Szemerédi-type clustering of peer-to-peer streaming system

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MODELING, ANALYSIS, AND CONTROL OF COMPLEX NETWORKS

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agenda:

review of Szemerédi's Regularity Lemma
Peer-to-peer streaming system
Szemerédi-type clustering of p2p-system

Szemerédi's Regularity Lemma (SzRL)

> a fundamental result in graph theory



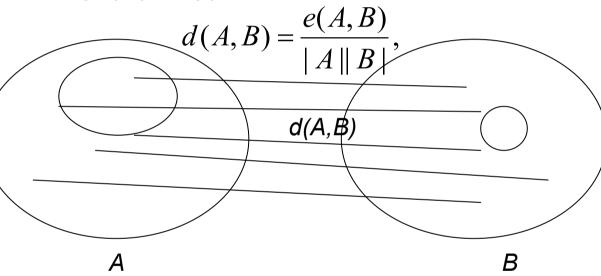
Szemerédi, E.:'Regular partitions of graphs', 1978

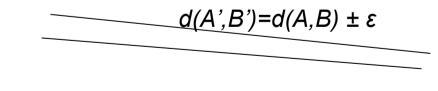
an 'ɛ-regular pair' (A,B) is a graph:

A and B are disjoint node sets

A'

link density (d(A,B)) between pair is almost uniform





B'

for all: $|A'| \ge \varepsilon |A|$, $|B'| \ge \varepsilon |B|$

SzRL

for every $\epsilon > 0$, and for every natural number m

→ \exists N=N(ϵ ,m) and M=M(ϵ ,m), natural numbers

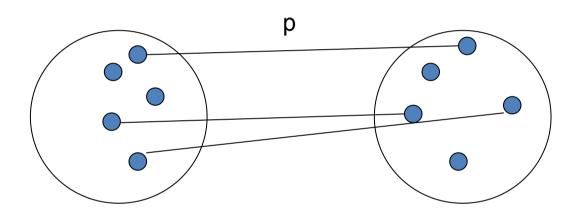
such that:

- any graph, G, with at least N nodes can be partitioned into k classes of almost equal sizes: (V₁, V₂, ..., V_k)
- ▶ m≤ k ≤ M and all but at most $εk^2$ pairs (V_i,V_s) are ε-regular

(then $(V_1, V_2, ..., V_k)$ is called an <u> ϵ -regular partition</u> of G)

note:

> a large random bipartite graph is 'quite' regular



- Ink is drawn with prob. p, and not drawn with prob. 1-p
- link density is close to p
- ➤ regularity ≈ uniformity (of links)

SzRL ≈> a large enough graph can be well approximated by bounded number of pseudo-random bipartite graphs $d_{2,3}$ $d_{1.2}$ G $d_{2,k}$ ≻G => Sz. graph G => > space of all graphs is totally bounded or precompact

some remarks on SzRL

- most effective for large and dense graphs
- sparse version (Scott 2010)?
- 'must' in extremal graph theory
- interesting new results:
 - > O(n) time algorithm to find reg. partitions (E. Fischer et al, 2010)
 - Sz. graph can be found in constant time! (E. Fischer et al, 2010, T. Tao, 2009)
 - <u>Theorem</u>: a graph property, *P*, is effectively testable verified from corresponding Sz. graphs (N. Alon, et al 2005)
 - Sz. graph tolerates substantial noise, spectral methods, M. Bolla 2005

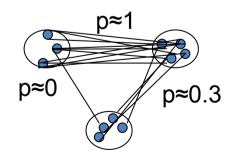
although SzRL does not guarantee reg. partitioning for any 'real life' graph:

because of terrific upper-bounds (M) like:

regular partition can still be meaningful
a reasonable model in many cases:

- Sz. graph as a model
 - fit real life graph with such model in max likelihood sense

- Itry to find partitioning where pairs look as random as possible => as regular as it can get(?)
- try to cluster nodes in a way that pairs are like random bipartite graphs:

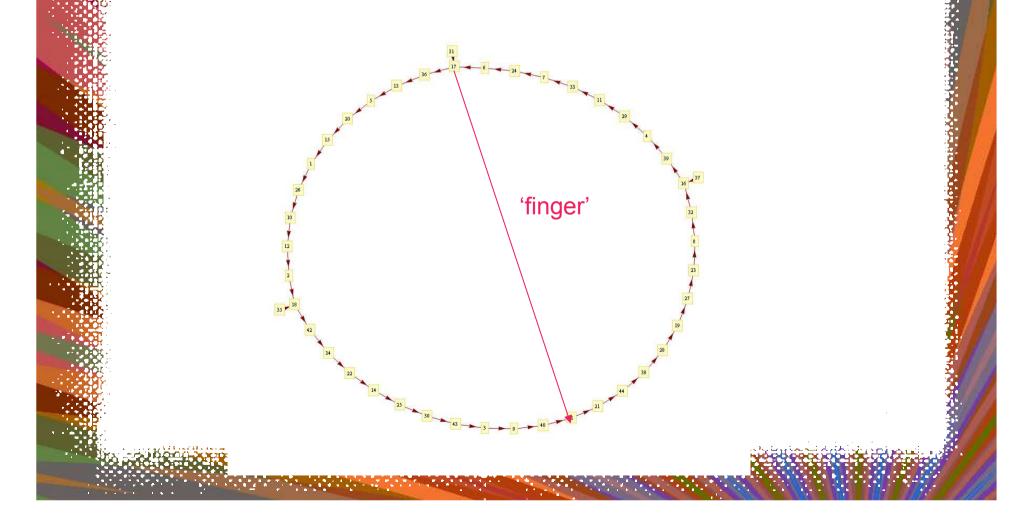


we adopted the last approach from:

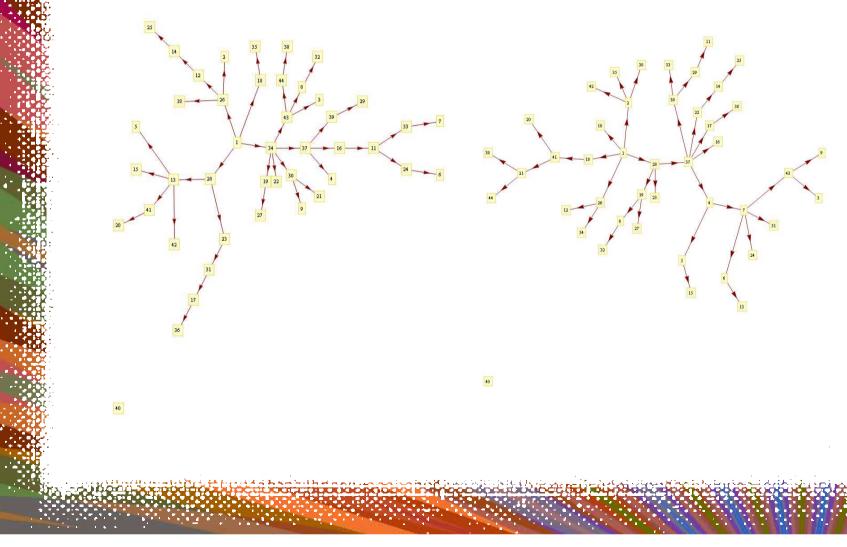
- Nepusz-Bazsó-Négyessy-Tusnády, "Reconstructing cortial networks..." Springer 2008.
- network of brain areas with neural connections as links
- with uncertain links
- Sz. clustering => predictions about such uncertain links

our experimental peer-to-peer streaming system, tests in PlanetLab

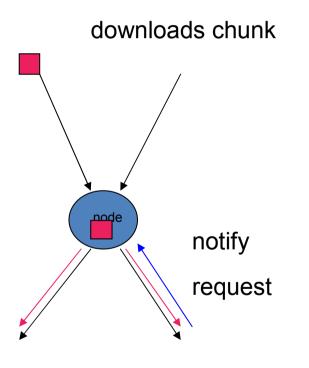
≻48 nodes organizing a Chord-network



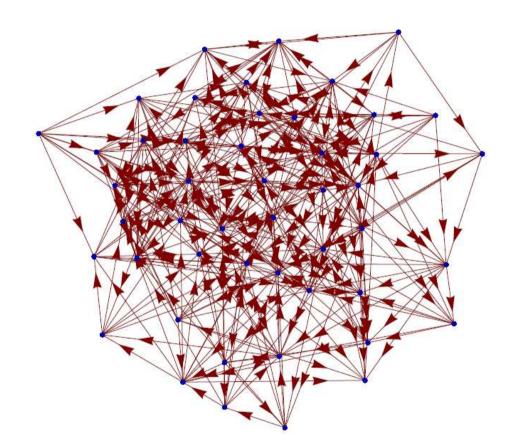
one node, the 'seed', streams chunks of the file to its antifinger neighbours -> new neighbours... a case of some two chunks:



a chunk is downloaded, neighbours are notified, one of them requests the chunk, and a peer accepts request and uploads the chunk further – a push scheme



after thousands of chunks were streamed: 'who downloaded from whom'-graph



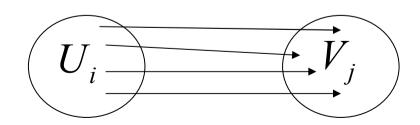
we would like to find peers downloading-uploading patterns

similar peers in the same class similarity in some Sz. clustering sense (?) without any aprior assumptions what the classes should be in directed case we want to find two partitions ('in' and 'out' clusters)

- in clusters: $Y = (V_1, V_2, ..., V_{Kin})$,
- and out clusters $\Omega = (U_1, U_2, ..., U_{Kout})$

$$V = V_1 + ... + V_{Kin} = U_1 + ... + U_{Kout}$$

• consider links from U_i to V_j

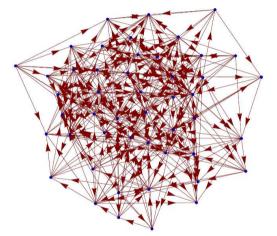


define a directed and weighted graph:

for link (i, j)

weight:

 $w_{i,j} = \frac{n_{i,j}}{\sum_{\alpha} n_{i,\alpha}}$



with $n_{i,j}$ = number of chunks node i gets from j during a long session

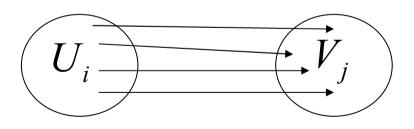
define a random graph model, \boldsymbol{g}_{W}

