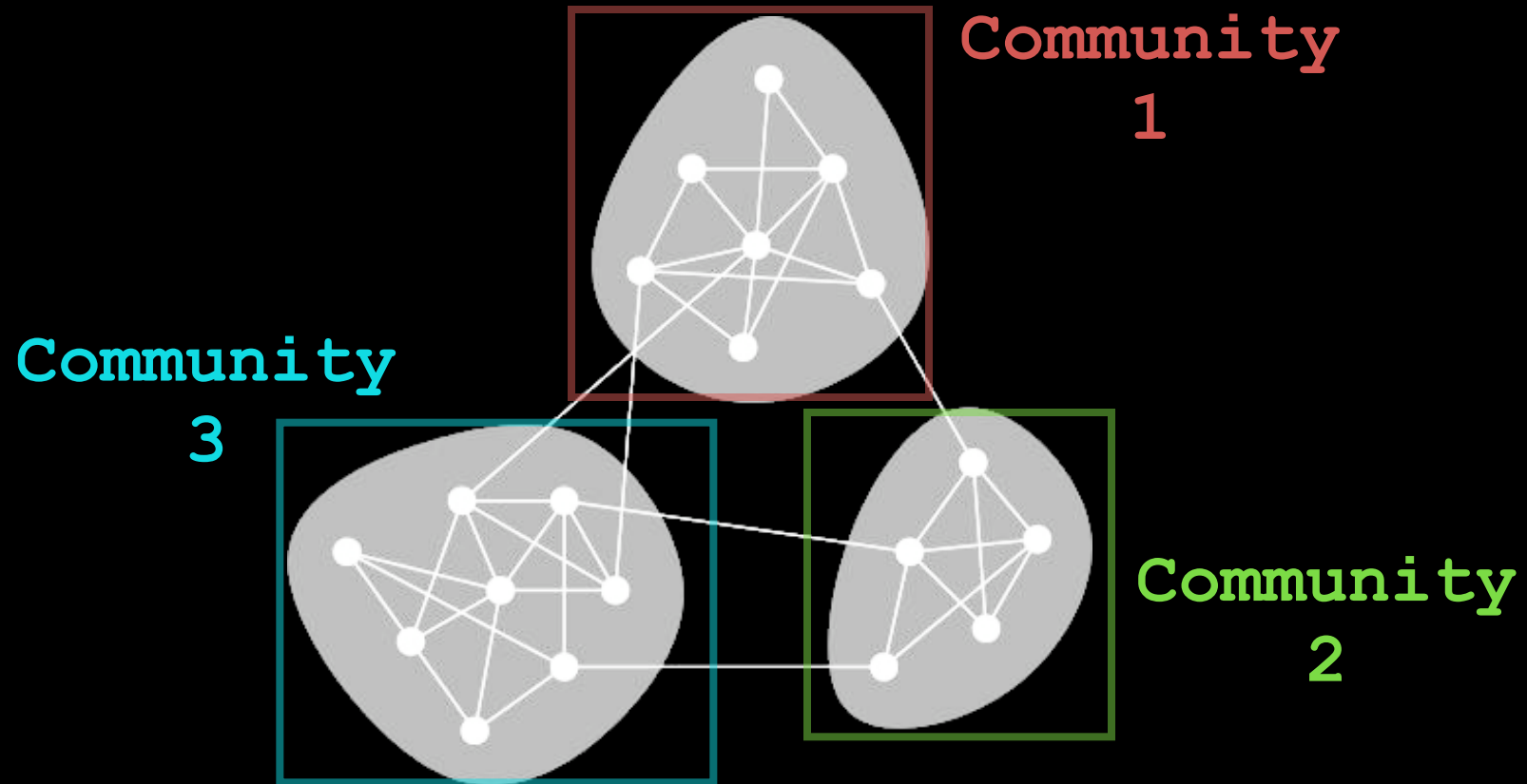


# Community Detection



# Goals

- **Tools and Datasets**
- **Community Detection Applications**
- **Measure the quality of a community**
  - ★ Modularity
  - ★ Node Similarity
- **Divide a graph into communities**
  - ★ Girvan and Newman (Divisive Algorithm)
  - ★ Hierarchical Clustering (Agglomerative Algorithm)

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# Python Library

## Small Graphs

- **networkx**
  - <https://networkx.github.io/>
  - Suitable for small graphs (~10,000 nodes)
  - Too slow for big graphs

