

Community Detection in Social Networks

Reference: 1) Network Science by Barabasi
2) Social Network Analysis by Tanmoy Chakraborty

Basics of Communities

- Fundamental Hypotheses
 - *A network's community structure is uniquely encoded in its wiring diagram.*
- What do we really mean by a community?
- How many communities are in a network?
- How many different ways can we partition a network into communities?

Defining Communities

- **Connectedness and Density Hypothesis**

A community is a locally dense connected subgraph in a network.

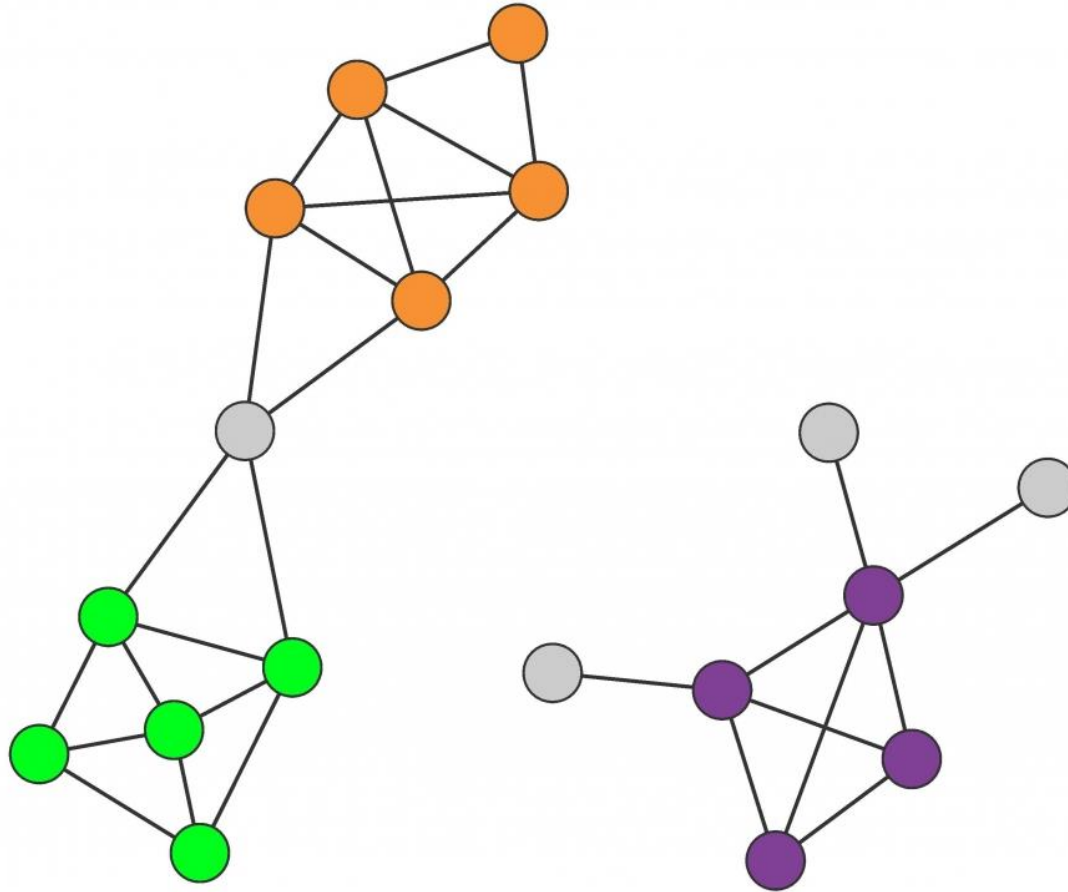
- **Connectedness Hypothesis**

- Each community corresponds to a connected subgraph

- **Density Hypothesis**

- Nodes in a community are more likely to connect to other members of the same community than to nodes in other communities.

Connectedness and Density Hypothesis



Maximum Cliques

- Community as a group of individuals whose members all know each other
 - i.e. a complete graph or a Clique
- Does a Clique satisfy our hypothesis of connectedness and density ???
- However, there are drawbacks
 - While triangles are frequent in networks, **larger cliques are rare.**
 - Requiring a community to be a complete subgraph may **be too restrictive**, missing many other legitimate communities.

Strong and Weak Communities

- Consider a connected subgraph C of N_C nodes in a network
- k_i^{int} : *internal degree* of node i
 - number of links that connect i to other nodes in C .
- k_i^{ext} : *external degree* of node i
 - number of links that connect i to the rest of the network.
- If $k_i^{ext}=0$, each neighbor of i is within C , hence C is a good community for node i .
- If $k_i^{int}=0$, then node i should be assigned to a different community.

Strong and Weak Communities...

- **Strong Community**

- C is a *strong community* if each node within C has more links **within the community** than with the rest of the graph
- a subgraph C forms a strong community if for each node $i \in C$,
 - $k_i^{int}(C) > k_i^{ext}(C)$

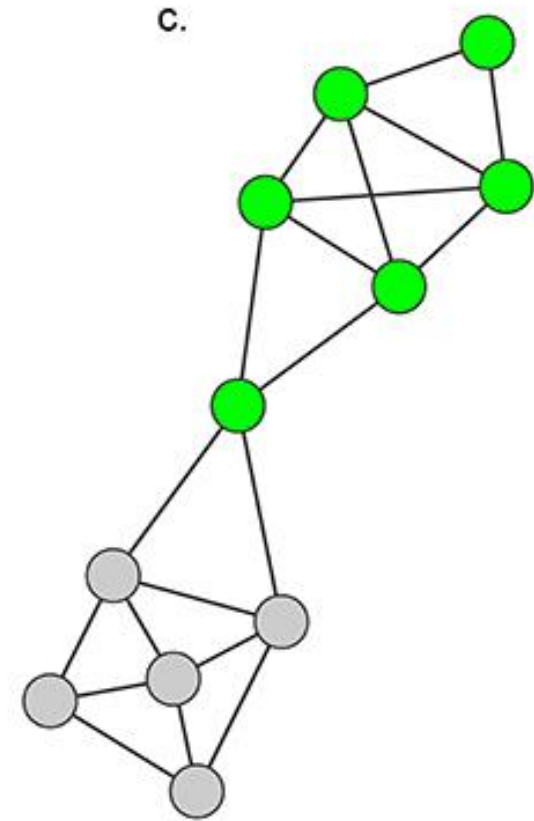
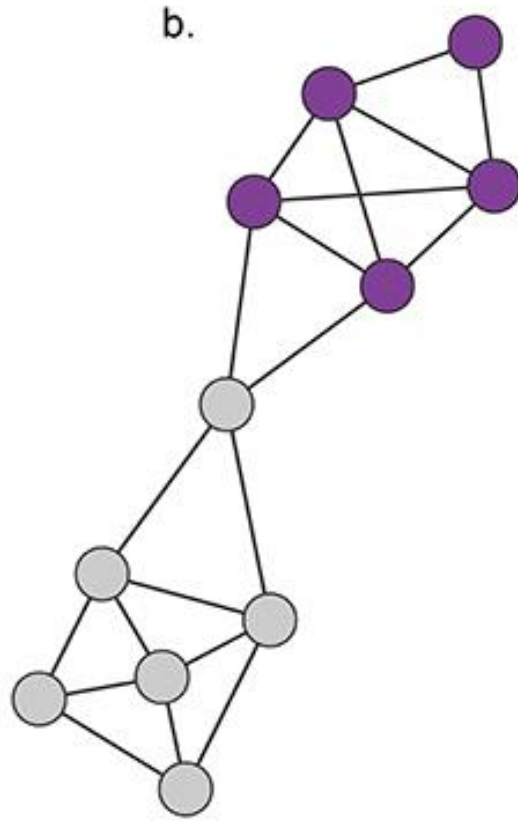
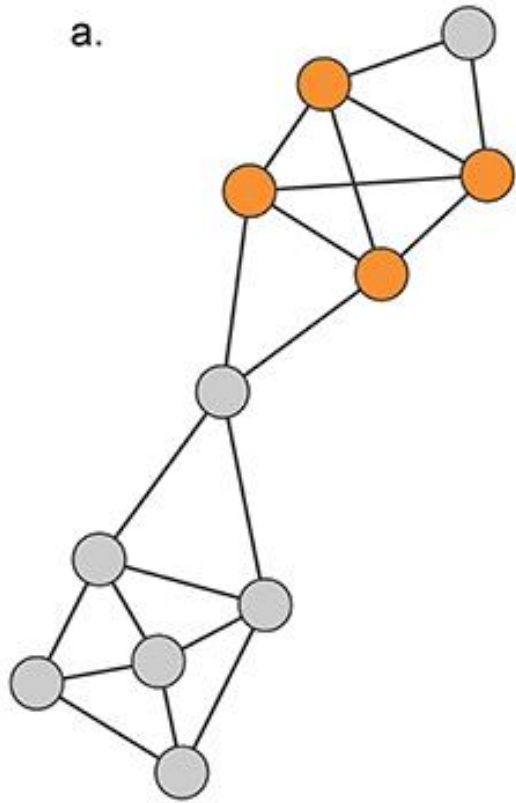
- **Weak Community**

