



NATIONAL RESEARCH
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Network communities

Network Science

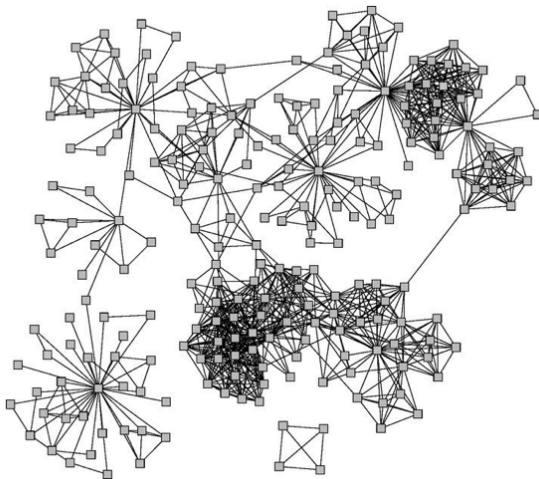
Lecture 8

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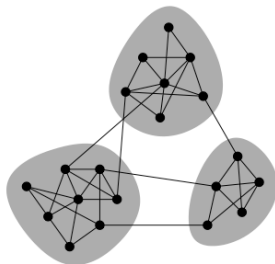
www.leonidzhukov.net/hse/2022/networks

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Definition

Network communities are groups of vertices such that vertices inside the group connected with many more edges than between groups.



- Community detection is an assignment of vertices to communities.

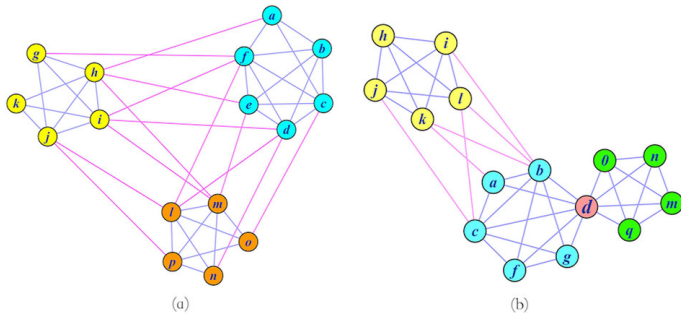


image from W. Liu, 2014

Community is a cohesive subgroup:

- Mutuality of ties. Almost everyone in the group has ties (edges) to one another
- Compactness. Closeness or reachability of group members in small number of steps, not necessarily adjacency
- Density of edges. High frequency of ties within the group
- Separation. Higher frequency of ties among group members compared to non-members

Wasserman and Faust

- Graph $G(V, E)$, $n = |V|$, $m = |E|$
- Community - set of nodes S
 n_s - number of nodes in S , m_s - number of edges in S
- Graph density

$$\rho = \frac{m}{n(n-1)/2}$$

- community internal density

$$\delta_{int} = \frac{m_s}{n_s(n_s-1)/2}$$

- external edges density

$$\delta_{ext} = \frac{m_{ext}}{n_s(n - n_s)}$$

- community (cluster): $\delta_{int} > \rho$, $\delta_{ext} < \rho$

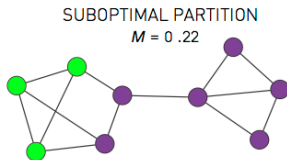
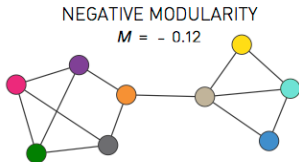
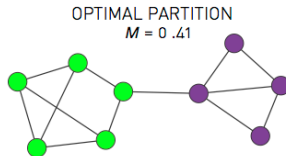
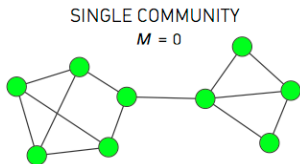
- Compare fraction of edges within the cluster to expected fraction in random graph with identical degree sequence
- Modularity score

$$Q = \frac{1}{2m} \sum_{ij} \left(A_{ij} - \frac{k_i k_j}{2m} \right) \delta(c_i, c_j), = \sum_u \left(\frac{m_u}{m} - \left(\frac{k_u}{2m} \right)^2 \right)$$

