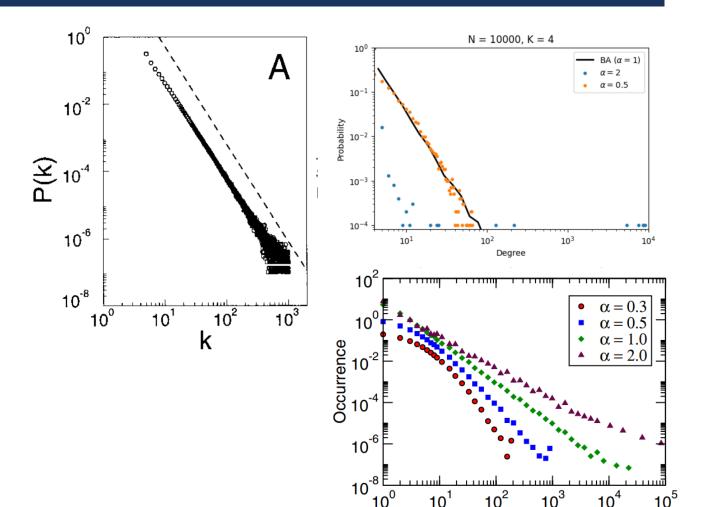
# NETWORK SCIENCE OF ONLINE INTERACTIONS

Chapter 6: Communities

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## **SUMMARY**

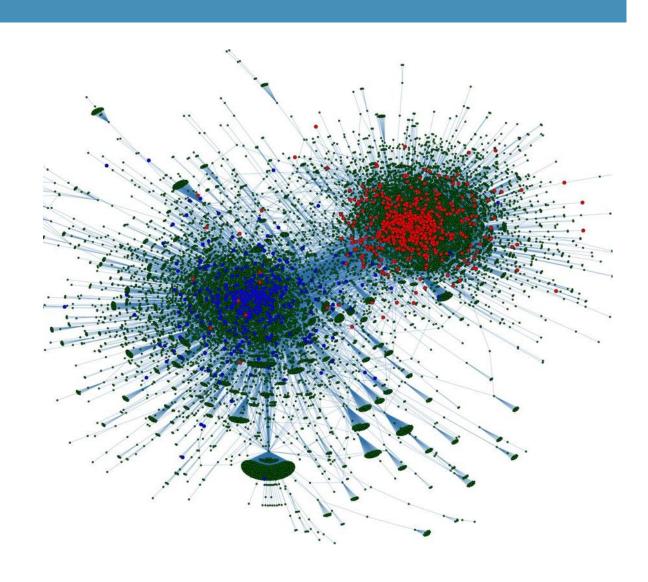
- Many models focusing on emulating certain properties
  - Degree distribution, clustering, triadic closure, etc
- No single "best model"
- Preferential attachment is a key mechanism
  - Can create heavy-tailed distributions
  - If unbalanced, can create hyper-concentrated hubs
- Variations of PA models can create a variety of degree distributions



k (degree)

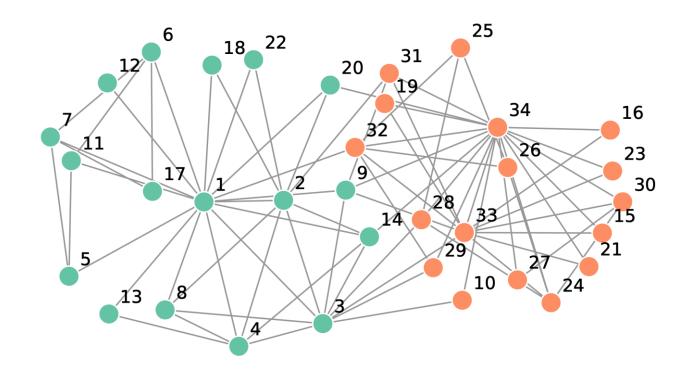
# INTRODUCTION

- Community detection is one of the main fields of network theory
- Many reasons:
  - Uncover network structure
  - Identify node affiliation
  - Find missing links ↔ predict links
- Different algorithms for different definitions of a community
- Two categories
  - Descriptive algorithms
  - Inferential algorithms

