

Algorithms and Data Structures 2, 340300 Lecture – 2023W Univ.-Prof. Dr. Alois Ferscha, teaching@pervasive.jku.at

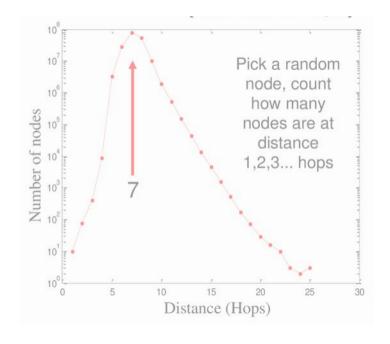
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Remember: Small-World Effect

- Six Degrees of Separation
- A famous experiment conducted by Travers and Milgram (1969)
 - Subjects were asked to send a chain letter to his acquaintance in order to reach a target person
 - The average path length is around **5.5**

Verified on a planetary-scale IM network (Microsoft Messenger) of 240 million people (Leskovec and Horvitz 2008) and 30 billion conversations

- 180 million nodes
- 1.3 billion undirected edges
- The average path length is 6.6



J. Leskovec and E. Horvitz. Planetary-scale views on a large instant-messaging network. In Proceedings of the 17th international conference on World Wide Web, WWW '08, pages 915–924, New York, NY, USA, 2008. ACM

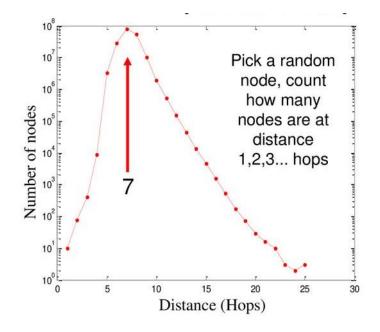


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Communities in Social Media

Community

Community: It is formed by individuals such that those **within** a group interact with each other **more frequently** than with those **outside** the group

a.k.a. group, cluster, cohesive subgroup, module in different contexts

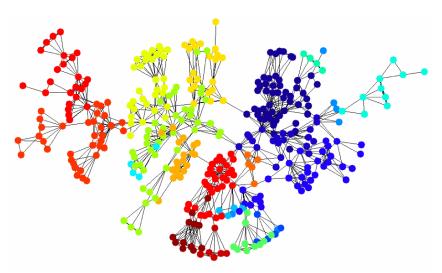
Community detection: discovering groups in a network where individuals' group **memberships are not explicitly given**

Why communities in social media?

- Human beings are social
- Easy-to-use social media allows people to extend their social life in unprecedented ways
- Difficult to meet friends in the physical world, but much easier to find friend online with similar interests

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Interactions between nodes can help determine communities



Example: community network in social media



Communities in Social Media

Communities in Social Media

Two types of groups in social media

- Explicit Groups: formed by user subscriptions
- Implicit Groups: implicitly formed by social interactions

Some social media sites allow people to join groups,

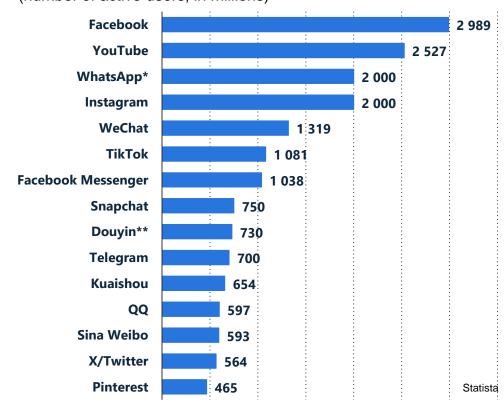
- extract groups based on network topology?
 - Not all sites provide a community platform
 - Not all people want to make effort to join groups
 - Groups can change dynamically

Network interaction provides rich information about the relationship between users

- Can complement other kinds of information
- Help network visualization and navigation
- Provide basic information for other tasks

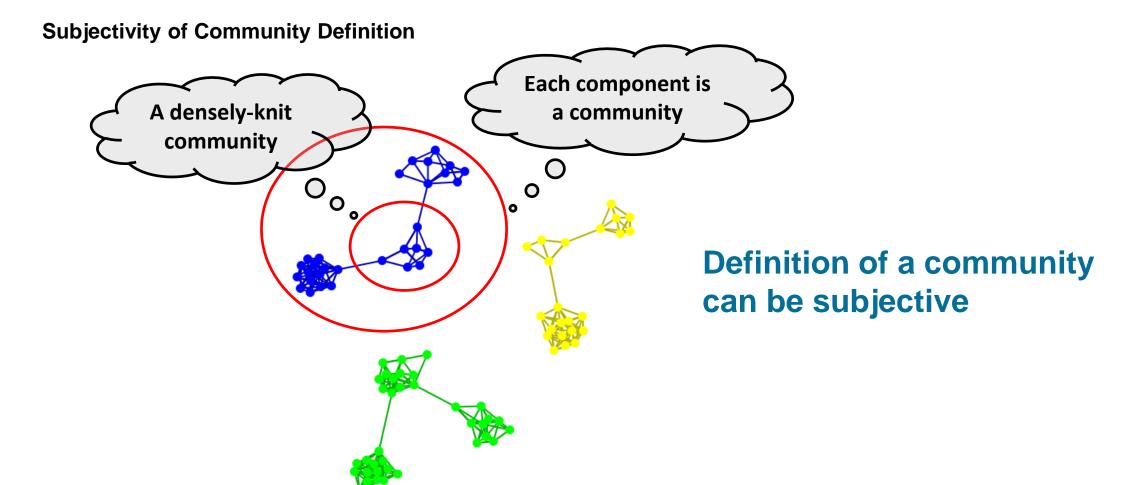
Example of explicit groups:

Most popular social networks worldwide, July 2023 (number of active users, in millions)





Community Definition





Community Detection Methods

Taxonomy of Community Criteria

Criteria vary depending on the tasks

Roughly, **community detection methods** can be divided into 4 categories (not exclusive):

Node-Centric Community

Each node in a group satisfies certain properties

Group-Centric Community

Consider the connections within a group as a whole.
 The group has to satisfy certain properties without zooming into node-level

Network-Centric Community

Partition the whole network into several disjoint sets

Hierarchy-Centric Community

Construct a hierarchical structure of communities



Community Metrics

Measures used to calibrate the small world effect:

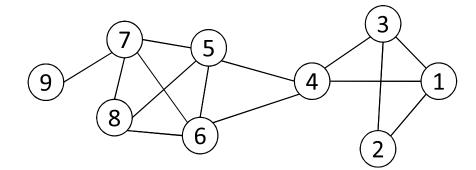
Diameter

Measures used to calibrate the small world effect

- **Diameter**: the (maximum) longest shortest path in a network
- Average shortest path length



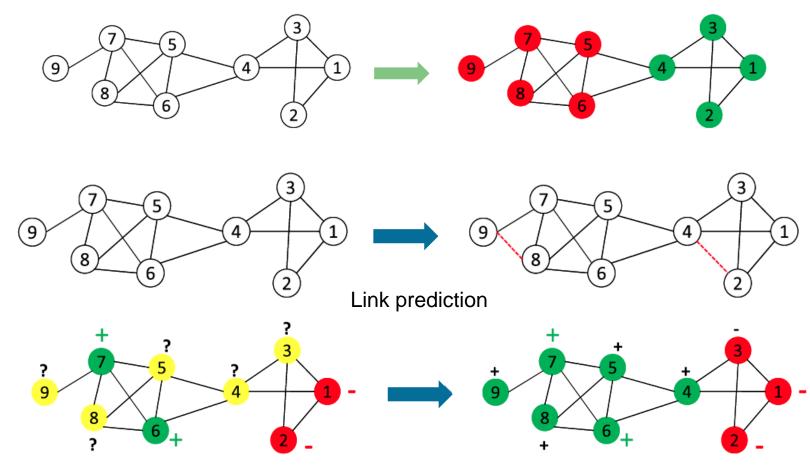
- The number of **hops** in the geodesic is the **geodesic distance**.
- The **geodesic distance** between node 1 and node 9 is **4**.
- The **diameter** of the network is **5**, corresponding to the geodesic distance between nodes 2 and 9.