- C is a *weak community* if the total internal degree of a subgraph exceeds its total external degree.
- a subgraph C forms a weak community if,
- $$\sum_{i \in C} k_i^{int}(C) > \sum_{i \in C} k_i^{ext}(C)$$

Strong and Weak Communities...

- Is clique a strong community?
- Is a strong community also a weak community?
- What about vice versa ?

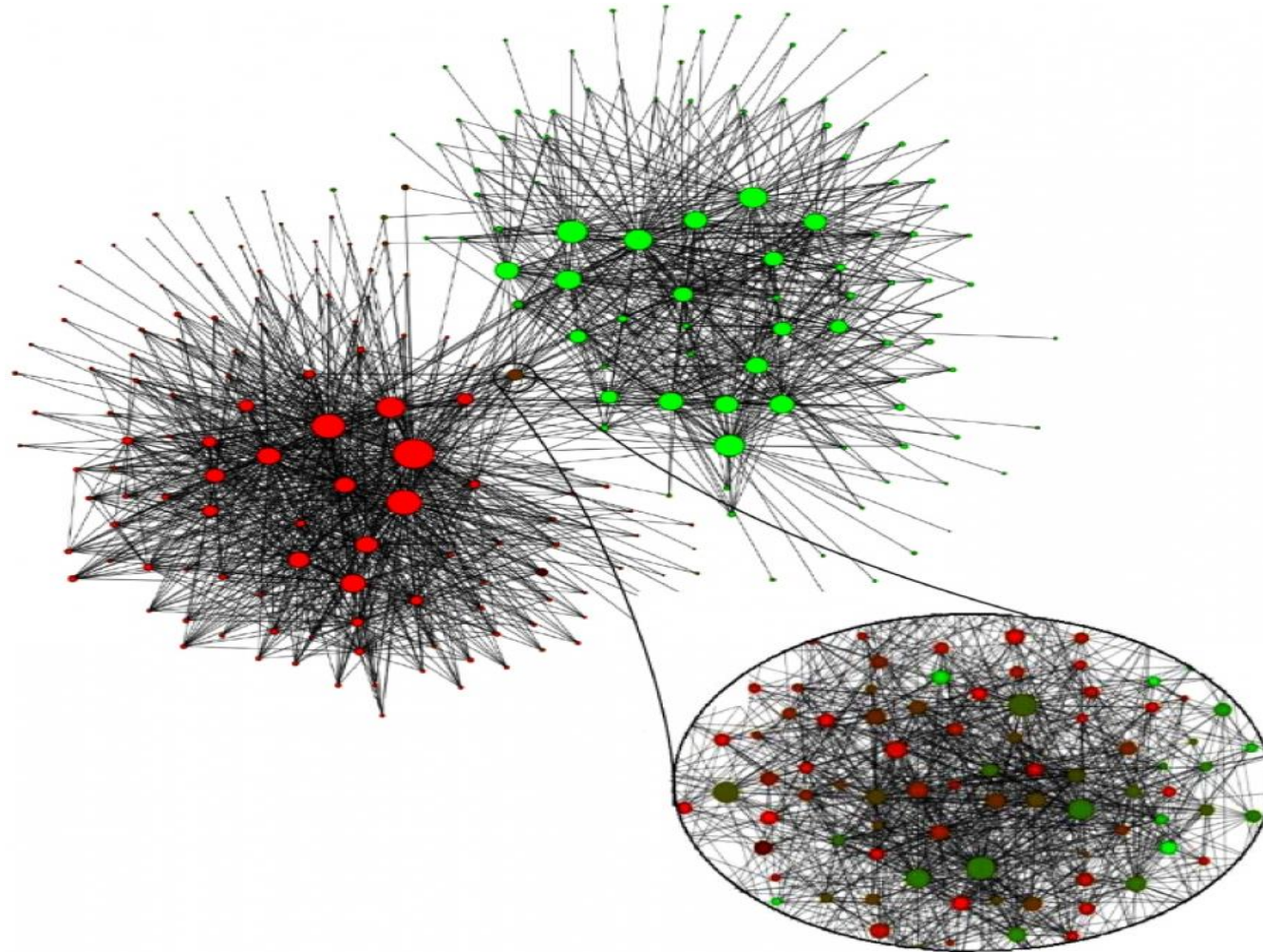
Strong and Weak Communities...