- with link probability: $P((i, j) \in E_w) = w_{i,j}$
- with independent links
- with the same nodes set as the original graph, V

with respect to in- and out-clusters define the weight density matrix:

$$(P)_{i,j} \coloneqq p_{i,j} = \frac{\sum_{\alpha \in U_i, \beta \in V_j} W_{\alpha,\beta}}{\left| U_i \right| \left| V_j \right|}$$

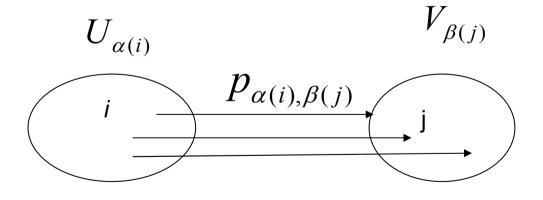
 $1 \le i \le Kout$, $1 \le j \le Kin$,



two random graphs: g_W and g_P

with link probabilities:

w(i,j) and $p_{lpha(i),eta(j)}$ with



find partitions (Ω , Y)

in such a way that the expected 'log-likelihood' is maximal:

$$l(\Omega, \mathbf{Y}) \coloneqq E_{\boldsymbol{g}_{\boldsymbol{w}}} \log P(G | \boldsymbol{g}_{P})$$

HARD STREET

where prob. of a graph G(E,V):

 $P(G \mid \boldsymbol{\mathcal{G}}_P) = \prod_{(i,j) \in V \times V: (i,j) \in E} p(i,j) \prod_{(i,j) \in V \times V: (i,j) \notin E} (1 - p(i,j))$

$$l(\Omega, \mathbf{Y}) = \sum_{(i,j) \in V \times V} \left(w_{i,j} \log p_{u(i),v(j)} + (1 - w_{i,j}) \log(1 - p_{u(i),v(j)}) \right)$$

where $i \in U_{u(i)}, j \in V_{v(j)}$

with fixed $|\Omega| = Kout, |Y| = Kin$

$$\max_{\Omega, Y} l(\Omega, Y) \coloneqq l^*(Kout, Kin)$$

(use EM-algorithm)

$$|\Omega| = Kout, |Y| = Kin???$$

from AIC, Akaike information criterion:

minimze:
$$AIC = -2l^*(Kout, Kin) + 2K$$

where *K* is number of parameters: $K = Kin \times Kout + 2|V| + 2$

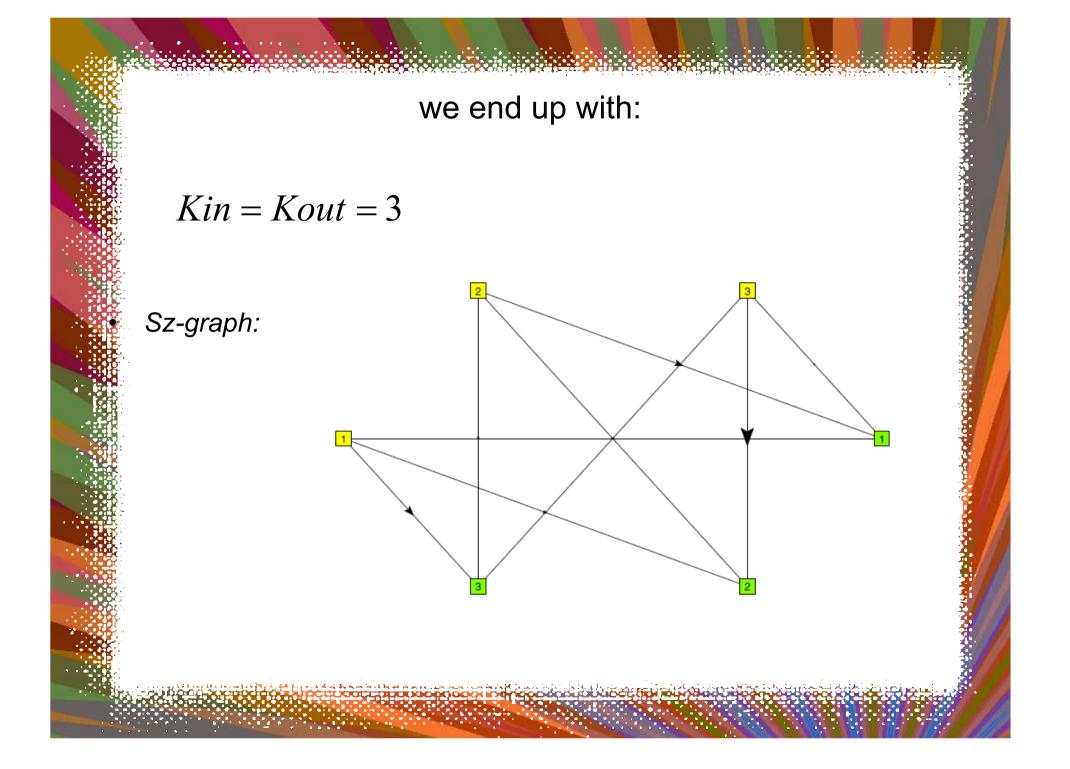
$$\Rightarrow AIC = -2l^*(Kout, Kin) + 2Kin \times Kout + 4|V| + 4$$