Community Dynamics

A community are people in a group interacting with each other more frequently than those outside the group





Density of connections

- Friends of a friend are likely to be friends as well
 - density of connections among one's friends

d ... number of neighbours

k ... connections among neighboured friends

Example:

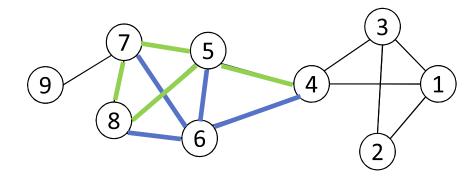
$$d_6$$
=4, N_6 = {4, 5, 7,8}
 k_6 =4 as e(4,5), e(5,7), e(5,8), e(7,8)
 C_6 = 4/(4*3/2) = 2/3

Average clustering coefficient

$$C = (C_1 + C_2 + ... + C_n)/n$$

C = 0.61 for the given network

$$C_{i} = \begin{cases} \frac{k_{i}}{d_{i} \times (d_{i}-1)/2} & d_{i} > 1\\ 0 & d_{i} = 0 \text{ or } 1 \end{cases}$$



Centrality

Centrality in Graphs assign numbers or rankings to nodes within a graph corresponding to their network position.

Applications

- identifying the most influential person(s) in a social network
- **key infrastructure nodes** in the Internet or urban networks
- super-spreaders of disease in pandemics
- cell networks in brains

Degree Cetrality

number of links incident to a node

Betweenness Cetrality (of a node)

• quantifies the **number of times** a node acts as a **connector/bridge along the shortest** path between two other nodes (the **number** of **shortest paths** that **pass** one node)

Eigenvector Cetrality (of a node)

assigns relative scores to all nodes in the network based on the concept that connections to high-scoring nodes
contribute more to the score of the node in question than equal connections to low-scoring nodes

Cross-Clique Cetrality

- determines the connectivity of a node to different cliques
- a node with **high cross-clique connectivity** facilitates the **propagation** of information (or diseases) in a graph
- (Cliques are subgraphs in which every node is connected to every other node in the clique).



Degree of Centrality

The **importance** of a node is determined by the **number** of **nodes adjacent** to it

- The larger the degree, the more import the node is
- Only a small number of nodes have high degrees in many real-life networks

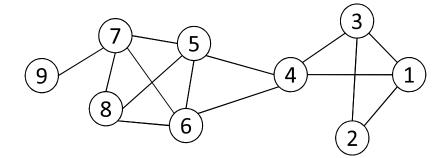
Degree Centrality

$$C_D(v_i) = d_i = \sum_j A_{ij}$$

Normalized Degree Centrality

$$C_D'(v_i) = d_i/(n-1)$$

- For node 1, degree centrality is 3;
- Normalized degree centrality is 3/(9-1)=3/8



Closeness Centrality

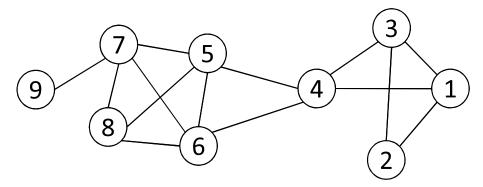
- "Central" nodes are important, as they can reach the whole network more quickly than non-central nodes
- Importance measured by how close a node is to other nodes

$$D_{avg}(v_i) = \frac{1}{n-1} \sum_{j \neq i}^{n} g(v_i, v_j)$$

$$C_C(v_i) = \left[\frac{1}{n-1} \sum_{j \neq i}^n g(v_i, v_j) \right]^{-1} = \frac{n-1}{\sum_{j \neq i}^n g(v_i, v_j)}$$



Closeness Centrality: Example



$$C_c(3) = \frac{9-1}{1+1+1+2+2+3+3+4} = \frac{8}{17} = 0.47$$

$$C_c(4) = \frac{9-1}{1+2+1+1+1+2+2+3} = \frac{8}{13} = 0.62$$

Table 2.1: Pairwise geodesic distance									
Node	1	2	3	4	5	6	7	8	9
1	0	1	1	1	2	2	3	3	4
2	1	0	1	2	3	3	4	4	5
3	1	1	0	1	2	2	3	3	4
4	1	2	1	0	1	1	2	2	3
5	2	3	2	1	0	1	1	1	2
6	2	3	2	1	1	0	1	1	2
7	3	4	3	2	1	1	0	1	1
8	3	4	3	2	1	1	1	0	2
9	4	5	4	3	2	2	1	2	0

Node 4 is more central than node 3



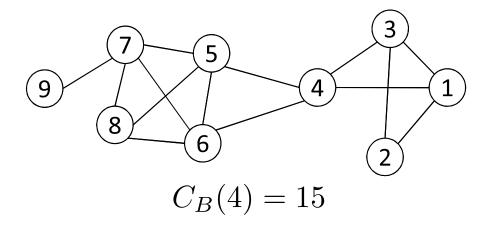
Betweenness Centrality

- Node betweenness counts the number of shortest paths that pass one node
- Nodes with high betweenness are important in communication and information diffusion

- Betweenness Centrality
$$C_B(v_i) = \sum_{v_s \neq v_i \neq v_t \in V, s < t} \frac{\sigma_{st}(v_i)}{\sigma_{st}}$$

 σ_{st} : The number of shortest paths between s and t

 $\sigma_{st}(v_i)$: The number of shortest paths between s and t that pass v_i



	$\sigma_{st}(4)/\sigma_{st}$						
	s = 1	s = 2	s = 3				
t = 5	1/1	2/2	1/1				
t = 6	1/1	2/2	1/1				
t = 7	2/2	4/4	2/2				
t = 8	2/2	4/4	2/2				
t = 9	2/2	4/4	2/2				

What's the betweenness centrality for node 5? (= 6)

 σ_{st} : The number of shortest paths between s and t

 $\sigma_{st}(v_i)$: The number of shortest paths between s and t that pass v_i

$$C_B(v_i) = \sum_{v_s \neq v_i \neq v_t \in V, s < t} \frac{\sigma_{st}(v_i)}{\sigma_{st}}$$



Eigenvalue (or Eigenvector) Centrality

- One's importance is determined by his friends'
- If one has many important friends, he should be important as well.