Communities in Belgium: A Case Study

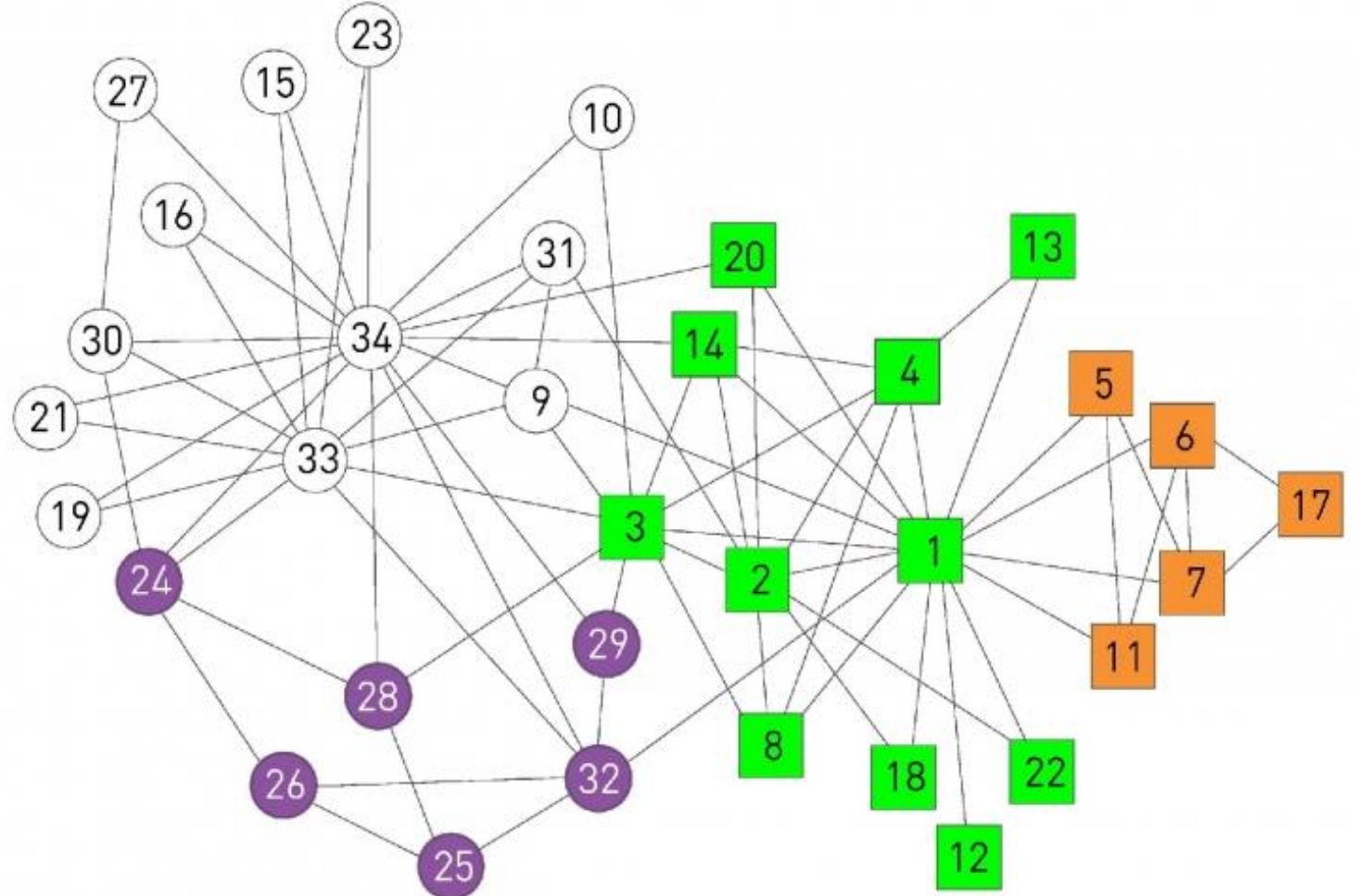
- Belgium, the model bicultural society: 59% of its citizens are Flemish, speaking Dutch and 40% are Walloons who speak French.
- What is the reason for the peaceful coexistence of these two ethnic groups since 1830 ?
 - Is it densely knitted society ?
 - Or we have two nations with the same borders, that learned to minimize contact with each other?
- Answer to the above question:
 - Research by Vincent Blondel and his students in 2007, who developed an algorithm to identify the **country's community structure**.

Communities extracted from the call pattern of the consumers of the largest Belgian mobile phone company



Community Detection : Major Application Areas

- Social Networks
 - E.g Zachary's Karate Club
- Biological Networks



Number of Communities

- How many ways can we group the nodes of a network into communities?
- Graph Bisection
 - Divide a network into two non-overlapping subgraphs, such that the number of links between the nodes in the two groups, called the *cut size*, is minimized
- Graph Partitioning
 - inspecting all possible divisions into two groups and choosing the one with the smallest cut size
 - number of distinct ways we can partition a network of N nodes into groups of N_1 and N_2 nodes is,

Number of Communities...

Number of distinct ways we can partition a network of N nodes into groups of N_1 and N_2 nodes is,

$$\frac{N!}{N_1! N_2!}$$

Using Stirling's formula

$$n! \approx 2\pi n (n/e)^n$$

For two equal sizes of N_1 and N_2 ,

$$\frac{N!}{N_1! N_2!} = e^{(N+1) \ln 2 - \frac{1}{2} \ln N}$$

The number of bisections increases exponentially with the size of the network.

Number of Communities...

- Consider a network with 10 nodes which we bisect into two subgraphs of size $N_1 = N_2 = 5$
 - What are the possible number of bisections?
- Now consider a network with 100 nodes with two subgraphs of size $N_1 = N_2 = 50$
 - What are the possible number of bisections?
- What are your observations from the above results???
- Is the Brute force approach feasible to compute graph bisection even for a modest size of network?

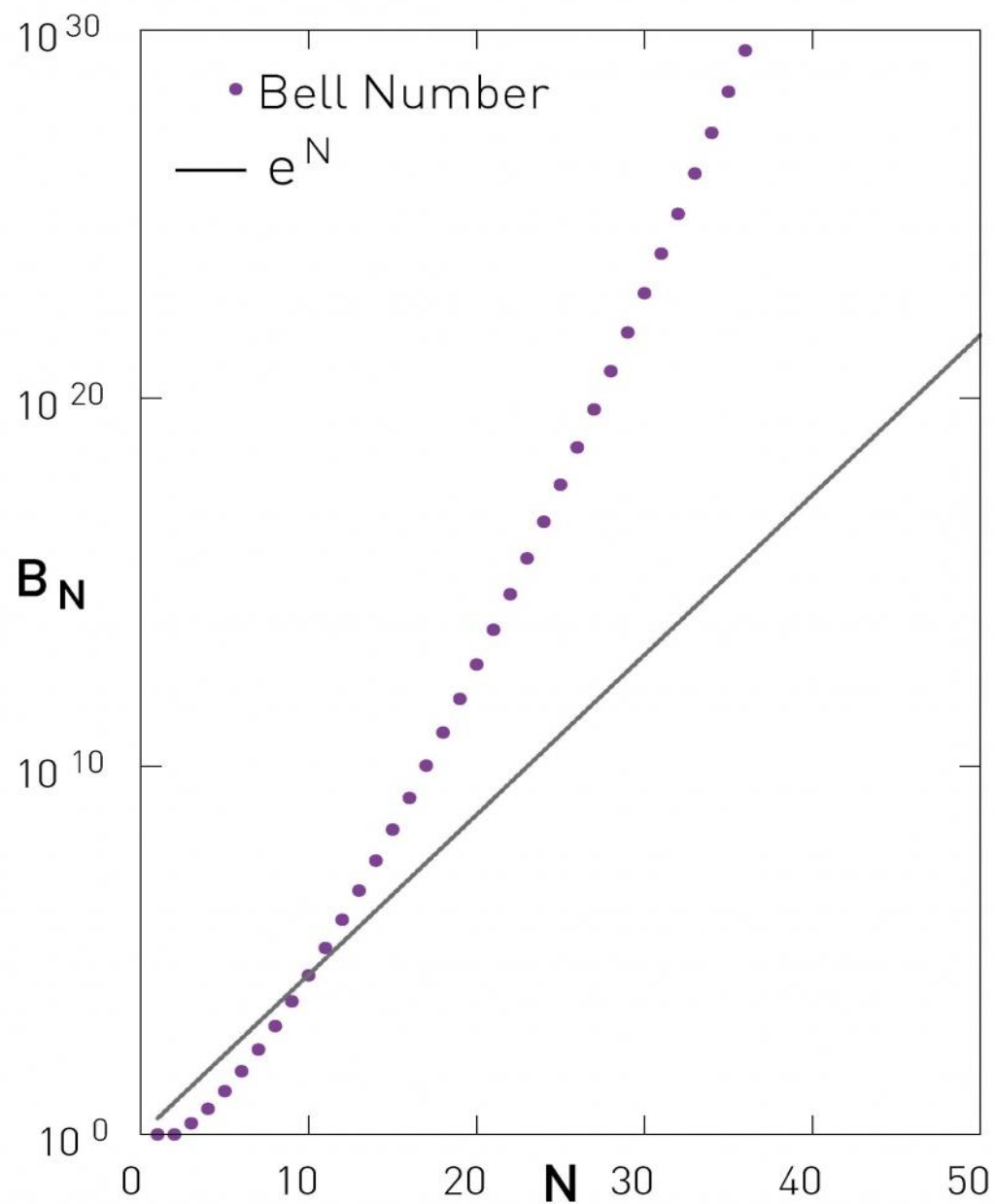
Difference between Graph Partitioning and Community Detection

- Graph partitioning divides a network into a **predefined** number of smaller subgraphs.
- In contrast community detection aims to **uncover** the inherent community structure of a network.
- Consequently in most community detection algorithms the **number and the size of the communities is not predefined**, but needs to be discovered by inspecting the network's wiring diagram.

