

Problems in Social Networks

PageRank and RandomWalks

Problem Definition Challenges Link Spam

Literature Review

Community Detection

Problem Definition

Taxonomy of Methods

Conclusion and Future Work

The Importance of Individuals and Groups in Social Networks

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Problem Definition Challenges

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Taxonomy of Methods

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- 4 Community Detection Problem Definition Taxonomy of Methods
- 5 Conclusion and Future Work



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- Social networks are becoming more popular each day!
- Need for studying in a scientific way
- Because of the size:
 - $\diamond~$ Solve for specific cases of input,
 - Working good practically,
 - Approximation algorithms or,
 - ◊ A mixture.





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- A Social Network: G = (V, E)
- Categories of problems:
 - ◊ Static vs. Dynamic
 - ◊ Content based vs. Structural

Table: Instances of Social Networks.

	Weighted	Unweighted
Directed	Email Network	Twitter
Undirected	DBLP	Facebook



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Conclusion and Future Work There are many problems in this area:

- Statistical Analysis:
 - ◊ On a large scale, how does a SN look like?
 - ◊ Extracting statistical features, such as:
 - · Degree distribution,
 - · Diameter,
 - · Clustering behavior,
 - · Behavior of connected components,
 - · Small-world phenomenon,
 - · Power-law degree distribution.



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Conclusion and Future Work • Centrality indices:

- ◊ How to rank nodes or edges?
- Many well-known indices:
 - · Degree,
 - · Closeness,
 - · Betweenness.
- ♦ Due to the size, *estimate* the value.



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Conclusion and Future Work • Centrality indices:

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 - · Degree,
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- ♦ Due to the size, *estimate* the value.

• PageRank and RandomWalks:

- ♦ Initiate a random surfer in the network,
- ◇ It will end up in most *important* nodes.



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• Community Detection:

- ♦ Members of a community have:
 - $\cdot\,$ Strong connections within the community,
 - $\cdot\,$ Loose connections to those outside.



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• Community Detection:

- ◊ Members of a community have:
 - $\cdot\,$ Strong connections within the community,
 - · Loose connections to those outside.
- Influence Maximization:
 - ♦ How does a message spread throughout the network?
 - ◊ 2 important research directions:
 - $\cdot\,$ Model the influence of individuals on each other,
 - $\cdot\,$ Select the best individuals for initiating the spread.



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- Link Prediction:
 - ♦ Which edges are likely to appear in the future?
 - · Friends suggestion,
 - · Product recommendation,
 - · expert hiring.



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Problems in Social Networks - cont.

- Link Prediction:
 - ♦ Which edges are likely to appear in the future?
 - · Friends suggestion,
 - · Product recommendation,
 - · expert hiring.
- Other problems:
 - ◊ Node classification
 - ◊ Expert discovery
 - ♦ Privacy issues
 - ◊ Visualizing
 - ◊ Data mining
 - $\diamond \cdots$



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Problems in Social Networks - cont.

- Link Prediction:
 - ♦ Which edges are likely to appear in the future?
 - · Friends suggestion,
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- Other problems:
 - ◊ Node classification
 - ◊ Expert discovery
 - ♦ Privacy issues
 - ◊ Visualizing
 - ◊ Data mining
 - $\diamond \cdots$
- We choose 2 of them: PageRank and Community Detection.



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• Motivation: Search Engines



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Concordia University - Wikipedia

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C-K officer found not liable after man dislocates shoulder - The ...

1 day ago - ... in Anthropology and majored in Communications at Concordia University. After finishing her Master of Journalism at Carleton University in ...



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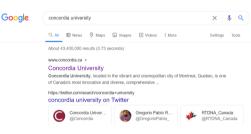
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Concordia University is a public comprehensive research university located in Montreal, Quebec, Canada. Founded in 1974 following the merger of Loyola ...

• IR + PR

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Problem Definition - cont.

