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# 15-826: Multimedia Databases and Data Mining

## Graph mining - Part 2&3


*Christos Faloutsos*

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## Thanks

- Deepayan Chakrabarti (CMU)
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- Michalis Faloutsos (UCR)
- George Siganos (UCR)



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## PART 2: PageRank, HITS, and eigenvalues

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## Outline

- Part 1: Topology, 'laws' and generators
- ➡ Part 2: PageRank, HITS and eigenvalues
- Part 3: Influence, communities

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## Part 2: PageRank, HITS and eigenvalues

- How important is a node?
- Who is the best customer to advertise to?

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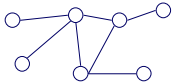
## Outline

- Part 1: Topology, 'laws' and generators
- ➡ Part 2: PageRank, HITS and eigenvalues
  - Eigenvalues and PageRank
  - SVD and HITS
- Part 3: influence, virus prop., communities

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## Motivating problem

Given a graph, find its most interesting/central node



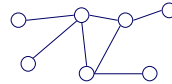
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## Motivating problem

Given a graph, find its most interesting/central node



A node is important, if it is connected with important nodes (recursive, but OK!)

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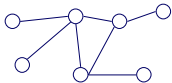
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## Motivating problem – pageRank solution

Given a graph, find its most interesting/central node

Proposed solution: Random walk; spot most 'popular' node (-> steady state prob. (ssp))



A node has high **ssp**, if it is connected with **high ssp** nodes (recursive, but OK!)

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## Notational conventions

- bold capitals -> matrix (eg. **A**, **U**, **A**, **V**)
- bold lower-case -> column vector (eg., **x**, **v**<sub>1</sub>, **u**<sub>3</sub>)
- regular lower-case -> scalars (eg.,  $\lambda_1$ ,  $\lambda_r$ )

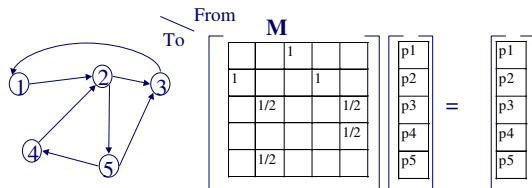
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## (Simplified) PageRank algorithm

- Let **A** be the transition matrix (= adjacency matrix); let  $M = A^T$  and column-normalized - then



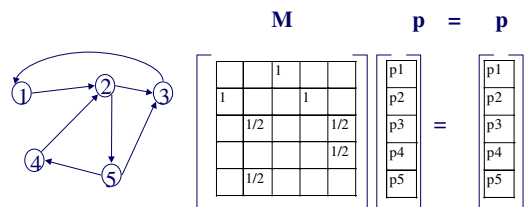
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## (Simplified) PageRank algorithm

- $M p = p$



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## (Simplified) PageRank algorithm

- $\mathbf{M} \mathbf{p} = \mathbf{1} * \mathbf{p}$
- thus,  $\mathbf{p}$  is the eigenvector that corresponds to the highest eigenvalue ( $=1$ , since the matrix is column-normalized)

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## (Simplified) PageRank algorithm

- In short: imagine a particle randomly moving along the edges
- compute its steady-state probabilities (ssp)

Full version of algo: with occasional random jumps – see later

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## Formal definition

If  $\mathbf{M}$  is a  $(n \times n)$  square matrix  
 $(\lambda, \mathbf{x})$  is an **eigenvalue/eigenvector** pair of  $\mathbf{M}$  if

$$\mathbf{M} \mathbf{x} = \lambda \mathbf{x}$$

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## (Published) PageRank

- Do a random walk, but
- with probability  $c$ , fly-out to a random node
- Then, the ssp vector  $\mathbf{v}$  obeys:

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## (Published) PageRank

$$\vec{\mathbf{v}} = (1-c) * \mathbf{M} \times \vec{\mathbf{v}} + c/n * \vec{\mathbf{1}}$$

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## (Published) PageRank

$$\vec{\mathbf{v}} = (1-c) * \mathbf{M} \times \vec{\mathbf{v}} + c/n * \vec{\mathbf{1}}$$

