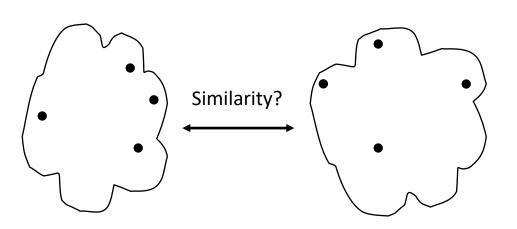
Where to cut?



Agglomerative Clustering Algorithm

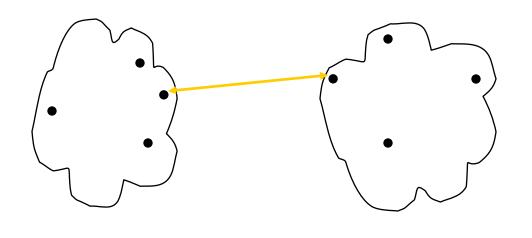
Key operation is the *computation of the proximity of two*

Different approaches to defining the distance between clusters distinguish the different algorithms



	p1	p2	р3	p4	р5	<u> </u>
<u>p1</u>						
<u>p2</u> <u>p3</u>						
<u>p4</u> <u>p5</u>						_

Proximity Matrix



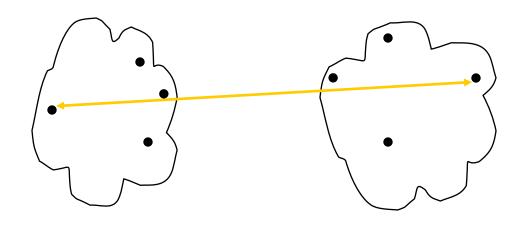
	p1	p2	р3	p4	р5	<u> </u>
p1						
p2						
<u>p2</u> <u>p3</u>						
<u>p4</u> <u>p5</u>						
<u> </u>						

MIN or single link

based on the two most similar (closest) points in the different clusters

Proximity Matrix

sensitive to outliers



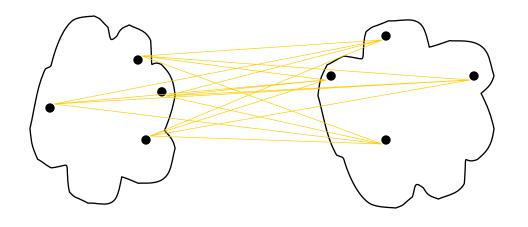
	p1	p2	р3	p4	р5	<u> </u>
p1						
p2						
<u>p2</u> <u>p3</u>						
<u>p4</u> <u>p5</u>						
<u> </u>						

MAX or complete linkage

Similarity of two clusters is based on the two least similar (most distant) points in the different clusters

Proximity Matrix

Tends to break large clusters
Biased towards globular clusters



	р1	p2	р3	p4	р5	<u> </u>
р1						
p2						
<u>p2</u> p3						
<u>p4</u> <u>p5</u>						
•						

Group Average

Proximity of two clusters is the average of pairwise proximity between points in the two clusters.

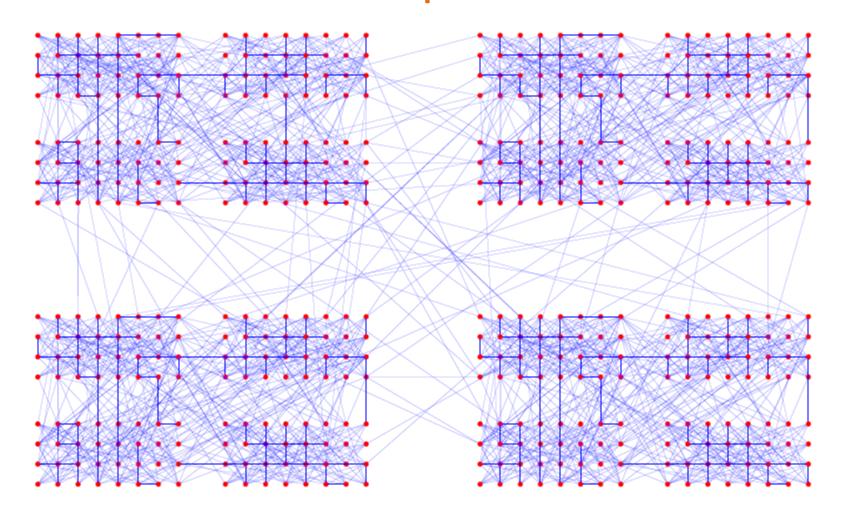
Proximity Matrix

Outline

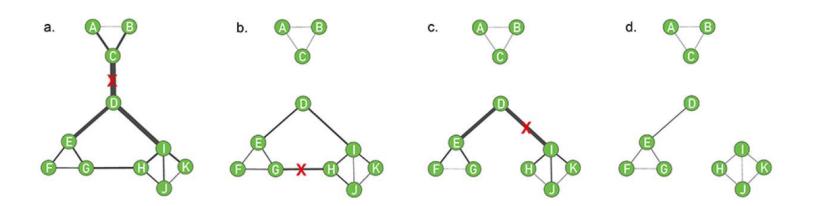
PART I

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Example of a Hierarchically Structured Graph