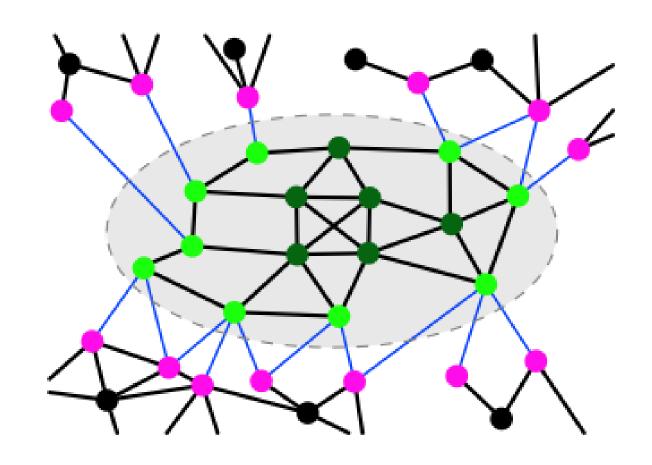


# INTRODUCTION

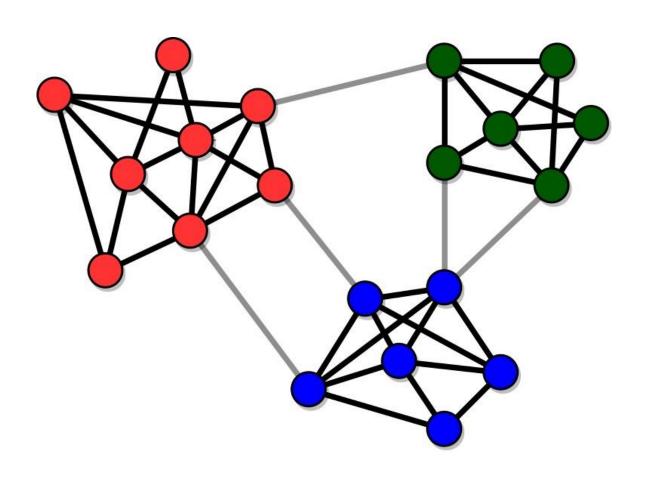
- Basic example: Zachary's Karate club
  - 34 individuals
  - Mapped external relationships
  - Disagreement between 2 instructors
  - Break in two groups
  - Standard test of community detection methods
    - Find two communities
    - Assign nodes to the correct community



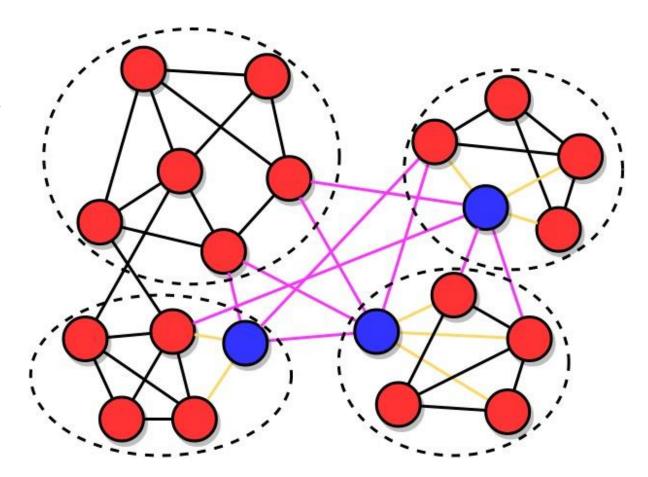
- Community variables
  - Node degree
    - Internal links  $k_i^{int}$  (black)
    - External links  $k_i^{ext}$  (blue)
  - Internal links  $L_C$ , internal nodes  $N_C$
  - Internal link density  $\delta_C^{int} = 2L_C/N_C(N_C 1)$
  - Community degree  $k_C = \sum_{j \in C} (k_j^{int} + k_j^{ext})$
- Valid for undirected, unweighted networks
  - Undirected, weighted: degree → strength
  - Directed: split in in- and out-links, harder to interpret