- Intuition: The more incoming edges a node has, the more important it is.
 - $\diamond\,$ Justification: Good websites do not link many pages, while recieve many links.
- Page P_i with importance r_i has *n* outgoing edges; each edge gets $\frac{r_i}{n}$.
- Importance of page P_j is the sum of the votes [15]:

$$r_j = \sum_{i \to j} \frac{r_i}{d_i} \tag{1}$$

• Stochastic adjacency matrix *M* [15]:

$$M_{j,i} = egin{cases} rac{1}{d_i} & ext{if } i o j \ 0 & ext{otherwise} \end{cases}$$

• Assume that $\sum_{i} r_i = 1$, and $r = [1/N, 1/N, \cdots, 1/N]^T$

(2)



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Problem Definition - cont.

• Main equation [15]:

$$= M.r \tag{3}$$

r

• Power Iteration method [15]:

Algorithm 1 Power Iteration

input : Graph *G* with *N* nodes, ε **output**: PageRank vector *r* Initialize $r^{(0)} = [\frac{1}{N}, \frac{1}{N}, \dots, \frac{1}{N}]^T$ while $|r^{(t+1)} - r^{(t)}|_1 < \varepsilon$ do $| r^{(t+1)} = M \cdot r^{(t)}$ end

return r



Problem Definition - cont.

Example:

M =

r =

ln+	rod		10
iii u			

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a b y m	$\begin{bmatrix} 0\\ \frac{1}{2}\\ 0\\ \frac{1}{2} \end{bmatrix}$	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	m 1 0 0 0	(4))		y the second sec	b
_	a b y m	$\left[\begin{array}{c} 1/4 \\ 1/4 \\ 1/4 \\ 1/4 \\ 1/4 \end{array} \right]$	\rightarrow	1/2 1/8 1/8 1/4	\rightarrow	$\left[\begin{array}{c} 6/16\\ 1/4\\ 1/16\\ 5/16\end{array}\right]$	$\rightarrow \cdots \rightarrow$	$\left[\begin{array}{c} 0.42\\ 0.21\\ 0.11\\ 0.26\end{array}\right]$

(5)



Challenges - Dead Ends

 a_1 b 0 0

0

 $\frac{1}{21}$ 0

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a_1 M =b₁ V_1

 m_1 Solutions:

- \diamond Removing them,
- Teleporting. \diamond

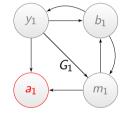
$$\begin{array}{cccc} y_1 & m_1 \\ \frac{1}{3} & \frac{1}{2} \\ \frac{1}{3} & \frac{1}{2} \\ 0 & 0 \\ \frac{1}{2} & 0 \end{array}$$

0

• Surfer gets stuck in this page \rightarrow Page rank will drain out.

(6)

• A page with no out going edge is a dead end.





Challenges - Spider Traps

an b_2

0

• A set of page with no edges outside the set.

1313013

• Surfer gets stuck in these pages \rightarrow They attract all the PageRank.

(7)

 m_2

 $\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$

Ō

0

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- a_2 $\begin{array}{c}
 0 \\
 \frac{1}{2} \\
 \frac{1}{2} \\
 \frac{1}{2}
 \end{array}$ 0 0 M =b₂ *Y*2 m_2
- Solutions:
 - \diamond Removing them.
 - Teleporting. \diamond

*Y*2 a_2

bo

 m_2

Gà





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Dealing with challenges: Teleporting

The random surfer has 2 options:

- With probability β follow an edge,
- With probability 1β jump to a random page.

$$r_j = \sum_{i \to j} \beta \frac{r_i}{d_i} + (1 - \beta) \frac{1}{N}$$
(8)

• Matrix *M* is modified:

$$A = \beta M + (1 - \beta) \left[\frac{1}{N} \right]_{N imes N}$$

- β is normally between 0.8 and 0.9.
- + Always teleport from a dead end!

(9)



Link Spam



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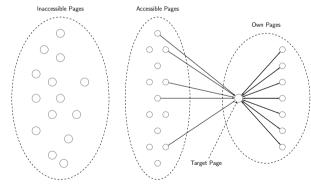
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- Not all individuals behave well in the WWW network.
- Designing a structure to increase the PR of a page, artificially.
- Architecture of a Spam Farm:





Link Spam - cont.

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- How to deal with Spam Farm?
 - ◊ Detect these structures,
 - ♦ Modify Page Rank:
 - Gyongyi et al. [10], Trust Rank: Teleport to trustworthy pages,
 - · Gyongyi et al. [9], Spam Mass: $rac{r-t}{r}pprox 1
 ightarrow$ SPAM!