Divisive Algorithms



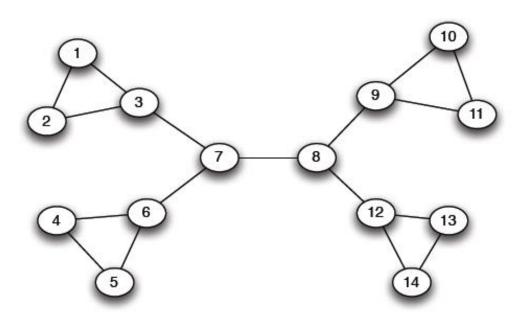


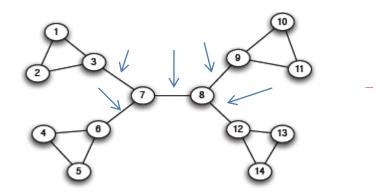
Which edge to remove?

Hierarchical divisive method

- Start with the whole graph
- Find edges whose removal "partitions" the graph
- Repeat with each subgraph until single vertices

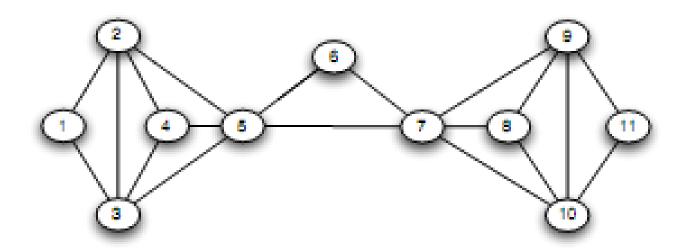
Which edge?



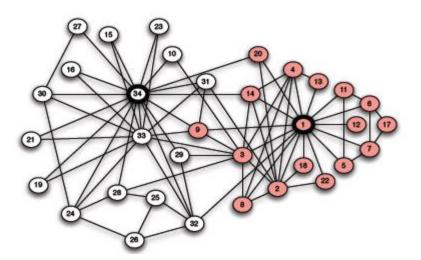


Use bridges or cut-edge (if removed, the nodes become disconnected)

Which one to choose?



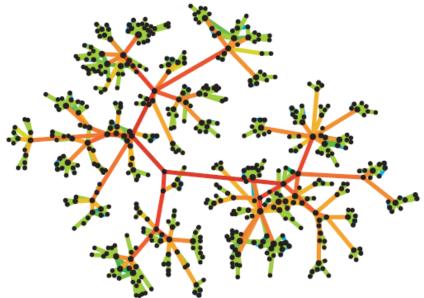
There may be none!



Strength of Weak Ties

- Edge betweenness: Number of shortest paths passing over the edge
- Intuition:

Assuming communication through shortest paths, captures *traffic*



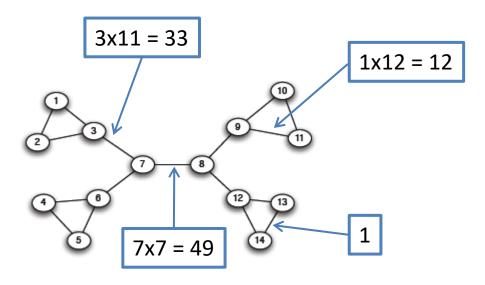
Edge betweenness in a real network

Edge Betweenness

Betweenness of an edge (a, b): number of pairs of nodes x and y such that the edge (a, b) lies on their shortest path

There can be multiple shortest paths, take the fraction that includes (a, b)

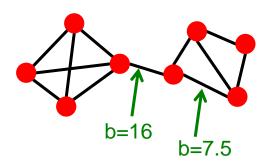
$$betweeness(a,b) = \sum_{(x,y) \in E} \frac{\#shortest_paths(x,y)_through(a,b)}{\#shortest_paths(x,y)}$$



edges that have a high probability to occur on a randomly chosen shortest path between two randomly chosen nodes

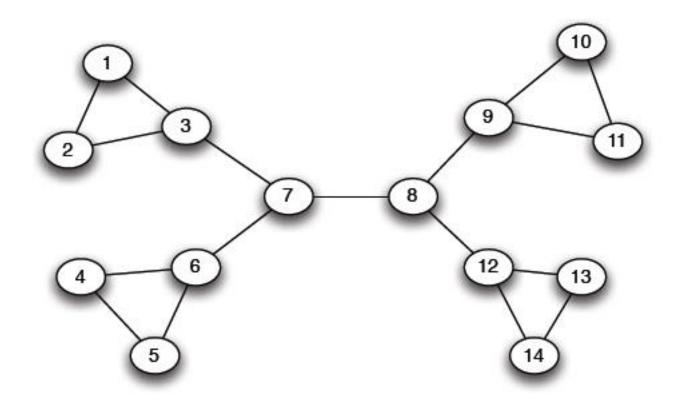
Edge Betweenness

$$betweeness(a,b) = \sum_{(x,y) \in E} \frac{\#shortest_paths(x,y)_through(a,b)}{\#shortest_paths(x,y)}$$