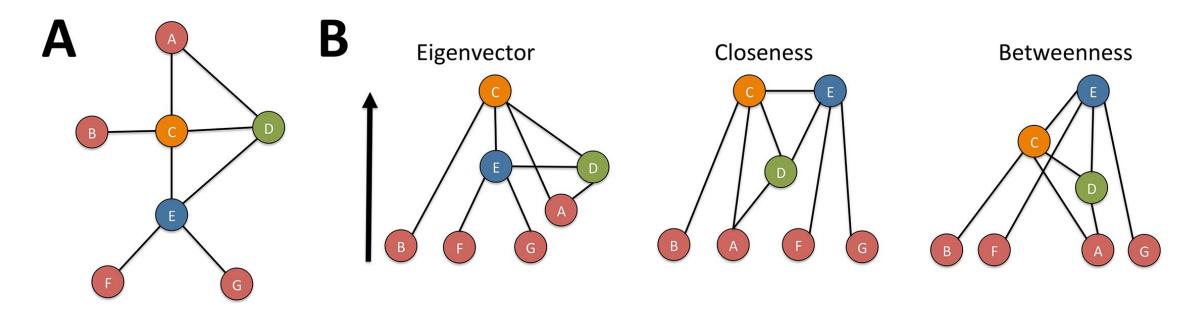
$$C_E(v_i) \propto \sum_{v_j \in N_i} A_{ij} C_E(v_j)$$

$$\mathbf{x} \propto A\mathbf{x}$$
 $A\mathbf{x} = \lambda \mathbf{x}$.

- The centrality corresponds to the top **eigenvector** of the **adjacency matrix A**.
- A variant of this eigenvector centrality is the PageRank score.



Community Analysis :: Importance (Closeness)



- (A) Example of a hypothetical social network illustrating the individual level metrics. Letters label individuals in the network.
- (B) Network is **restructured** in a **hierarchy** such that the **node with the highest relevant centrality measure** is on **top** following the arrow). For example, node **C** has the highest Eigenvector centrality, but node **E** had the highest betweenness centrality.



Community Analysis: Ties and Influence

Weak and Strong Ties

- In practice, connections are not of the same strength
- Interpersonal social networks are composed of strong ties (close friends) and weak ties (acquaintances).
- Strong ties and weak ties play different roles for community formation and information diffusion
- strong ties are transitive with high probability
- weak ties are not transitive (or with low probability)
- The Strength of Weak Ties (Granovetter, 1973)
 - Occasional encounters with distant acquaintances can provide important information about new opportunities for job search

Weak Ties Strong Ties

M. Granovetter. The Strength of Weak Ties. The American Journal of Sociology, 78(6):1360-1380, 1973.



Connections in Social Media

Social Media allows users to connect to each other more easily than ever

- One user might have thousands of friends online
- Who are the most important ones among your Facebook friends?

Imperative to **estimate** the **strengths** of **ties** for advanced analysis

- Analyze network topology
- Learn from User Profiles and Attributes
- Learn from User Activities

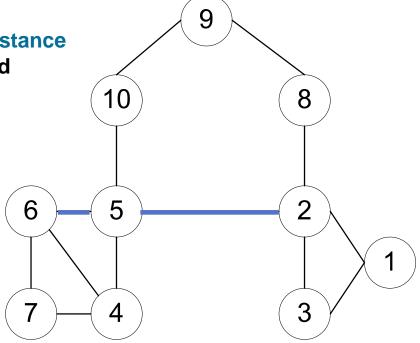


Community Analysis :: Network Topology

"shortcut" Bridge

- Bridges are rare in real-life networks
- Alternatively, one can relax the definition by checking if the distance between two terminal nodes increases if the edge is removed
- The larger the distance, the weaker the tie is

- d(2,5) = 4 if e(2,5) is removed
- d(5,6) = 2 if e(5,6) is removed
- e(5,6) is a **stronger** tie than e(2,5)



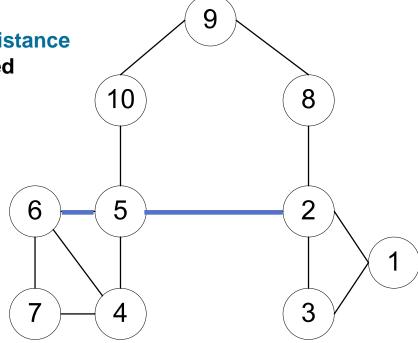


Community Analysis :: Network Topology

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Community Analysis :: Network Topology

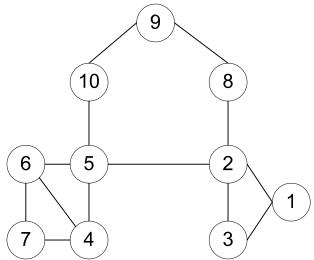
Neighbourhood Overlap

Tie strength can be measured based on neighborhood overlap; the larger the overlap, the stronger the tie is.

$$\begin{array}{ll} \textit{overlap}(v_i, v_j) & = & \frac{\text{number of shared friends of both } v_i \text{ and } v_j}{\text{number of friends who are adjacent to at least } v_i \text{ or } v_j} \\ & = & \frac{|N_i \cap N_j|}{|N_i \cup N_j| - 2}. \end{array}$$

-2 in the denominator is to exclude v_i and v_j

$$overlap(2, 5) = 0$$
,
 $overlap(5, 6) = \frac{|\{4\}|}{|\{2, 4, 5, 6, 7, 10\}| - 2} = 1/4$



Community Analysis :: User Activities

Learning from User Activities

- One might learn how one influences his friends if the user activity log is accessible
- Depending on the adopted influence model
 - Independent cascading model
 - Linear threshold model

Maximizing the likelihood of user activity given an influence model



Influence Modeling

Influence modeling is one of the fundamental questions in order to understand the information diffusion, spread of new ideas, and word-of-mouth (viral) marketing

Well known methods:

- Linear threshold model (LTM)
- Independent cascade model (ICM)

Common properties of influence modeling methods

- A social network is represented by a directed graph, with each actor being one node;
- Each node is started as active or inactive;
- A node, once activated, will activate his neighboring nodes;
- Once a node is activated, this node cannot be deactivated.



Linear Threshold Model (LTM)

An actor would take an action if the number of his friends who have taken the **action exceeds** (reaches) a certain **threshold**

- Each node *v* chooses a threshold Θ_v randomly from a uniform distribution in an interval between 0 and 1.
- In each discrete step, all nodes that were active in the previous step remain active
- The nodes satisfying the following condition will be activated

$$\sum_{w \in N_v, w \text{ is active}} b_{w,v} \ge \theta_v$$

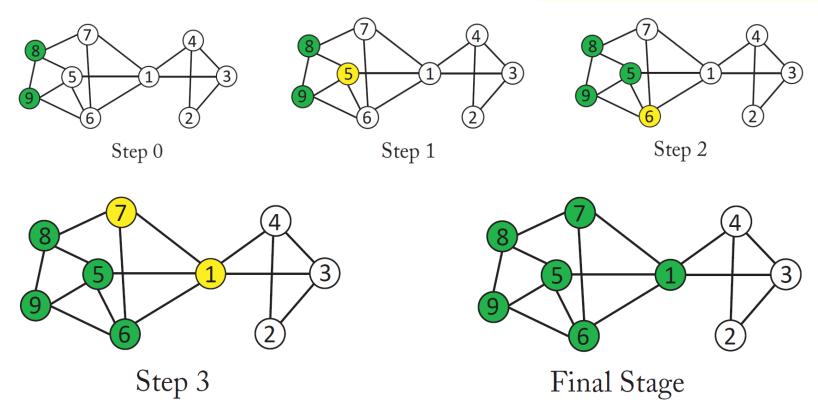


activation "receiver driven"