# Python Library

## Large Graphs

- **igraph** (C code)

- ★ <http://igraph.org/python/doc/igraph.clustering-module.html>

- **graph-tool** (Heavily optimized C code)

- ★ Takes ~1/2 hour to install

- ★ <https://graph-tool.skewed.de/static/doc/community.html>

# Benchmarks

**N = 39796 vertices**

**E = 301498 edges**

Algorithm	graph-tool (4 cores)	graph-tool (1 core)	igraph	NetworkX
Single-source shortest path	0.0064 s	0.0063 s	0.012 s	0.127 s
PageRank	0.193 s	0.555 s	0.781 s	34.26 s
K-core	0.0205 s	0.0250 s	0.0181 s	0.9586 s
Minimum spanning tree	0.0268 s	0.0296 s	0.0397 s	0.413 s
Betweenness	579.7 s (~9.6 mins)	1977.6 s (~33 mins)	1182.6 s (~19.7 mins)	53716.692 s (~14.9 hours)

# Visualization and Database

- **Neo4j (Database)**

- ★ <http://console.neo4j.org/>

- **Gephi (Visualization)**

- ★ <http://gephi.github.io/>



# Network Data Sources

- <http://snap.stanford.edu/class/cs224w-2012/resources.html>
- <http://konect.uni-koblenz.de/>