» Undirected unweighted networks

- Repeat until no edges are left:
 - Calculate betweenness of edges
 - Remove edges with highest betweenness
- Connected components are communities
- Gives a hierarchical decomposition of the network

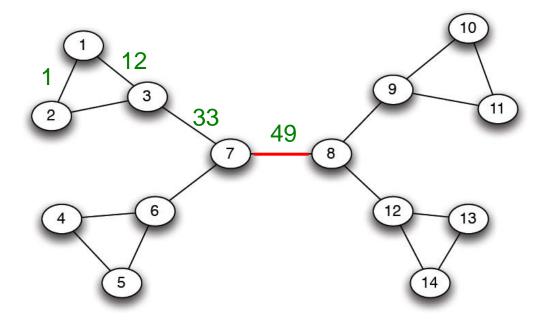


Betweenness(7, 8) = 7x7 = 49

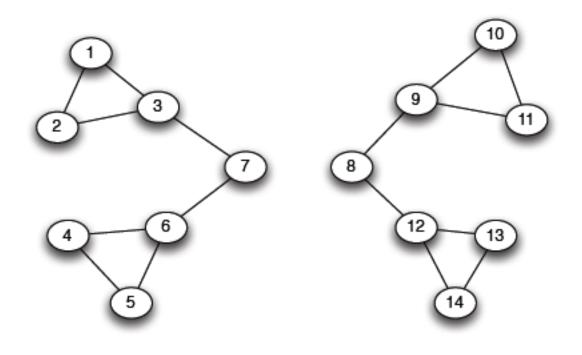
Betweenness(3, 7)=Betweenness(6, 7)=Betweenness(8, 9) = Betweenness(8, 12)= 3x11=33

Betweenness(1, 3) = 1x12=12

Girvan-Newman: Example



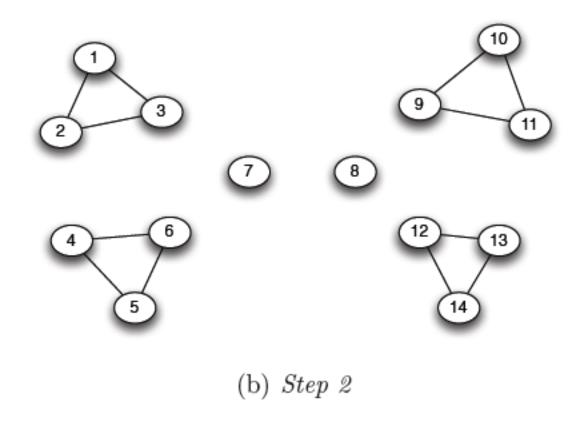
Need to re-compute betweenness at every step



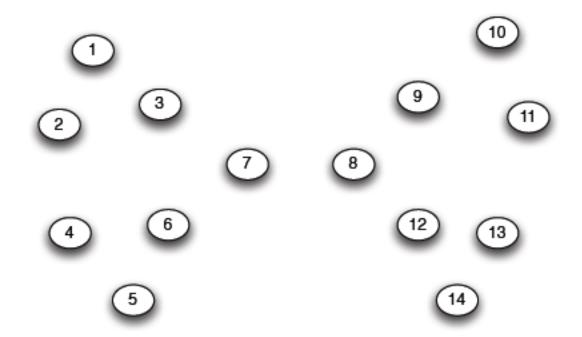
(a) Step 1

Betweenness(1, 3) = 1x5=5

Betweenness(3,7)=Betweenness(6,7)=Betweenness(8,9) = Betweenness(8,12)= 3x4=12

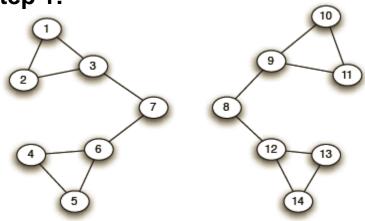


Betweenness of every edge = 1

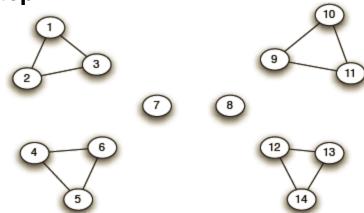


Girvan-Newman: Example

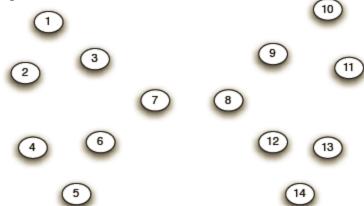
Step 1:



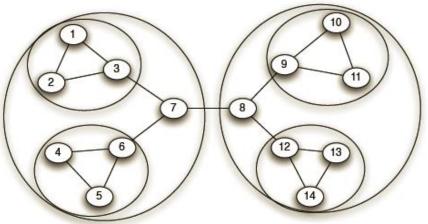
Step 2:



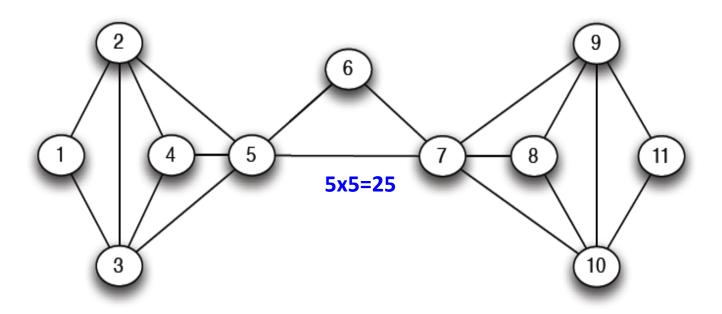
Step 3:



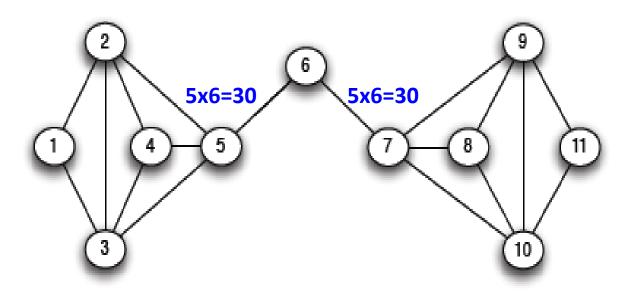
Hierarchical network decomposition:



Another example

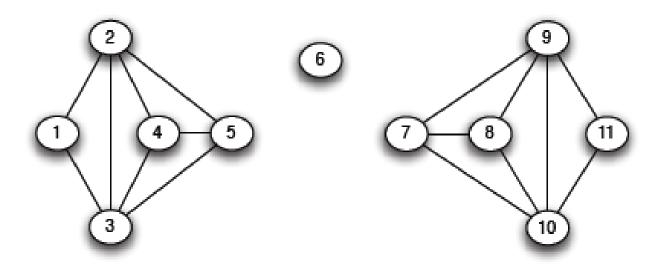


Another example



(a) Step 1

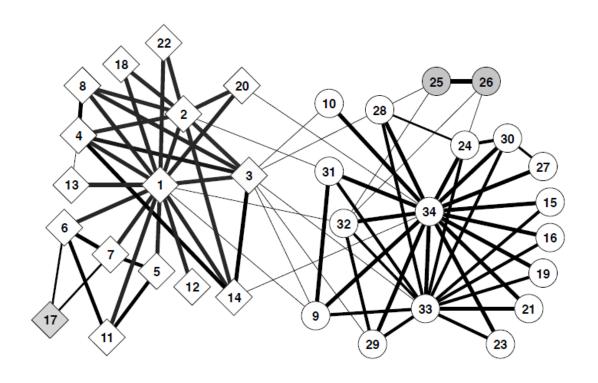
Another example

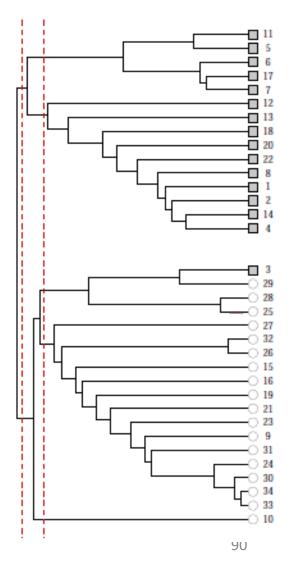


(b) Step 2

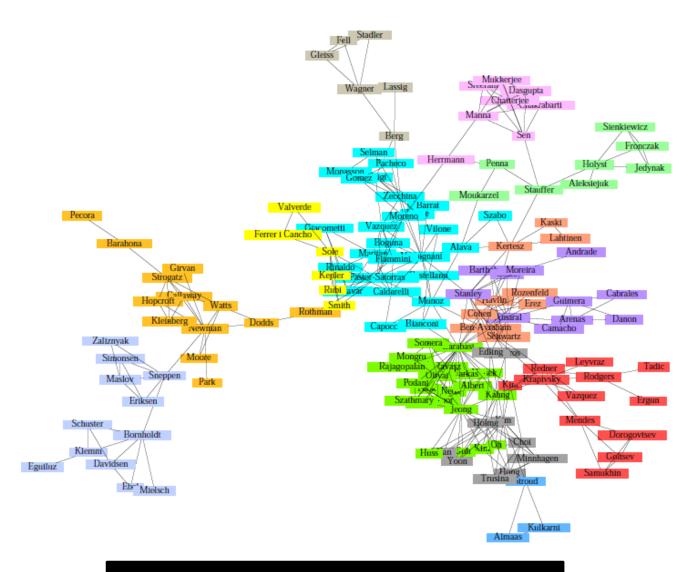
Girvan-Newman: Results

 Zachary's Karate club: Hierarchical decomposition



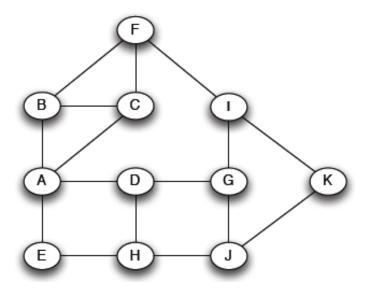


Girvan-Newman: Results



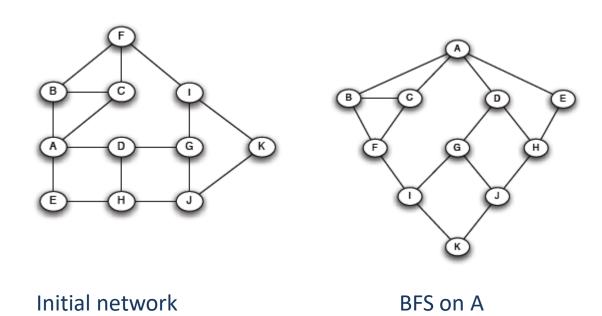
How to Compute Betweenness?

 Want to compute betweenness of paths starting at node A

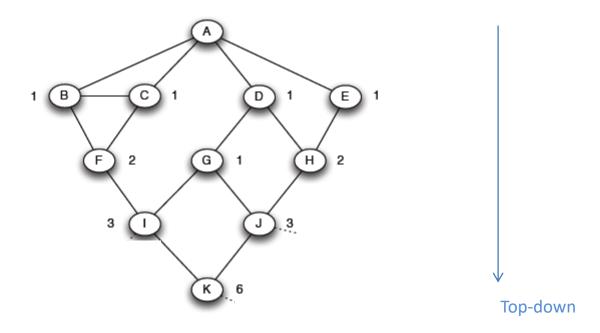


Computing Betweenness

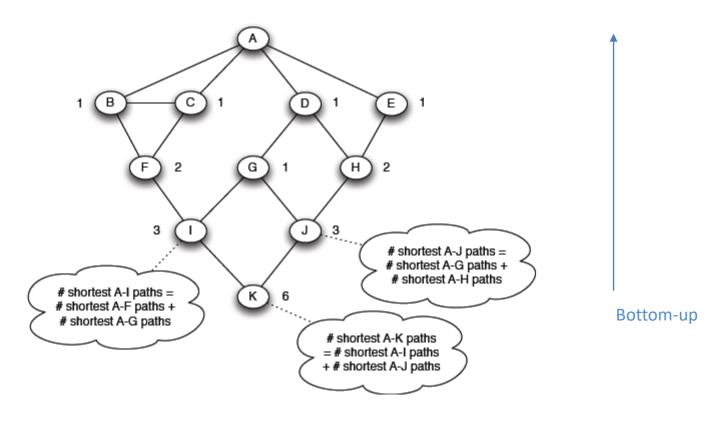
- 1. Perform a *BFS* starting from A
- 2. Determine the *number of shortest path* from A to each other *node*
- 3. Based on these numbers, determine the amount of *flow* from A to all other nodes that uses each edge



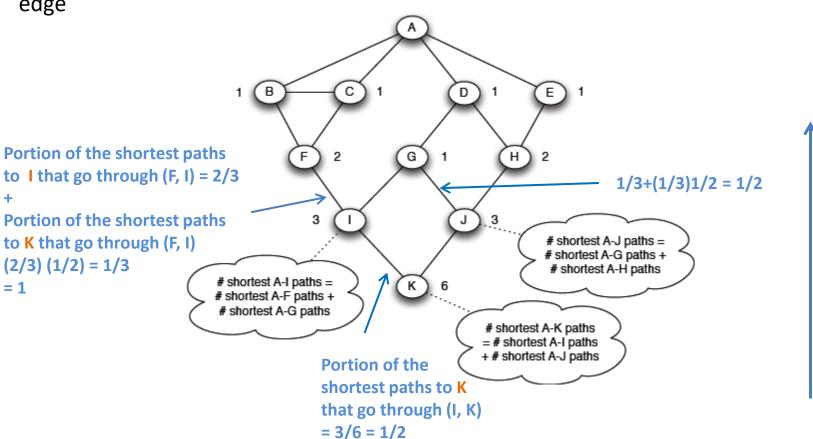
Count how many shortest paths from A to a specific node

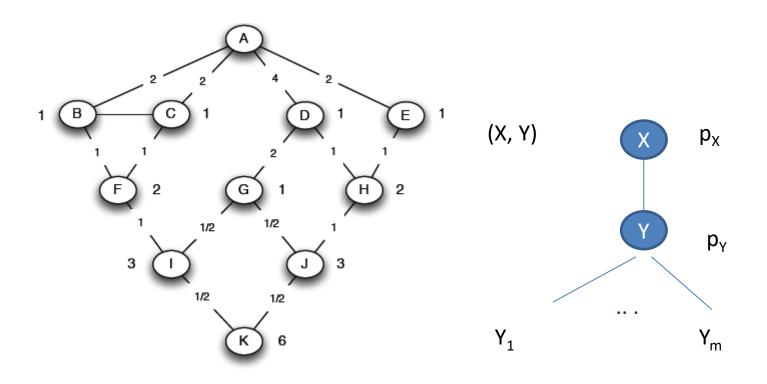


Compute betweenness by working up the tree: If there are multiple paths count them fractionally



