Clique and Link Percolation

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Overview

- Classifications of Communities
- Overlapping Communities
- Clique Percolation
- Link Clustering
- Applications

Definitions

- Connectedness Hypothesis: A community is a connected subgraph
- Density Hypothesis: Nodes in a community are more likely to connect to each other than to outside nodes



Classifications of Communities

- Clique: A completely connected subgraph
- Strong Community: A connected subgraph whose nodes have more links to nodes within the same community than nodes outside the community
- Weak Community: A connected subgraph whose total internal degree exceeds the total external degree









Overlapping Communities

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- A node is rarely confined to a single community
- How can we detect overlapping communities?



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1) A community is the union of overlapping, or adjacent, cliques	2) Two k-cliques are considered adjacent if they share k-1 nodes.
3) A k-clique community is the largest connected subgraph obtained by the union of all adjacent k-cliques	4) K-cliques that cannot be reached from a particular k-clique belong to another k-clique community











Can these emerge by chance?

 Random networks can produce very large cliques if they have a very high density

$$p_c(k) = rac{1}{\left[(k-1)N
ight]^{1/(k-1)}}$$

k = clique size, N = number of nodes

- Subcritical Communities: $p < p_c(k)$
- Supercritical Communities: $p > p_c(k)$



p = 0.22, p_c(3) = 0.16



Link Clustering

- Nodes can belong to many communities but links typically are community specific
- Algorithm finds link communities by exploring the similarity between their neighbors and themselves

Link Clustering Algorithm

 Link Similarity can be defined as S:





$$S((i,k),(j,k)) = \frac{|n_{+}(i) \cap n_{+}(j)|}{|n_{+}(i) \cup n_{+}(j)|} d.$$

 Identify link clusters using a matrix



Clique Percolation in Biological Networks

- CFinder: Locating cliques and overlapping modules in biological networks by Balázs Adamcsek, Gergely Palla, Illés J. Farkas, Imre Derényi, and Tamás Vicsek
- Identify which groups of proteins interact with each other

network of yeast PPI modules node: module of proteins, link: overlap of modules (b) vps17 Vps29 (a) Vps35 Vps26 Vps5 retromer complex (C) Rad10 Rad14 Rad1 Msh2 nucleotide-excision repair factor 1 complex + Msh2 (d) | vps11 Vps39 Vps16 Vps18 Vps33 Vps41 Vps8 HOPS complex + Vps8

enlarged portions of the network of modules



References

Adamcsek, Balazs, et al. *CFinder: Locating Cliques and Overlapping Modules in Biological Networks*.

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