

Propagation and Immunization on Networks: Theory and Tools

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Guest Lecture

15-826 – Multimedia Databases and Data Mining



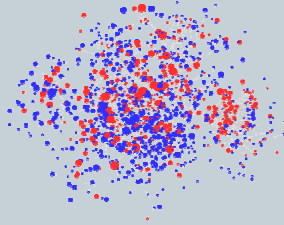
15 September 2011

Motivation

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One liner: "Contagions spreading over a network"

- Disease Propagation
- Blog Cascades
- Viral Marketing
- Influence Propagation



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Reading Material

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- **Primary:** Chapter 21, Networks, Crowds, and Markets: Reasoning about a Highly Connected World.

by David Easley and Jon Kleinberg. Cambridge University Press, 2010.

<http://www.cs.cornell.edu/home/kleinber/networks-book/networks-book-ch21.pdf>

- **Additional:** The Mathematics of Infectious Diseases

by H. W. Hethcote. SIAM Review Volume 42, Issue 4, 2000.

<http://www.maths.usyd.edu.au/u/marym/populations/hethcote.pdf>

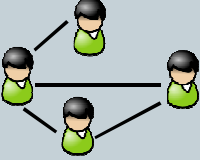
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Underlying phenomenon


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- Contagions over a network

Human contact-networks



Biological Viruses
e.g. H1N1




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Contagions over networks


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- examples...

Computer Networks



Computer Viruses
e.g. trojans



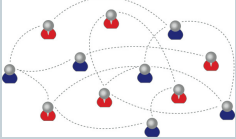
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Contagions over a network


6

- examples...

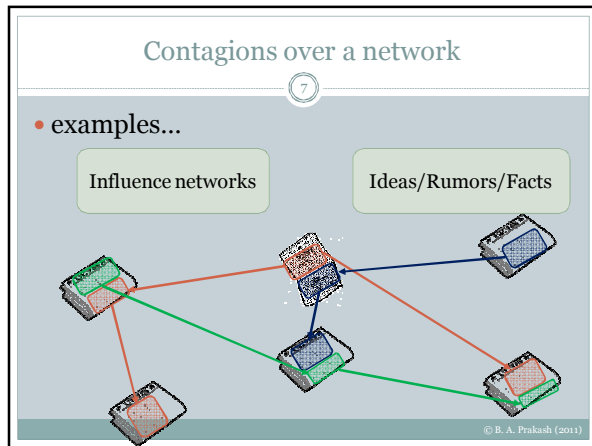
Influence networks

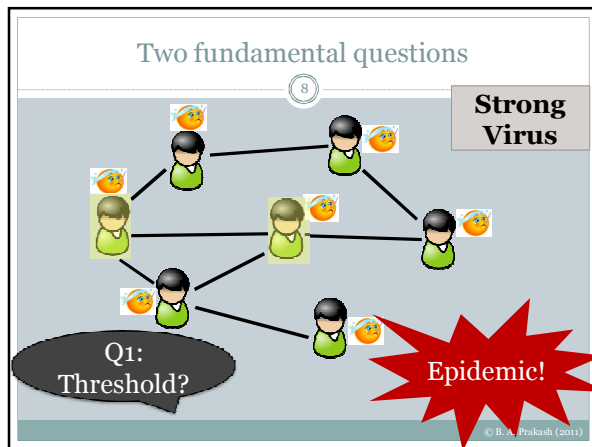


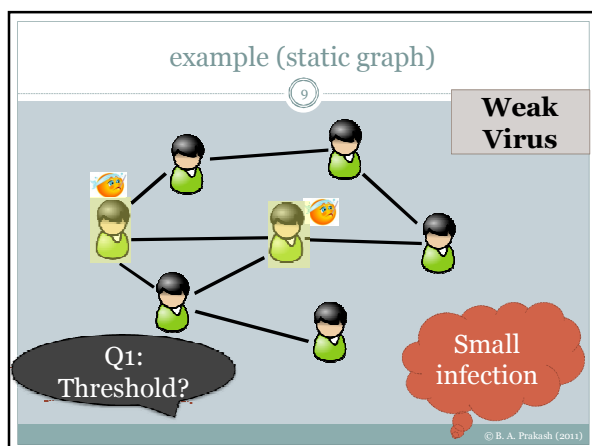
Products



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Questions...

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Q2: Immunization

SARS costs 700+ lives; \$40+ Bn; H1N1 costs Mexico \$2.3bn

This lecture...

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- Focus on **propagation** processes on **arbitrary static** and **dynamic** networks and **temporal evolution** of data.
- Theory:
 - Tipping point? Footprint? Etc.
- Algorithms:
 - Which nodes to immunize? Etc.

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Lecture: Overview

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	Theory: Observations and Results	Algorithms: Tools
Propagation Models on Static Graphs		
Propagation Models on Dynamic Graphs		

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Lecture: Overview

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Propagation Models on Static Graphs Propagation Models on Dynamic Graphs	Theory: Observations and Results Tipping point?	Algorithms: Tools Immunization Algorithms
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Outline

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- Introduction
- Terminology
- Theory
- Algorithms
- Open Problem
- Conclusion

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“SIR” model (mumps-like)

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- Each node in the graph is in one of three states
 - **S**usceptible (i.e. healthy)
 - **I**nfected
 - **R**emoved (i.e. can't get infected again)

$t = 1$

$t = 2$

$t = 3$

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Terminology: continued

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- Other virus models
 - SIS : susceptible-infected-susceptible, flu-like
 - SIRS : temporary immunity, like pertussis
 - SEIR : mumps-like with virus incubation (E = Exposed)
- Underlying contact-network – ‘who-can-infect-whom’
 - Nodes (*people/computers*)
 - Edges (*links between nodes*)