Count the flow through each edge





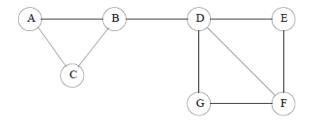
$$flow(X,Y) = \frac{p_X}{p_y} + \sum_{Y_i child \ of \ Y} \frac{p_X}{p_y} flow(Y,Y_i)$$

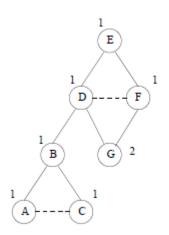
Computing Betweenness

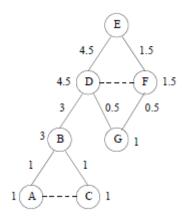
Repeat the process for all nodes

Sum over all BFSs

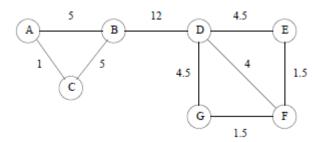
Example







Example





Computing Betweenness

Issues

- Test for connectivity?
- Re-compute all paths, or only those affected
- Parallel computation
- Sampling

Centrality measures

Degree centrality

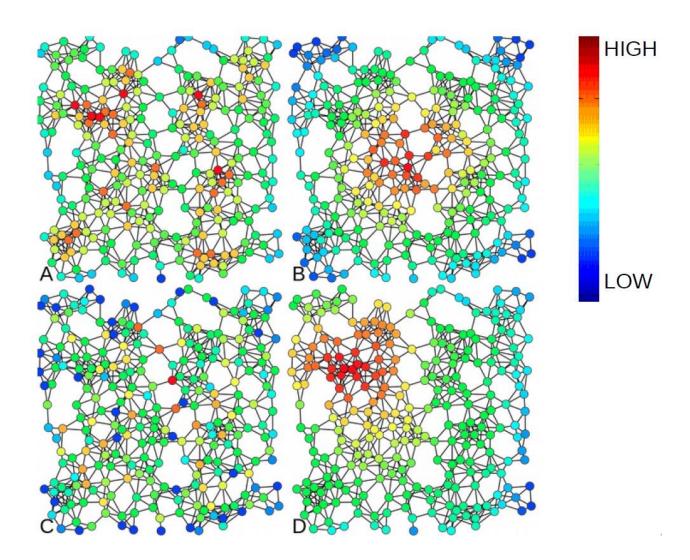
closeness
$$(u) = \frac{1}{\sum_{v \in V, v \neq u} d(u, v)}$$

A: Degree

B: Closeness

C: Betweenness

D: PageRank



Outline

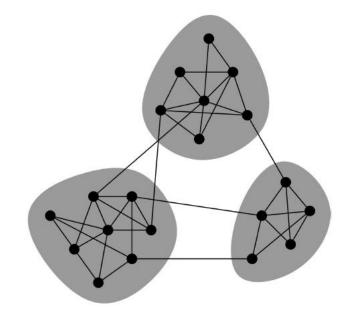
PART I

- 1. Introduction: what, why, types?
- 2. Cliques and vertex similarity
- 3. Background: Cluster analysis
- 4. Betweeness centrality
- 5. Modularity, label propagation
- 6. How to evaluate

Modularity

- Communities: sets of tightly connected nodes
- Define: Modularity Q
 - A measure of how well a network is partitioned into communities
 - Given a partitioning of the network into groups $s \in S$:

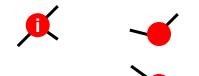
 $Q \propto \sum_{s \in S} [$ (# edges within group s) – (expected # edges within group s)]



Need a null model!

Null Model: Configuration Model

- Given real G on n nodes and m edges,
 construct rewired network G'
 - Same degree distribution but random connections



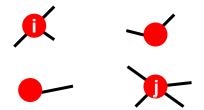
- Consider G' as a multigraph
- The expected number of edges between nodes i and j of degrees d_i and d_j equals to: $d_i \cdot \frac{d_j}{2m} = \frac{d_i d_j}{2m}$

For any edge going out of i randomly, the probability of this edge getting connected to node j is $\frac{d_j}{2m}$ Because the degree for i is d_i , we have d_i number of such edges

Note:

$$\sum_{u \in N} d_u = 2m$$

Null Model: Configuration Model



• The expected number of edges in (multigraph) G':

$$- = \frac{1}{2} \sum_{i \in N} \sum_{j \in N} \frac{d_i d_j}{2m} = \frac{1}{2} \cdot \frac{1}{2m} \sum_{i \in N} d_i \left(\sum_{j \in N} d_j \right) =$$
$$- = \frac{1}{4m} 2m \cdot 2m = m$$