Literature Review - Spam Farm

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- Wu et al. [22]: Pages in a Spam Farm are densely connected;
 - \diamond Seed set of bad pages: more than T_{IO} common in-out links,
 - \diamond Pages that link to more than T_{PP} bad pages: potential spammers,
 - Give 0 weight to links between bad pages,
 - ♦ Recalculate PR.



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Literature Review - Spam Farm - cont.

• Ghosh et al. [7], Collusionrank: Penalize the ones who are following bad users:

Algorithm 2 Collusionrank

input : Graph G, Set of known spammers S, β output: Collusionrank vector c

$$d(n) = egin{cases} rac{-1}{|S|} & ext{if } n \in S, \ 0 & ext{otherwise} \end{cases}$$

 $c \leftarrow d$ while c not converged do $\left|\begin{array}{c} \text{foreach node } v \text{ in } G \text{ do} \\ 1 \text{ foreach node } v \text{ in } G \text{ do} \\ 1 \text{ tmp } \leftarrow \sum_{n \in following(v)} \frac{c(n)}{[followers(n)]} \\ 1 \text{ c}(n) = \beta \times tmp + (1 - \beta) \times d(n) \\ 1 \text{ end} \\ \text{end} \\ \text{return } c$



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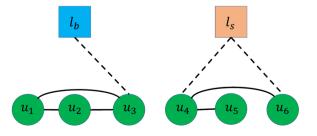
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Literature Review - Spam Farm - cont.

- Jia et al. [13]:
 - \diamond Two artificial nodes: real I_b , fake I_s ,
 - \diamond Add edge from every fake known node to I_s ,
 - \diamond Add edge from every real known node to I_b ,
 - ◊ For each unknown node, initiate a RandomWalk,
 - \diamond Badness score = the probability of reaching I_s sooner than I_b .





Literature Review - Applications of PageRank

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- In link prediction problem:
 - ♦ Tong et al. [21]:
 - \cdot Initiate a random walk with restart from v,
 - $\cdot\,$ Nodes with highest PR score will form an edge with v.
 - ♦ Backstrom and Leskovec [2]:
 - \cdot Initiate a random walk from v,
 - $\cdot\,$ Learn to visit the nodes that will have a potential edge.



Literature Review - Applications of PageRank - cont.

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- In Influence Maximization problem:
 - ♦ Java et al. [11]:
 - $\cdot\,$ Nodes with high PR are good seed sets
 - ◊ Bar-Yossef et al. [3], Reverse PageRank:
 - · Change the direction of edges
 - · Run PageRank
 - $\cdot\,$ Nodes with high PR are good seed sets



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- In graph G = (V, E), divide V into c subsets, C₁, C₂, ..., C_c in a way that:
 Members of each community have dense connections inside,
 - $\diamond~$ And loose connections to those outside
- In 2 phases:
 - Detecting communities with an algorithm,
 - ♦ *Evaluating* the appropriateness of communities.



Taxonomy of Methods

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- 4 Categories of methods:
 - ♦ Disjoint or non-overlapping communities,
 - Overlapping communities,
 - ♦ Hierarchical communities and,
 - Local communities.
- Evaluation Metrics



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Conclusion and Future Work • Girvan and Newman [8] $(O(VE^2) \text{ or } O(V^3))$:

Calculate edge betweenness for all edges,

- Remove the edge with highest betweenness,
- ◊ Recalculate the edge betweenness for all edges,

Taxonomy of Methods - Disjoint Communities

◊ Repeat step 2 and 3 until no edge remains.



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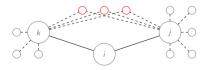
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Taxonomy of Methods - Overlapping Communities

- Ahn et al. [1], Link algorithm:
 - ♦ Similarity of two edges:

$$S(e_{ij}, e_{ik}) = \frac{|n_+(j) \cap n_+(k)|}{|n_+(j) \cup n_+(k)|}$$
(10)

Merge the edges with highest similarity into 1 community,
For e_{ij} and e_{ik}, if k and j belong to different communities, i is an overlapping node.