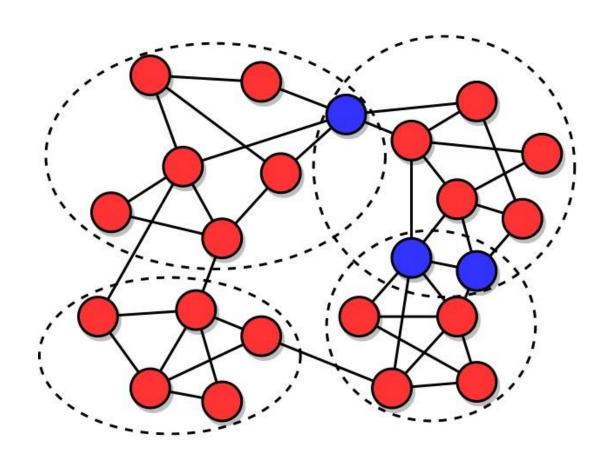
- Intuitive idea:
  - Many internal links (cohesion)
  - Few external links (separation)
- A community should have more internal than external links
- Definition #1
  - **Strong community**: subnetwork where  $k_i^{int} > k_i^{out}$  for each node
  - Weak community:  $\sum_{i} k_{i}^{int} > \sum_{i} k_{i}^{out}$
  - Problem: compares the community to the entire network



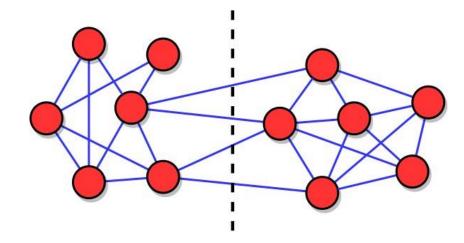
- Definition #2
  - Compares communities to other communities
  - Strong community: internal degree of each node higher than to any other single community (red)
  - Weak community: sum of internal degrees greater than sum of external to any other community (red + blue)
- Current definitions: counting links
  - Bias against small communities
- Alternative: compare link probabilities
  - Requires a network model
  - Inferential methods (more later)



- Communities can be
  - Non-overlapping: partition
  - Overlapping: cover
- Number of communities
  - Number of possible partitions grows super-exponentially (Bell's number)
  - $N = 15 \rightarrow 1.3$  billion possible partitions
  - Number of possible covers is even worse
  - Need for a heuristic algorithm to detect communities
- Graph partitioning is well-studied
  - Community detection in networks
  - Task parallelization in computer science

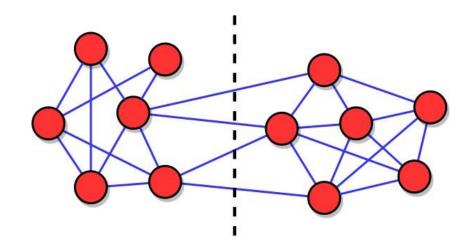


- Network partitioning with fixed group size
  - Cut size: number of links between partitions
  - Good partitions: minimum cut size
  - Graph bisection: two partitions with equal size
    - How to do it?
- Kerninghan-Lin algorithm
  - The idea: minimize cut size
  - The algorithm
    - I. Start from a random bisection
    - 2. For each pair of nodes from the two groups, compute the swap that would result in the largest decrease in cut size
    - 3. Swap those nodes and **lock** them in place
    - 4. Repeat from 2 until cut size cannot be decreased



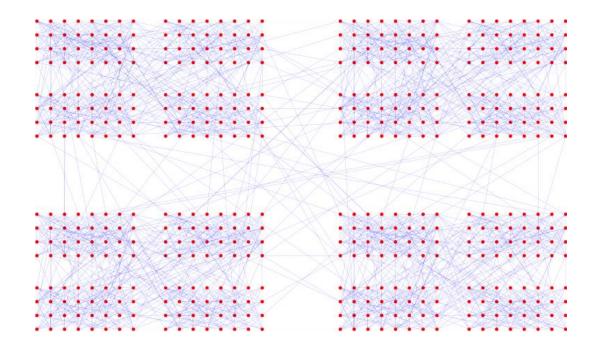
#### Kerninghan-Lin algorithm

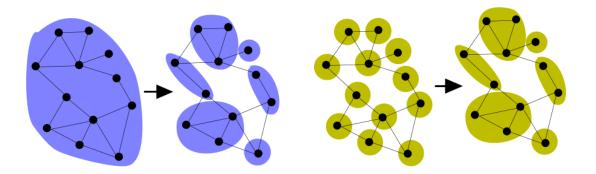
- Greedy algorithm: always tries to minimize/maximize by step
- May get stuck in local minima/maxima
  - Variants implement random changes to avoid this problem
- Depends on initial condition
  - Run many in parallel and choose the one with lowest cut size
- Widely used as post-processing to improve partitioning from other methods (e.g. expert knowledge)
- Limitations of partitioning
  - Partitioning maximizes separation, not internal link density: not necessarily good communities
  - Requires giving the number of communities



partition = nx.community.kernighan\_lin\_bisection(G)