Modularity

- Modularity of partitioning S of graph G:
 - Q $\propto \sum_{s \in S}$ [(# edges within group *s*) − (expected # edges within group *s*)]

$$-Q(G,S) = \frac{1}{2m} \sum_{S \in S} \sum_{i \in S} \sum_{j \in S} \left(A_{ij} - \frac{d_i d_j}{2m} \right)$$
Normalizing cost.: -1ij = 1 if i \rightarrow j o else

- Modularity values take range [-1, 1]
 - It is positive if the number of edges within groups exceeds the expected number
 - 0.3-0.7 < Q means significant community structure</p>

Modularity

Greedy method of Newman (one of the many ways to use modularity)

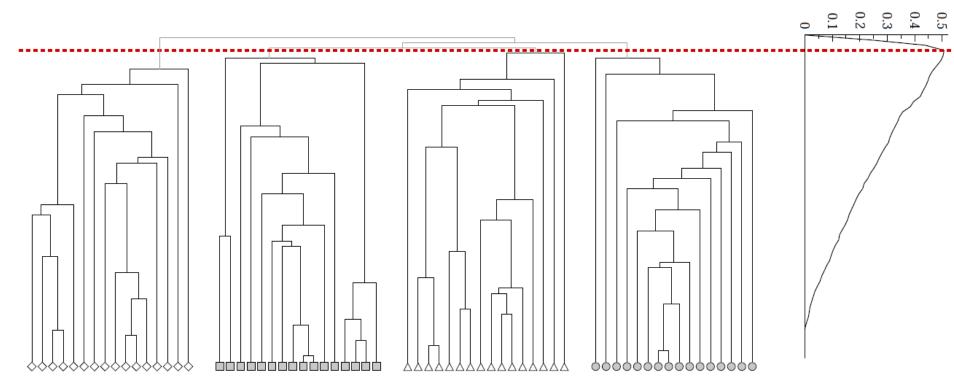
Agglomerative hierarchical clustering method

- 1. Start with a state in which each vertex is the sole member of one of *n* communities
- Repeatedly join communities together in pairs, choosing at each step the join that results in the greatest increase (or smallest decrease) in Q.

Since the joining of a pair of communities between which there are no edges can never result in an increase in modularity, we *need only consider* those pairs between which there are edges, of which there will at any time be at most m

Modularity: Number of clusters

 Modularity is useful for selecting the number of clusters:



modularity

Modularity: Cluster quality

When a given clustering is "good"?

Also, it is both a local (per individual cluster) and global measure

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Label propagation

Vertices are initially given unique labels (e.g., their vertex labels).

At each iteration,

sweep over all vertices, in random sequential order:

each vertex takes the label shared by the majority of its neighbors.

If no unique majority, one of the majority label is picked at random.

Stop (convergence) when each vertex has the majority label of its neighbors

Communities: groups of vertices having identical labels at convergence

Label propagation

- Labels propagate across the graph: most labels will disappear, others will dominate.
- By construction, each vertex has more neighbors in its community than in any other community.
- Due to many possible ties, different partitions
 - Perform many propagations from the same initial condition, with different random seeds
 - Aggregate partition label each vertex with the set of all labels it has in different partitions → overlapping communities

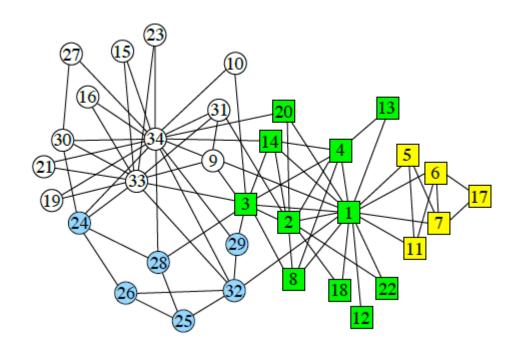
Outline

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Community Evaluation

- With ground truth
- Without ground truth

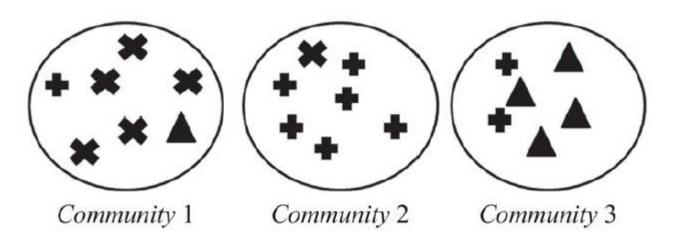


Zachary's Karate Club
Club president (34) (circles) and instructor (1) (rectangles)

Metrics: purity

the fraction of instances that have labels equal to the label of the community's majority

$$Purity = \frac{1}{N} \sum_{i=1}^{k} \max_{j} |C_i \cap L_j|$$

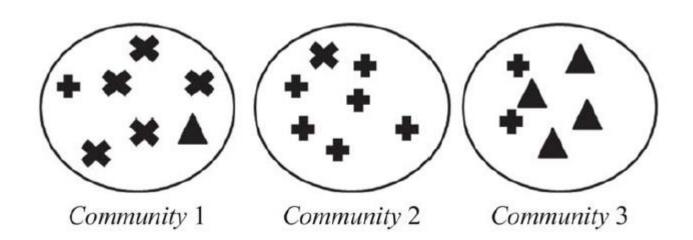


(5+6+4)/20 = 0.75

Metrics

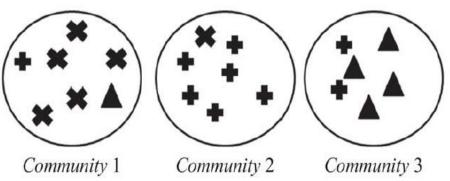
Based on pair counting: the number of pairs of vertices which are classified in the same (different) clusters in the two partitions.