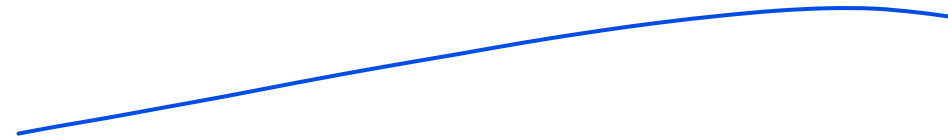
Community Detection : Non-overlapping

- Divide a network into an arbitrary number of groups, such that each node belongs to one and only one group
- The number of possible partitions are given by **Bell Number** as shown below:

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



We therefore need polynomial time algorithms that can identify communities without inspecting all partitions



Hierarchical Clustering

- 1) Calculate *Similarity matrix*, whose elements x_{ij} indicate the distance of node i from node j .
- 2) Iteratively identify groups of nodes with high similarity
 - 1) **Agglomerative algorithms** merge nodes with high similarity into the same community
 - 2) **Divisive algorithms** isolate communities by removing low similarity links that tend to connect communities.
- 3) Outcome: a hierarchical tree, called a dendrogram, that predicts the possible community partitions

Agglomerative Procedures: the Ravasz Algorithm

- Step 1: Define the Similarity Matrix
- Step 2: Decide Group Similarity
- Step 3: Apply Hierarchical Clustering
- Step 4: Dendrogram

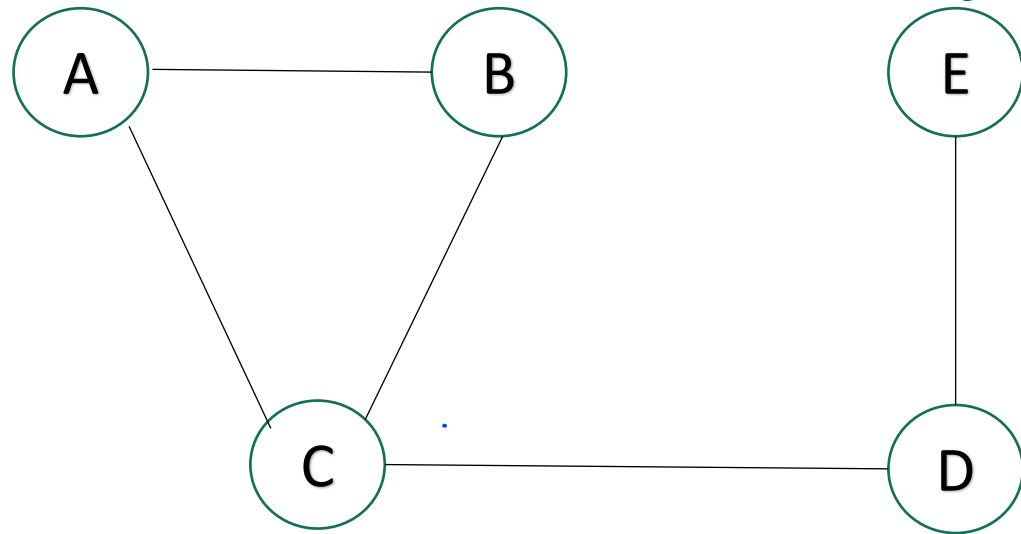
Ravasz Algorithm : Step 1-Similarity Matrix

- The topological overlap matrix,

$$x_{ij}^0 = \frac{j(i, j)}{\min(k_i, k_j) + 1 - \theta(A_{ij})}$$

- $\Theta(x)$ is the Heaviside step function, which is zero for $x \leq 0$ and one for $x > 0$;
- $J(i, j)$ is the number of common neighbors of node i and j , to which we add one (+1) if there is a direct link between i and j ;
- $\min(k_i, k_j)$ is the smaller of the degrees k_i and k_j

Ravasz Algorithm : Step 1-Similarity Matrix...



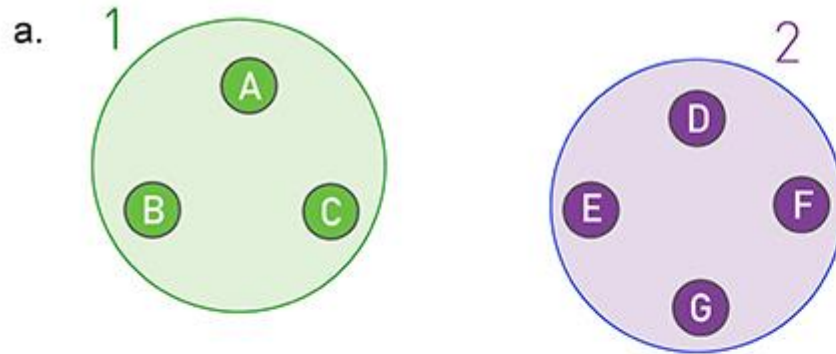
$$x_{ij}^0 = \frac{j(i,j)}{\min(k_i, k_j) + 1 - \theta(A_{ij})}$$

$$\begin{matrix}
 & \text{B} & \text{C} & \text{D} & \text{E} \\
 \begin{matrix} \text{A} \\ \text{B} \\ \text{C} \\ \text{D} \\ \text{E} \end{matrix} & \begin{bmatrix} - & 1 & 1 & 1/3 & 0 \\ & - & 1 & 1/3 & 0 \\ & & - & 1/2 & 1/2 \\ & & & - & 1 \\ & & & & - \end{bmatrix}
 \end{matrix}$$

Ravasz Algorithm : Step 2-Decide Group Similarity

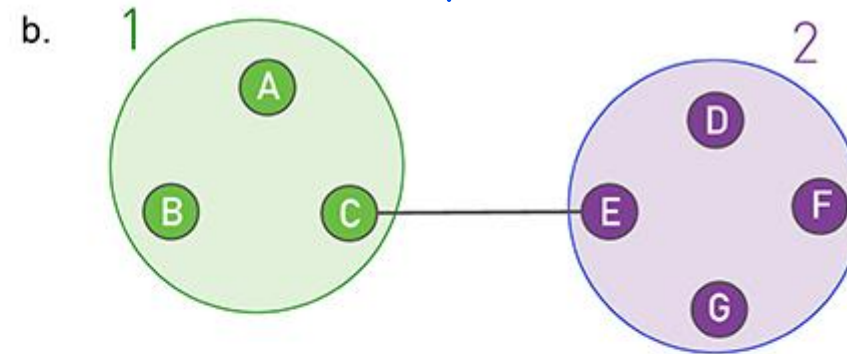
- We need to determine the similarity of two communities from the node similarity matrix x_{ij}
- Single Linkage Clustering
 - The similarity between communities 1 and 2 is the smallest of all x_{ij} , where i and j are in different communities.
- Complete Linkage Clustering
 - The similarity between two communities is the maximum of x_{ij} , where i and j are in distinct communities.
- Average Linkage Clustering
 - The similarity between two communities is the average of x_{ij} over all node pairs i and j that belong to different communities.
 - Ravasz algorithm uses this procedure