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Related Work

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- ◻ R. M. Anderson and R. M. May. Infectious Diseases of Humans. Oxford University Press, 1991.
- ◻ A. Barrat, M. Barthélemy, and A. Vespignani. Dynamical Processes on Complex Networks. Cambridge University Press, 2010.
- ◻ F. M. Bass. A new product growth for model consumer durables. Management Science, 19(3):215–227, 1969.
- ◻ D. Chakrabarti, Y. Wang, C. Wang, J. Leskovec, and C. Faloutsos. Epidemic thresholds in real networks. ACM TISSEC, 10(4), 2008.
- ◻ D. Basley and J. Kleinberg. Networks, Crowds, and Markets: Reasoning About a Highly Connected World. Cambridge University Press, 2010.
- ◻ A. Ganesh, L. Massoulié, and D. Towsley. The effect of network topology in spread of epidemics. IEEE INFOCOM, 2005.
- ◻ Y. Hayashi, M. Minoura, and J. Matsukubo. Recoverable prevalence in growing scale-free networks and the effective immunization. arXiv:cond-mat/0305549 v2, Aug. 6 2003.
- ◻ H. W. Hethcote. The mathematics of infectious diseases. SIAM Review, 42, 2000.
- ◻ H. W. Hethcote and J. A. Yorke. Gonorrhea transmission dynamics and control. Springer Lecture Notes in Biomathematics, 46, 1984.
- ◻ J. O. Kephart and S. R. White. Directed-graph epidemiological models of computer viruses. IEEE Computer Society Symposium on Research in Security and Privacy, 1991.
- ◻ J. O. Kephart and S. R. White. Measuring and modeling computer virus prevalence. IEEE Computer Society Symposium on Research in Security and Privacy, 1993.
- ◻ R. Pastor-Santorrans and A. Vespignani. Epidemic spreading in scale-free networks. Physical Review Letters 86, 14, 2001.
- ◻

All are about *either*:

➤ **Structured topologies** (cliques, block-diagonals, hierarchies, random)

➤ **Specific virus models**

➤ **Static graphs**

Outline

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- Introduction
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- **Theory**
- Algorithms
- Open Problem
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Lecture: Overview		
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	Theory: Observations and Results	Algorithms: Tools
Propagation Models on Static Graphs	(TS1) G2-threshold theorem	(AS1) Full Symmetric Static Immunization (NetShield)
Propagation Models on Dynamic Graphs	(TD1) Thresholds	(AD1) Full Symmetric Dynamic Immunization

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(TS1) Static Graphs: G2-threshold Theorem

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Problem Statement

- Given:
 - An undirected unweighted graph G , and
 - A virus propagation model (VPM) and its parameters (e.g., β and δ for SIR)
- Find:
 - A condition for virus extinction/invasion



In Prakash+ ICDM 2011. Also posted as *arXiv:1004.0060*

Intuitively

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- Answer should depend on:
 - Graph
 - Virus Propagation Model (VPM)
- But how??
 - Graph – average degree? max. degree? diameter?
 - VPM – which parameters?
 - How to combine – linear? quadratic? exponential?

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Static Graphs: G2-threshold Theorem

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- Main result (informal):

For,

- any arbitrary topology (adjacency matrix A)
- any virus propagation model (VPM) in standard literature

the epidemic threshold depends only

1. on the λ , first eigenvalue of A , and
2. some constant C_{VPM} , determined by the virus propagation model

“G2”: two orthogonal generalizations

 λ C_{VPM}

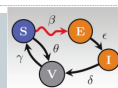
No epidemic if $\lambda * C_{VPM} < 1$

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Thresholds for some models

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- s = effective strength
- $s < 1$: below threshold



Models	Effective Strength (s)	Threshold (tipping point)
SIS, SIR, SIRS, SEIR	$s = \lambda \cdot \left(\frac{\beta}{\delta} \right)$	$s = 1$
SIV, SEIV	$s = \lambda \cdot \left(\frac{\beta\gamma}{\delta(\gamma + \theta)} \right)$	
SI ₁ I ₂ V ₁ V ₂ (H.I.V.)	$s = \lambda \cdot \left(\frac{\beta_1 v_2 + \beta_2 \epsilon}{v_2(\epsilon + v_1)} \right)$	

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Largest Eigenvalue (λ)

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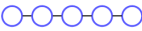

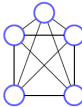
- “Official” definition:
Let A be the adjacency matrix. Then λ is the root with the largest magnitude of the characteristic polynomial of A [$\det(A - xI)$].
- Yeah, doesn’t give too much intuition! ☺

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Largest Eigenvalue (λ)

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better connectivity \rightarrow higher λ

(a)Chain
(b)Star
(c)Clique



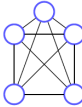
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Largest Eigenvalue (λ)

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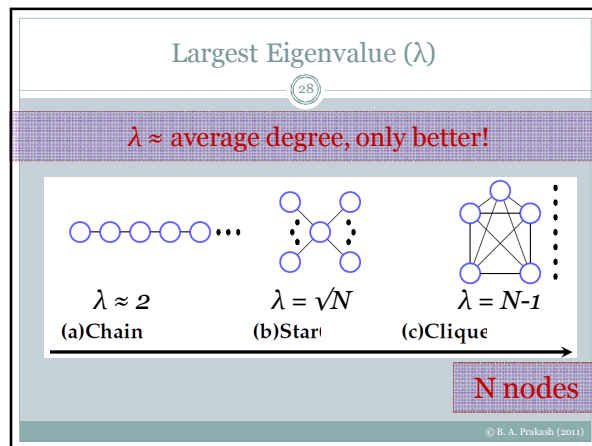
better connectivity \rightarrow higher λ