- True Positive (TP) Assignment: when similar members are assigned to the same community. This is a correct decision.
- True Negative (TN) Assignment: when dissimilar members are assigned to different communities. This is a correct decision.
- False Negative (FN) Assignment: when similar members are assigned to different communities. This is an incorrect decision.
- False Positive (FP) Assignment: when dissimilar members are assigned to the same community. This is an incorrect decision.



For TP, we need to compute the number of pairs with the same label that are in the same community

$$TP = \underbrace{\binom{5}{2}}_{Community 1} + \underbrace{\binom{6}{2}}_{Community 2} + \underbrace{\binom{4}{2} + \binom{2}{2}}_{Community 3} = 32$$



$$TN = \underbrace{(5 \times 6 + 1 \times 1 + 1 \times 6 + 1 \times 1)}_{Communities 1 and 2}$$

×,+

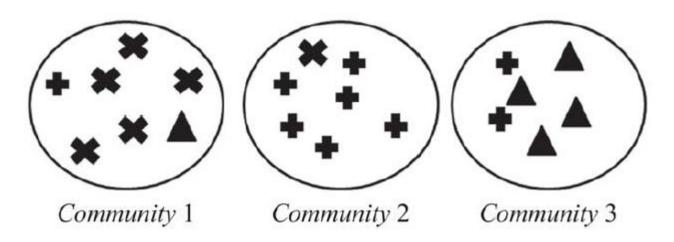
For TN: compute the number of dissimilar pairs in dissimilar communities

$$+\underbrace{(5\times4+5\times2+1\times4+1\times2)}^{\times,\wedge}$$

Communities 1 and 3

+
$$\underbrace{(6 \times 4 + 1 \times 2 + 1 \times 4)}_{+,\triangle} = 104.$$

Communities 2 and 3



For FP, compute dissimilar pairs that are in the same community.

$$FP = \underbrace{(5 \times 1 + 5 \times 1 + 1 \times 1)}_{Community 1} + \underbrace{(6 \times 1)}_{Community 2} + \underbrace{(4 \times 2)}_{Community 3} = 25$$

For FN, compute similar members that are in different communities.

$$FN = \underbrace{(5 \times 1)}_{\times} + \underbrace{(6 \times 1 + 6 \times 2 + 2 \times 1)}_{+} + \underbrace{(4 \times 1)}_{\triangle} = 29$$

Precision (P): the fraction of pairs that have been correctly assigned to the same community.

Recall (R): the fraction of pairs assigned to the same community of all the pairs that should have been in the same community.

F-measure

$$2PR/(P+R)$$

- Cluster Cohesion: Measures how closely related are objects in a cluster
- Cluster Separation: Measure how distinct or well-separated a cluster is from other clusters
- Example: Squared Error
 - Cohesion is measured by the within cluster sum of squares (SSE)

$$WSS = \sum_{i} \sum_{x \in C} (x - m_i)^2$$

 $i \ x \in C_i$ — Separation is measured by the between cluster sum of squares

$$BSS = \sum_{i} |C_{i}| (m - m_{i})^{2}$$

- Where $|C_i|$ is the size of cluster i

$$cohesion(C) = \frac{\#internal\ edges\ of\ C}{N_C(N_C - 1)/2}$$

$$seperation(C) = \frac{\#inter - cluster \ edges \ of \ C}{N_C(N - N_C)}$$

Modularity

Both as a local (per individual community) and as a global measure

With semantics:

- (ad hoc) analyze other attributes (e.g., profile, content generated) for coherence
- human subjects (user study) Mechanical Turk
 Visual representation (similarity/adjacency matric, word clouds, etc)





(a) U.S. Constitution

(b) Sports

Basic References

- Jure Leskovec, Anand Rajaraman, Jeff Ullman, Mining of Massive Datasets,
 Chapter 10, http://www.mmds.org/
- Reza Zafarani, Mohammad Ali Abbasi, Huan Liu, Social Media Mining: An Introduction, Chapter 6, http://dmml.asu.edu/smm/
- Santo Fortunato: Community detection in graphs. CoRR abs/0906.0612v2 (2010)
- Pang-Ning Tan, Michael Steinbach, Vipin Kumar, Introduction to Data Mining, Chapter 8, http://www.users.cs.umn.edu/~kumar/dmbook/index.php
- Albert-László Barabasi, Network Science, Chapter 9, http://networksciencebook.com/

Questions?