m_u - number of internal edges in a community u ,

k_u - sum of node degrees within a community

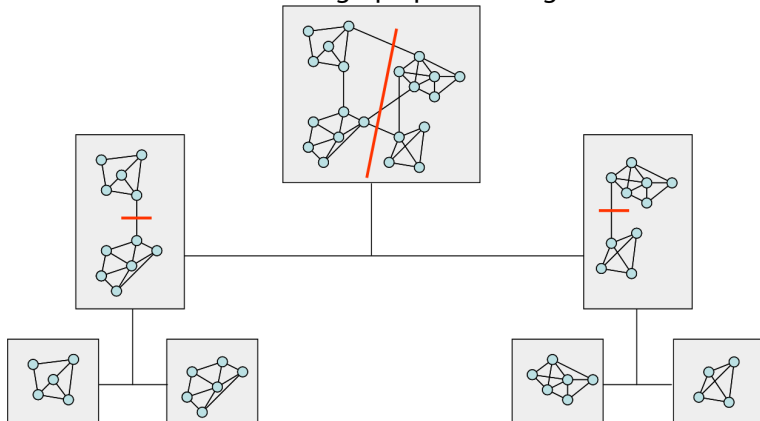
- Modularity score range $Q \in [-1/2, 1)$, single community $Q = 0$



- The higher the modularity score - the better are communities

from A.L. Barabasi 2016

Recursive graph partitioning



Graph $G(E, V)$ partition: $V = V_1 + V_2$

- Graph cut

$$Q = \text{cut}(V_1, V_2) = \sum_{i \in V_1, j \in V_2} e_{ij}$$

- Ratio cut:

$$Q = \frac{\text{cut}(V_1, V_2)}{||V_1||} + \frac{\text{cut}(V_1, V_2)}{||V_2||}$$

- Normalized cut:

$$Q = \frac{\text{cut}(V_1, V_2)}{\text{Vol}(V_1)} + \frac{\text{cut}(V_1, V_2)}{\text{Vol}(V_2)}$$

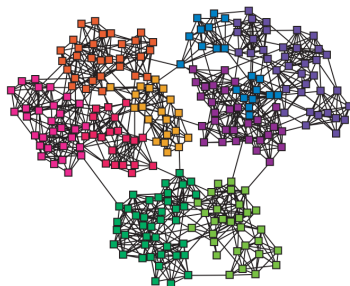
- Quotient cut (conductance):

$$Q = \frac{\text{cut}(V_1, V_2)}{\min(\text{Vol}(V_1), \text{Vol}(V_2))}$$

where: $\text{Vol}(V_1) = \sum_{i \in V_1, j \in V} e_{ij} = \sum_{i \in V_1} k_i$

Edge betweenness - number of shortest paths $\sigma_{st}(e)$ going through edge e

$$C_B(e) = \sum_{s \neq t} \frac{\sigma_{st}(e)}{\sigma_{st}}$$



Recover communities by progressively removing edges

Newman-Girvan, 2004

Algorithm: Edge Betweenness

Input: graph $G(V,E)$

Output: Dendrogram

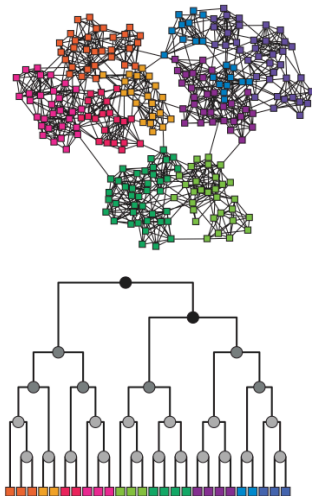
repeat

 For all $e \in E$ compute edge betweenness $C_B(e)$;
 remove edge e_i with largest $C_B(e_i)$;

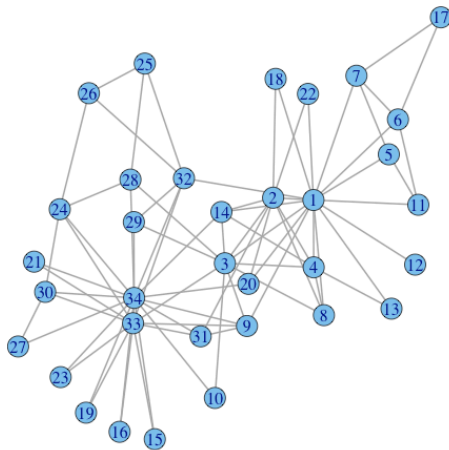
until *edges left*;

If bi-partition, then stop when graph splits in two components
(check for connectedness)