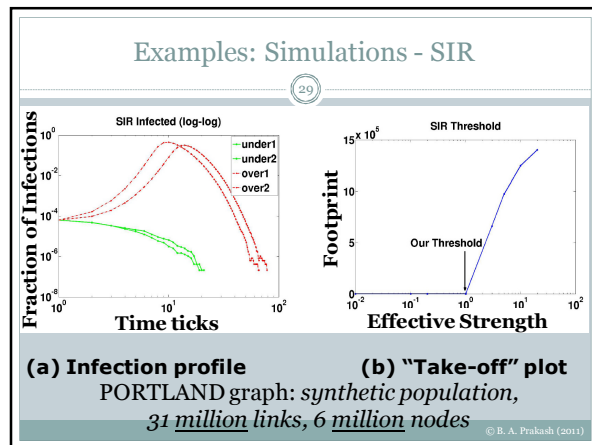




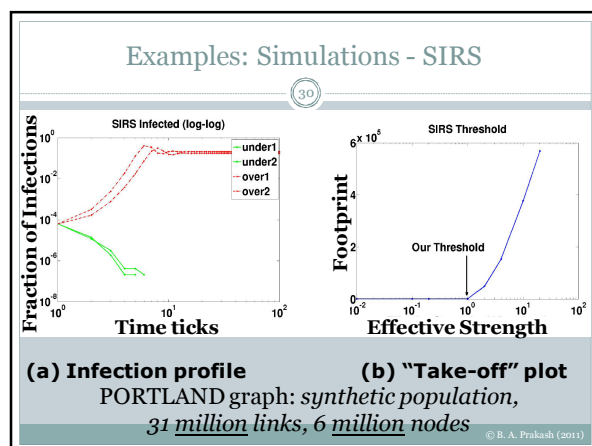
(a)Chain($\lambda = 1.73$)
(b)Star($\lambda = 2$)
(c)Clique($\lambda = 4$)

\rightarrow

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So how did we do that?

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Some trivia

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- > first person in the US identified as a healthy carrier of the pathogen associated with typhoid fever.
- > infected some 53 people, over the course of her career as a cook!
- > forcibly quarantined by public health authorities

Two "Infected" States?