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Taxonomy of Methods - Hierarchical Communities

• Mann et al. [18]:

 $\diamond~$ Using the idea of sparsest cut:

$$(S, T) - \text{cut density} = \frac{|(S, T)|}{(|S| \cdot |T|)}$$
(11)

- The cut with minimum density is suitable for partitioning the graph,
 Find minimum density cut and repeat it for bigger sub-graph.
 Finding sparsest cut is NP-Hard:
 - · Closely related to maximum concurrent flow,
 - $\cdot\,$ Could be solved efficiently with linear programming.



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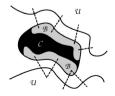
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Taxonomy of Methods - Local Communities

- Clauset [5]:
 - $\diamond~$ Divide V into three sets: C, B, and U.
 - ♦ Local modularity:

- $R = \frac{I}{T} \tag{12}$
- *· I*: the number of those edges with neither end point in U *· T*: the number of edges with one or more end points in B
 ◊ Start with C = v₀ and discover k vertices that are in the same community as v₀,
- $\diamond\,$ In each step, add the one with highest difference in terms of R.





Taxonomy of Methods - Evaluation Metrics

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- $M = \frac{E_{in}}{E_{out}}$ $\diamond E_{in}$: Number of edges within the community, $\diamond E_{out}$: Number of crossing edges.
- Chen et al. [4]:

 $\diamond L_{in} = \frac{E_{in}}{|\mathcal{C}|}$ $\diamond L_{out} = \frac{E_{out}}{|\mathcal{B}|}$

• Luo et al. [16]:

 $L = \frac{L_{in}}{L_{out}}$

(13)

(14)



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- $\bullet\,$ 2 important problems in social networks were considered in this study:
 - ◊ PageRank and RandomWalks
 - Community Detection
- We're interested in the following directions for future work:
 - ◊ Connection between PageRank and Broadcasting
 - ◊ Local community detection
 - With PageRank
 - · Evaluation Metric



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- 1. Connection between PageRank and Broadcasting:
 - Broadcasting:
 - A message is transmitted throughout the network,
 - ♦ Starting from a single *originator*,
 - $\diamond\,$ All informed vertices may initiate a call in each time step,

$$\diamond \ b(G) = \max_{v \in V} \{b(v, G)\}$$

$$\diamond \ \lceil \log_2 n \rceil \le b(G) \le n-1$$

- ♦ Research directions:
 - mbgs
 - · Find broadcast time of any graph: NP-Complete
 - \checkmark Find Center nodes
 - ✓ Find Worst originators
- Experiment: Run Power Iteration method with $\beta=$ 0.85 and $\varepsilon=$ 0.00005



Conclusion and Future Work - cont.

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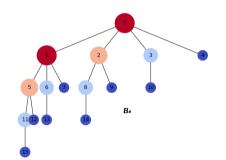
Community Detection

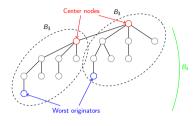
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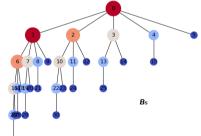
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Conclusion and Future Work **1**. Connection between PageRank and Broadcasting:

• Binomial Tree:









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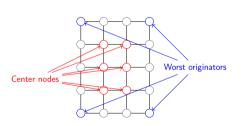
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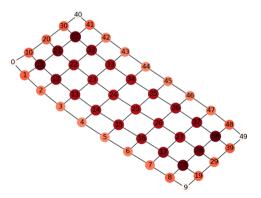
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- 1. Connection between PageRank and Broadcasting:
 - Grids:







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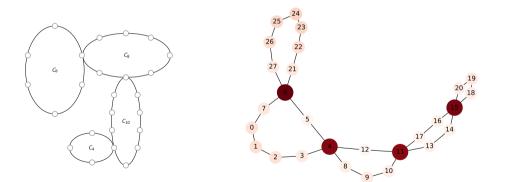
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- 1. Connection between PageRank and Broadcasting:
 - Necklace graph:





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2.1. Local Community Detection - Method:

- Source vertex v_0 ,
- Discover k vertices that are in the same local community as v_0
- Possible algorithm:
 - \diamond Start a random walk from v_0 (with TrustRank modification)
 - $\diamond \ C = v_0, \ \mathsf{Trusted} = v_0$
 - ♦ Until |C| = k:
 - · Add v_i with the highest page rank to the community: $C = C \cup v_i$ · Trusted = Trusted $\cup v_i$
- Why it works?
 - \diamond The vertices with highest PageRank have good similarity with v_0
 - $\diamond~\ln$ 1 β fraction of times, teleport to the discovered community