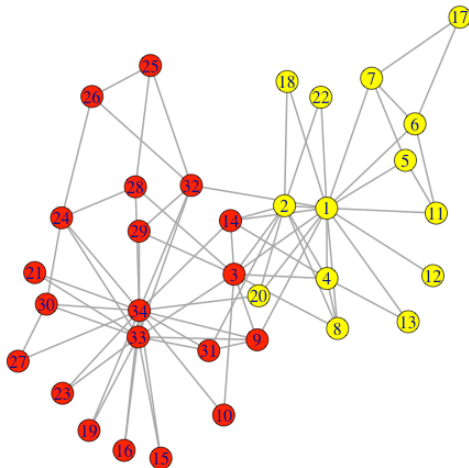
Hierarchical algorithm, dendrogram



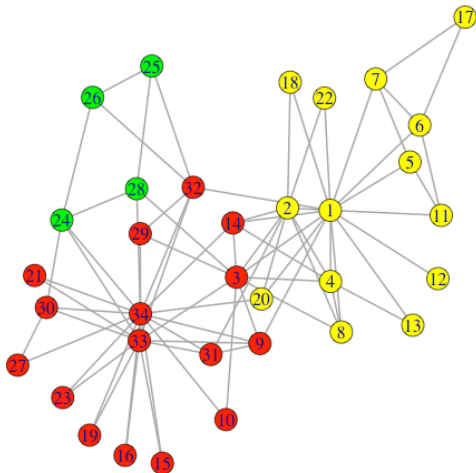
Zachary karate club

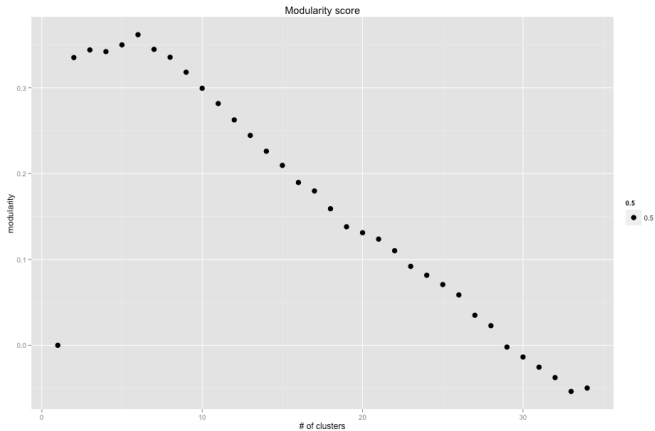


Zachary karate club

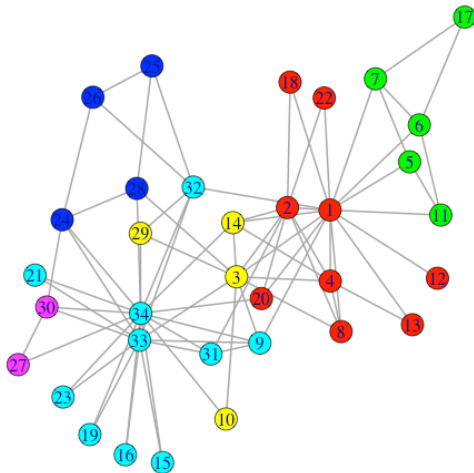


Zachary karate club

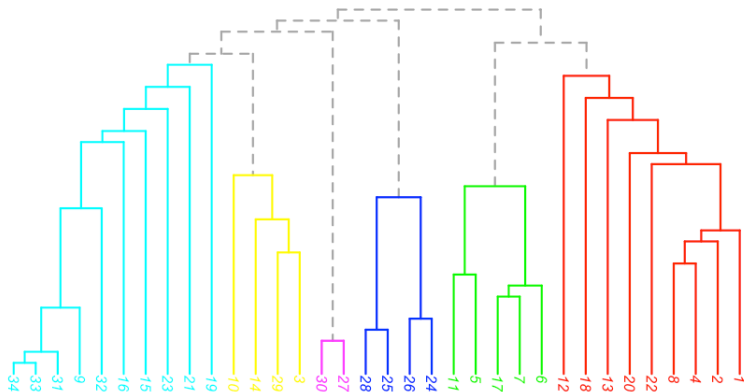




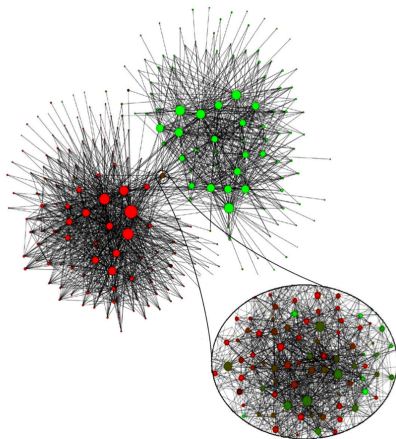
best: clusters = 6, modularity = 0.345



Zachary karate club



Multi-resolution scalable method



2 mln mobile phone network

V. Blondel et.al., 2008

"The Louvain method"

- Heuristic method for greedy modularity optimization
- Find partitions with high modularity
- Multi-level (multi-resolution) hierarchical scheme