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SICR:
with a
carrier

Asymptomatic

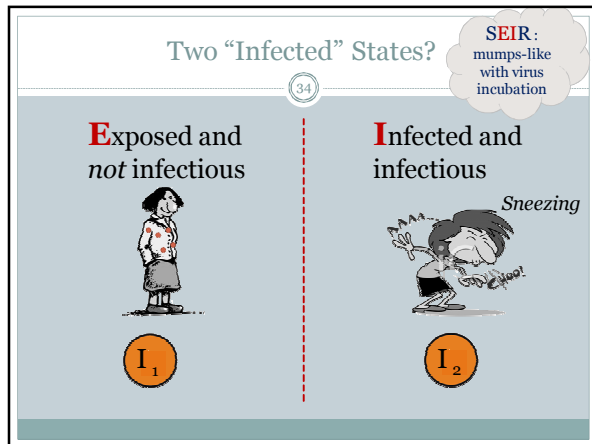


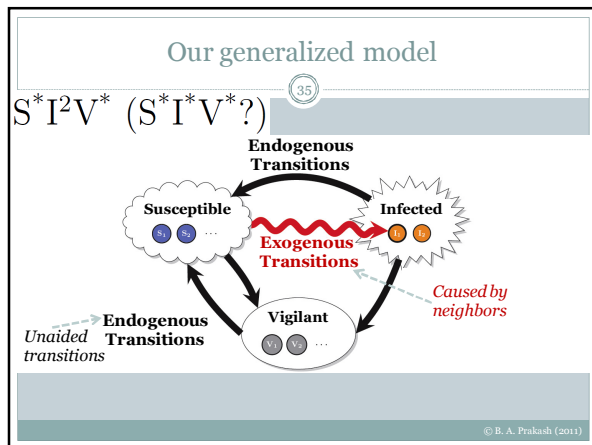
I_1

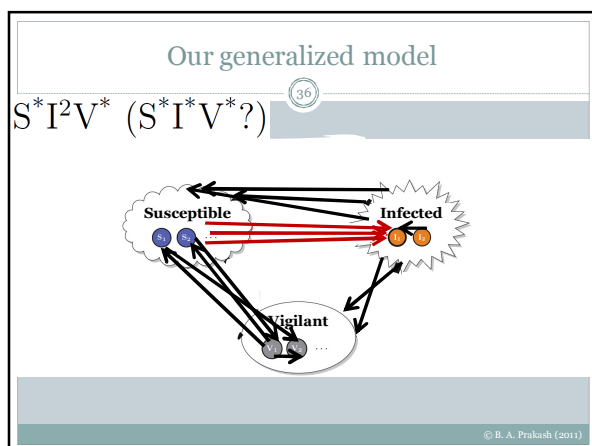
Symptomatic

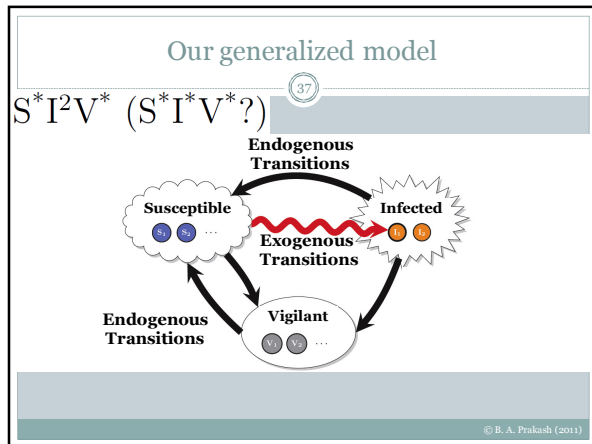


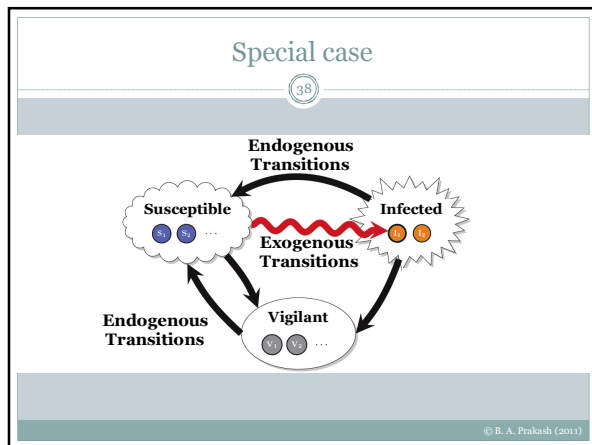
I_2

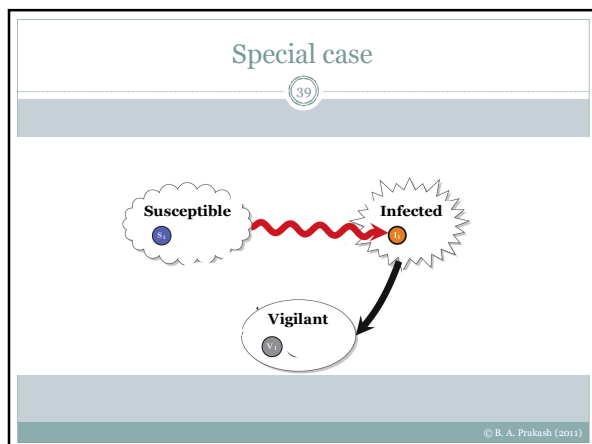


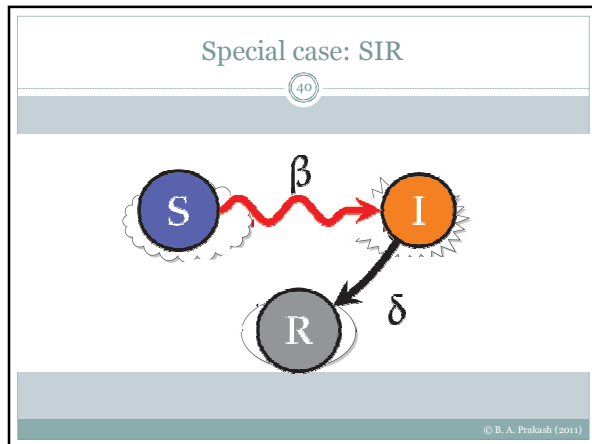


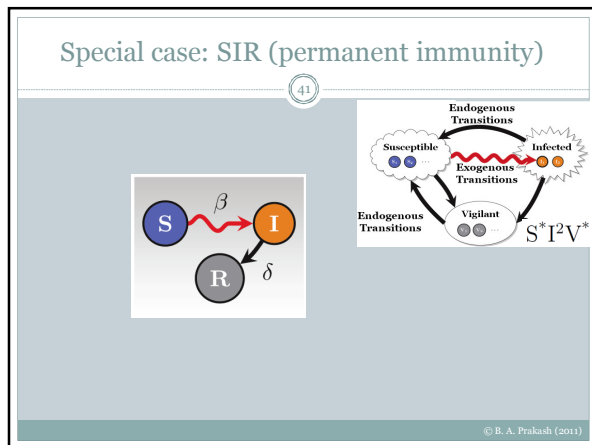


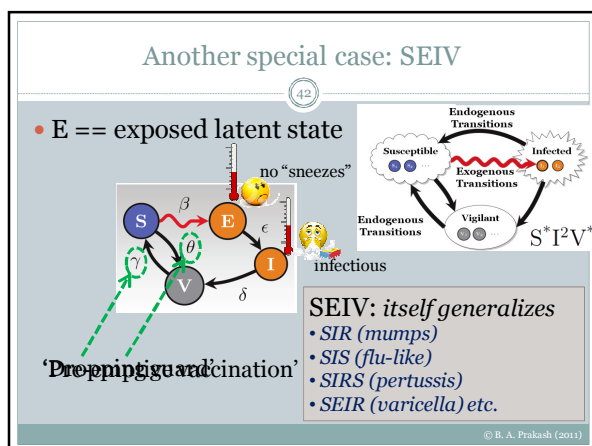


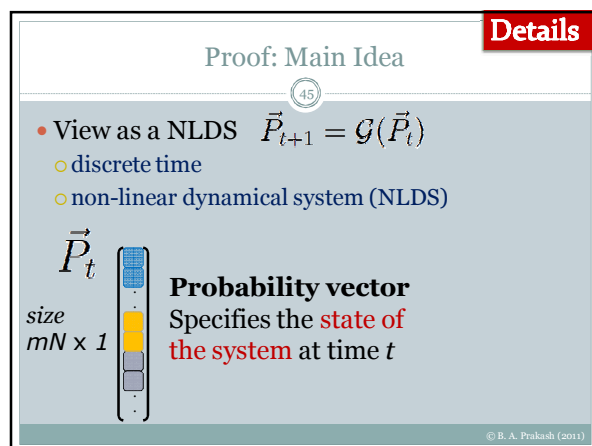
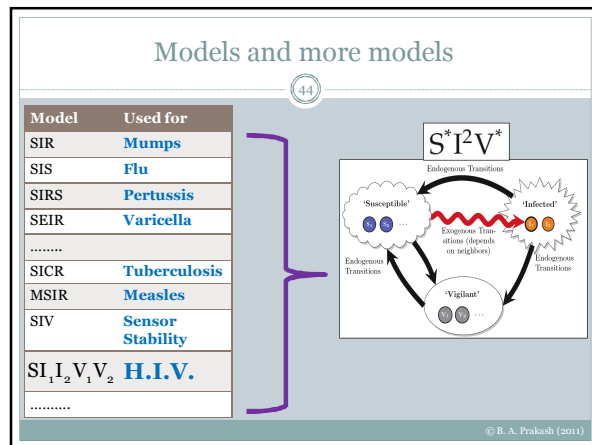
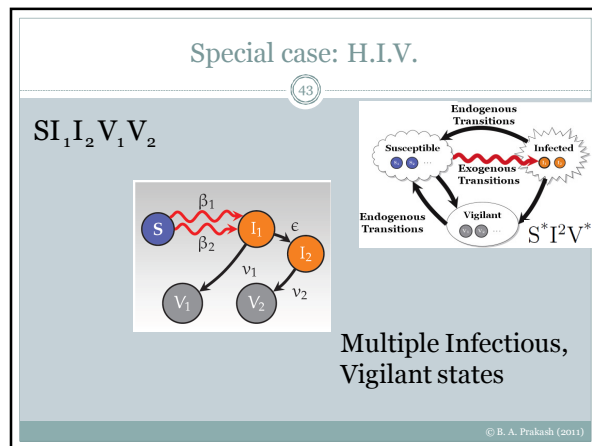