Linear Threshold Model – Diffusion Process (Threshold = 50%)

assume every v_i chooses 0.5

i.e. 50% of neighbours have to be active





Independent Cascade Model (ICM)

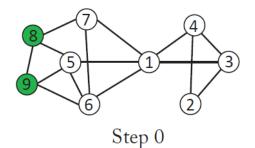
The independent cascade model focuses on the **sender's** rather than the receiver's **view**

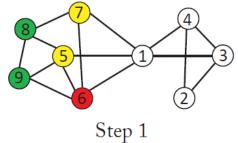
- A node w, once activated at step t, has one chance to activate each of its neighbors randomly
 - For a neighboring node (say, v), the activation succeeds with probability $p_{w,v}$ (e.g. p = 0.5)
- If the activation succeeds, then v will become active at step t + 1
- In the **subsequent** rounds, **w** will **not** attempt to activate **v** anymore.
- The diffusion process, starts with an initial activated set of nodes, then continues until no further activation is possible

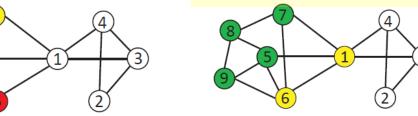


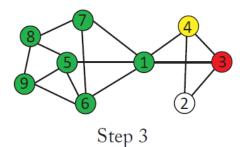
Community Analysis :: Influence Modeling activation "sender driven"

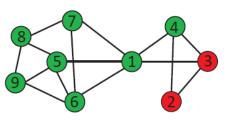
Independent Cascade Model (ICM)

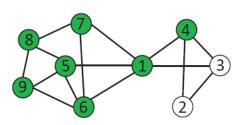












Step 2

with a certain probability

yellow ... sucessful red ... not successful

only active odes can set friends active ...

Step 4

Final Stage

Influence Maximization

- Given a network and a parameter k:
 which k nodes should be selected to be in the activation set B
 In order to maximize the influence in terms of active nodes at the end?
- Let σ(B) denote the expected number of nodes that can be influenced by B, the optimization problem can be formulated as follows:

$$\max_{B \subseteq V} \sigma(B) \ s.t. \ |B| \le k$$



Influence Maximization – A greedy approach

Maximizing the influence is an NP-hard problem but it is proven that the greedy approach gives a solution that is
 63 % of the optimal.

A greedy approach

- Start with B = Ø
- Evaluate $\sigma(v)$ for each node, and pick the node with maximum σ as the first node v_1 to form $B = \{v_1\}$
- Select a node which will increase $\sigma(B)$ most if the node is included in B

Essentially, we greedily find a node $v \in V \setminus B$ such that

$$v = \arg\max_{v \in V \setminus B} \sigma(B \cup \{v\})$$



Community Analysis :: Distinguishing Between Influence & Correlation

Correlation

It has been widely observed that user attributes and behaviors tend to correlate within their social networks

- Suppose we have a binary attribute with each node (say, whether or not being smoker)
- If the attribute is correlated with the network, we expect actors sharing the same attribute value to be positively correlated with social connections
- That is, smokers are more likely to interact with other smokers, and non-smokers with non-smokers



Community Analysis :: Distinguishing Between Influence & Correlation

Test For Correlation

If the **fraction** of **edges linking nodes** with **different attribute values** are **significantly less** than the expected probability, then there is **evidence** of **correlation**

Example if connections are independent of the smoking behavior:

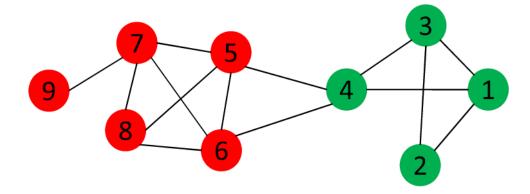
p fraction are smokers (1-p non-smoker)

- one edge is expected to connect two smokers with probability: p x p
- two non-smokers with probability: $(1 p) \times (1 p)$
- a smoker and a non-smoker: 2 p × (1-p)



Community Analysis :: Distinguishing Between Influence & Correlation

Test For Correlation



- Red nodes denote non-smokers, and green ones are smokers. If there is no correlation, then the probability of one edge connecting a smoker and a non-smoker is 2 × 4/9 × 5/9 = 49%.
- In this example the fraction is **2/14** = 14% < 49%, so this network demonstrates some degree of **correlation** with respect to the smoking behavior.
- A more formal way is to conduct a χ2 test for independence of social connections and attributes [1]

[1] T. La Fond and J. Neville. Randomization tests for distinguishing social influence and homophily effects. In Proceedings of the 19th international conference on World wide web, WWW '10, pages 601–610, New York, NY, USA, 2010. ACM.



Community Analysis :: Clustering

Node Centric clustering

Nodes satisfy different properties:

Complete Mutuality

Cliques

Reachability of members

k-clique, k-clan, k-club

Nodal degrees

k-plex, k-core

Relative frequency of Within-Outside Ties

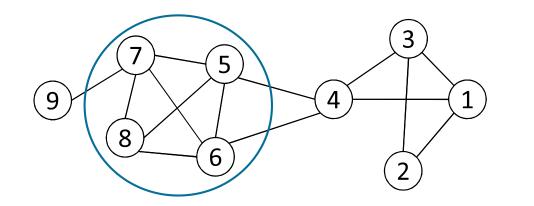
- LS sets, Lambda sets
- Commonly used in traditional social network analysis
- Here, we discuss some representative ones



Community Analysis :: Clustering

Clique

A maximum complete subgraph in which all nodes are adjacent to each other



Nodes 5, 6, 7 and 8 form a clique

NP-hard to find the **maximum clique** in a network

Straightforward implementation to find cliques is very expensive in time complexity



Finding the Maximum Clique

In a clique of size k, each node maintains degree >= k-1

Nodes with degree < k-1 will **not be included** in the maximum clique

Recursively apply the following **pruning** procedure

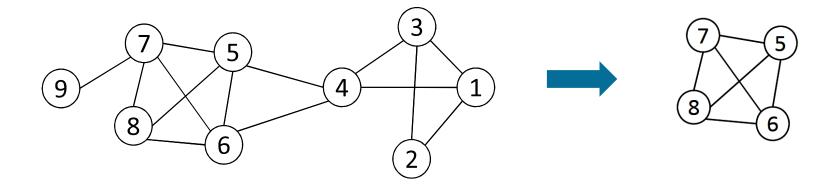
- Sample a sub-network from the given network, and find a clique in the sub-network, say, by a greedy approach
- Suppose the clique above is size k, in order to find out a larger clique, all nodes with degree <= k-1 should be removed.