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# Community Detection

## Learning to Discover Social Circles in Ego Networks

**Julian McAuley**

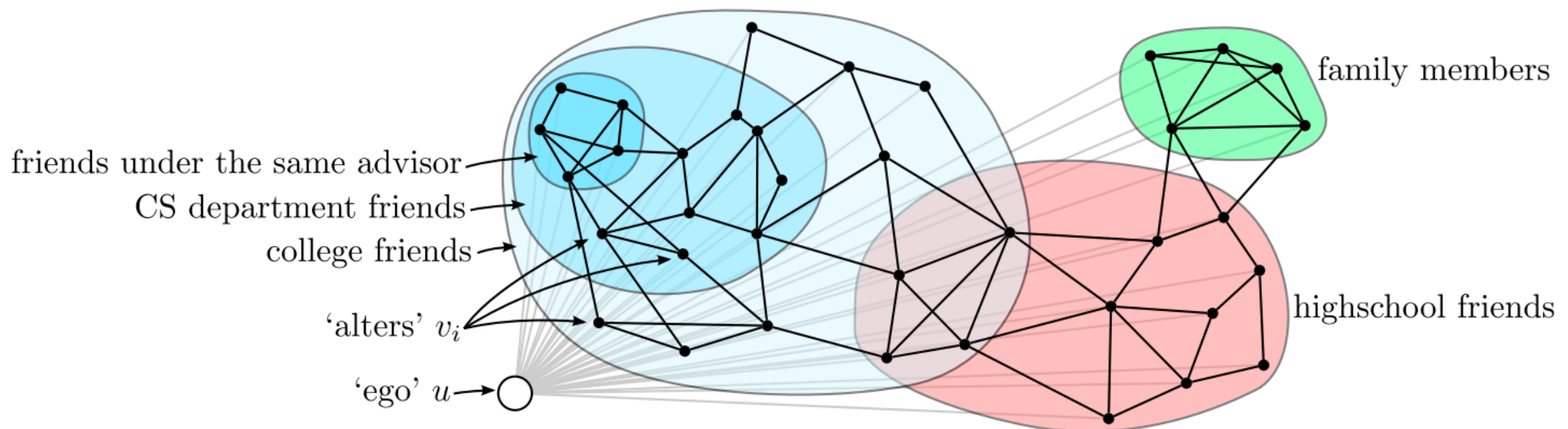
Stanford, USA

`jmcauley@cs.stanford.edu`

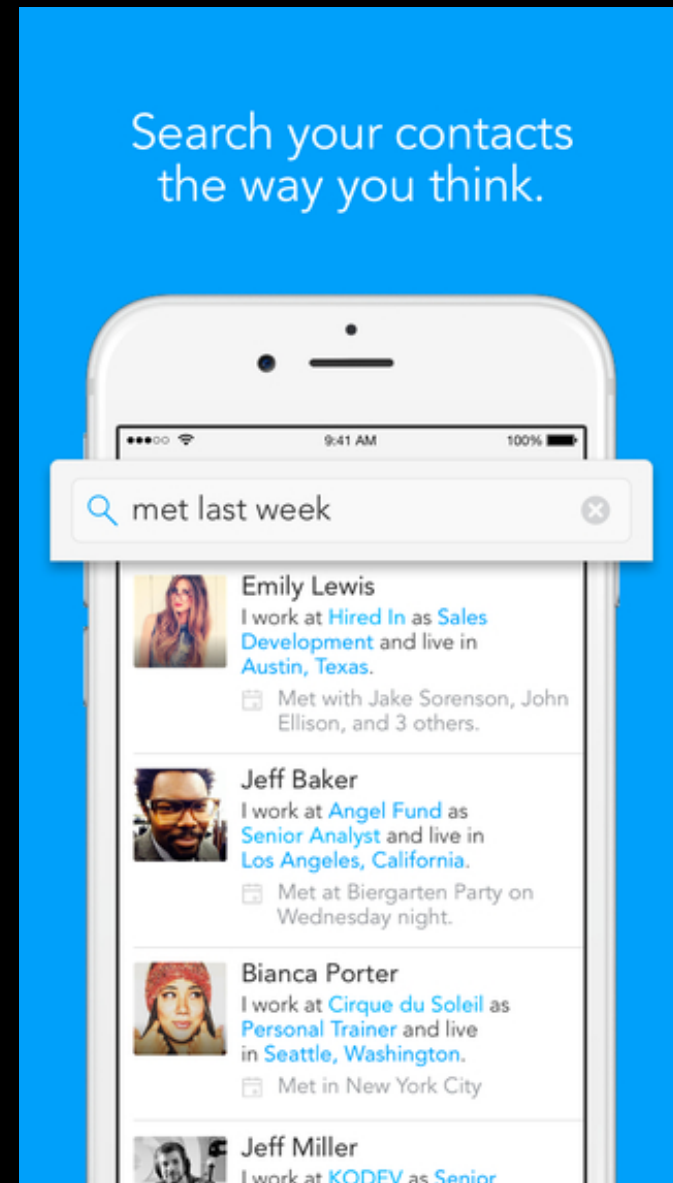
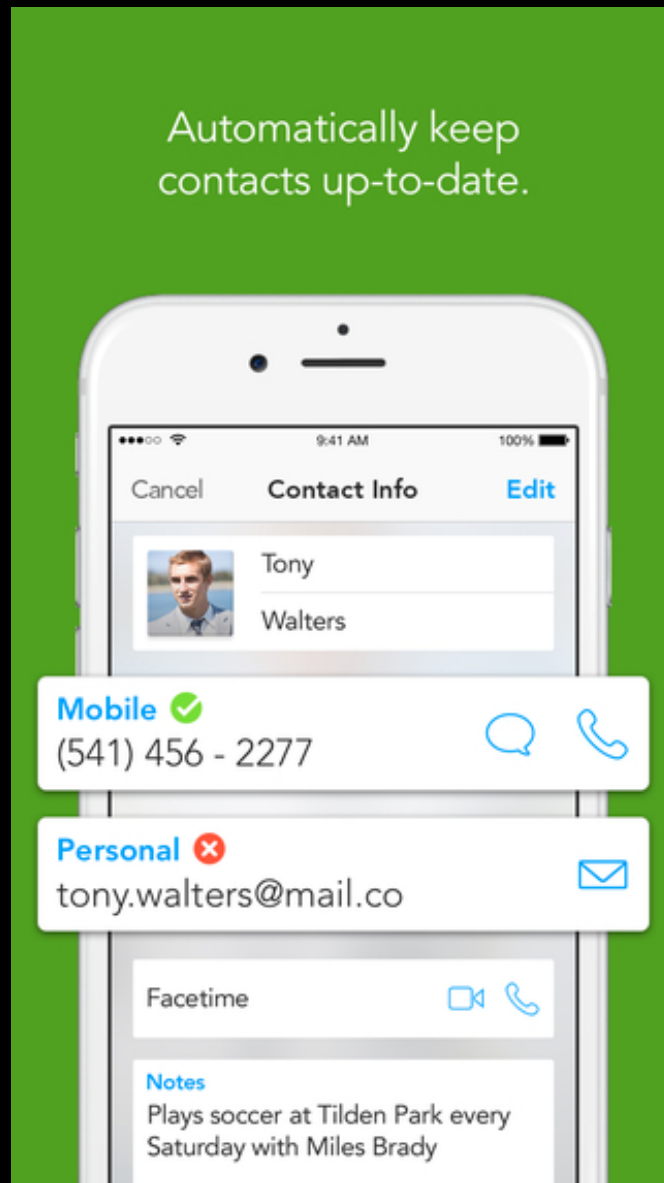
**Jure Leskovec**