Details

Proof: Main Idea


(46)

- View as a NLDS $\vec{P}_{t+1} = \mathcal{G}(\vec{P}_t)$
 - discrete time
 - non-linear dynamical system (NLDS)

\vec{P}_t size $mN \times 1$

Non-linear function
Explicitly gives the evolution of system

$\mathcal{G} : \mathbb{R}^{mN} \rightarrow \mathbb{R}^{mN}$




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Proof: Main Idea

(47)

- View as a NLDS $\vec{P}_{t+1} = \mathcal{G}(\vec{P}_t)$
 - discrete time
 - non-linear dynamical system (NLDS)
- Threshold \rightarrow Stability of NLDS



(A) Unstable (B) Stable (C) Neutral (at threshold)

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Details

Special case: SIR

(48)

\vec{P}_{t+1} size $3N \times 1$

$\mathcal{G} : \mathbb{R}^{3N} \rightarrow \mathbb{R}^{3N}$

$P_{S,i,t+1} = P_{S,i,t} \zeta_{i,t}(I)$
 $P_{L,i,t+1} = P_{S,i,t} (1 - \zeta_{i,t}(I)) + (1 - \delta) P_{L,i,t}$
 $P_{R,i,t+1} = \delta P_{L,i,t} + P_{R,i,t}$

$\zeta_{i,t}(I)$ = probability that node i is not attacked by any of its infectious neighbors

NLDS

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Fixed Point

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Details

$$\vec{P}^* = \begin{bmatrix} 1 \\ 1 \\ 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

State when no node is infected

NLDS

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Stability for SIR

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Stable
under threshold

Unstable
above threshold

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General Model Structure

See paper for full proof

+

→

Model-based

Graph-based

$\lambda^* C_{VPM} < 1$

NLDS stability

(A) Unstable

(B) Stable

(C) Neutral (at threshold)

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Lecture: Overview		
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	Theory: Observations and Results	Algorithms: Tools
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(TD1) Dynamic Graphs: Thresholds

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- What about time-varying dynamic graphs?

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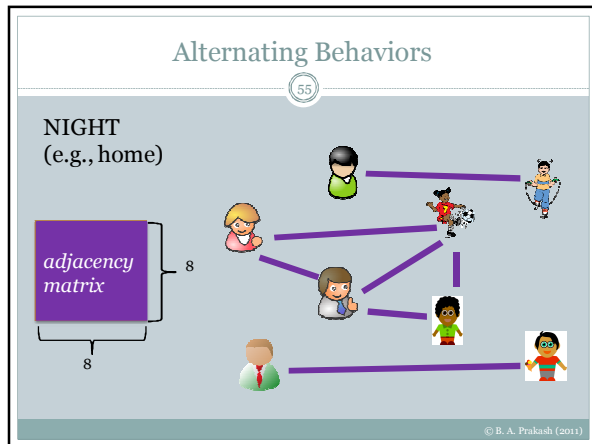
(TD1) Dynamic Graphs: Thresholds

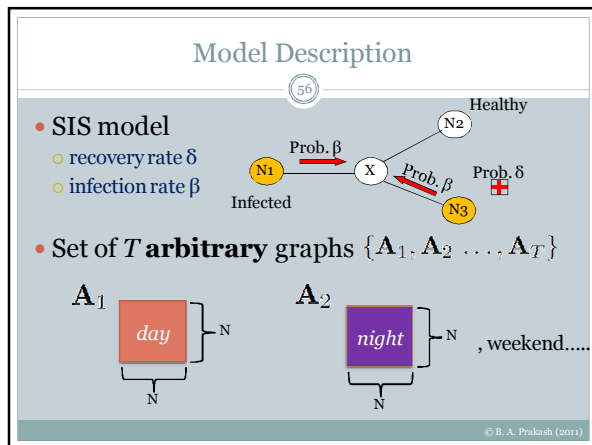
54

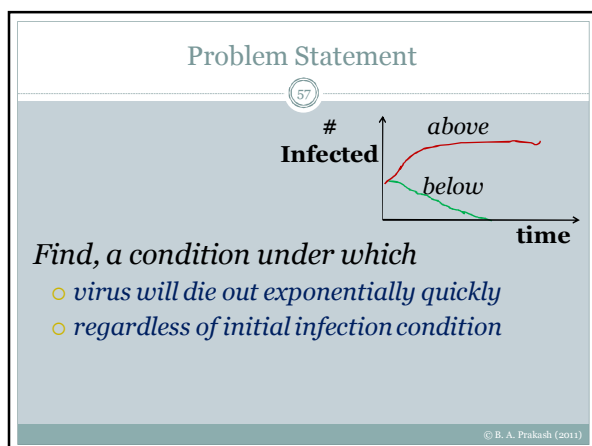
DAY
(e.g., work)

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Should we,

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- Add the adjacency matrices?  + 
- Multiply?  x 
- “Superimpose”? 
- or some other crazy combination?