- Scalable

Modularity:

$$Q = \frac{1}{2m} \sum_{i,j} \left(A_{ij} - \frac{k_i k_j}{2m} \right) \delta(c_i, c_j)$$

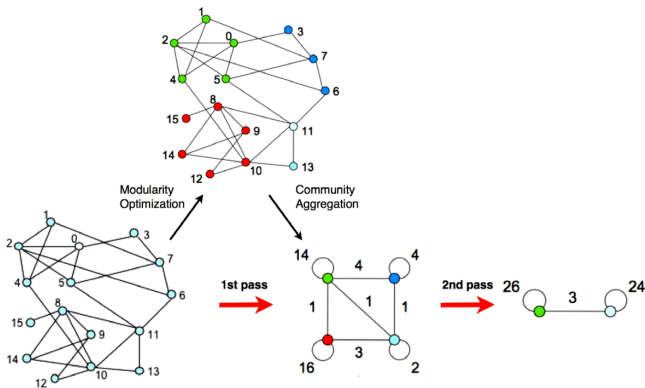
V. Blondel et.al., 2008

Algorithm

- Assign every node to its own community
- Phase I
 - For every node evaluate modularity gain from removing node from its community and placing it in the community of its neighbor
 - Place node in the community maximizing modularity gain
 - repeat until no more improvement (local max of modularity)
- Phase II
 - Nodes from communities merged into "super nodes"
 - Weight on the links added up
- Repeat until no more changes (max modularity)