Stanford, USA

`jure@cs.stanford.edu`



# Community Detection Cont.



# Goals

- **Tools and Datasets**
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# Community Quality

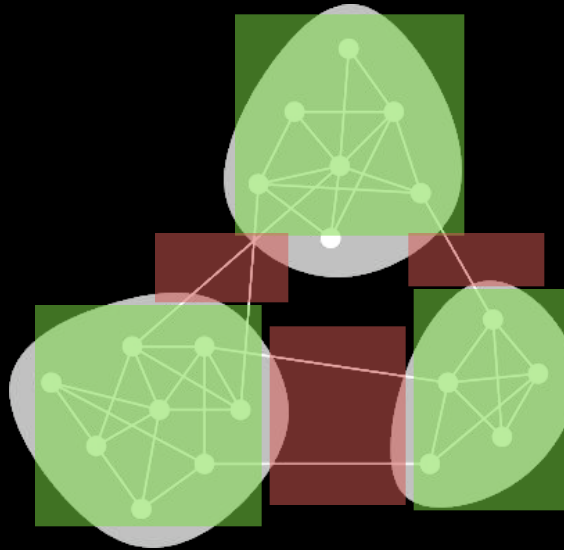
- Quantitative definition of a community
- Used to guide the division of communities

# A General Idea

# Edges  
"inside"  
community A

>

# Edges  
"between"  
community A  
and others



# Modularity

- Density of edges within a subgraph
- Minus baseline edge density of the same subgraph randomized

**High modularity = “Better” community**



Fraction of edges  
within a subgraph

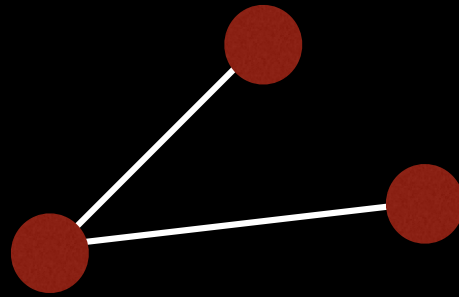
—

Fraction of edges if  
edges were randomly  
distributed in the  
subgraph

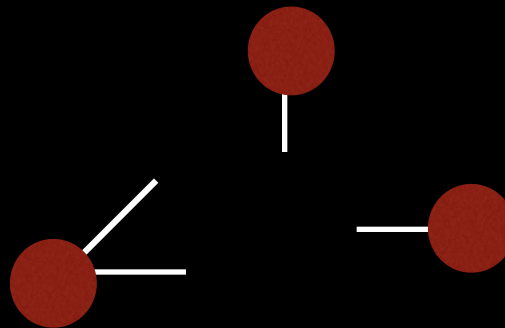
- Edge density must be *higher than expected at random* to be regarded as “high”

# Randomizing Subgraph

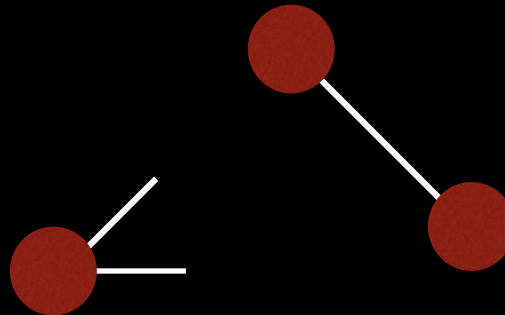
1. Halve each edge (stub) in the subgraph
2. Rewire stubs randomly to other nodes / self node
3. Degree distribution of randomized subgraph remains the same



Original  
Subgraph



Intermediate



Randomized  
Subgraph

$$Q = \frac{1}{4m} \sum_{\substack{i,j \\ \text{in same} \\ \text{module}}} \left( A_{ij} - \frac{k_i k_j}{2m} \right)$$

Normalization  
 Adjacency between  $i$  and  $j$  (0/1)  
 Prob. of Edge between  $i$  and  $j$

