Uncovering the overlapping modular structure of complex networks

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Why modules (densely interconnected parts)?

The internal organization of large networks is responsible for their function.

Complex systems/networks are typically *hierarchical*.

The units organize (become more closely connected) into groups which can themselves be regarded as units on a higher level.

We call these densely interconnected groups of nodes as modules/communities/cohesive groups/clusters etc. They are the "building blocks" of the complex networks on many scales.

For example:

Person->group->department->division->company->industrial sector

Letter->word->sentence->paragraph->section->chapter->book

Questions:

How can we recover the hierarchy of overlapping groups/modules/communities in the network if only a (very long) list of links between pairs of units is given?

What are their main characteristics?

Outline

- Basic facts and principles
- Community finding *versus k*-clique percolation
- Results for protein interaction, word association, phone calls, school friendship and collaboration networks

Basic observations:

A large complex network is bounded to be highly structured (has modules; function follows from structure)

The internal organization is typically hierarchical

(i.e., displays some sort of self-similarity of the structure)

An important new aspect: Overlaps of modules are essential





Complexity is between randomness and regularity

Graduate School

Role of overlaps















Hierarchical versus template rolling clustering

Common clustering methods lead to a partitioning in which someone (a node) can belong to a single community at a time only.

For example, I can be located as a member of the community "physicists", but not, at the same time, be found as a member of my community "family" or "friends", etc.

k-clique template rolling allows large scale, systematic (deterministic) analysis of the network of overlapping communities

k-CLIQUE PERCOLATION

with I. Derényi and G. Palla



Definitions

k-clique: complete subgraph of *k* vertices

k-clique adjacency: two *k*-cliques share a *k*-1 – clique

k-clique walk: series of steps to adjacent *k*-cliques

k-clique cluster: set of vertices of all *k*-clique walks from a given *k*-clique



(E-R percolation is the *k=2* case)

I. D, G. P. and T.V., Phys. Rev. Lett. 2005



Order parameter for clique percolation, k=4

Percolation threshold at

 $p_c(k) = [(k-1)N]^{(-1/(k-1))}$

The scaling of the relative size of the giant cluster of k=3,4 and 5-cliques at p_c

For $k \le 3$, $N_k^* / N_k(p_c) \sim N^{-k/6}$ For $k \ge 3$ $N_k^* / N_k(p_c) \sim N^{1-k/2}$

UNCOVERING THE OVERLAPPING COMMUNITY STRUCTURE OF COMPLEX NETWORKS IN NATURE AND SOCIETY

with G. Palla, I. Derényi, and I. Farkas

Definitions An order *k* community is a *k*-clique percolation cluster

Such communities/clusters obviously can overlap This is why a lot of new interesting questions can be posed

New fundamental quantities (cumulative distributions) defined:

P(d ^{com})	community degree distribution
<i>P(m)</i>	membership number distribution
$P(S^{ov})$	community overlap distribution
P(s)	community size distribution (not new)

G.P,I.D,I.F,T.V Nature 2005

DATA

cond-mat (electronic preprints, about 30,000 authors) protein-protein (DIP database, yeast, 2,600 nodes) word association (sets of words associated with given words, questionnaire, 10,600 words) mobile phone (~ 4,000,000 users calling each other) school friendship (84 schools from USA)

large data sets: efficient algorithm is needed! Our method is the fastest known to us for these type of data

Steps:

determine: cliques (not k-cliques!) clique overlap matrix components of the corresponding adjacency matrix

Do this for "optimal" k and w, where optimal corresponds to the "richest" (most widely distributed cluster sizes) community structure





"Web of networks"

Each node is a community

Nodes are weighted for community size Links are weighted for overlap size

DIP "core" data base of protein interactions (S. cerevisiase, a yeast)

The other networks we analysed are much larger!!







Community overlap size

membership number

Case studies + dynamics

Protein interaction (prediction of function)

School friendship (disassortativity of communities, role of races)

Social group evolution in a co-authorship and a mobile phone network



B.A,G.P,I.D,I.F,T.V.: BIOINFROMATICS 2006

enlarged portions of the network of modules Marked:

single proteins (function prediction) and groups (anticipated new modules)



Three schools from the Add-Health school friendship data set

Grades 7-12







Network of school friendship communities

Distribution functions (for k=3)

O communities □ individuals



P(k) – degree distribution C(k) – clustering coefficient $< k_n > (k)$ – degree of neighbour (individuals: assortative communities: diassortative)

Quantifying social group evolution

with G. Palla and A-L Barabási (Nature, April 2007)



Small part of the phone call network (surrounding the circled yellow node up to the fourth neighbour) Small part of the collaboration network (surrounding the circled green node up to the fourth neighbour

Callers with the same zip code or age are over-represented in the communities we find



Examples for tracking individual communities.



Lifetime (τ) of a social group as a function of stability (steadiness, ζ) and size (s)



Probability of disintegrating (p_d) and the lifetime (τ^*) of a community whose members have a total amount of "commitments" to other communities equal to W_{out}



Home page

of

CFinder





Social network of the 3000 employees of an European company determined from an on-line survey. Visualization of the betweenness centrality



Visualization of the communities for the same company shown here using an adaptation of our CFinder-Firmnet software.

Theridion provides organizational development services based on network analysis.

Outlook:

Networks of networks

- hierarchical aspects
- correlations, clustering, etc., i.e., everything you can do for vertices
- applications, such as protein function prediction or organizational development



This will also become a commercial product by **Firmlinks** with **GORDIO**, a Budapest based HR company

Internal organization of large complex networks in terms of their modular structure

- Research on modules/communities is a very active field (Amaral, Barabási, Newman + many further groups)
- How does a large complex network may look like?



To find overlapping communities we

consider: connected groups (clusters) of motifs e.g. a 4-clique

define: a cluster of adjacent complete subgraps (cliques) is a community (simple assumption)

Two aspects

- I) *k*-clique percolation
- II) communities in large real networks:

overlaps and their statistics

Evolution of a single large community of collaborators

s – size (number of authors), t – time (in months)



Dedicated home page (software, papers, data) http://angel.elte.hu/clustering/

<u>Home</u>

Screen shots

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Communities in a "tiny" part of a phone calls network of 4 million users (with A-L Barabási and G. Palla

Nature, April 5 2007)





Information about the age distribution of users in communities of size *s* (Ratio of the standard deviation in a randomized set over actual)

Information about the Zip code (spatial) distribution of users in communities of size *s*

(Ratio of the standard deviation in a randomized set over actual)



A.-L. B., H.J, Z.N., E.R., A. S., T. V. (Physica A, 2002)



Data: collaboration graphs in (M) Mathematics and (NS) Neuroscience

Cumulative data, 1991 - 98

Degree distribution:

power-law with

$$\gamma_{\rm M} = 2.1, \ \gamma_{\rm NS} = 2.4$$
 due to growth and





Internal preferential attachment:



communities of collaborators are formed

The scaling of the relative size of the giant cluster of k-cliques at p_c



For $k \le 3$, $N_k^* / N_k(p_c) \sim N^{-k/6}$ For $k \ge 3$ $N_k^* / N_k(p_c) \sim N^{1-k/2}$



Distribution of community sizes

Over-representation of the usage of a given service as a function of the number of users in a community

Community dynamics

with P. Pollner and G. Palla

Dynamics of community growth: the preferential attachment principle applies on the level of communities as well



The probability that a previously unlinked community joins a community larger than *s* grows approximately linearly (for the cond-mat coauthorship network)

P.P,G.P,T.V Europhys Lett. 2006