Repeat until the network is small enough

Many nodes will be pruned as social media networks follow a power law distribution for node degrees



Maximum Clique Example



Suppose we sample a sub-network with nodes {1-5} and find a clique {1, 2, 3} of size 3

In order to find a clique >3, remove all nodes with degree <=3-1=2 (**prune recursively**)

- Remove nodes 2 and 9
- Remove nodes 1 and 3
- Remove node 4



Clique Percolation Method (CPM)

Clique is a very strict definition, unstable

Normally use cliques as a core or a seed to find larger communities

CPM is such a method to find overlapping communities

Input

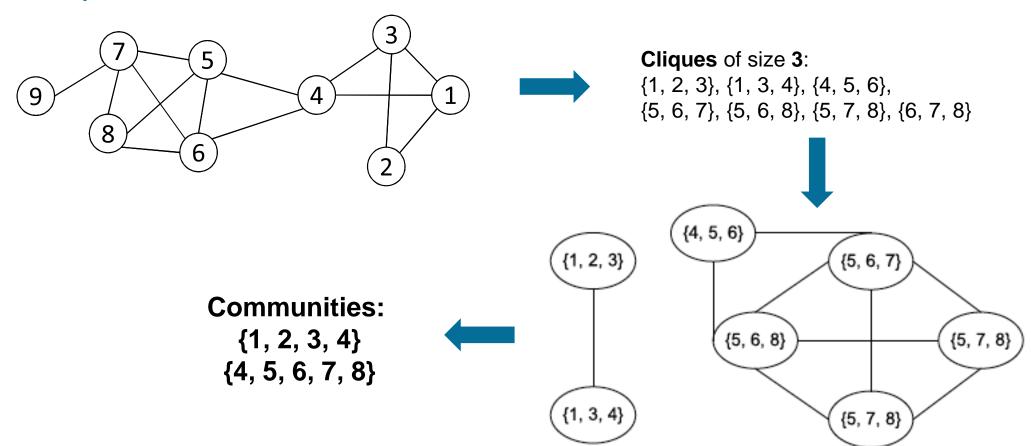
A parameter k, and a network

Procedure

- Find out all cliques of size k in a given network
- Construct a clique graph. Two cliques are adjacent if they share k-1 nodes
- Each connected components in the clique graph form a community



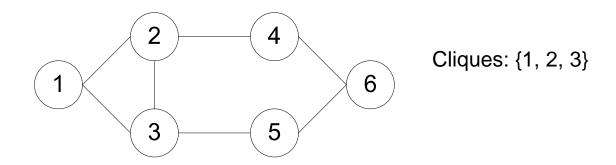
CPM Example





Reachability: k-clique, k-club

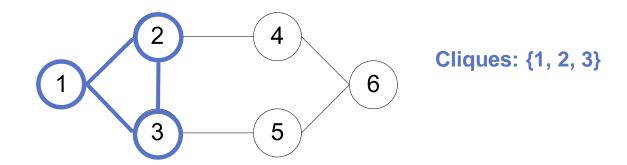
Any node in a group should be reachable in k hops





Reachability: k-clique, k-club

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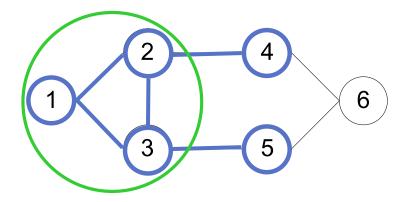




Reachability: k-clique, k-club

Any node in a group should be reachable in k hops

k-clique: a maximal subgraph in which the largest geodesic distance between any nodes <= k



Cliques: {1, 2, 3}

2-cliques: {1, 2, 3, 4, 5}, {2, 3, 4, 5, 6}

A **k-clique** might have **diameter larger than k** in the subgraph

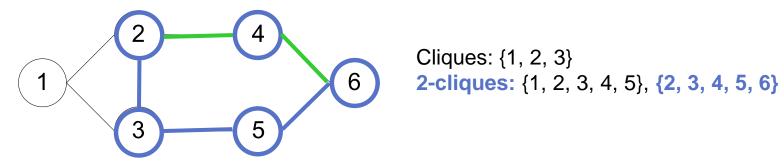
Commonly used in traditional SNA



Reachability: k-clique, k-club

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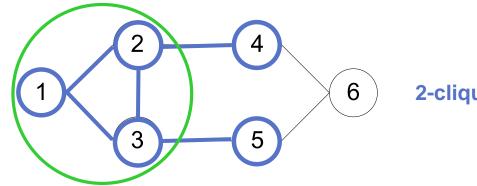
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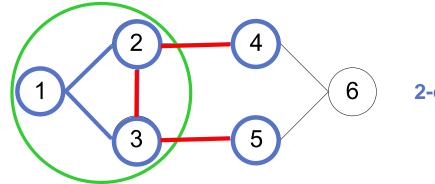
Commonly used in traditional SNA





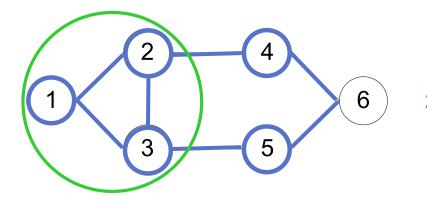
2-cliques: {1, 2, 3, 4, 5}, **{2, 3, 4, 5, 6}**





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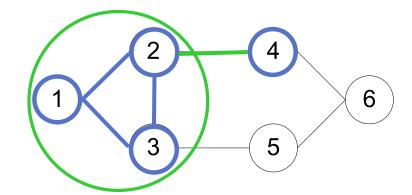


Reachability: k-clique, k-club

Any node in a group should be reachable in k hops

k-clique: a **maximal subgraph** in which the **largest geodesic** distance between any nodes **<= k**

k-club (also: **k-clan**): a **substructure** of **diameter <= k**



Cliques: {1, 2, 3}

2-cliques: {1, 2, 3, 4, 5}, {2, 3, 4, 5, 6}

2-clubs: {1,2,3,4}, {1, 2, 3, 5}, {2, 3, 4, 5, 6}

A **k-clique** might have **diameter larger than k** in the subgraph

Commonly used in traditional SNA

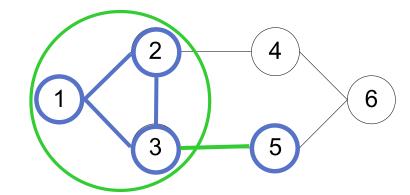


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Commonly used in traditional SNA

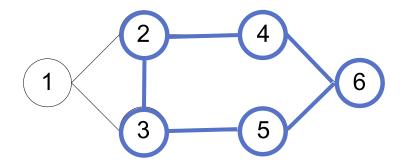


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Cliques: {1, 2, 3}

2-cliques: {1, 2, 3, 4, 5}, {2, 3, 4, 5, 6}

2-clubs: {1,2,3,4}, {1, 2, 3, 5}, **{2, 3, 4, 5, 6}**

A **k-clique** might have **diameter larger than k** in the subgraph

Commonly used in traditional SNA



Group-Centric clustering: Density Based Groups

Quasi-cliques are dense incomplete subgraphs of a graph that generalize the notion of cliques.