$k_i$  = Degree of Node  $i$   
(Randomized Graph)

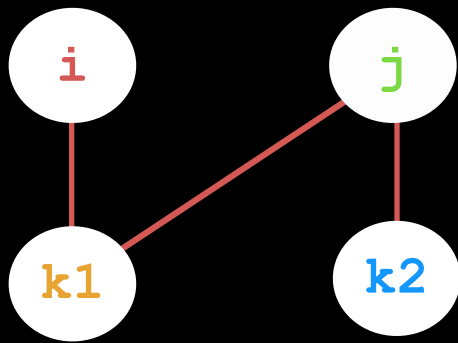
$m$  = # of edges

# Node similarity

- The number of neighbors 2 nodes share
- A lot of shared neighbors = High similarity

$$n_{ij} = \sum_k A_{ik} A_{kj}$$

# Node similarity Example



$$\mathbf{A}_{ik1} = 1$$

$$\mathbf{A}_{jk1} = 1$$

$$\mathbf{A}_{ik2} = 0$$

$$\mathbf{A}_{jk2} = 1$$

$$\begin{aligned} n_{ij} &= ( \mathbf{A}_{ik1} * \mathbf{A}_{jk1} ) + ( \mathbf{A}_{ik2} * \mathbf{A}_{jk2} ) \\ &= ( 1 * 1 ) + ( 0 * 1 ) \\ &= 1 \end{aligned}$$

# Remarks about Node similarity

- Node similarity is not normalized
- According to the degree of the nodes
- Usually measure **similarity** with **Cosine Similarity**
- Or **(dis)similarity** with **Euclidean Distance**

# Node (Dis)Similarity: Euclidean Distance

$$d_{ij} = \sum_k (A_{ik} - A_{jk})^2$$

$$normal(d_{ij}) = \frac{d_{ij}}{\boxed{k_i} + \boxed{k_j}}$$

Degree of node *i*      Degree of node *j*



# Node Similarity: Cosine Similarity

# of shared neighbors

$$\sigma_{ij} = \frac{n_{ij}}{\sqrt{k_i k_j}}$$

Degree of node  $i$       Degree of node  $j$

# Goals

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# How to divide a graph

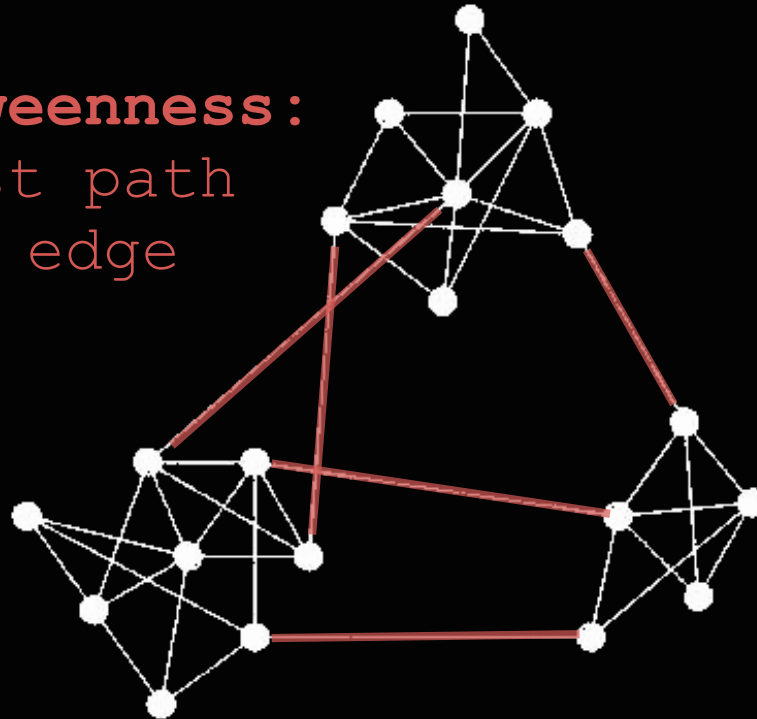
	Divisive	Agglomerative
Approach	Top-down	Bottom-up
Starting Point	Graph	Individual Nodes
Community Formation	Removing edges	Iteratively merging
Technique	Girvan and Newman	Hierarchical Clustering
More popular	Yes	No