$\vec{\mathbf{v}}$  is an  $n \times 1$  ssp vector  
 $\mathbf{M}$  is a column-normalized matrix to-from adjacency matrix  
 $c$  is the fly-out probability  
 $n$  is the number of nodes  
 $\vec{\mathbf{1}}$  is a vector full of 'ones'

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## Personalized PageRank

- [Haveliwala+]

$$\vec{v}_i = (1-c) * \mathbf{M} \times \vec{v}_i + c * \vec{e}_i$$

ssp, when we restart from node 'i'

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## Personalized PageRank

- [Haveliwala+]

$$\vec{v}_i = (1-c) * \mathbf{M} \times \vec{v}_i + c * \vec{e}_i$$

ssp, when we restart from node 'i'

i-th row →

0
...
1
0
...

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## Personalized PageRank

- [Haveliwala+]

$$\vec{v}_i = (1-c) * \mathbf{M} \times \vec{v}_i + c * \vec{e}_i \quad \text{new}$$

$$\vec{v} = (1-c) * \mathbf{M} \times \vec{v} + c / n * \vec{1} \quad \text{original}$$

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## Personalized PageRank

- [Haveliwala+]

$$\vec{v}_i = (1-c) * \mathbf{M} \times \vec{v}_i + c * \vec{e}_i$$

- then  $s_{i,j}$  = prob( a random walker with restarts from node  $i$ , will find itself at node  $j$ )

$$[s_{i,j}] = \mathbf{S} = c * [\mathbf{I} - (1-c) * \mathbf{M}]^{-1}$$

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## Our wish list:

- ✓ How important is a node?
- ✓ Who is the best customer to advertise to?

ssp values answer these questions

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## Outline

Part 1: Topology, 'laws' and generators

Part 2: PageRank, HITS and eigenvalues

- Eigenvalues and PageRank
- ➡ SVD and HITS

Part 3: influence, virus prop., communities

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## Kleinberg's algorithm ('HITS')

- Problem dfn: given the web and a query
- find the most 'authoritative' web pages for this query

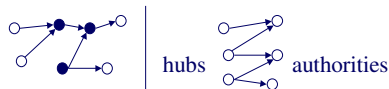
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## Kleinberg's algorithm

- give high score (= 'authorities') to nodes that many important nodes point to
- give high importance score ('hubs') to nodes that point to good 'authorities'



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## Kleinberg's algorithm

### Observations

- recursive definition!
- each node (say, ' $i$ '-th node) has both an authoritativeness score  $a_i$  and a hubness score  $h_i$

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## Kleinberg's algorithm

Let  $\mathbf{A}$  be the adjacency matrix:

the  $(i,j)$  entry is 1 if the edge from  $i$  to  $j$  exists

Let  $\mathbf{h}$  and  $\mathbf{a}$  be  $[n \times 1]$  vectors with the 'hubness' and 'authoritativeness' scores.

Then:

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## Kleinberg's algorithm

In conclusion, we want vectors  $\mathbf{h}$  and  $\mathbf{a}$  such that:

$$\mathbf{h} = \mathbf{A} \mathbf{a}$$

$$\mathbf{a} = \mathbf{A}^T \mathbf{h}$$

That is:

$$\mathbf{a} = \mathbf{A}^T \mathbf{A} \mathbf{a}$$

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## Kleinberg's algorithm

$\mathbf{a}$  is a right-singular vector of the adjacency matrix  $\mathbf{A}$  (by dfn!)

$\Rightarrow$  eigenvector of  $\mathbf{A}^T \mathbf{A}$

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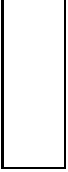



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## SVD & HITS

•  $A = U \Lambda V^T$  - example:

A	U	Lambda	V <sup>T</sup>
Nxn	Nxr	rxr	rxn
			

=

v1: author. scores

u1: hubness scores

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## Conclusions

eigenvalues/eigenvectors: vital for

- PageRank,
- (virus propagation - coming up next!)
- (graph partitioning - not mentioned here)

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## Conclusions, cont'd

SVD

- closely related: HITS/Kleinberg
- (and also LSI, KLT, PCA, Least squares, ...)