The **group-centric criterion** requires the **whole group** to satisfy a certain condition

• E.g., the group density >= a given threshold

$$G_s(V_s,E_{\dot{s}})$$
 a

A subgraph
$$G_s(V_s, E_s)$$
 a $\gamma - depse i-clique$ if

$$\frac{|E_s|}{|V_s|(|V_s|-1)/2} \ge \gamma$$

number of edges in the quasi-clique number of edges in a (full) clique of the same size

A **similar strategy** to that of cliques can be used

- •Sample a subgraph, and find a maximal $\gamma d_{\text{CD-SP-clique}}$ (say, of size k)
- •Remove nodes with degree $< k\gamma$



Network-Centric clustering (as opposed to **Node-Centric clustering** so far)

Network-centric criterion needs to consider the connections within a network globally

Goal: partition nodes of a network into disjoint sets

Approaches:

- Clustering based on vertex similarity
- Latent space models
- Block model approximation
- Spectral clustering
- Modularity maximization



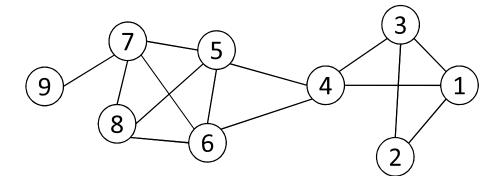
Clustering based on Vertex Similarity

Apply **k-means** or **similarity-based clustering** to nodes

Vertex similarity is defined in terms of the similarity of their neighborhood

Structural equivalence: two nodes are structurally equivalent if they are connecting to the same set of actors

Nodes 1 and 3 are structurally equivalent;



Structural equivalence is too restrictive for practical use.



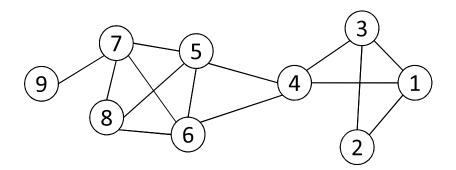
Vertex Similarity

Jaccard Similarity

$$Jaccard(v_i, v_j) = \frac{|N_i \cap N_j|}{|N_i \cup N_j|}$$

Cosine similarity

$$cosine(v_i, v_j) = \frac{|N_i \cap N_j|}{\sqrt{|N_i| \cdot |N_j|}}$$



$$Jaccard(4,6) = \frac{|\{5\}|}{|\{1,3,4,5,6,7,8\}|} = \frac{1}{7}$$
$$cosine(4,6) = \frac{1}{\sqrt{4 \cdot 4}} = \frac{1}{4}$$

Latent Space Models

Map nodes into a low-dimensional space such that the proximity between nodes based on network connectivity is preserved in the new space, then apply k-means clustering

Multi-dimensional scaling (MDS)

- Given a network, construct a proximity matrix P representing the pairwise distance between nodes (e.g., geodesic distance)
- Let $S \hat{I} R^{n'\ell}$ denote the coordinates of nodes in the low-dimensional space

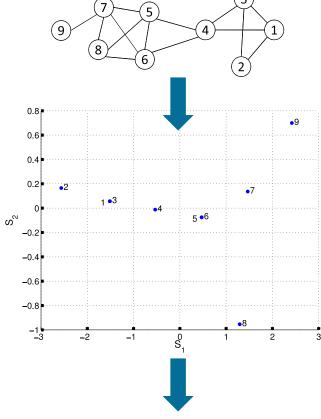
$$SS^T \approx -\frac{1}{2}(I - \frac{1}{n}\mathbf{1}\mathbf{1}^T)(P \circ P)(I - \frac{1}{n}\mathbf{1}\mathbf{1}^T) = \widetilde{P}$$

- Objective function: $\min \|SS^T \widetilde{P}\|_F^2$
- Solution: $S = V\Lambda^{\frac{1}{2}}$
- lacktriangledown V is the top ℓ eigenvectors of \widetilde{P} , and Λ is a diagonal matrix of top eigenvalues $\Lambda=diag(\lambda_1,\lambda_2,\cdots,\lambda_\ell)$

Sosa, J. and Buitrago, L., 2021. A Review of Latent Space Models for Social Networks. Revista Colombiana de Estadística, 44(1), pp.171-200.



MDS Example



Two communities:

{1, 2, 3, 4} and {5, 6, 7, 8, 9}

$$\widetilde{P} = \begin{bmatrix} 2.46 & 3.96 & 1.96 & 0.85 & -0.65 & -0.65 & -2.21 & -2.04 & -3.65 \\ 3.96 & 6.46 & 3.96 & 1.35 & -1.15 & -1.15 & -3.71 & -3.54 & -6.15 \\ 1.96 & 3.96 & 2.46 & 0.85 & -0.65 & -0.65 & -2.21 & -2.04 & -3.65 \\ 0.85 & 1.35 & 0.85 & 0.23 & -0.27 & -0.27 & -0.82 & -0.65 & -1.27 \\ -0.65 & -1.15 & -0.65 & -0.27 & 0.23 & -0.27 & 0.68 & 0.85 & 1.23 \\ -0.65 & -1.15 & -0.65 & -0.27 & -0.27 & 0.23 & 0.68 & 0.85 & 1.23 \\ -2.21 & -3.71 & -2.21 & -0.82 & 0.68 & 0.68 & 2.12 & 1.79 & 3.68 \\ -2.04 & -3.54 & -2.04 & -0.65 & 0.85 & 0.85 & 1.79 & 2.46 & 2.35 \\ -3.65 & -6.15 & -3.65 & -1.27 & 1.23 & 1.23 & 3.68 & 2.35 & 6.23 \end{bmatrix}$$

$$V = \begin{bmatrix} -0.33 & 0.05 \\ -0.55 & 0.14 \\ -0.33 & 0.05 \\ -0.11 & -0.01 \\ 0.10 & -0.06 \\ 0.10 & -0.06 \\ 0.32 & 0.11 \\ 0.28 & -0.79 \\ 0.52 & 0.58 \end{bmatrix}, \quad \Lambda = \begin{bmatrix} 21.56 & 0 \\ 0 & 1.46 \end{bmatrix}, \quad S = V\Lambda^{1/2} = \begin{bmatrix} -1.51 & 0.06 \\ -2.56 & 0.17 \\ -1.51 & 0.06 \\ -0.53 & -0.01 \\ 0.47 & -0.08 \\ 0.47 & -0.08 \\ 1.47 & 0.14 \\ 1.29 & -0.95 \\ 2.42 & 0.70 \end{bmatrix}$$



Block Models

Table 3.1: Adjacency Matrix										
-	1	1	1	0	0	0	0	0		
1		1	0	0	0	0	0	0		
1	1		1	0	0	0	0	0		
1	0	1		1	1	0	0	0		
0	0	0	1		1	1	1	0		
0	0	0	1	1		1	1	0		
0	0	0	0	1	1		1	1		
0	0	0	0	1	1	1		0		
0	0	0	0	0	0	1	0	-		

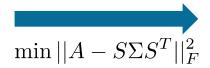


Table 3.2: Ideal Block Structure								
1	1	1	1	0	0	0	0	0
1	1	1	1	0	0	0	0	0
1	1	1	1	0	0	0	0	0
1	1	1	1	0	0	0	0	0
0	0	0	0	1	1	1	1	1
0	0	0	0	1	1	1	1	1
0	0	0	0	1	1	1	1	1
0	0	0	0	1	1	1	1	1
0	0	0	0	1	1	1	1	1

S is the community indicator matrix

Relax S to be numerical values, then the optimal solution corresponds to the top eigenvectors of A