# Divisive Algorithm

## Girvan and Newman

**High edge betweenness:**

Many shortest path  
passes the edge



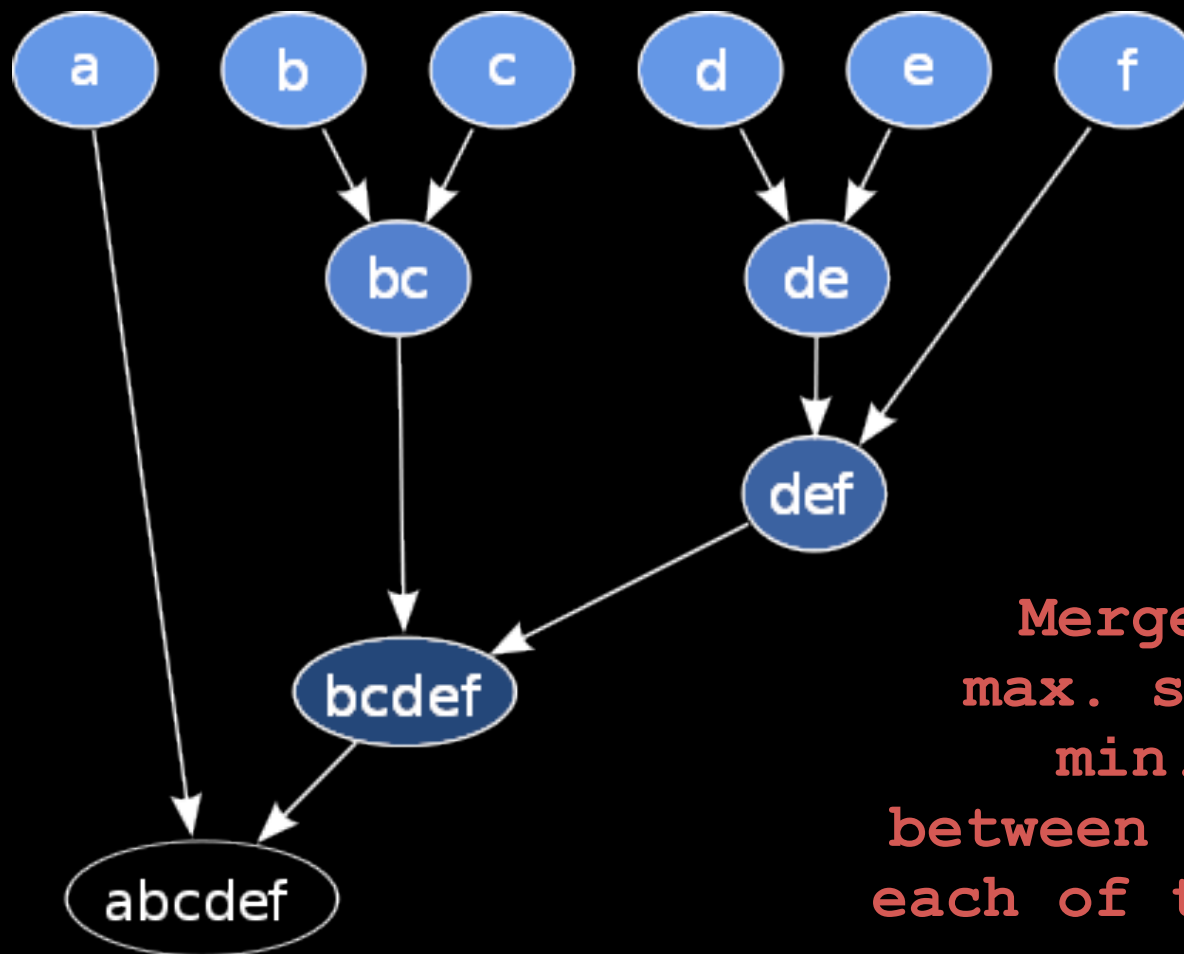
# Girvan and Newman

1. Compute betweenness for all edges
2. Remove edge with largest betweenness
3. Recalculate betweenness

4. Calculate modularity if new communities formed
5. Stop if average modularity is maximized  
(i.e. Further iteration would reduce modularity)
6. Otherwise repeat from step 2

# Agglomerative Algorithm

## Hierarchical Clustering



Merge based on  
max. similarity /  
min. distance  
between elements from  
each of the 2 clusters