Both are **extremely useful, well understood** tools for graphs / matrices.

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## PART 3: Influence, virus propagation, communities

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## Problem definition

- Q1: How does a virus spread across an arbitrary network?
- Q2: will it create an epidemic?

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## Framework

- Susceptible-Infected-Susceptible (SIS) model
  - Cured nodes immediately become susceptible

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## The model

- (virus) Birth rate  $\beta$ : probability that an infected neighbor attacks
- (virus) Death rate  $\delta$ : probability that an infected node heals

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## The model

- Virus 'strength'  $s = \beta/\delta$

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## Other models:

- SIR: Susceptible - infected & infectious - recovered/removed
  - eg., mumps, chickenpox; black plague

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## Other models:

- and many more:
- SEIR: Susceptible; Exposed (= infected, but not infectious yet); I; R
- variations:
  - M: passively immune, like infants
  - with births/newcomers
  - ...

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## Epidemic threshold $\tau$

of a graph, defined as the value of  $\tau$ , such that  
if strength  $s = \beta/\delta < \tau$   
an epidemic can not happen

Thus,

- given a graph
- compute its epidemic threshold

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## Epidemic threshold $\tau$

What should  $\tau$  depend on?

- avg. degree? and/or highest degree?
- and/or variance of degree?
- and/or third moment of degree?



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## Epidemic threshold

- [Theorem] We have no epidemic, if

$$\beta/\delta < \tau = 1/\lambda_{1,A}$$

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## Epidemic threshold

- [Theorem] We have no epidemic, if recovery prob.

epidemic threshold

$$\beta/\delta < \tau = 1/\lambda_{1,A}$$

attack prob. largest eigenvalue of adj. matrix  $A$

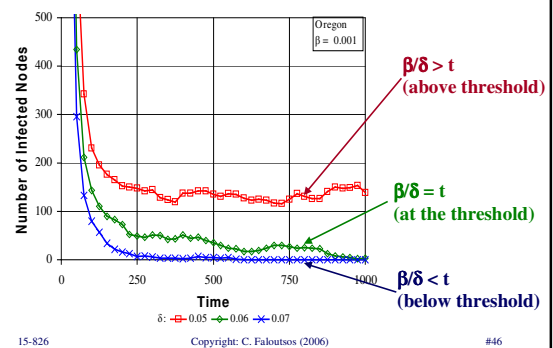
Proof: [Wang+03]

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## Experiments (Oregon)



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## Our wish list:

- Who is the best person/computer to immunize against a virus?

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#47

## Our wish list:

- ✓ Who is the best person/computer to immunize against a virus? Highest diff in  $\lambda_1$

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## Graph clustering & mining

- Q1: which edges/nodes are 'abnormal'?
- Q2: split a graph in  $k$  'natural' communities  
- but how to determine  $k$ ?

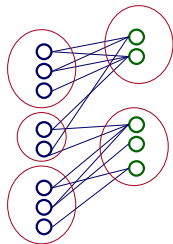
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## Graph partitioning

- Documents x terms
- Customers x products
- Users x web-sites



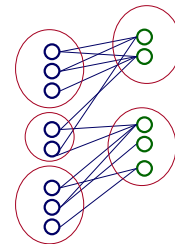
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## Graph partitioning

- Documents x terms
- Customers x products
- Users x web-sites



- Q: HOW MANY PIECES?

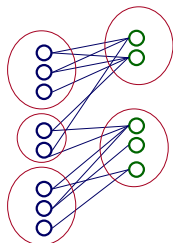
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## Graph partitioning

- Documents x terms
- Customers x products
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- Q: HOW MANY PIECES?