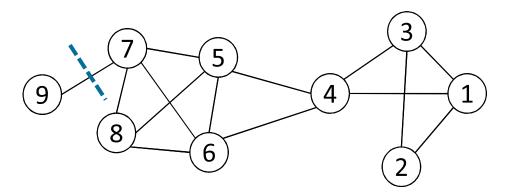
$$S = \begin{bmatrix} 0.20 & -0.52 \\ 0.11 & -0.43 \\ 0.20 & -0.52 \\ 0.38 & -0.30 \\ 0.47 & 0.15 \\ 0.47 & 0.15 \\ 0.41 & 0.28 \\ 0.38 & 0.24 \end{bmatrix}, \Sigma = \begin{bmatrix} 3.5 & 0 \\ 0 & 2.4 \end{bmatrix}.$$

$$\{1, 2, 3, 4\} \text{ and } \{5, 6, 7, 8, 9\}$$



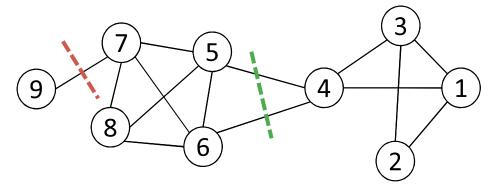
Cut

- Most interactions are within a group, whereas interactions between groups are few
- community detection (clustering) → minimum cut problem
- Cut: A partition of vertices of a graph into two disjoint sets
- Minimum cut problem: find a graph partition such that the number of edges between the two sets is minimized





Ratio Cut & Normalized Cut



- Minimum cut often returns an imbalanced partition, with one set being a singleton (e.g. node 9)
- Change the objective function to consider **community size**

Ratio Cut
$$(\pi) = \frac{1}{k} \sum_{i=1}^{k} \frac{cut(C_i, \bar{C}_i)}{|C_i|},$$

Normalized
$$\operatorname{Cut}(\pi) = \frac{1}{k} \sum_{i=1}^{k} \frac{\operatorname{cut}(C_i, \bar{C}_i)}{\operatorname{vol}(C_i)}$$
 $|C_i|$: number of nodes in C_i (=community size) $\operatorname{vol}(C_i)$: sum of degrees in C_i

C_{i,}: a community

cut(A, B): number of edges induced by cut

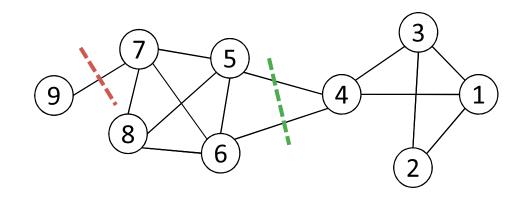


Ratio Cut & Normalized Cut Example

For partition in red: π_1

Ratio
$$Cut(\pi_1) = \frac{1}{2} \left(\frac{1}{1} + \frac{1}{8} \right) = 9/16 = 0.56$$

Normalized $Cut(\pi_1) = \frac{1}{2} \left(\frac{1}{1} + \frac{1}{27} \right) = 14/27 = 0.52$



For partition in green: π_2

Ratio
$$Cut(\pi_2) = \frac{1}{2} \left(\frac{2}{4} + \frac{2}{5} \right) = 9/20 = 0.45 < Ratio $Cut(\pi_1)$$$

Normalized Cut(
$$\pi_2$$
) = $\frac{1}{2} \left(\frac{2}{12} + \frac{2}{16} \right) = 7/48 = 0.15 < \text{Normalized Cut}(\pi_1)$

Both ratio cut and normalized cut prefer a balanced partition



Spectral Clustering

Both ratio cut and normalized cut can be reformulated as

$$\min_{S \in \{0,1\}^{n \times k}} Tr(S^T \widetilde{L}S)$$

Where

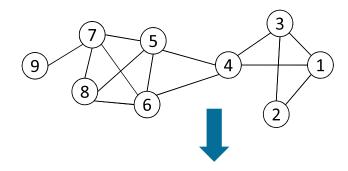
$$\widetilde{L} = \left\{ \begin{array}{ll} D-A & \text{graph Laplacian for ratio cut} \\ I-D^{-1/2}AD^{-1/2} & \text{normalized graph Laplacian} \end{array} \right.$$

$$D = diag(d_1, d_2, \cdots, d_n) \qquad \text{diagonal matrix of degrees}$$

Spectral relaxation: $\min_{S} Tr(S^T \widetilde{L}S)$ s.t. $S^T S = I_k$

Optimal solution: top eigenvectors with the smallest eigenvalues

Spectral Clustering Example



$$D = diag(3, 2, 3, 4, 4, 4, 4, 3, 1)$$

$$\widetilde{L} = D - A = \begin{bmatrix} 3 & -1 & -1 & -1 & 0 & 0 & 0 & 0 & 0 \\ -1 & 2 & -1 & 0 & 0 & 0 & 0 & 0 & 0 \\ -1 & -1 & 3 & -1 & 0 & 0 & 0 & 0 & 0 \\ -1 & 0 & -1 & 4 & -1 & -1 & 0 & 0 & 0 \\ 0 & 0 & 0 & -1 & 4 & -1 & -1 & -1 & 0 \\ 0 & 0 & 0 & -1 & -1 & 4 & -1 & -1 & 0 \\ 0 & 0 & 0 & 0 & -1 & -1 & 4 & -1 & -1 \\ 0 & 0 & 0 & 0 & 0 & -1 & -1 & 3 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & -1 & 0 & 1 \end{bmatrix}$$

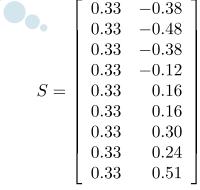
Two communities:

{1, 2, 3, 4} and {5, 6, 7, 8, 9}



k-means

The 1st eigenvector means all nodes belong to the same cluster, no use



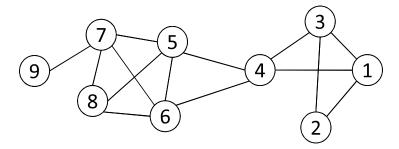


Modularity Maximation

Modularity measures the strength of a community partition by taking into account the degree distribution

Given a network with m edges, the expected number of edges between two nodes with d_i and d_i is

 $d_i d_j / 2m$



Strength of a community:

$$\sum_{i \in C, j \in C} A_{ij} - d_i d_j / 2m$$

Modularity

$$Q = \frac{1}{2m} \sum_{\ell=1}^{k} \sum_{i \in C_{\ell}, j \in C_{\ell}} (A_{ij} - d_i d_j / 2m)$$

A larger value indicates a good community structure



Modularity Maximation

Modularity matrix:
$$B = A - \mathbf{dd}^T/2m \quad (B_{ij} = A_{ij} - d_i d_j/2m)$$

Similar to spectral clustering, Modularity maximization can be reformulated as

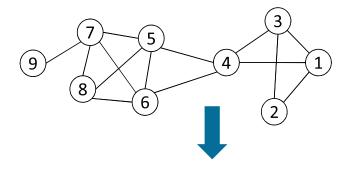
$$\max Q = \frac{1}{2m} Tr(S^T B S) \quad s.t. \ S^T S = I_k$$