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Our result: Threshold

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- Informally, **NO** epidemic if

$$\text{eig}(\mathbf{S}) = \lambda_{\mathbf{S}} < 1$$

Single number!
Largest eigenvalue of
the “system matrix” (dfn. next)

In Prakash+, ECML-PKDD 2010

NO epidemic if

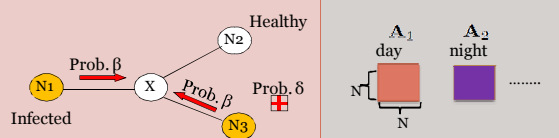
$$\text{eig}(\mathbf{S}) = \lambda_{\mathbf{S}} < 1$$

$$\mathbf{S} = \prod_i \mathbf{S}_i$$

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$$\mathbf{S}_i = (1 - \delta)\mathbf{I} + \beta\mathbf{A}_i$$

Recovery rate Infection rate Adjacency matrices



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Simulation Examples

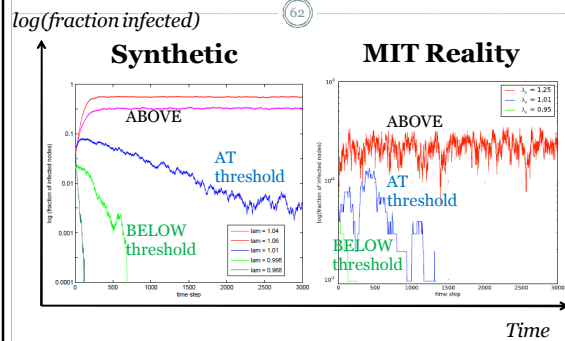
61

- Synthetic
 - 100 nodes
 - A_1 - Clique A_2 - Chain
- MIT Reality Mining
 - 104 mobile devices
 - September 2004 – June 2005
 - 12-hr adjacency matrices
 - A_1 (day) A_2 (night)

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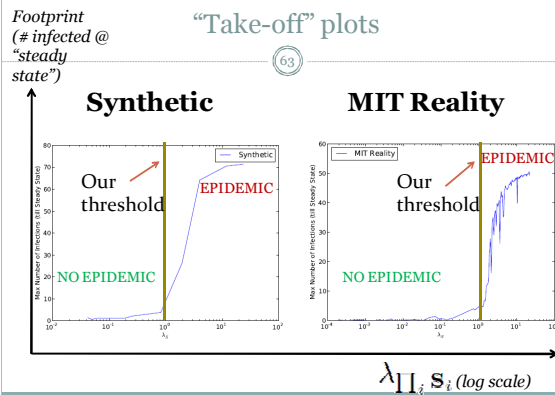
Infection-profile

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“Take-off” plots

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Outline

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- Introduction
- Terminology
- Theory
- Algorithms
- Open Problem
- Conclusion

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Lecture: Overview

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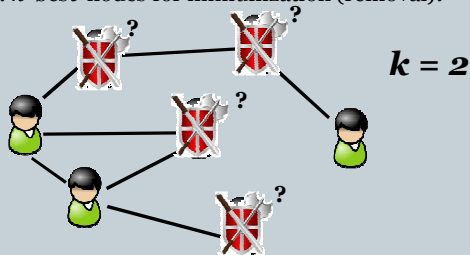
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(AS1): Full Static Immunization

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Given: a graph A , virus prop. model and budget k ;
Find: k 'best' nodes for immunization (removal).



In Tong, Prakash+ ICDM 2010

Challenges

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- Given a graph A , budget k ,

Q1 (Metric) How to measure the 'shield-value' for a set of nodes (S)?

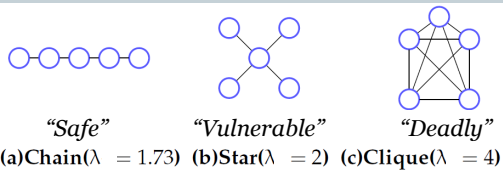
Q2 (Algorithm) How to find a set of k nodes with highest 'shield-value'?

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Proposed vulnerability λ

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λ is the epidemic threshold!



Increasing λ
Increasing vulnerability

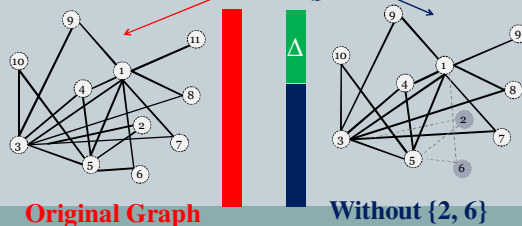
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A1: Eigen-Drop: an ideal shield value

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Eigen-Drop(S)

$$\Delta \lambda = \lambda - \lambda_s$$



(Q2) - Direct Algorithm too expensive!

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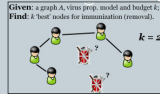
- Immunize k nodes which maximize $\Delta \lambda$

$$S = \operatorname{argmax} \Delta \lambda$$

- Combinatorial!
- Complexity: $O\left(\binom{n}{k} \cdot m\right)$

- Example:

- 1,000 nodes, with 10,000 edges
- It takes 0.01 seconds to compute λ
- It takes **2,615 years** to find 5-best nodes!