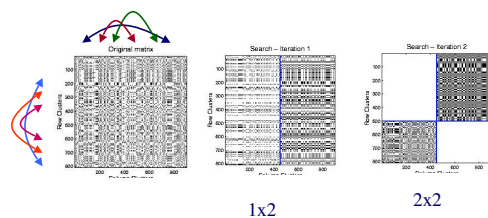
- A: MDL/ compression

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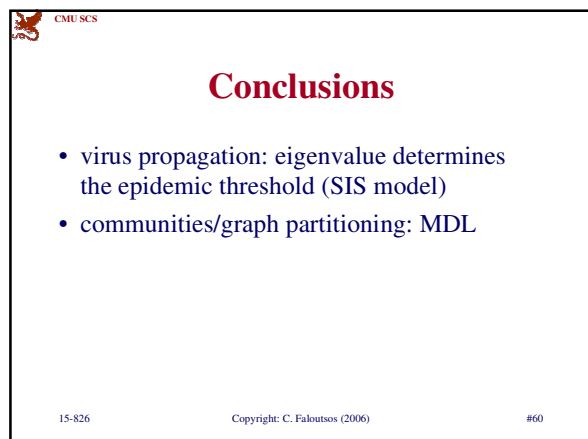
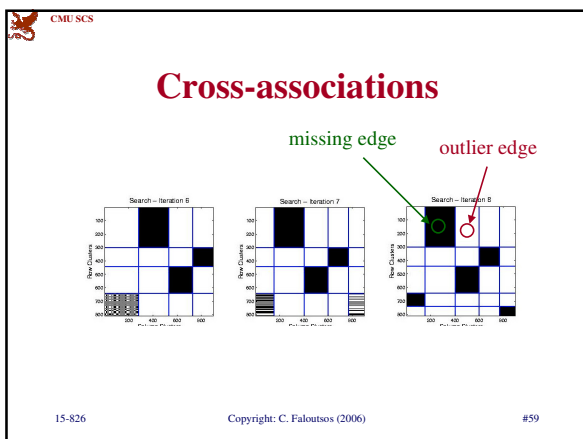
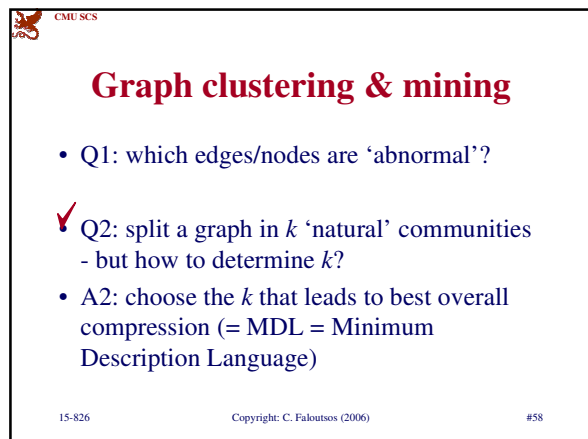
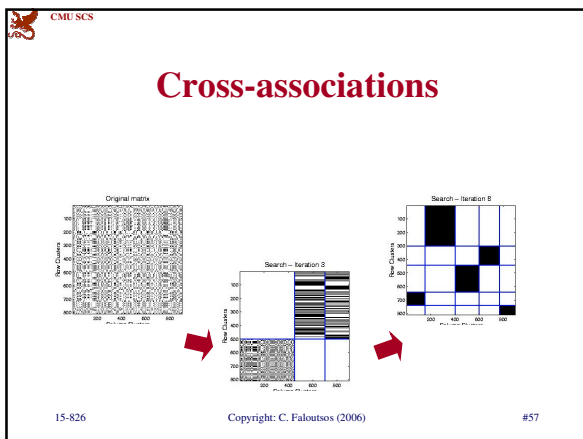
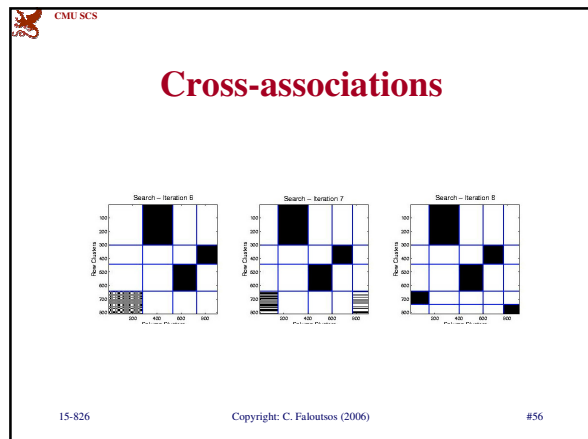
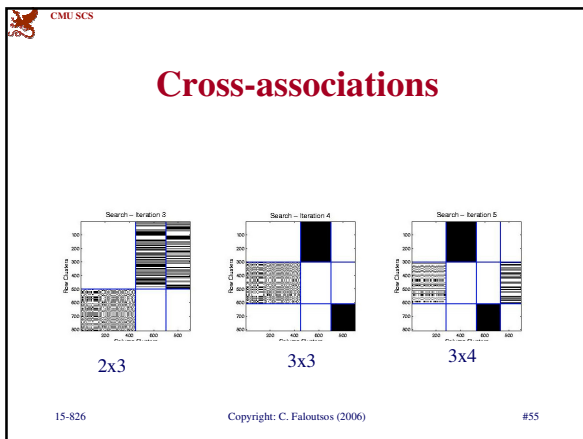
## Cross-associations