# Single Linkage Clustering

Merge based on the minimum distance between  
elements of 2 clusters

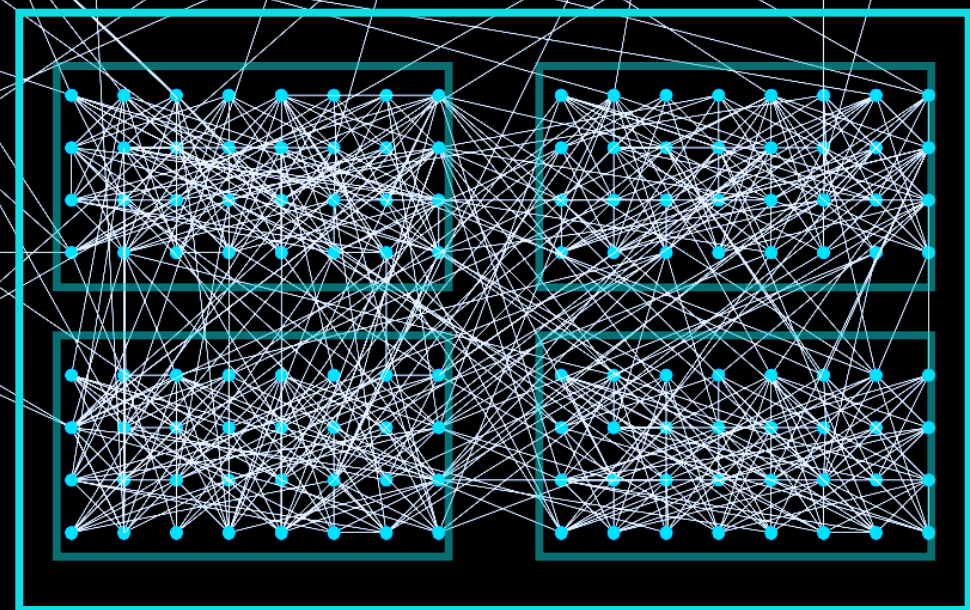
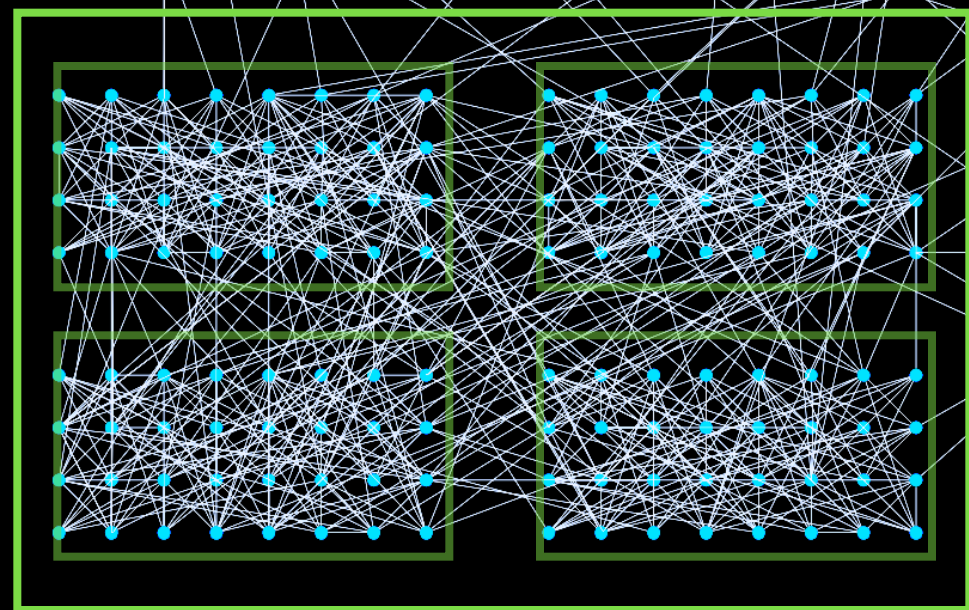
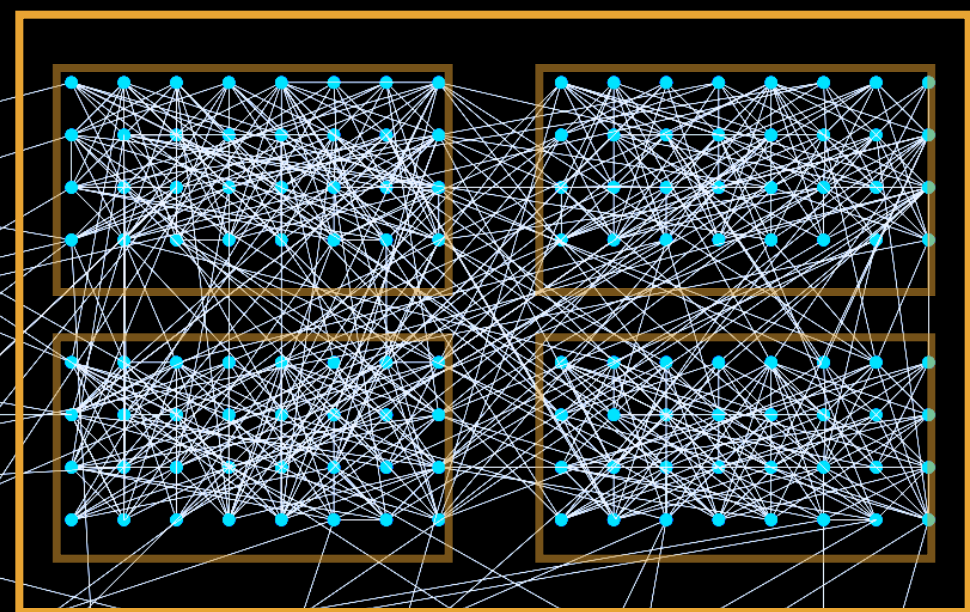
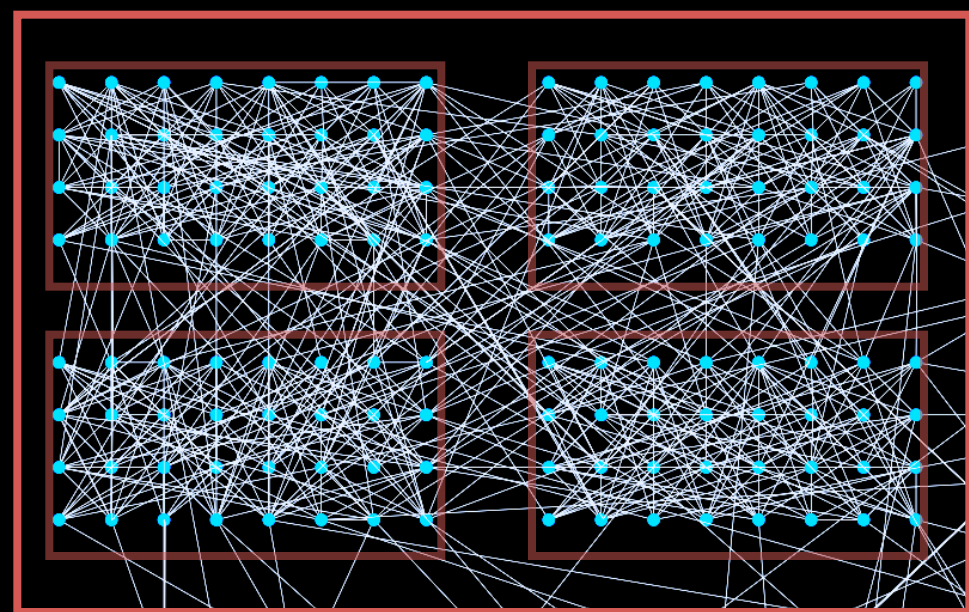
$$\min\{ d(x, y) : x \in \mathcal{A}, y \in \mathcal{B} \}.$$

*A and B are 2 separate clusters*



# Remarks about Hierarchical Clustering

- Have to decide cut-off
- Hierarchy is by construction, not always sensible
- Good for networks that are hierarchical  
(social / biological)



# Summary

- Applications to detect social community
- Different metric to evaluate communities
- Different algorithms to create communities from graph

# Next Steps

- **Cliques** and **k-core** as a measure of similarity

★ p. 10 – 11 (<http://arxiv.org/pdf/0906.0612.pdf>)

- **Kernighan-Lin (Minimum bisection)** algorithm

★ p. 17 – 18 (<http://arxiv.org/pdf/0906.0612.pdf>)

- **Partitional Clustering**

★ p. 19 – 20 (<http://arxiv.org/pdf/0906.0612.pdf>)

- **igraph** and **graph-tool**