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2.2. Local Community Detection - Evaluation Metric:

- Idea of using Geodesic Distance (GD),
- Length of the shortest path between two nodes,
- Possible metric:
 - Sum of GD for all vertices within a local community,
 - ◊ The smaller the sum, the better the community.
- Why it works?
 - $\diamond\,$ More edges in a community \rightarrow length of shortest path will decrease,
 - $\diamond~$ More edges in a community \rightarrow the community is more dense.
- Also, design a heuristic algorithm that uses this metric:
 - Add the vertices with highest difference in terms of the metric to local community.



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Important References I

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Thanks a bunch!



Why power iteration method works?

- Recall that when $Ax = \lambda x$, x is the eigenvector and λ is the eigenvalue.
- In equation r = M.r: r is the principal eigenvector of M with eigenvalue of 1 (largest eigenvalue of M)
 - ♦ Because *M* is column stochastic.
- Eventually, we want to find the dominant eigenvector of M.

•
$$r^{(1)} = M.r^{(0)}$$

•
$$r^{(2)} = M.r^{(1)} = M(M.r^{(0)}) = M^2.r^{(0)}$$

• • • •

•
$$r^{(k)} = M^k . r^{(0)}$$

Claim

The sequence of $M.r^{(0)}, M^2.r^{(0)}, \cdots, M^k.r^{(0)}$ approaches the dominant eigenvector of M.



Claim

The sequence of $M.r^{(0)}, M^2.r^{(0)}, \cdots, M^k.r^{(0)}$ approaches the dominant eigenvector of M.

Proof.

Suppose *M* has *n* eigenvectors x_1, x_2, \dots, x_n with corresponding eigenvalues $\lambda_1, \lambda_2, \cdots, \lambda_n$ in a way that: $\lambda_1 > \lambda_2 > \cdots > \lambda_n$ [15]. We can write $r^{(0)} = c_1 x_1 + c_2 x_2 + \cdots + c_n x_n$, now: $M.r^{(0)} = M(c_1x_1 + c_2x_2 + \cdots + c_nx_n) \rightarrow$ $M.r^{(0)} = c_1(Mx_1) + c_2(Mx_2) + \cdots + c_n(Mx_n) \xrightarrow{M.x_i = \lambda_i.x_i}$ $M.r^{(0)} = c_1(\lambda_1 x_1) + c_2(\lambda_2 x_2) + \dots + c_n(\lambda_n x_n)$ repeat multiplication on both sides $M^k \cdot r^{(0)} = c_1(\lambda_1^k x_1) + c_2(\lambda_2^k x_2) + \cdots + c_n(\lambda_n^k x_n) \rightarrow$ $M^{k}.r^{(0)} = \lambda_{1}^{k}(c_{1}x_{1} + c_{2}(\frac{\lambda_{2}}{\lambda_{1}})^{k}x_{2} + \dots + c_{n}(\frac{\lambda_{n}}{\lambda_{1}})^{k}x_{n}) \xrightarrow{\lambda_{1} > \lambda_{2} > \dots > \lambda_{n}}{\lim_{k \to \infty} (\frac{\lambda_{i}}{\lambda_{1}})^{k} = 0}$ $M^{k} r^{(0)} = c_{1}(\lambda_{1}^{k} x_{1})$



A note on β :

- We want to simulate the user's behavior. With $\beta = 0.85$, we are giving a chance of entering a new URL in $\frac{1}{6}$ of times.
- As β increases, the PageRank becomes more and more sensitive to small changes in ${\it M}$ matrix.
- The smaller the $\beta,$ the faster the convergence, but the structure of the graph is not used so much!
- A trade off!
- Langville [14] showed that a rough estimate of the number of iterations needed to converge to a tolerance level ε is log₁₀ε/log₁₀β. So:
 - $\diamond~$ for $\beta=0.85~{\rm and}~\varepsilon=10^{-6}$ it takes roughly $\frac{-6}{\log_{10}0.85}\approx 85$ (A very common situation),
 - $\diamond~{\rm Or}$ for $\beta=0.85$ and $\varepsilon=10^{-8}$ it takes almost 114 iterations,
 - $\diamond\,$ While for $\beta=$ 0.99 and $\varepsilon=10^{-8},$ it takes 1833 iterations!