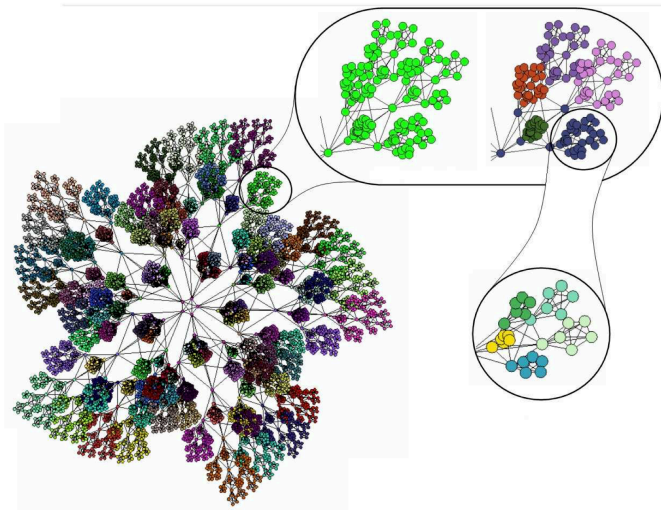
V. Blondel et.al., 2008

Fast community unfolding



V. Blondel et.al., 2008

Fast community unfolding

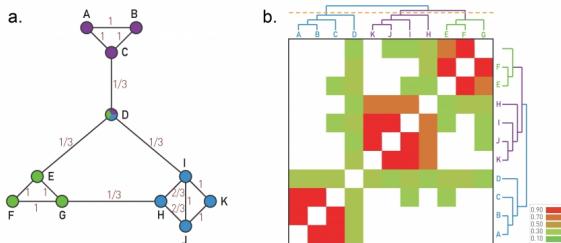


V. Blondel et.al., 2008

Similarity matrix

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

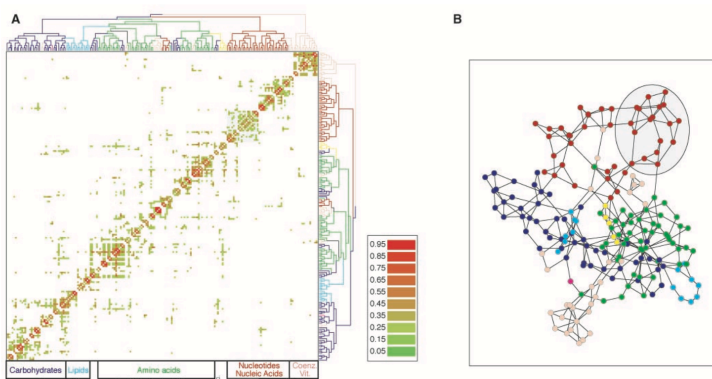
- N_{ij} - number of common neighbors
- $\theta()$ - step function, 0 for $x \leq 0$ and 1 for $x > 0$



The Ravasz algorithm:

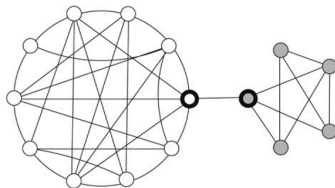
- Assign each node to a community of its own and evaluate x_{ij} for all node pairs.
- Find the node (community) pair with the highest similarity and merge them into a single community.
- Calculate the similarity between the new community and all other communities.
- Repeat Steps 2 and 3 until all nodes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

E. Ravasz et.al., 2002



Reduced E.coli metabolic network

E. Ravasz et.al., 2002



- Random walks on a graph tend to get trapped into densely connected parts corresponding to communities.

Walktrap

- Consider random walk on graph
- At each time step walk moves to NN uniformly at random
 $P_{ij} = \frac{A_{ij}}{d(i)}, P = D^{-1}A, D_{ii} = \text{diag}(d(i))$
- P_{ij}^t - probability to get from i to j in t steps, $t \ll t_{\text{mixing}}$
- Assumptions: for two i and j in the same community P_{ij}^t is high
- if i and j are in the same community, then $\forall k, P_{ik}^t \approx P_{jk}^t$
- Distance between nodes:

$$r_{ij}(t) = \sqrt{\sum_{k=1}^n \frac{(P_{ik}^t - P_{jk}^t)^2}{d(k)}} = \|D^{-1/2}P_i^t - D^{-1/2}P_j^t\|$$

Computing node distance r_{ij}

- Direct (exact) computation: $P_{ij}^t = (P^t)_{ij}$ or $P_i^t = P^t p_i^0, p_i^0(k) = \delta_{ik}$
- Approximate computation (simulation):
 - Compute K random walks of length t starting from node i
 - Approximate $P_{ik}^t \approx \frac{N_{ik}}{K}$, number of walks end up on k

Distance between communities:

$$P_{Cj}^t = \frac{1}{|C|} \sum_{i \in C} P_{ij}^t$$

$$r_{C_1 C_2}(t) = \sqrt{\sum_{k=1}^n \frac{(P_{C_1 k}^t - P_{C_2 k}^t)^2}{d(k)}} = \|D^{-1/2} P_{C_1}^t - D^{-1/2} P_{C_2}^t\|$$

Algorithm (hierarchical clustering)

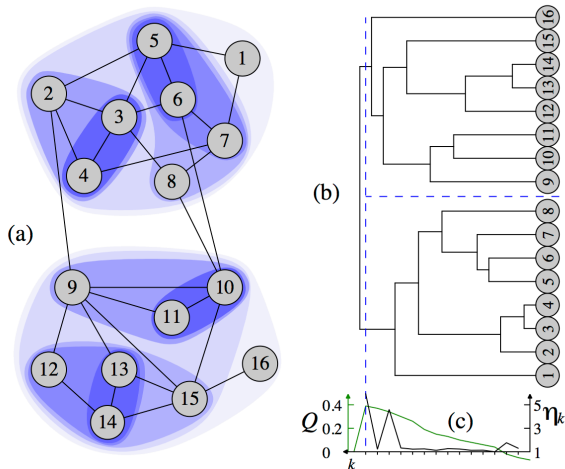
- Assign each vertex to its own community $S_1 = \{\{v\}, v \in V\}$
- Compute distance between all adjacent communities $r_{C_i C_j}$
- Choose two "closest" communities that minimizes (Ward's methods):