Ravasz Algorithm : Step 2-Decide Group Similarity

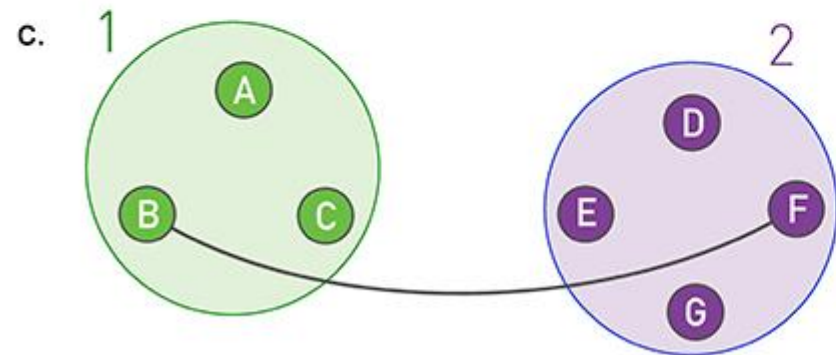


$$x_{ij} = r_{ij} =$$

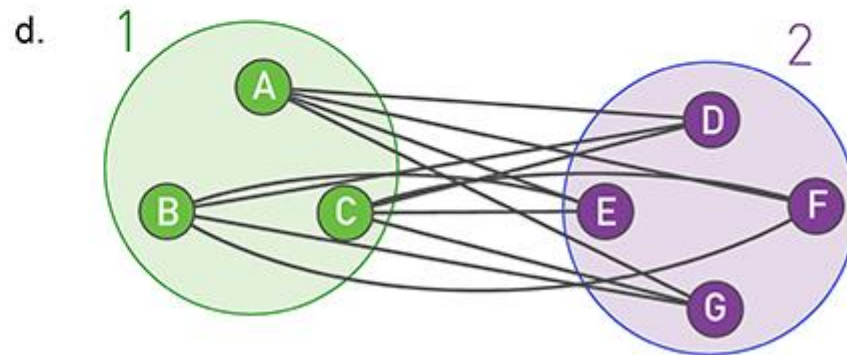
	D	E	F	G
A	2.75	2.22	3.46	3.08
B	3.38	2.68	3.97	3.40
C	2.31	1.59	2.88	2.34



Single Linkage: $x_{12} = 1.59$



Complete Linkage: $x_{12} = 3.97$



Average Linkage: $x_{12} = 2.84$

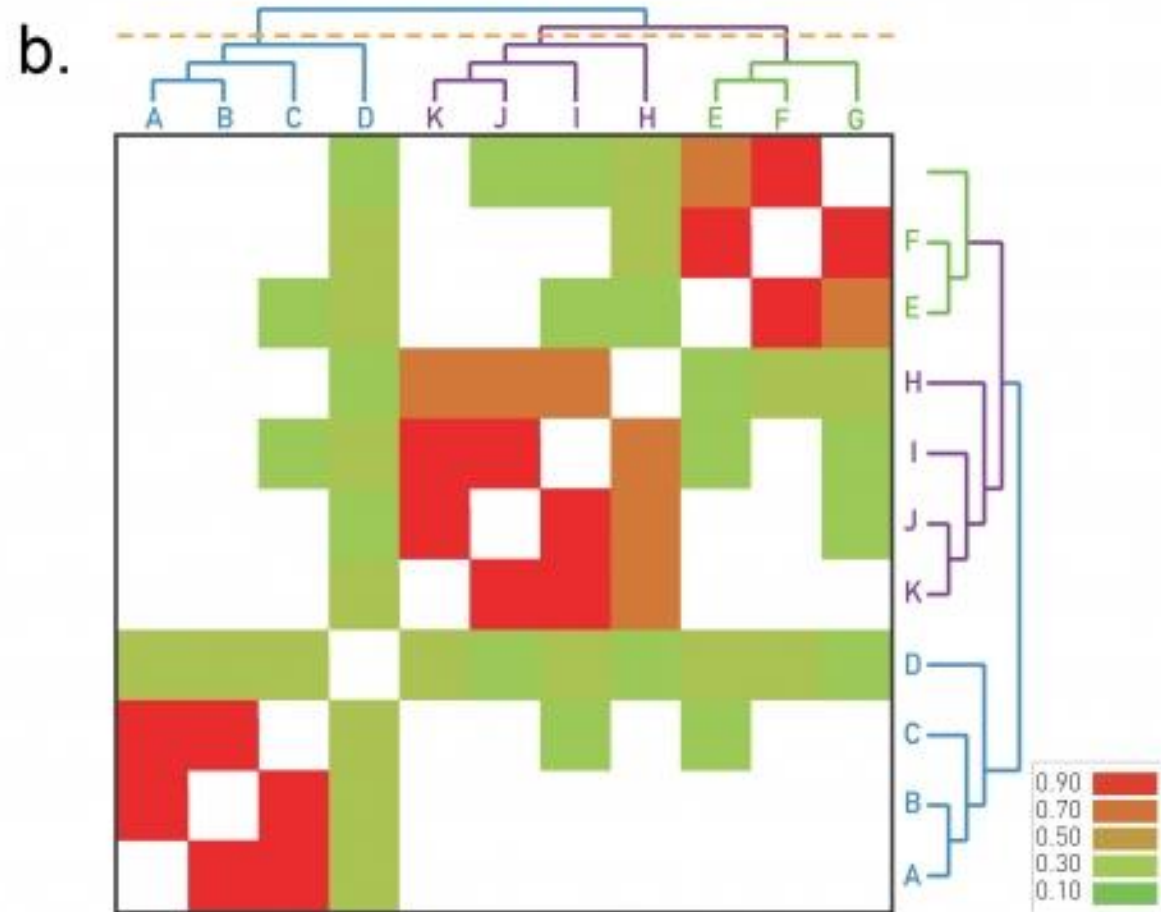
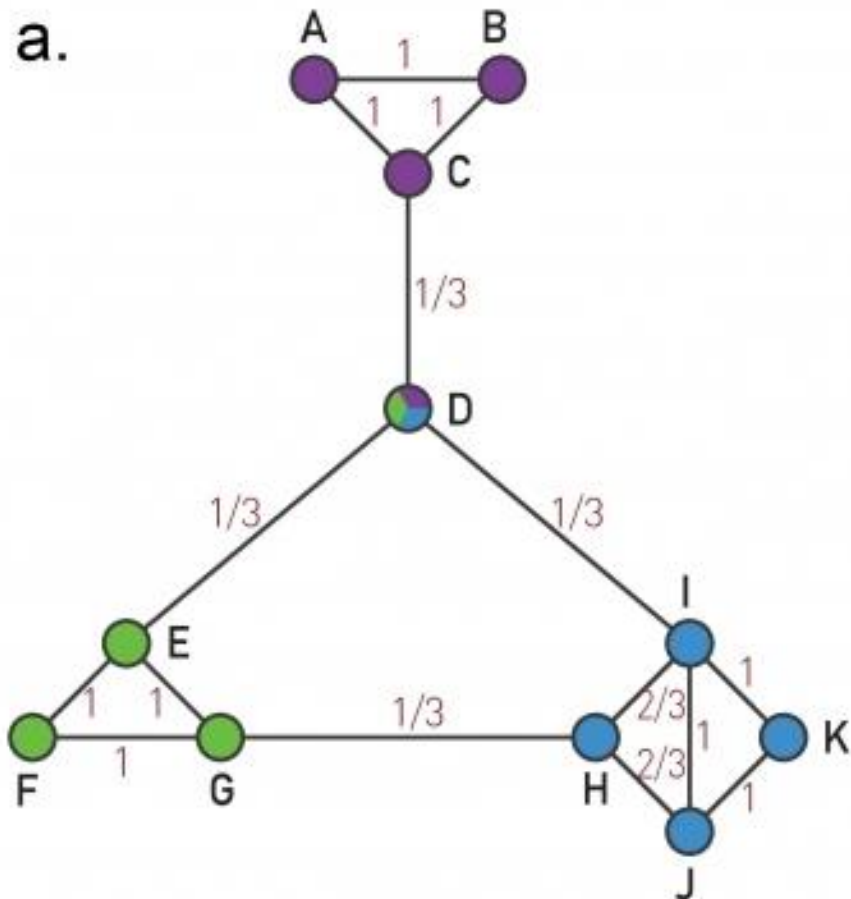
Ravasz Algorithm : Step 3: Apply Hierarchical Clustering

1. Assign each node to a community of its own and evaluate x_{ij} for all node pairs.
2. Find the community pair or the node pair with the highest similarity and merge them into a single community.
3. Calculate the similarity between the new community and all other communities.
4. Repeat Steps 2 and 3 until all nodes form a single community.