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## Resources: Software and urls

- SVD packages: in **many** systems (matlab, mathematica, LINPACK, LAPACK)
- stand-alone, free code: SVDPACK from Michael Berry  
<http://www.cs.utk.edu/~berry/projects.html>

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## Books

- Faloutsos, C. (1996). *Searching Multimedia Databases by Content*, Kluwer Academic Inc.
- Jolliffe, I. T. (1986). *Principal Component Analysis*, Springer Verlag.

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## Books

- [Press+92] William H. Press, Saul A. Teukolsky, William T. Vetterling and Brian P. Flannery: *Numerical Recipes in C*, Cambridge University Press, 1992, 2nd Edition. (Great description, intuition and code for SVD)

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- Berry, Michael: <http://www.cs.utk.edu/~lsi/>
- [Brin+98] Brin, S. and L. Page (1998). *Anatomy of a Large-Scale Hypertextual Web Search Engine*. 7th Intl World Wide Web Conf.
- [Chakrabarti'04] D. Chakrabarti, *AutoPart: Parameter-Free Graph Partitioning and Outlier Detection*, PKDD 2004 (pages 112-124), Pisa, Italy

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- [Chakrabarti+,04a] D. Chakrabarti, S. Papadimitriou, D. Modha and C. Faloutsos, *Fully Automatic Cross-Associations*, KDD 2004 (pp. 79-88), Washington, USA
- [Haveliwala02] Taher H. Haveliwala, *Topic-Sensitive PageRank* World Wide Web Conference, 2002

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- Fukunaga, K. (1990). *Introduction to Statistical Pattern Recognition*, Academic Press.
- Kleinberg, J. (1998). *Authoritative sources in a hyperlinked environment*. Proc. 9th ACM-SIAM Symposium on Discrete Algorithms.

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- [Wang+03] Yang Wang, Deepayan Chakrabarti, Chenxi Wang and Christos Faloutsos: *Epidemic Spreading in Real Networks: an Eigenvalue Viewpoint*, SRDS 2003, Florence, Italy.

## Discussion

A lot of recent interest - topics we didn't cover:

- Relational learning, e.g., [David Jensen; Daphne Koller; Saso Dzeroski]
- Frequent sub-graphs, e.g., [Jiawei Han, Jian Pei; George Karypis, Vipin Kumar; Mohammed Zaki]

## Discussion cont'd

- Graph partitioning, e.g., [METIS (Karypis)]
- Social networks, e.g., [Kathleen Carley; Wasserman+Faust]
- Web mining, e.g., [Soumen Chakrabarti]

## Overall conclusions

- Surprising patterns in graphs
- Powerful tools exist:
  - Self-similarity, fractals, Kronecker
  - SVD, eigenvalues
  - MDL for partitioning