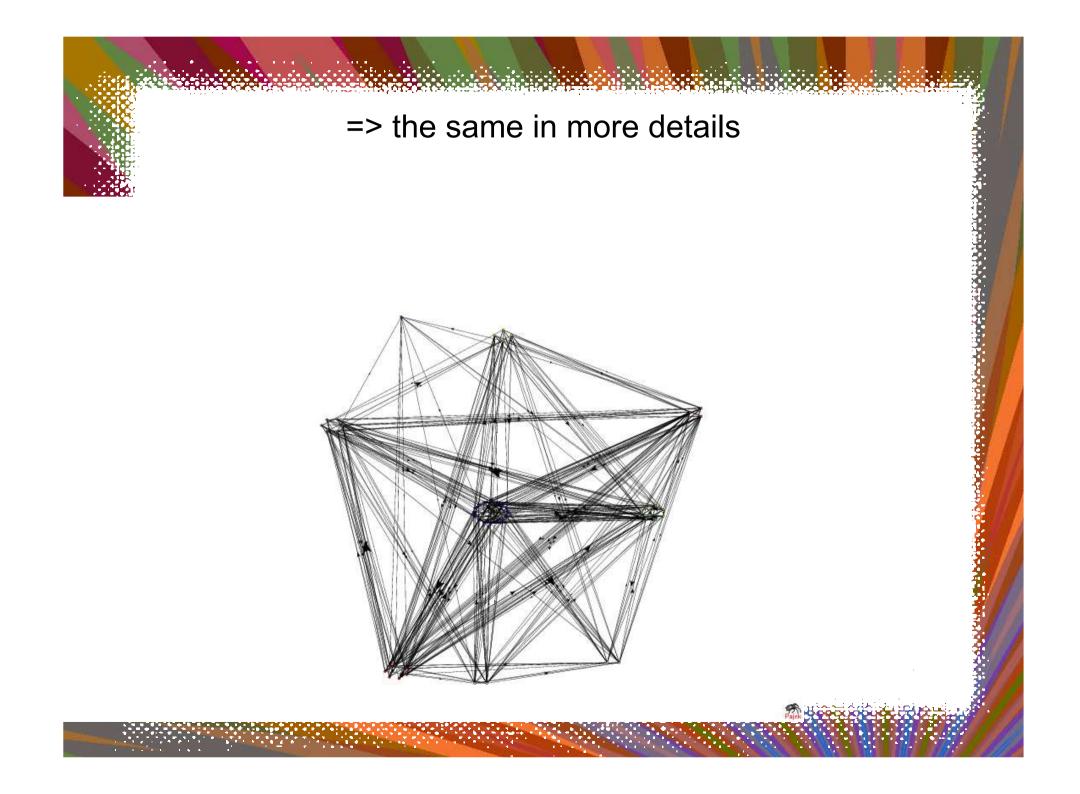
Min A/C => $|\Omega| = Kout, |Y| = Kin$

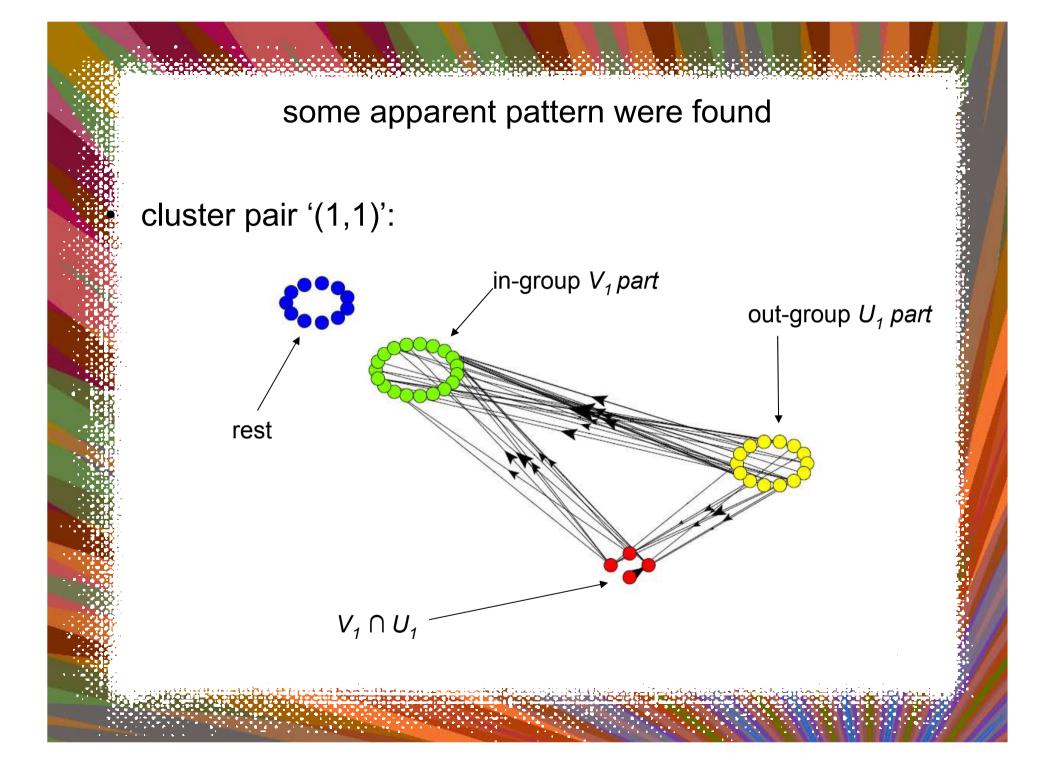
clustering algorithm:

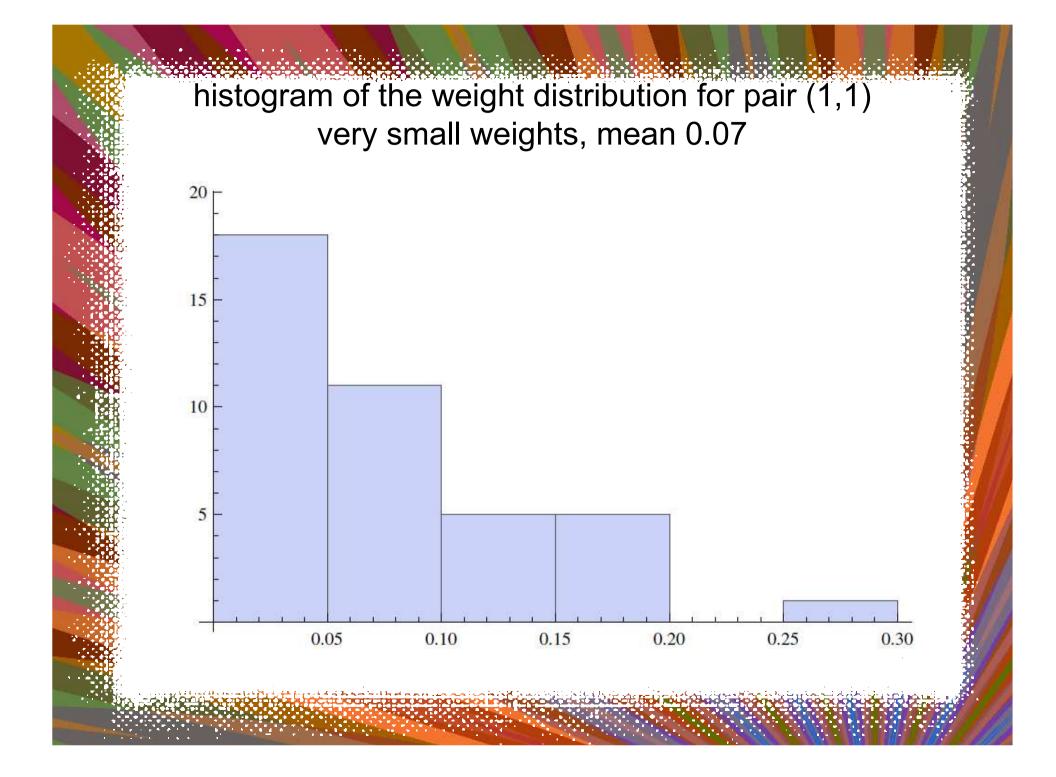
- use EM (expectation-maximisation)
- FOR:

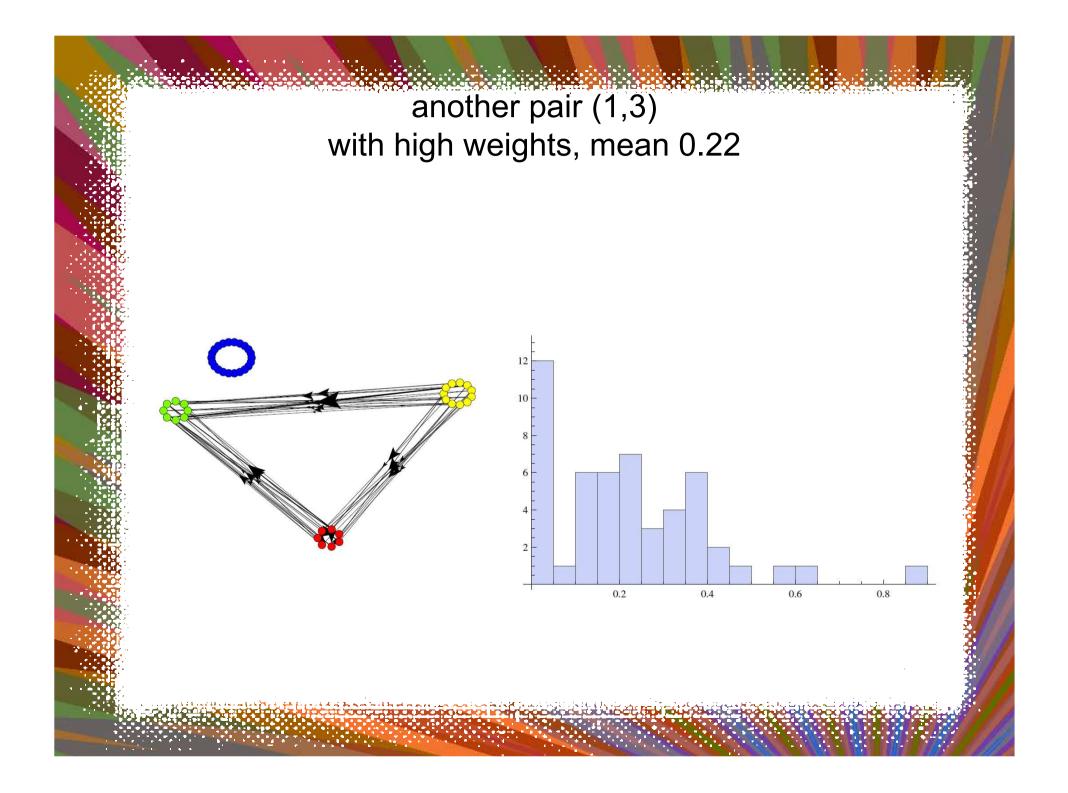
- start from random partitions
- A: calculate *P-matrix*
- for each node: calculate exp. log-likelihood considering that the node belongs to a given pair of in- and out-cluster; find max over all such cluster pairs; (*P* is fixed)
- redistribute nodes in clusters, replace node to cluster pair that gives the max of expected log-likelihood
- Go To A and iterate until the clusters saturate
- calculate AIC
 - continue FOR, take clustering corresponding to min AIC

