



Finding Overlapping Communities in Social Networks: Toward a Rigorous Approach

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Introduction

- What is a **community in a social network**?
 - a group of nodes more densely connected with each other than with the rest of the network
- Communities **overlap** each other
- Direct approach → **NP-hard** problems
- Heuristic or generative model approach → **egg & chicken** problem
- Instead: Assumptions are based on **ego-centric networks**
 - Studied in sociology
 - Suggested algorithms also have ego-centric analysis feel

Assumptions

0. Each person participates in **up to d communities**
 - d is constant or small
1. **Expected degree model**
 - Each node u in community C has an *affinity* $p_u \in [0, 1]$
 - The edge (u, v) exists with probability $p_u p_v$
2. **Maximality** with gap ε
 - If for $u, v \in C$, (u, v) exists with probability α ,
then $w \notin C$ has edges to $\leq \alpha - \varepsilon$ fraction of nodes in C
3. **Communities explain γ fraction** of each person ties

First Step: Communities are Cliques

- Another Assumption: $\forall C : \delta k \leq |C| \leq k$
- Output each community with prob. $\geq 2/3$
 - in time $O(nk) \cdot 2^{\tilde{O}(\log^2 d)}$
- Algorithm Description
 1. Pick **starting nodes** uniformly at random
 2. For each starting node v , **randomly sample** $S \subseteq \Gamma(v)$
 3. Look at **cliques** U in $G(S)$
 4. Let V' be the set of nodes in $\Gamma(v)$ which are **connected to all** nodes in U
 5. Return **high degree vertices** from $G(V')$

Communities are Dense Subgraphs

- **Setup 1:** $p_u = \sqrt{\alpha}$
 - Find each community
 - With high probability over G randomness
 - With prob. $2/3$ over algorithm randomness
 - In time $O(nk) \cdot 2^{\tilde{O}(\log^2 d)}$
- **Setup 2:** $p_u \geq \sqrt{\alpha}$
 - Need to **loop over** all $S \subseteq \Gamma(v)$ of size T
 - Sample $G(\Gamma(S))$ for each S
 - Worse running time: $O(n \cdot (kd)^T) \cdot 2^{\tilde{O}(\log^2 d)}$

Communities with Very Different Sizes

- Sampling may **miss small communities**
 - So previous ideas will not work
- **Definition:** A is a $(\alpha, \alpha - \varepsilon)$ -set if
 - Nodes in A have edges to $\geq \alpha$ fraction of nodes in A
 - Outside nodes have edges to $\leq \alpha - \varepsilon$ fraction of nodes in A
- **Algorithm** (assuming $p_u \geq \sqrt{\alpha_{min}}$)
 1. For $\alpha = 1$ downto α_{min} step $-\varepsilon/4$
 - 1.1. For all sets of nodes S of size T
 - 1.1.1. $U = \{v: \geq \alpha - \varepsilon/4 \text{ fraction of its edges are to } S\}$
 - 1.1.2. Return U if it is a $(\alpha, \alpha - \varepsilon/2)$ set
- **Running time:** $n^{C \log kd}$ (not polynomial)

Cliques with Very Different Sizes

- Looking for a **polynomial** algorithm for **cliques**
- **Extra assumptions** are needed:
 - Distinctness: For $u \in C$, at least a constant factor of C does not lie in any other community containing u
 - Duck assumption
 - Small communities are distinguishable from “noise” edges
- Polynomial algorithm description
 - Find large cliques first (sampled easily), then ignore their edges
 - Extra assumptions ensure smaller cliques can be found

Relaxing the Assumptions

- **Expected degree model** assumption can be relaxed if:
 - The following are **concentrated** near their expectation:
 - **# of edges** from any node u to any community C
 - **Degree** of each node
 - **Intersection** of two nodes in a community
- **Gap assumption**
 - Can be relaxed if:
 - $\forall C : \delta k \leq |C| \leq k$
 - Communities are cliques or $p_u = \sqrt{\alpha}$
 - The returned communities will be close to the real ones

Sparser Communities

- Different assumptions
 - (u,v) exists with probability B/\sqrt{k} (where $|C| = k$)
 - All edges belong to some community
 - Communities intersection size is limited
- Transform G to a dense graph G'
 - Nodes are the same
 - (u,v) exists in G' iff they have $\geq B^2/2$ length-2 path in G

Summary

case no.	extra / different assumptions?	probability of edges in communities	communities sizes must be similar?	running time
1	No	Cliques	Yes	Polynomial
2	No	$p_u = \sqrt{\alpha}$	Yes	Polynomial
3	No	$p_u \geq \sqrt{\alpha}$	Yes	Polynomial
4	No	$p_u \geq \sqrt{\alpha}$	No	Quasi-Poly
5	Extra	Cliques	No	Polynomial
6	Different	Sparse	Yes	Polynomial

Areas of Possible Further Research

- Releasing the assumptions in more cases
 - Expected degree model assumption
 - Maximality (gap) assumption
- Polynomial algorithm for dense communities with different sizes
- Fast implementation using heuristics
- Testing on real-world data
- Adapting the algorithms to a dynamic setting



Questions?