Tong et al. [21]:

- Proposing node-to-node proximity measure Prox based on RandomWalks,
- escape probability $e_{p_{i,j}}$: the probability that the random walk which starts from node *i* will visit node *j* before it returns to node *i*,
- generalized voltage $v_k(i, j)$: the probability that a random walk that starts from node k will visit node j before node i,
- $p_{i,k}$: probability of a direct transition from node *i* to node *j*,
- Prox measure: $Prox(i,j) \stackrel{\Delta}{=} ep_{i,j} = \sum_{k=1}^{n} p_{i,k} \cdot v_k(i,j)$
- Predict a link between i and j iff Prox(i, j) + Prox(j, i) > th,
 th is a given threshold.



Backstrom and Lescovec [2]:

- Combining two general approaches for link prediction:
 - ♦ Using graph structural information (with RandomWalk),
 - ♦ Using node and edge attributes (with ML).
- Assign each edge a RandomWalk transition probability (learn strength function for each edge),
- Initiate a RandomWalk with restart from source node s,
- Nodes with highest PageRank are the ones that *s* will form an edge with.
- Excellent results on co-authorship network \rightarrow suggest who to write a paper with!



Jeh et al [12]:

- Intuition: two objects are similar if they are related to similar objects,
- SimRank: if a = b then s(a, b) = 1. Otherwise:

$$s(a,b) = \frac{C}{|\mathsf{InDegree}(a)|.|\mathsf{InDegree}(b)|} \sum_{v \in \mathsf{InNeighbor}(a)} \sum_{u \in \mathsf{InNeighbor}(b)} s(u,v)$$
(15)

• They show that SimRank score s(a, b) measures how soon two random surfers are expected to meet at the same node if they started at nodes a and b and randomly walked the graph backwards.

$$m(a,b) = \sum_{t:(a,b) \rightsquigarrow (x,x)} P[t] I(t)$$
(16)

◊ t =< w₁, · · · , w_k > is a tour on G² graph with V² as the nodes and
< (a, b), (c, d) >∈ E(G²) means (a, c) ∈ E(G) and (b, d) ∈ E(G)
◊ P[t] is the probability of traveling P[t] = ∏^{k-1}_{i=1} 1 1 |OutDegree(w_i)|
◊ I(t) is the path length and is k - 1.



Bar-Yossef et al [3]:

- They show that local PageRank approximation is not efficient in graphs with high in-degree nodes (such as SNs).
- However, ReversePageRank can be approximated locally in the graph obtained by reversing the direction of all edges.
- They also argue that ReversePageRank is useful for selecting influential nodes in IM problem, and many other applications (such as crawler's seed set selection).



Fast Newman [19]:

- O(V(V + E)) or $O(V^2)$:
 - ♦ Modularity $Q = \sum_i (e_{ii} a_i^2)$:
 - $\cdot e_{ii}$ fraction of edges within the group i
 - $\cdot ~ e_{ij}$ one-half of the fraction of edges that connect a vertex from group i to j
 - · $a_i = \sum_j e_{ij}$ the fraction of all ends of edges that are attached to vertices in the group *i*
 - $\cdot a_i^2$ the value that it would take if edges were placed at random.
 - \diamond Two nodes are joined with biggest difference in Q.



Fast Newman [19]:

- Modularity $Q = \sum_i (e_{ii} a_i^2)$
- Why not optimize Q over all possible divisions to find the best one?
- It is very costly:
 - $\diamond\,$ Number of ways to divide n objects into g non-empty groups is the Stirling number of the second kind $S_n^{(g)},$
 - ♦ The sum is not known in closed form,
 - ♦ But we know that $S_n^{(1)} + S_n^{(2)} = 2^{n-1}$; thus, it is at least exponentially!
- Instead, use a greedy approximation optimization:
 - Each vertex is the sole member of a community,
 - Repeatedly join communities together,
 - $\diamond\,$ Choosing the join that results in the greatest increase (or smallest decrease) in Q.
 - ◊ Generates a dendrogram!