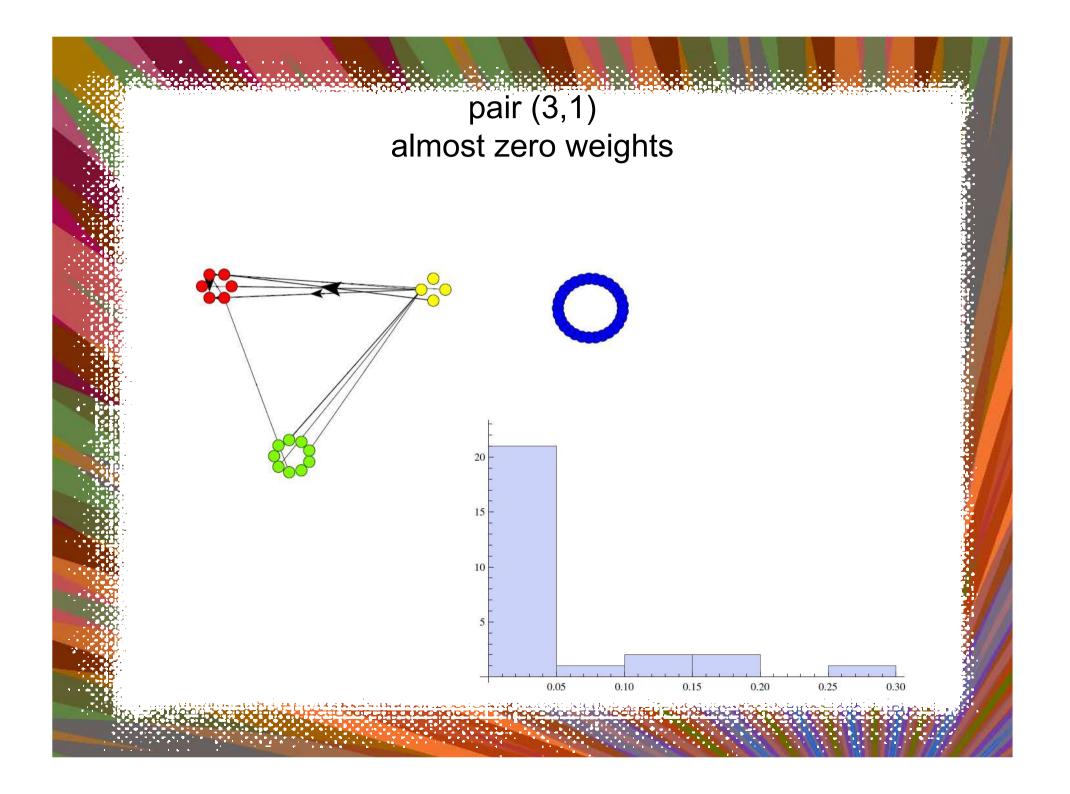


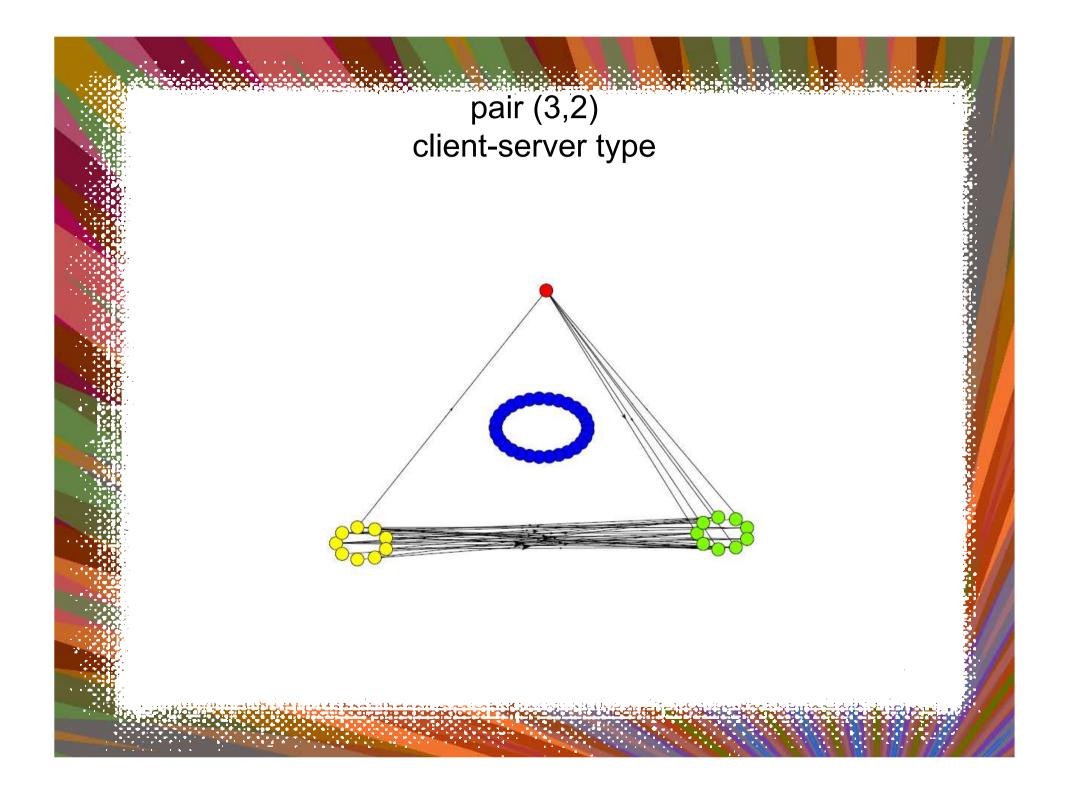


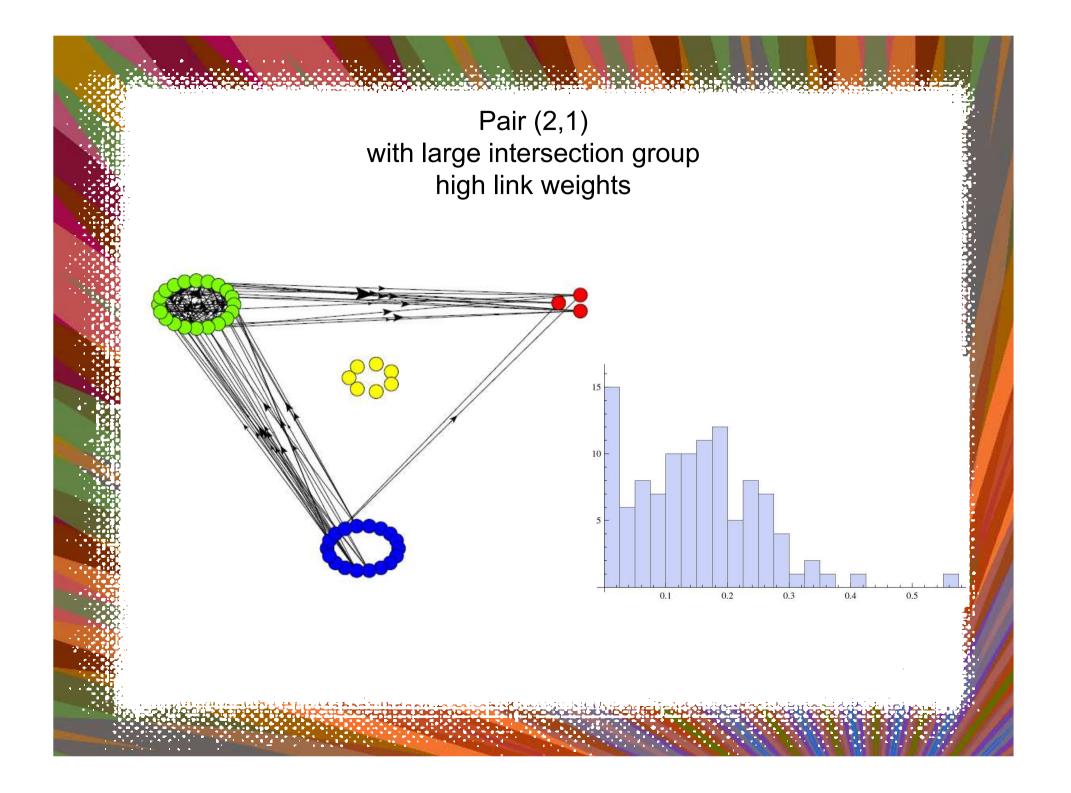