Optimal solution: top eigenvectors of the modularity matrix

Apply k-means to S as a post-processing step to obtain community partition



Modularity Maximation Example



$$B = \begin{bmatrix} -0.32 & 0.79 & 0.68 & 0.57 & -0.43 & -0.43 & -0.43 & -0.32 & -0.11 \\ 0.79 & -0.14 & 0.79 & -0.29 & -0.29 & -0.29 & -0.29 & -0.21 & -0.07 \\ 0.68 & 0.79 & -0.32 & 0.57 & -0.43 & -0.43 & -0.43 & -0.32 & -0.11 \\ 0.57 & -0.29 & 0.57 & -0.57 & 0.43 & 0.43 & -0.57 & -0.43 & -0.14 \\ -0.43 & -0.29 & -0.43 & 0.43 & -0.57 & 0.43 & 0.43 & 0.57 & -0.14 \\ -0.43 & -0.29 & -0.43 & 0.43 & 0.43 & -0.57 & 0.43 & 0.57 & -0.14 \\ -0.43 & -0.29 & -0.43 & 0.43 & 0.43 & -0.57 & 0.57 & 0.86 \\ -0.32 & -0.21 & -0.32 & -0.43 & 0.57 & 0.57 & 0.57 & -0.32 & -0.11 \\ -0.11 & -0.07 & -0.11 & -0.14 & -0.14 & -0.14 & 0.86 & -0.11 & -0.04 \end{bmatrix}$$

Modularity Matrix

Two communities:

{1, 2, 3, 4} and {5, 6, 7, 8, 9}



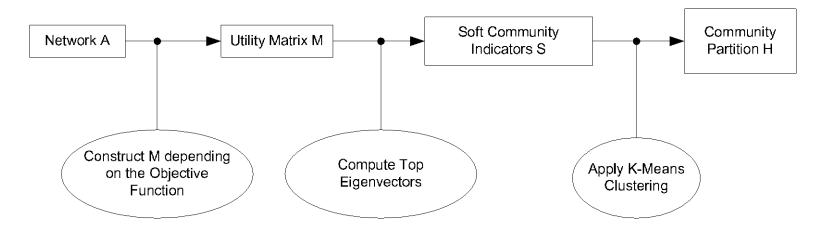
-0.00

$$S = \begin{bmatrix} 0.38 & 0.23 \\ 0.44 & -0.00 \\ 0.17 & -0.48 \\ -0.29 & -0.32 \\ -0.32 & -0.32 \\ -0.38 & 0.34 \\ -0.34 & -0.08 \\ 0.63 \end{bmatrix}$$



A Unified View for Community Platform

Latent space models, block models, spectral clustering, and modularity maximization can be unified as



$$\text{Utility Matrix } M = \left\{ \begin{array}{ll} \text{modified proximity matrix } \widetilde{P} & \textit{if latent space models} \\ \text{adjacency matrix } A & \textit{if block models} \\ \text{graph Laplacian } \widetilde{L} & \textit{if spectral clustering} \\ \text{modularity maximization } B & \textit{if modularity maximization} \end{array} \right.$$



Hierarchy-Centric Clustering

Goal: build a **hierarchical structure** of **communities** based on **network topology**

Allow the analysis of a network at different resolutions

Representative approaches:

- Divisive Hierarchical Clustering
- Agglomerative Hierarchical Clustering



Divisive Hierarchical Clustering

Divisive clustering

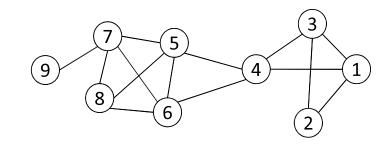
- Partition nodes into several sets
- Each set is further divided into smaller ones
- Network-partition can be applied for the partition

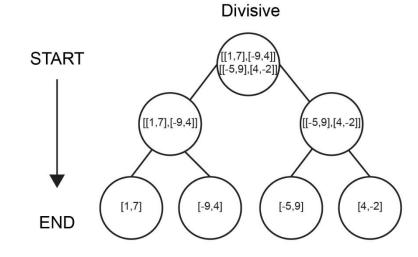
One particular example: recursively remove the "weakest" tie

- Find the edge with the least strength
- Remove the edge and update the corresponding strength of each edge

Recursively apply the above two steps until a network is discomposed into desired number of connected components.

Each component forms a community



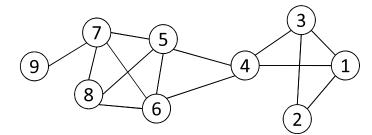




Edge Betweeness

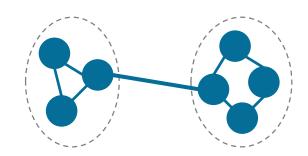
The **strength** of a **tie** can be measured by **edge betweenness Edge betweenness**: the number of shortest paths that pass along with the edge

edge-betweenness(e) =
$$\Sigma_{s < t} \frac{\sigma_{st}(e)}{\sigma_{s,t}}$$

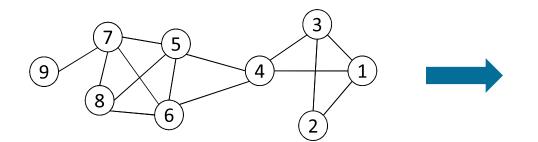


The edge betweenness of e(1, 2) is 4, as all the shortest paths from 2 to $\{4, 5, 6, 7, 8, 9\}$ have to either pass e(1, 2) or e(2, 3), and e(1,2) is the shortest path between 1 and 2

The edge with higher betweenness tends to be the bridge between two communities.

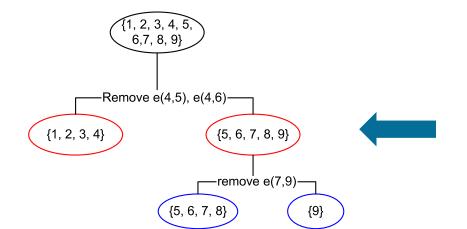


Divisive clustering based on Edge Betweenness



Initial betweenness value

	Table 3.3: Edge Betweenness									
	1	2	3	4	5	6	7	8	9	
1	0	4	1	9	0	0	0	0	0	
2	4	0	4	0	0	0	0	0	0	
3	1	4	0	9	0	0	0	0	0	
4	9	0	9	0	10	10	0	0	0	
5	0	0	0	10	0	1	6	3	0	
6	0	0	0	10	1	0	6	3	0	
7	0	0	0	0	6	6	0	2	8	
8	0	0	0	0	3	3	2	0	0	
9	0	0	0	0	0	0	8	0	0	



After remove e(4,5), the betweenness of e(4, 6) becomes 20, which is the highest;

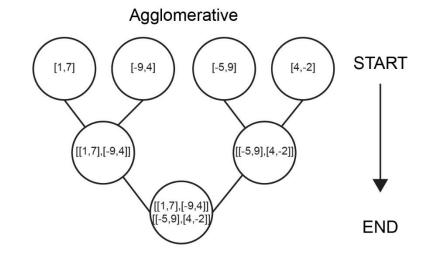
After remove e(4,6), the edge e(7,9) has the highest betweenness value 4, and should be removed.



Agglomerative Hierarchical Clustering

- Initialize each node as a community
- Merge communities successively into larger communities following a certain criterion

(E.g., based on modularity increase)







Social Computing:: Community Detection

Summary of Community Detection

Node-Centric Community Detection

cliques, k-cliques, k-clubs

Group-Centric Community Detection

quasi-cliques

Network-Centric Community Detection

- Clustering based on vertex similarity
- Latent space models, block models, spectral clustering, modularity maximization

Hierarchy-Centric Community Detection

- Divisive clustering
- Agglomerative clustering





Community Analysis



Algorithms and Data Structures 2, 340300 Lecture – 2023W Univ.-Prof. Dr. Alois Ferscha, teaching@pervasive.jku.at

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