Ravasz Algorithm : Step 4-Dendrogram

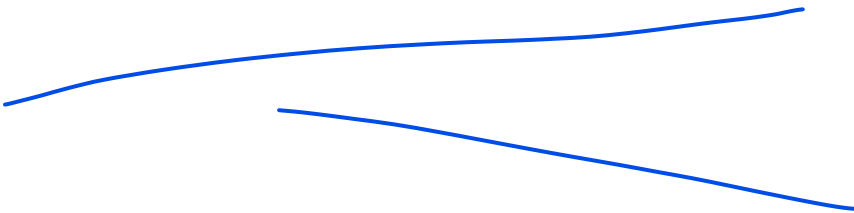
- To extract the underlying community organization
 - By cutting the Dendrogram
- The dendrogram visualizes the order in which the nodes are assigned to specific communities.

Agglomerative Procedures: the Ravasz Algorithm



Ravas Algorithm : Computational Complexity

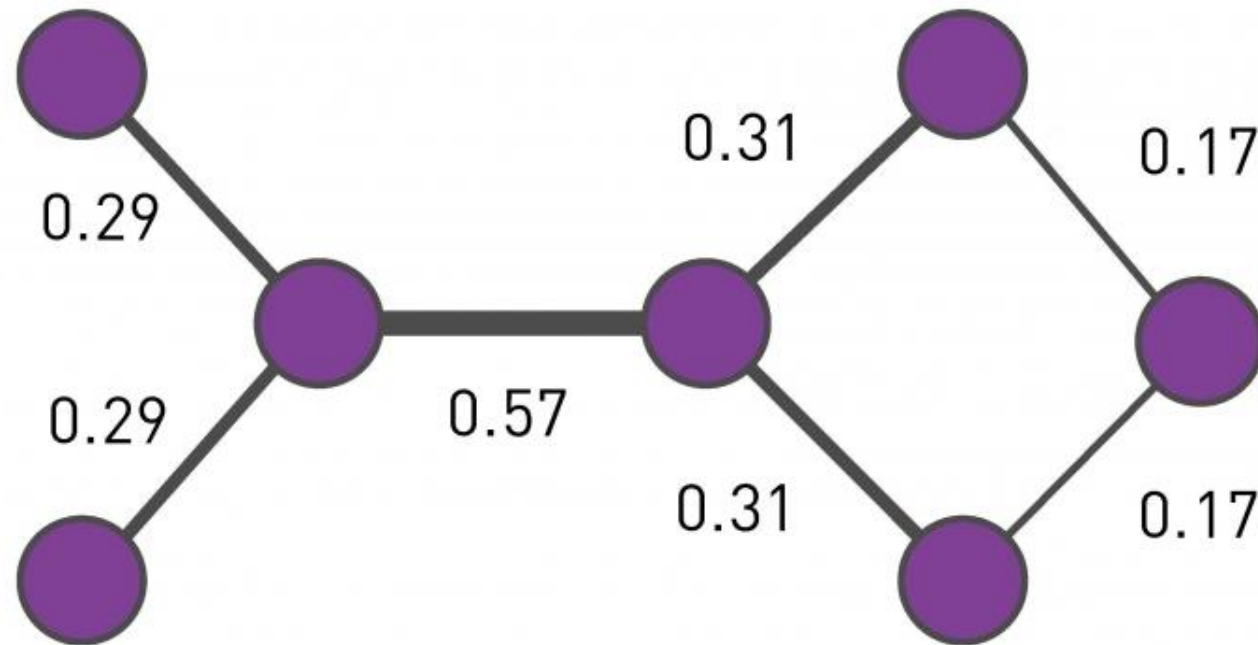
- Exercise: Is it a polynomial time algorithm?



Divisive Procedures: The Girvan-Newman algorithm

- Idea: Systematically remove the links connecting nodes that belong to different communities, eventually breaking a network into isolated communities.
- Step 1: Define Centrality X_{ij} using **Link Betweenness**
 - i.e the number of shortest paths that pass through link (i, j)
 - Large X_{ij} for the links connecting nodes in different communities
 - Small X_{ij} for the links connecting nodes in same community

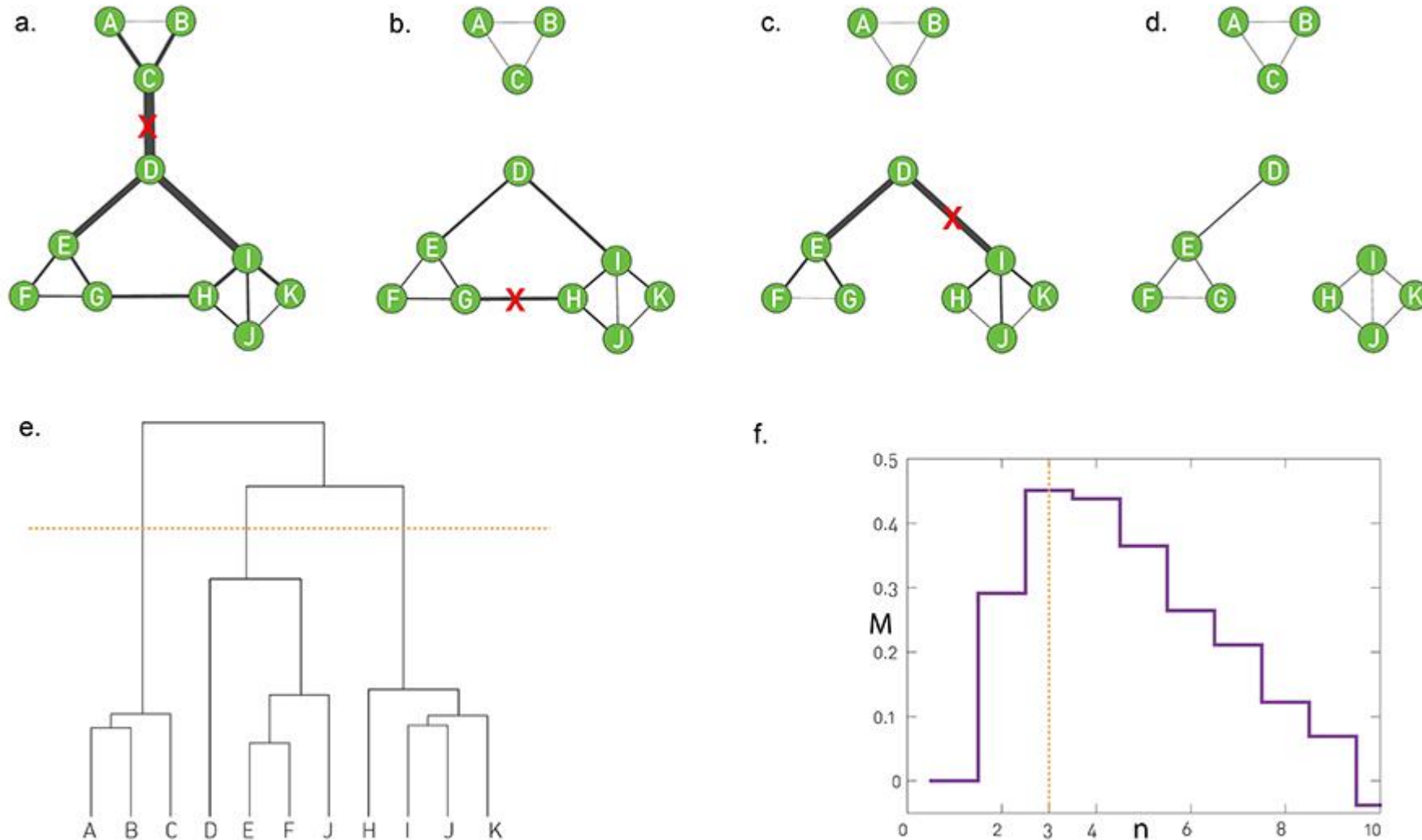
Divisive Procedures: The Girvan-Newman algorithm