- Partitions can be hierarchical
  - Subdivisions in companies
  - Classes in school
  - How to detected them?
- Hierarchical clustering
  - Main ingredient: similarity measure
  - Approaches:
    - Agglomerative hierarchical clustering
    - Divisive hierarchical clustering



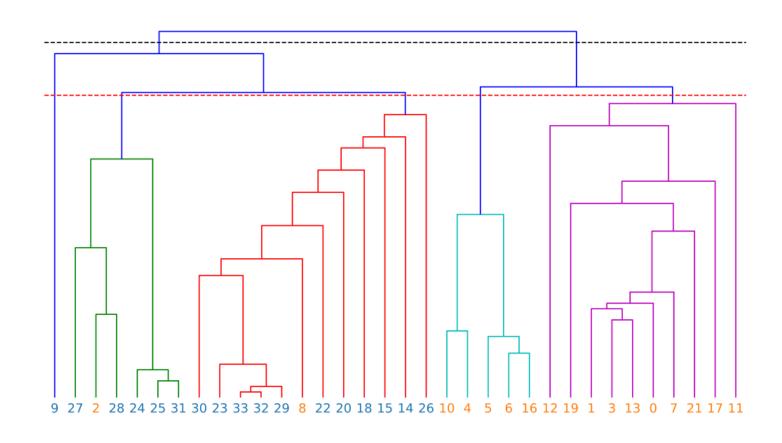


- Example similarity: structural equivalence
  - Idea: nodes are similar if their neighbours are similar

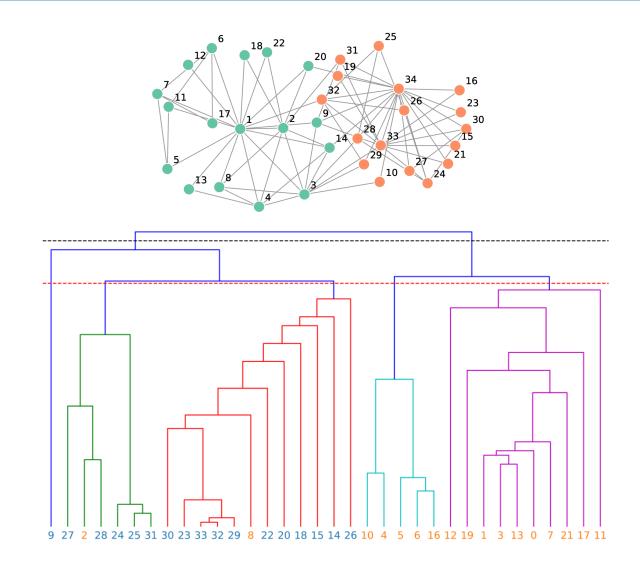
$$S_{ij}^{SE} = \frac{\text{number of neighbors shared by } i \text{ and } j}{\text{total number of nodes neighboring only } i, \text{ only } j, \text{ or both}}$$

- How do define similarity between partitions?
  - Single linkage: take the maximum pairwise similarity  $S_{G_1,G_2} = \max_{i,j} S_{i,j}$
  - **Complete linkage**: take the minimum pairwise similarity  $S_{G_1,G_2} = \min_{i,j} S_{i,j}$
  - Average linkage: take the average pairwise similarity  $S_{G_1,G_2} = \langle S_{ij} \rangle_{i,j}$

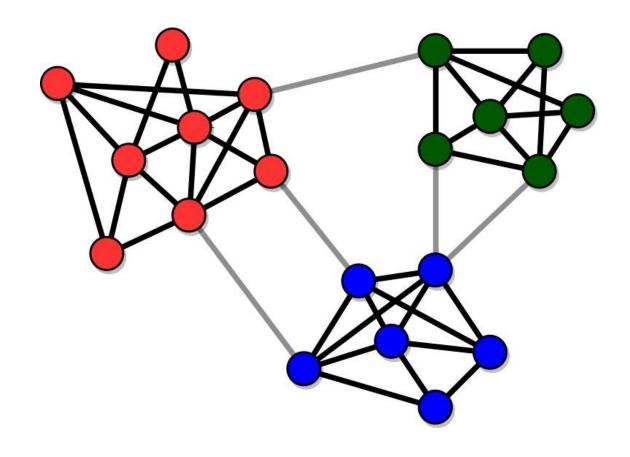
- Result of clustering: dendrogram
  - Summary of similarity between nodes
  - Shows the partitioning for each value of number of partitions (horizontal cuts)
  - Each partition includes the partitions lower in the hierarchy