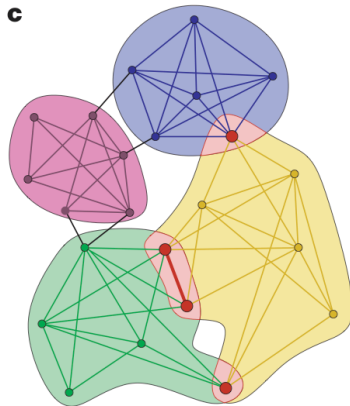
$$\Delta\sigma(C_1, C_2) = \frac{1}{n} \left(\sum_{i \in C_3} r_{iC_3}^2 - \sum_{i \in C_1} r_{iC_1}^2 - \sum_{i \in C_2} r_{iC_2}^2 \right)$$

and merge them $S_{k+1} = (S_k \setminus \{C_1, C_2\}) \cup C_3, C_3 = C_1 \cup C_2$

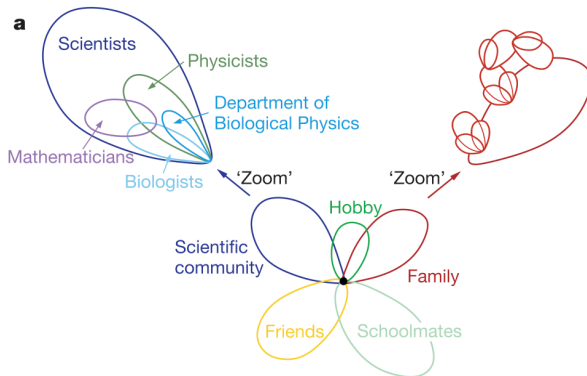
- update distance between communities

After $n - 1$ steps finish with one community $S_n = \{V\}$



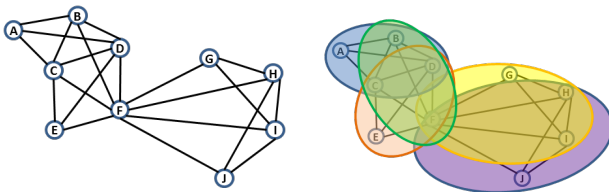


Palla, 2005

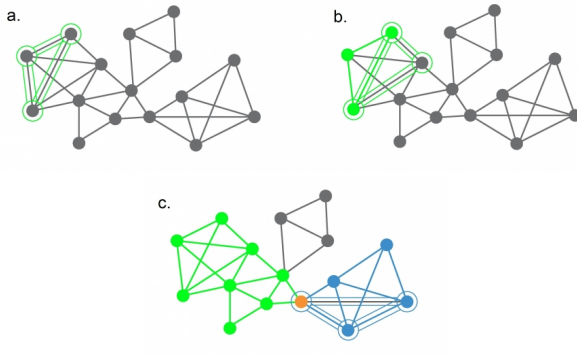


Palla, 2005

- k -clique is a clique (complete subgraph) with k nodes
- k -clique community a union of all k -cliques that can be reached from each other through a series of adjacent k -cliques
- two k -cliques are said to be adjacent if they share $k - 1$ nodes.



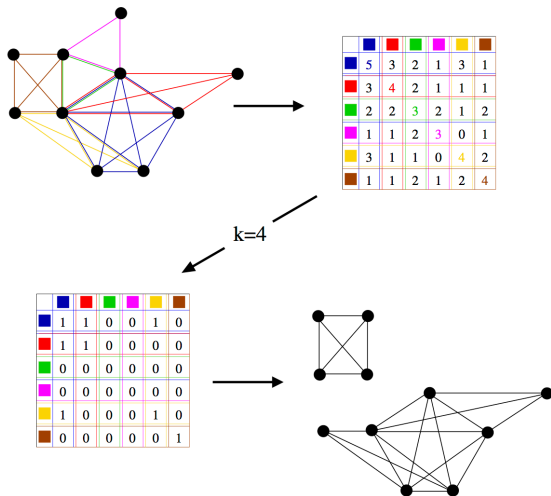
Adjacent 4-cliques



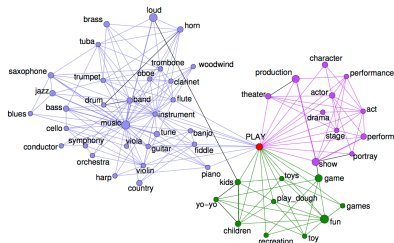
Palla, 2005

- Find all maximal cliques
- Create clique overlap matrix
- Threshold matrix at value $k - 1$
- Communities = connected components