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A2: Our Solution

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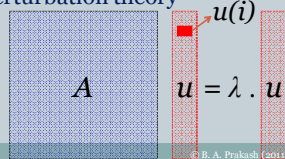
- Part 1
 - Approximate Eigen-drop ($\Delta \lambda$)
- Not enough!
- Part 2
 - Greedily pick best node at each step
- Algorithm: *NetShield*
 - $O(nk^2+m)$

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Our Solution: Part 1

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- Approximate Eigen-drop ($\Delta \lambda$)
- $\Delta \lambda \approx \widehat{SV}(S) = \sum_{i \in S} 2\lambda u(i)^2 - \sum_{i,j \in S} A(i,j)u(i)u(j)$
- Result using Matrix perturbation theory
- $u(i) == \text{'eigenscore'}$
- $\sim \text{pagerank}(i)$



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Details

Details

$$\sum_{i \in S} 2\lambda u(i)^2 - \sum_{i,j \in S} A(i,j)u(i)u(j)$$

P1: node importance P2: set diversity

Original Graph Select by P1 Select by P1+P2

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Details

Done?

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- $S = \operatorname{argmax} \widehat{SV}(S)$
- Still combinatorial
- Complexity: $O(\binom{n}{k} \cdot k^2)$
- Example:
 - 1,000 nodes, with 10,000 edges
 - It takes 0.00001 seconds to compute $\widehat{SV}(S)$
 - It takes **3 months** to find 5-best nodes

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Our Solution: Part 2: NetShield

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- We prove that:
 $\widehat{SV}(S)$ is sub-modular (& monotone non-decreasing)

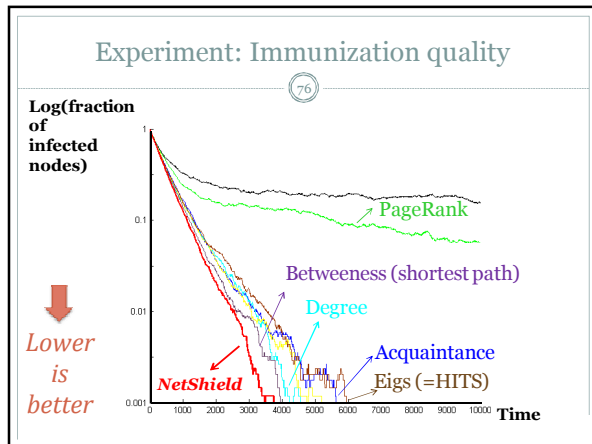
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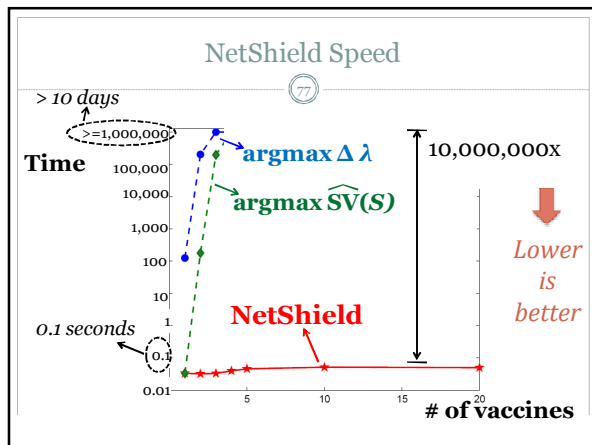
Corollary: Greedy algorithm works

1. NetShield is near-optimal (w.r.t. $\max \widehat{SV}(S)$)
2. NetShield is $O(nk^2+m)$

- NetShield: Greedily add best node at each step

Footnote: near-optimal means $\widehat{SV}(S^{\text{NetShield}}) \geq (1-1/e) \widehat{SV}(S^{\text{Opt}})$





Lecture: Overview

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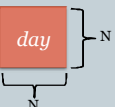
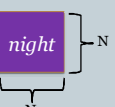
	Theory: Observations and Results	Algorithms: Tools
Propagation Models on Static Graphs	(TS1) G2-threshold Theorem	(AS1) Full Symmetric Static Immunization (NetShield)
Propagation Models on Dynamic Graphs	(TD1) Thresholds	(AD1) Full Symmetric Dynamic Immunization

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(AD1) Full Dynamic Immunization

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- Given:
 - Set of T arbitrary graphs $\{A_1, A_2, \dots, A_T\}$

A_1  A_2  , weekend.....



- Find:
 - k 'best' nodes to immunize (remove)

In Prakash+ ECML-PKDD 2010

(AD1) Full Dynamic Immunization

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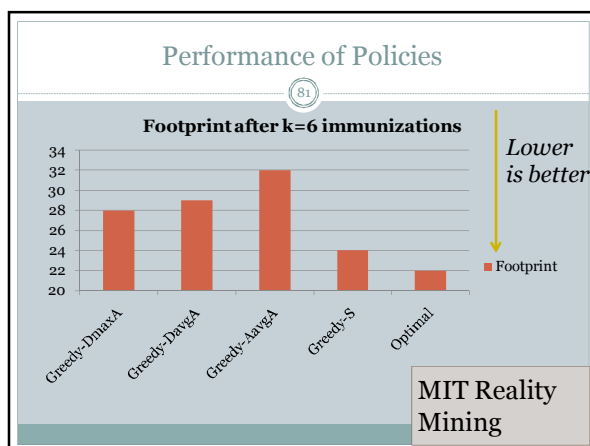
- Our solution
 - Recall theorem
 - Simple: reduce $\lambda_1(S_i) (= \lambda)$