Divisive Procedures: The Girvan-Newman algorithm

- Step 2: Hierarchical Clustering
 - Compute the centrality x_{ij} of each link.
 - Remove the link with the largest centrality. In case of a tie, choose one link randomly.
 - Recalculate the centrality of each link for the altered network.
 - Repeat steps 2 and 3 until all links are removed.

Divisive Procedures: The Girvan-Newman algorithm



Modularity

- How do we decide which of the many partitions predicted by a hierarchical method offers the best community structure?
- Selecting the one for which **modularity** is maximal.
- Measures the quality of each partition.
- Allows us to decide if a particular community partition is better than some other one.
- Modularity optimization offers a novel approach to community detection.

Modularity...

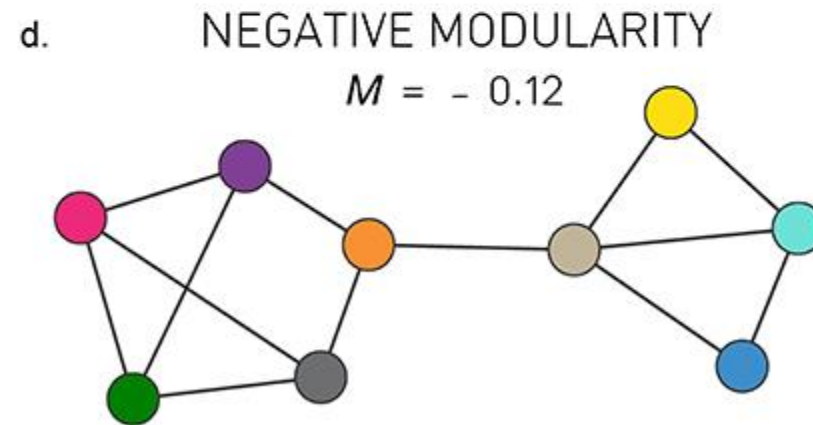
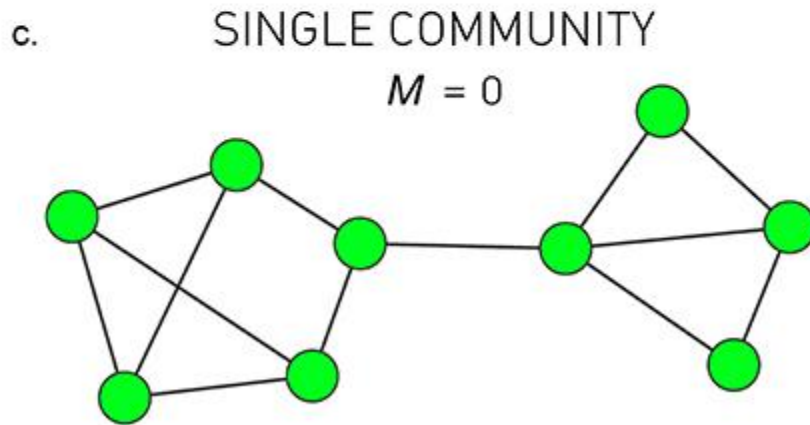
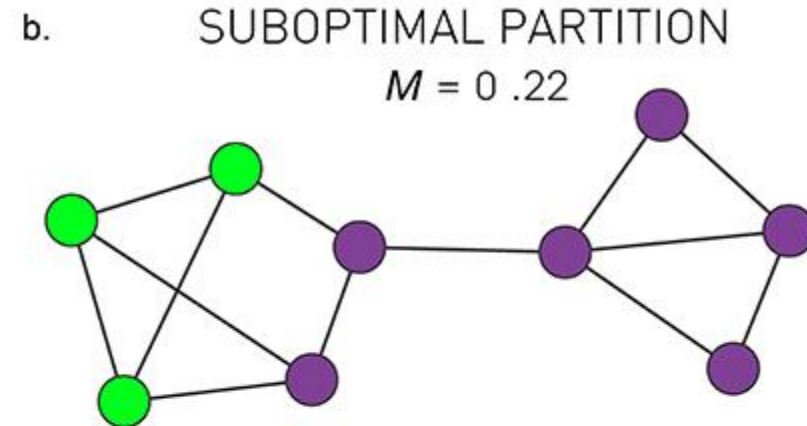
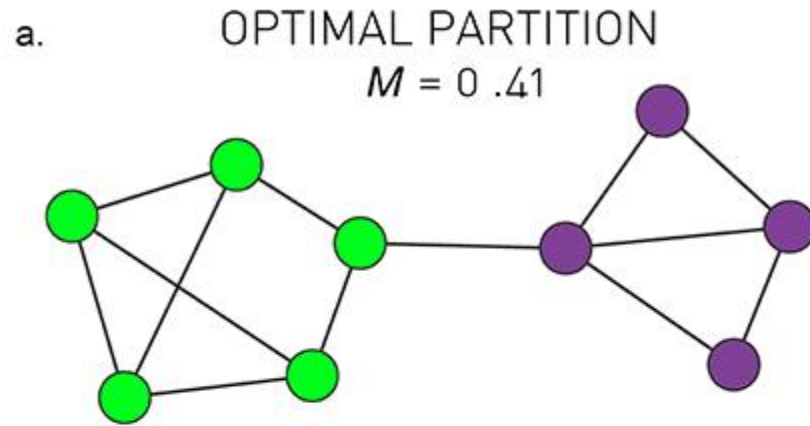
- Consider the following scenario
- A network with N nodes and L links
- Network is partitioned into n_c number of communities
- Each community has N_c nodes and L_c links
- If L_c is larger than the expected number of links between the N_c nodes given the network's degree sequence, then the nodes of the subgraph C_c could indeed be part of a true community.
- We therefore measure the difference between the network's real wiring diagram (A_{ij}) and the expected number of links between i and j if the network is randomly wired (p_{ij})

Modularity...

$$M = \sum_{c=1}^{n_c} \left[\frac{L_c}{L} - \left(\frac{k_c}{2L} \right)^2 \right]$$

- L : total number of links in Graph
- n_c : total number of communities in the graph
- L_c : total number of links in community c
- k_c : total degree of nodes in community c