Palla, 2005



k-clique percolation

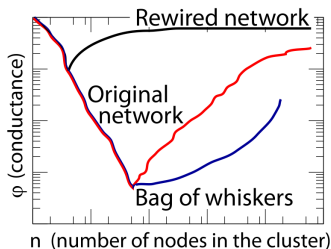


$k = 4$

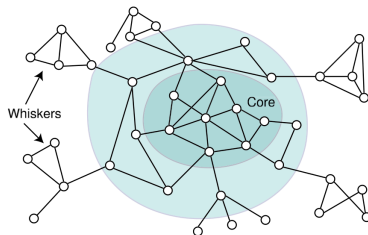


$k = 5$

Palla, 2005



(a) Typical NCP plot



(b) Caricature of network structure

Best conductance of a vertex set S of size k :

$$\Phi(k) = \min_{S \in V, |S|=k} \phi(S), \quad \phi(S) = \frac{\text{cut}(S, V \setminus S)}{\min(\text{vol}(S), \text{vol}(V \setminus S))}$$

where $\text{vol}(S) = \sum_{i \in S} k_i$ - sum of all node degrees in the set

Author	Ref.	Label	Order
Eckmann & Moses	(Eckmann and Moses, 2002)	EM	$O(m(k^2))$
Zhou & Lipowsky	(Zhou and Lipowsky, 2004)	ZL	$O(n^3)$
Latapy & Pons	(Latapy and Pons, 2005)	LP	$O(n^3)$
Clauset et al.	(Clauset <i>et al.</i> , 2004)	NF	$O(n \log^2 n)$
Newman & Girvan	(Newman and Girvan, 2004)	NG	$O(nm^2)$
Girvan & Newman	(Girvan and Newman, 2002)	GN	$O(n^2 m)$
Guimerà et al.	(Guimerà and Amaral, 2005; Guimerà <i>et al.</i> , 2004)	SA	parameter dependent
Duch & Arenas	(Duch and Arenas, 2005)	DA	$O(n^2 \log n)$
Fortunato et al.	(Fortunato <i>et al.</i> , 2004)	FLM	$O(m^3 n)$
Radicchi et al.	(Radicchi <i>et al.</i> , 2004)	RCCLP	$O(m^4/n^2)$
Donetti & Muñoz	(Donetti and Muñoz, 2004, 2005)	DM/DMN	$O(n^3)$
Bagrow & Boltt	(Bagrow and Boltt, 2005)	BB	$O(n^3)$
Capocci et al.	(Capocci <i>et al.</i> , 2005)	CSCC	$O(n^2)$
Wu & Huberman	(Wu and Huberman, 2004)	WH	$O(n + m)$
Palla et al.	(Palla <i>et al.</i> , 2005)	PK	$O(\exp(n))$
Reichardt & Bornholdt	(Reichardt and Bornholdt, 2004)	RB	parameter dependent

Author	Ref.	Label	Order
Girvan & Newman	(Girvan and Newman, 2002; Newman and Girvan, 2004)	GN	$O(nm^2)$
Clauset et al.	(Clauset <i>et al.</i> , 2004)	Clauset et al.	$O(n \log^2 n)$
Blondel et al.	(Blondel <i>et al.</i> , 2008)	Blondel et al.	$O(m)$
Guimerà et al.	(Guimerà and Amaral, 2005; Guimerà <i>et al.</i> , 2004)	Sim. Ann.	parameter dependent
Radicchi et al.	(Radicchi <i>et al.</i> , 2004)	Radicchi et al.	$O(m^4/n^2)$
Palla et al.	(Palla <i>et al.</i> , 2005)	Cfinder	$O(\exp(n))$
Van Dongen	(Dongen, 2000a)	MCL	$O(nk^2)$, $k < n$ parameter
Rosvall & Bergstrom	(Rosvall and Bergstrom, 2007)	Infomod	parameter dependent
Rosvall & Bergstrom	(Rosvall and Bergstrom, 2008)	Infomap	$O(m)$
Donetti & Muñoz	(Donetti and Muñoz, 2004, 2005)	DM	$O(n^3)$
Newman & Leicht	(Newman and Leicht, 2007)	EM	parameter dependent
Ronhovde & Nussinov	(Ronhovde and Nussinov, 2009)	RN	$O(m^{\beta} \log n)$, $\beta \sim 1.3$

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