- Result of clustering: dendrogram
- Benefits
  - Clear picture of node affiliation at different levels
  - Different view from the graph
- Caveats
  - No criteria to select ideal partitioning
  - Depends heavily on similarity measure
  - Rather useless for large networks (slow, too many lines)



- Many different methods
- Today
  - Bridge removal
  - Modularity maximization
- Wednesday:
  - Stochastic block modelling
  - Community validation

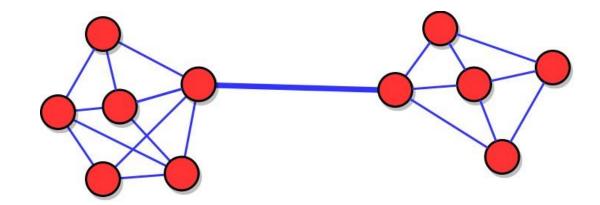


#### Bridge removal

- The idea: detects bridges between clusters, removes them until clusters are disconnected
- Those clusters are communities
- What is a good bridge metric?

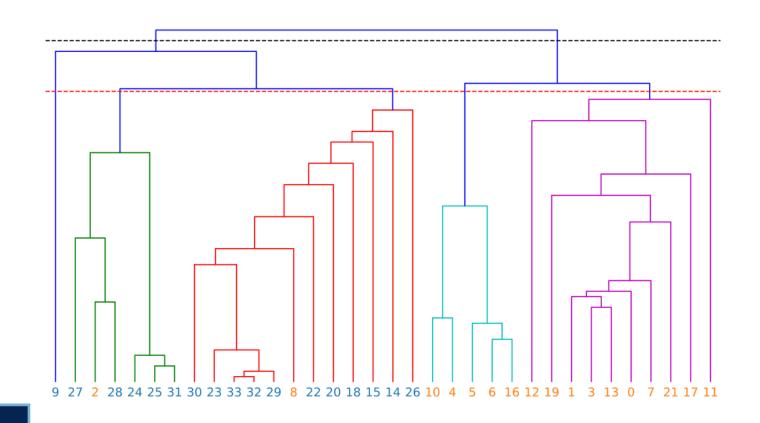
#### Girvan-Newman algorithm

- Measure: link betweenness centrality
- Algorithm
  - I. Compute betweenness for all links
  - 2. Removes the link with largest betweenness, recomputes betweenness for all links
  - 3. Iterate until all links are removed



#### Girvan-Newman: the bad

- Recalculate betweenness for all links at each step: very slow
- Not feasible for large networks (say, > 10000 nodes)
- Girvan-Newman: the good
  - Dendogram
  - If strong community structure, nodes quickly get disconnected
  - Faster variants
    - Computing similarity over a sample
    - Faster similarity than betweenness



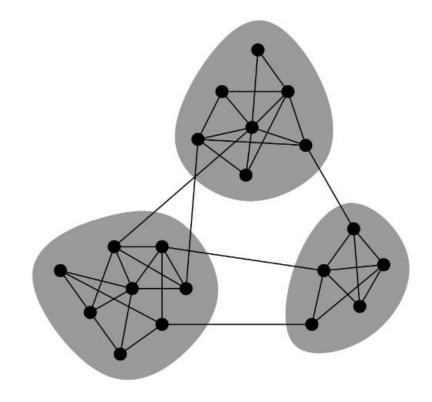
- Method finds a partition: how good is it?
  - Quality function
  - Assumption: random networks have no communities
  - Must distinguish between real communities and random fluctuations
- Most famous quality function: modularity
  - The idea: compares community structure against randomized networks with equal degree
  - For each community, computes the difference in number of internal links between the network and the ensemble of random networks
  - Value  $Q \in [-0.5,1]$

## Modularity and community structure in networks

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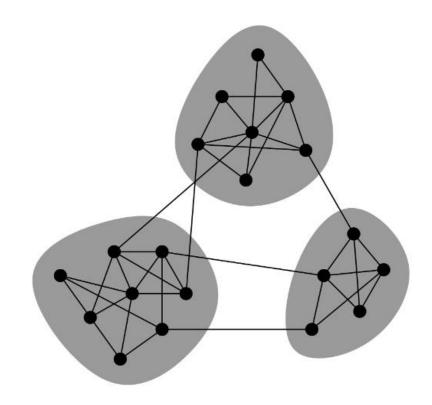
Edited by Brian Skyrms, University of California, Irvine, CA, and approved April 19, 2006 (received for review February 26, 2006)