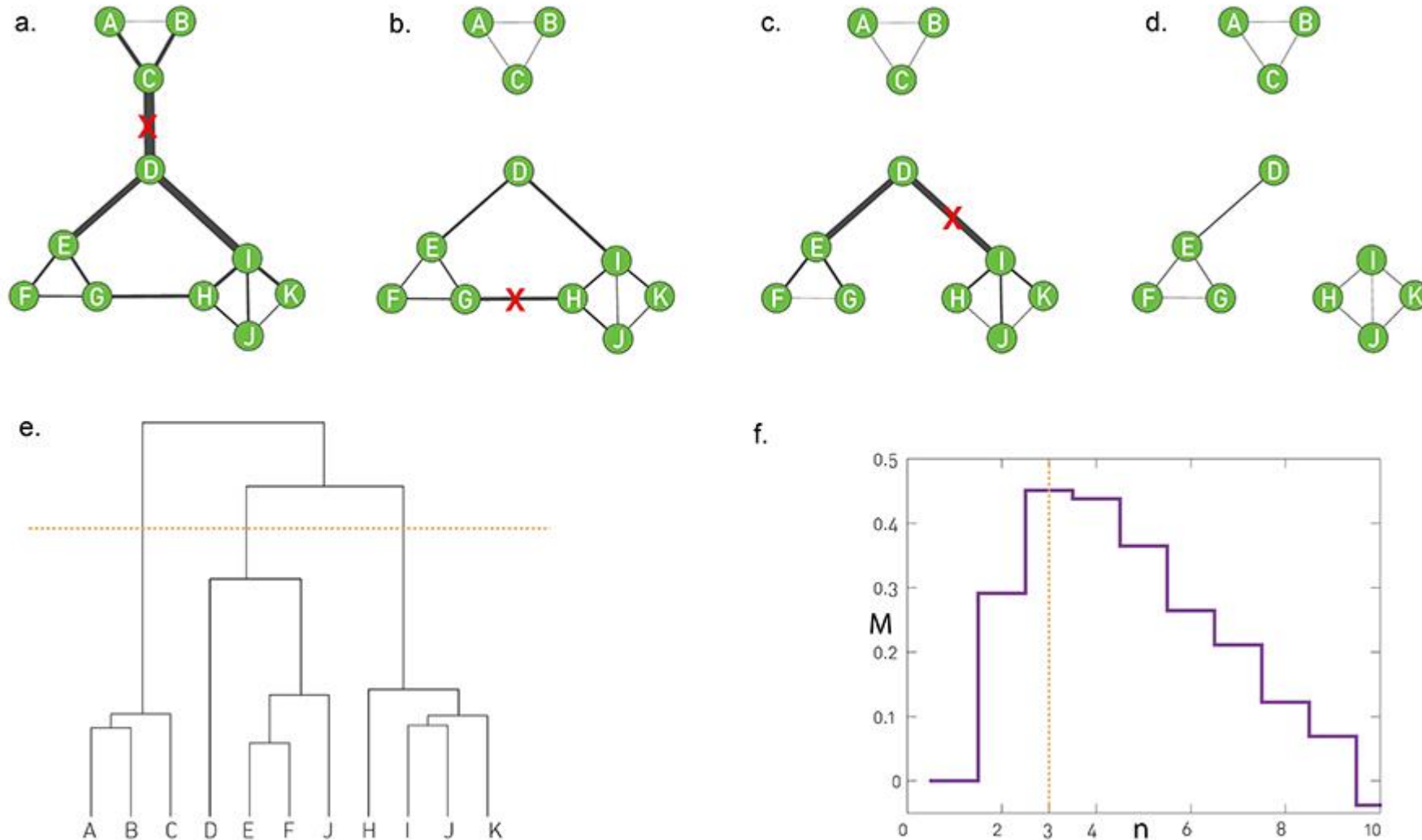
Modularity...



Modularity...

- Optimal Partition
 - The partition with maximal modularity $M=0.41$ closely matches the two distinct communities.
- Suboptimal Partition
 - A partition with a sub-optimal but positive modularity, $M=0.22$, fails to correctly identify the communities present in the network.
- Single Community
 - If we assign all nodes to the same community we obtain $M=0$, independent of the network structure.
- Negative Modularity
 - If we assign each node to a different community, modularity is negative, obtaining $M=-0.12$.

Divisive Procedures: The Girvan-Newman algorithm



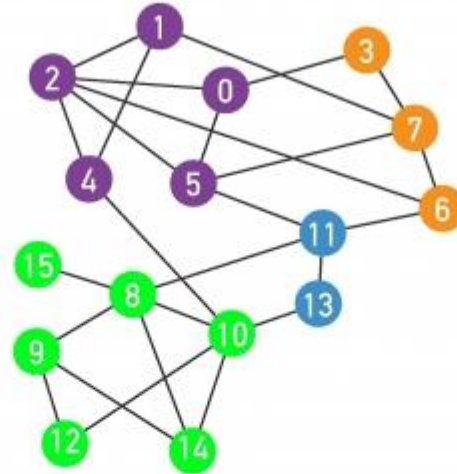
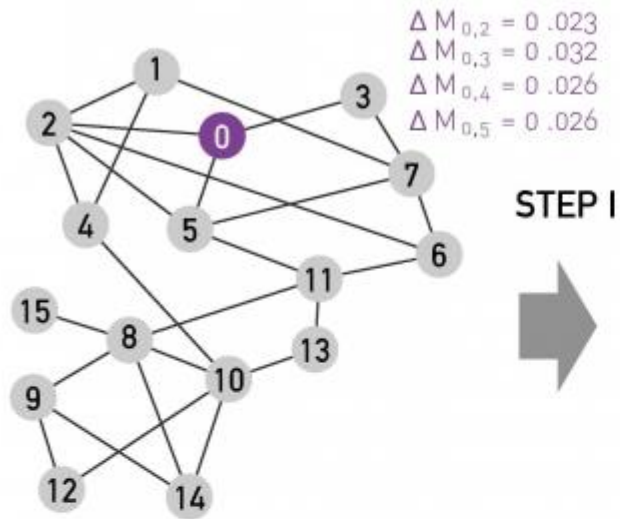
The Louvain Algorithm

- Greedy algorithm for Community Detection
 - $O(n \log n)$ run time
- Supports weighted graphs
- Provide hierarchical communities
- Widely utilized to study large networks because
 - Fast
 - Rapid Convergence
 - High modularity output (i.e. better communities)

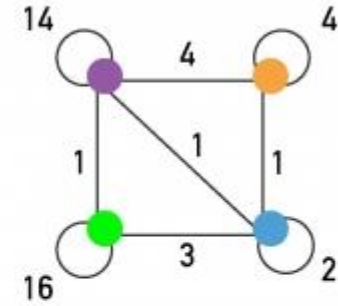
The Louvain Algorithm...

- Operates in several iterations
- Each iteration consists of 2 phases
 - **Phase 1:** Modularity is optimized by allowing only local changes to node-communities memberships
 - **Phase 2:** The identified communities are aggregated into super nodes to build a new network
- Goto Phase 1 until no increase in modularity is possible

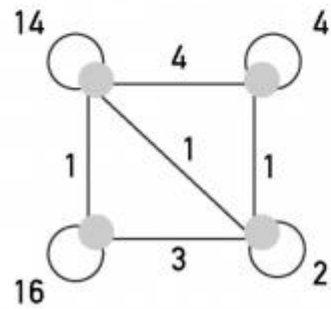
1ST PASS



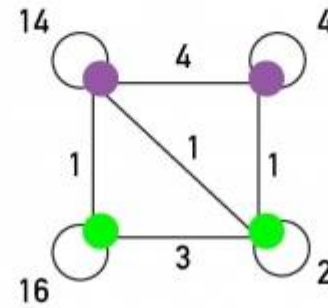
STEP II



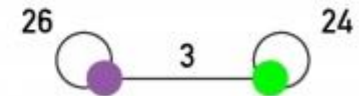
2ND PASS



STEP I



STEP II



Community Detection through Modularity Maximization: Limitations

1) Resolution limit:

- well-connected smaller communities tend to get merged with larger communities even if the resultant communities are not that dense
- fails to detect those communities which are well-separated with densely connected intra-community nodes but only a single inter-community edge with the rest of the network

2) Degeneracy of solutions:

- the case when there is an exponential number of community structures with same (maximum) modularity value