- Derènyi et al. [6], Clique Percolation Method (CPM):
 - ♦ Find all *k-cliques*,
 - \diamond Build hyper graph (two k-cliques are connected if they share k-1 vertices),
 - ◊ Connected parts are communities.



- Raghavan et al. [20], Label Propagation Algorithm (LPA):
 - \diamond The community of x is the same as majority of its neighbors,
 - ◊ Initiate labels,
 - ◊ Propagate until convergence.
 - $\diamond\,$ BUT it is likely to find many communities for the same graph.



Ahn et al. [1]:

- Where to cut the generated dendrogram?
- They proposed partition density D as follows:
 - ♦ For a graph with *M* edges and *N* nodes, $P = \{P_1, \dots, P_C\}$ is a partition of the links into *C* subsets.
 - \diamond The number of links in subset P_c is m_c ,
 - \diamond The number of adjacent nodes in subset P_c is $n_c = |\cup_{e_{i,j} \in P_c} \{i,j\}|$

◊ Density of community *c* is:

$$D_c = \frac{m_c - (n_c - 1)}{\frac{n_c(n_c - 1)}{2} - (n_c - 1)}$$
(17)

 \diamond The partition density *D* is the average of D_c weighted by the fraction of present links:

$$D = \frac{2}{M} \sum_{c} m_{c} \frac{m_{c} - (n_{c} - 1)}{(n_{c} - 2)(n_{c} - 1)}$$
(18)

• Cut dendrogram where maximum D happens.



Mann et al. [18]:

- Using the idea of the sparsest cut (highly related to MCFP).
- Ford-Fulkerson method:
 - While there is an augmenting path in the graph;
 - Augment the value of flow along the path,
 - Reduce the capacities along the path,
- If the augmenting path is found via BFS, the algorithm is called Edmonds-Karp.
- A good heuristic for finding the sparsest cut.



• Lue et. al [17]:

$$LQ = \frac{e_c}{S} - (\frac{d_c}{2S})^2$$
 (19)

- \diamond e_c : number of edges within the detected local community,
- $\diamond d_c$: summation of degrees of all nodes belonging to that local community,
- ◊ S: the number of edges with one or two endpoints in the local community.
- Zhen-Qing et al. [23]:

$$Q^{d} = \sum_{r=1}^{s} \left(\frac{L_{r}}{D_{r}} - \frac{\tilde{L_{r}}}{\tilde{D_{r}}}\right)$$
(20)

- \diamond L_r : Number of edges inside the community
- ◊ D_r: Average minimal path for all pairs of nodes within a given community,
- \diamond $\tilde{L_r}$ and $\tilde{D_r}$: Expected values for the graph that is generated randomly.



Zhen-Qing et al. [23]:

- We know what are L_r and D_r in this equation $Q^d = \sum_{r=1}^{s} \left(\frac{L_r}{D_r} \frac{\tilde{L}_r}{\tilde{D}_r}\right)$
- But how to calculate $\tilde{L_r}$ and $\tilde{D_r}$?

 $\diamond \ ilde{L}_r = d_r^2/4L$ where:

- \cdot d_r is the sum degree of nodes in community r,
- $\cdot \ L$ is the total number of edges for the underlying network.



 $\diamond \tilde{D}_r$:

- $\cdot K = (k_1, k_2, \cdots, k_n)$ is the degree distribution of the original graph,
- · L_{ij} The path length of certain vertices (i, j) can be approximately calculated based on

$$L_{ij}(k_i, k_j) = \frac{-\ln k_i k_j + \ln(\langle k^2 \rangle - \langle k \rangle) + \ln N - \gamma}{\ln(\langle k^2 \rangle / \langle k \rangle - 1)} + \frac{1}{2} \quad (21)$$

- $\cdot \ <.>$ indicates the average operation over the entire degree sequence,
- \cdot N is the number of vertices,
- $\cdot \ \gamma$ is a constant value of 0.5772

$$\tilde{D}_{r} = \frac{2}{n_{r}(n_{r}-1)} \sum_{i,j \in r, i < j} L_{ij}(k_{i},k_{j})$$
(22)

 \cdot *n_r* is the number of nodes in community *r*.