Modularity definition:

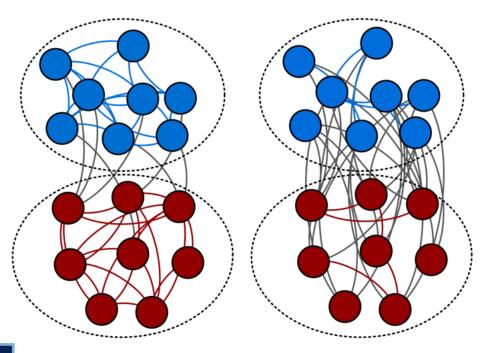
$$Q = \frac{1}{L} \sum_{C} \left( L_C - \frac{k_C^2}{4L} \right)$$

- where
  - L: number of links in the network
  - $L_C$ : number of internal links in community C
  - $k_C$ : degree of community C
  - $k_C^2/4L$ : expected number of internal links in community C



- Explaining  $k_C^2/4L$ :
  - Random links by pairing stubs (configuration model)
  - Total number of stubs in C is  $k_C$
  - Probability of selecting a stub is  $k_C/2L$
  - Probability of link is  $p_C = k_C^2/4L^2$
  - Expected number of internal links is  $Lp_C = k_C^2/4L$
- Properties of Q
  - Q > 0: some structure
  - Q = 0: no structure
  - Q < 0 if particularly bad partitioning

$$Q = \frac{1}{L} \sum_{C} \left( L_C - \frac{k_C^2}{4L} \right)$$



**Original network** 

Randomized network

- Modularity has straightforward extensions
- Directed:

$$Q_d = \frac{1}{L} \sum_{C} \left( L_C - \frac{k_C^{in} k_C^{out}}{L} \right)$$

Weighted:

$$Q_w = \frac{1}{W} \sum_C \left( W_C - \frac{s_C^2}{4W} \right)$$

Directed and weighted:

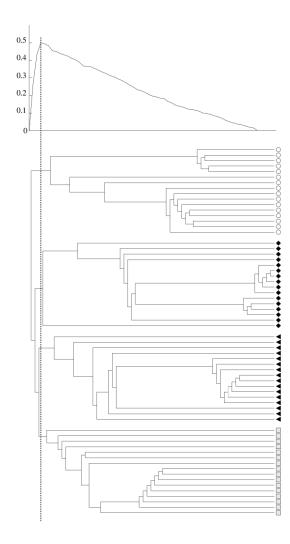
$$Q_{dw} = \frac{1}{W} \sum_{C} \left( W_C - \frac{s_C^{in} s_C^{out}}{W} \right)$$

- Modularity maximization
  - Finding partitioning that maximizes Q

#### Newman's greedy algorithm

- I. Each node in its community (agglomerative)
- 2. Merge nodes that yield the largest increase in Q
- 3. Continue until single community
- 4. Pick partition with largest *Q*

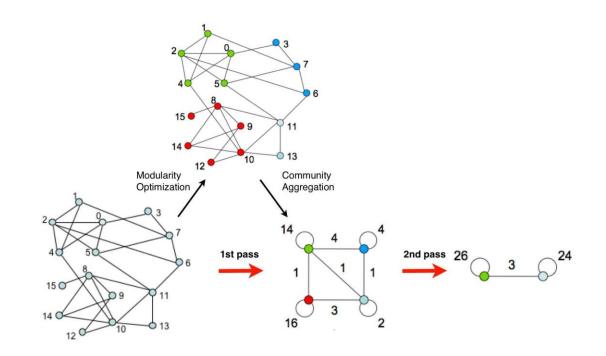
partition = nx.community.greedy\_modularity\_communities(G)



- Newman's greedy algorithm limitations
  - greedy: gets stuck on local minima
  - tends to generate unbalanced networks
  - Slow
- Standard modularity algorithm:

#### Louvain algorithm

- Similar to Newman's
- Very fast
- In python



- Modularity maximization is widely used
- Has a few problems
  - **Resolution limit**: small communities are undetected
  - Fails the random network test
- Solution: stochastic blockmodels

# **SUMMARY**

- Community detection is a huge part of network theory
- Best method depends on community definition
- Dendogram is useful for small networks
- Modularity is very popular, but comes with caveats