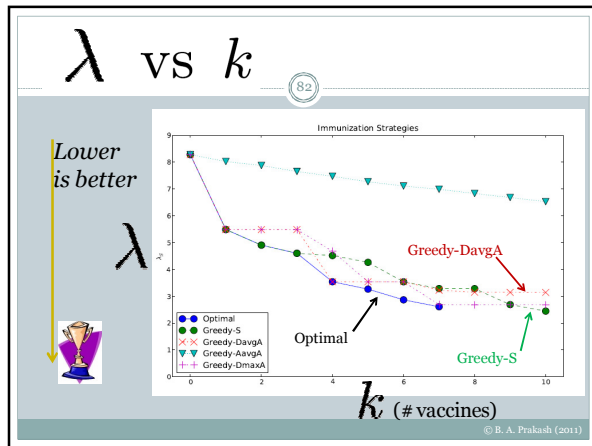
Matrix Product  

- Goal: max eigendrop $\Delta \lambda$

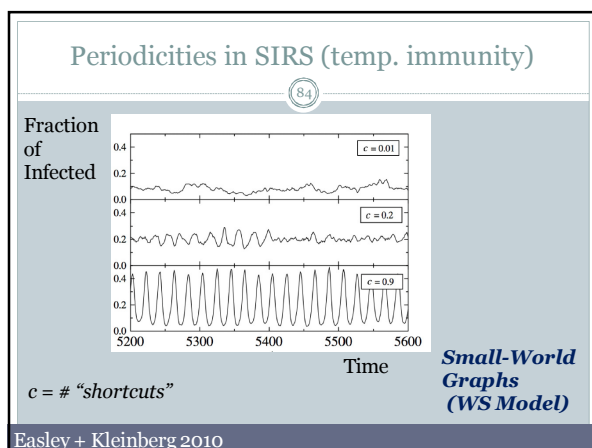
$$\Delta \lambda = \lambda_{\text{before}} - \lambda_{\text{after}}$$
- No competing policy for comparison
- We propose and evaluate many policies

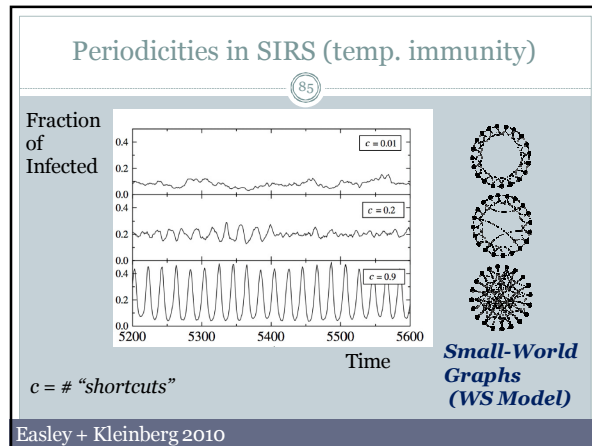
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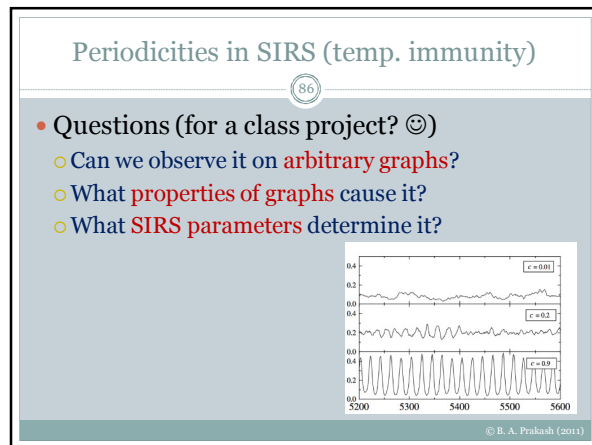


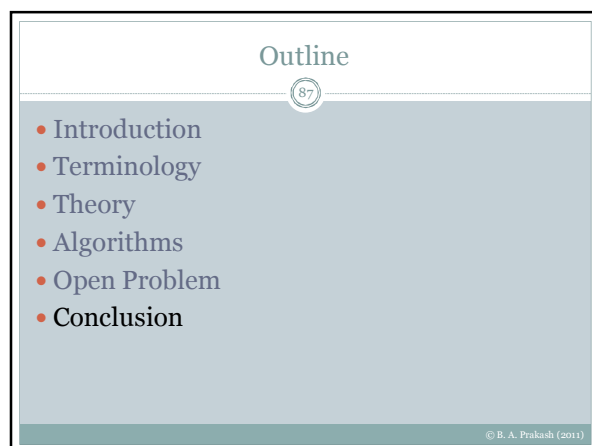


- Outline
- (83)
- Introduction
 - Terminology
 - Theory
 - Algorithms
 - Open problem
 - Conclusion
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Conclusion

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- Threshold depends (for any VPM)
 - on λ (largest eigenvalue) of suitable matrix
 - a constant C_{VPM}
- Use $\Delta\lambda$ to guide immunization policies
 - fast, scalable, provably near-optimal algorithms

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Conclusion

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- **Real, compelling** applications in **diverse** domains
- Lots of interesting research problems!
 - we saw only some of them
 - multiple competing viruses? (Iphone vs android)
 - information diffusion (Twitter) etc. etc.

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References

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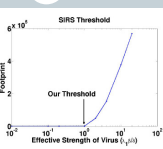
Christos Faloutsos, Roni Rosenfeld, Michalis Faloutsos, Lada Adamic, Jack Iwashyna, Deepayan Chakrabarti, Hanghang Tong, Varun Gupta, Polo Chau, Nicholas Valler

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Any questions?

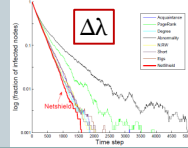


Models	Effective Strength (s)	Threshold (tipping point)
SIS, SIR, SIRS, SEIR	$S = \lambda \cdot \left(\frac{\beta}{\delta} \right)$	$S = 1$
SIV, SEIV	$S = \lambda \cdot \left(\frac{\beta \gamma}{\delta(\gamma + \theta)} \right)$	
SI, I, V, V ₂ (~H.I.V.)	$S = \lambda \cdot \left(\frac{\beta \gamma_1 + \beta \gamma_2}{\gamma_1(\gamma + \gamma_1)} \right)$	



Theory

Tools



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