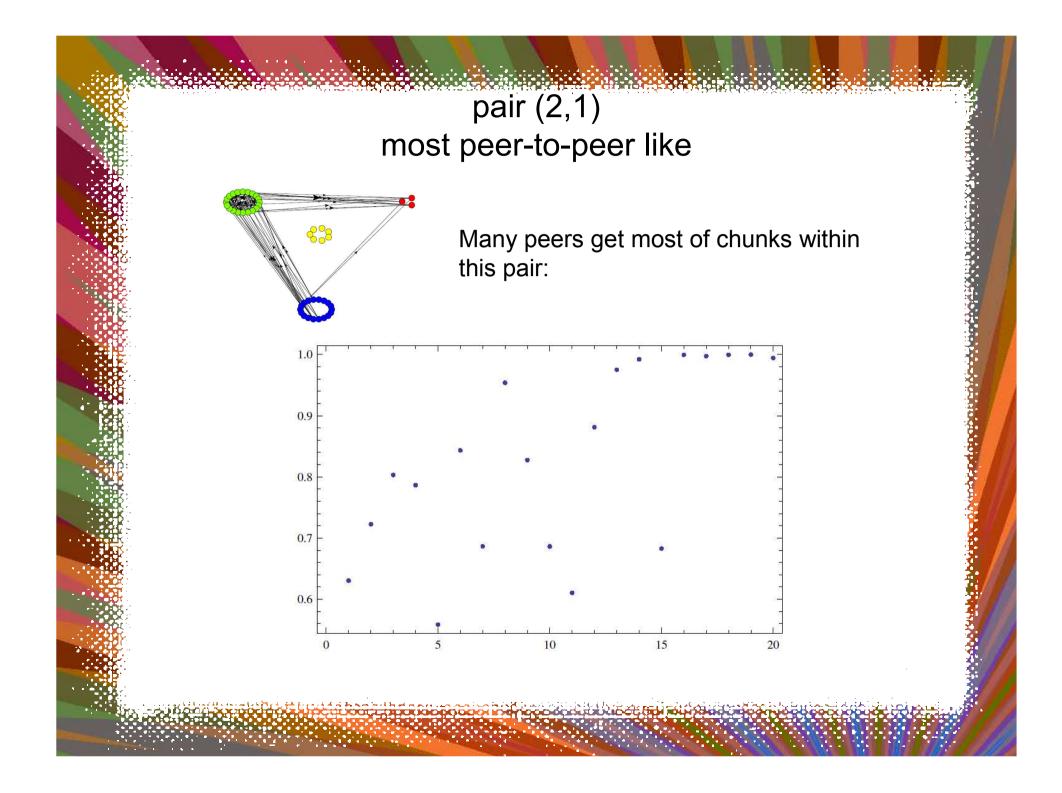


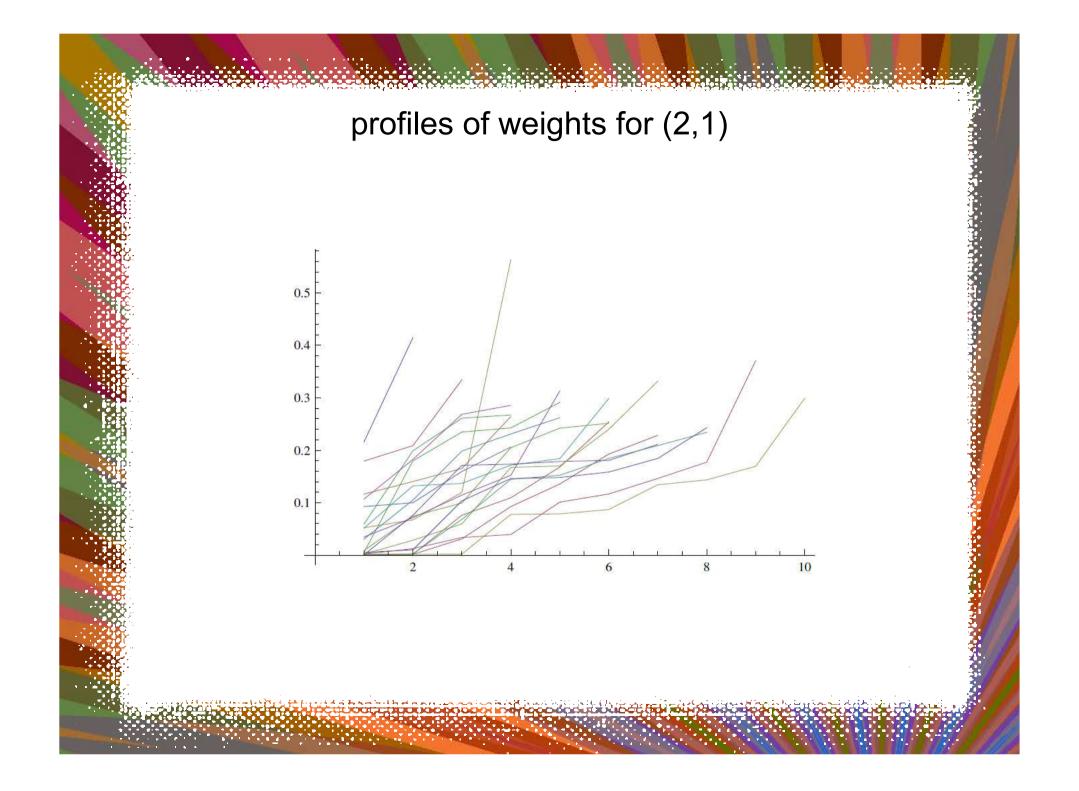












comments:

- more interesting results for larger networks?
- use in metabolic networks
- MC-algorithm would be a better choice
- seems to have some potential

