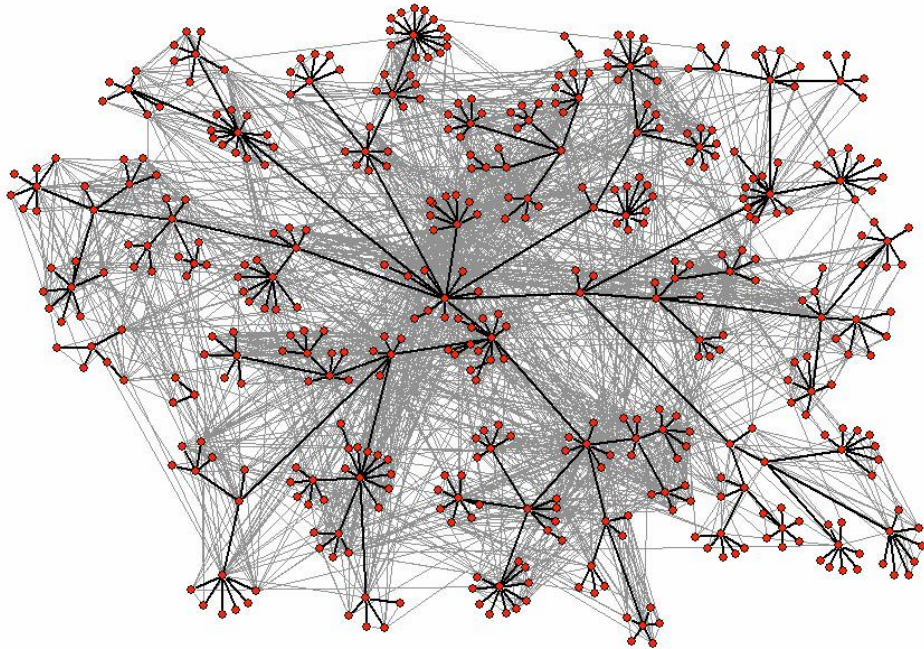


Epidemics in Social Networks



Epidemic Processes

- Viruses, diseases
- Online viruses, worms
- Fashion
- Adoption of technologies
- Behavior
- Ideas

Example: Ebola virus

- First emerged in Zaire 1976 (now Democratic Republic of Kongo)
- Very lethal: it can kill somebody within a few days
- A small outbreak in 2000
- From 10/2000 – 01/2009 173 people died in African villages

Example: HIV

- Less lethal than Ebola
- Takes time to act, lots of time to infect
- First appeared in the 70s
- Initially confined in special groups:
homosexual men, drug users, prostitutes
- Eventually escaped to the entire
population

Example: Melissa computer worm

- Started on March 1999
- Infected MS Outlook users
- The user
 - Receives email with a word document with a virus
 - Once opened, the virus sends itself to the first 50 users in the outlook address book
- First detected on Friday, March 26
- On Monday had infected >100K computers

Example: Hotmail

- Example of Viral Marketing: Hotmail.com
- Jul 1996: Hotmail.com started service
- Aug 1996: 20K subscribers
- Dec 1996: 100K
- Jan 1997: 1 million
- Jul 1998: 12 million

Bought by Microsoft for \$400 million

Marketing: At the end of each email sent there was a message to subscribe to Hotmail.com
“Get your free email at Hotmail”

The Bass model

- Introduced in the 60s to describe product adoption
- Can be applied for viruses
- No network structure

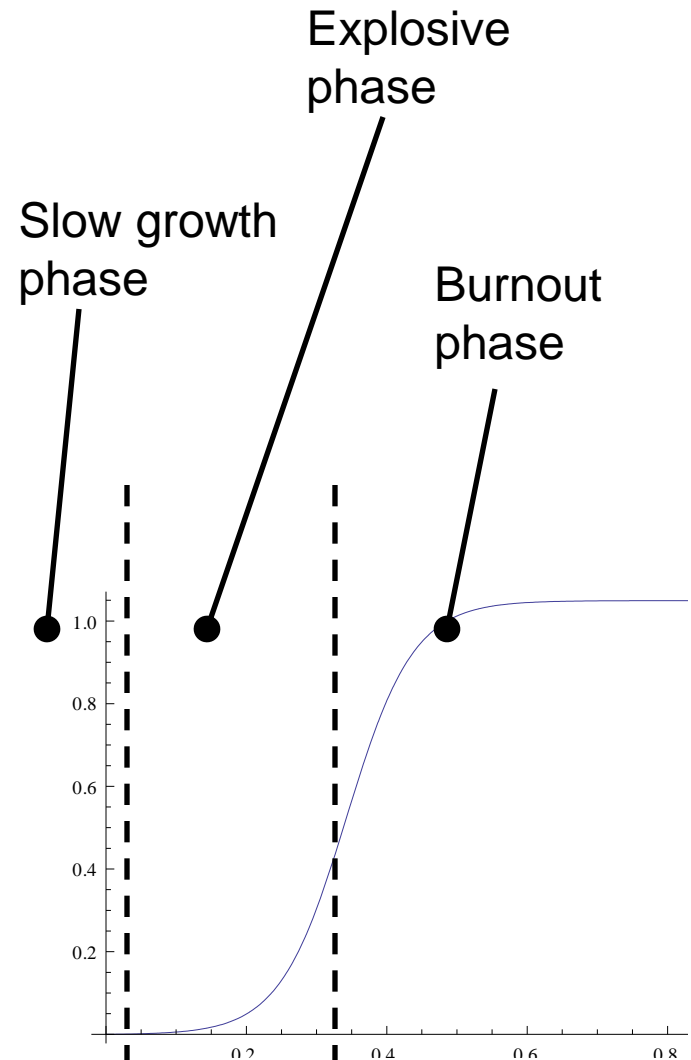
$$F(t + 1) = F(t) + p(1 - F(t)) + q(1 - F(t))F(t)$$

- $F(t)$: Ratio of infected at time t
- p : Rate of infection by outside
- q : Rate of contagion

The Bass model

- $F(t)$: Ratio of infected at time t
- p : Rate of infection by outside
- q : Rate of contagion

$$F(t) = \frac{1 - e^{-(p+q)t}}{1 + \frac{q}{p}e^{-(p+q)t}}$$

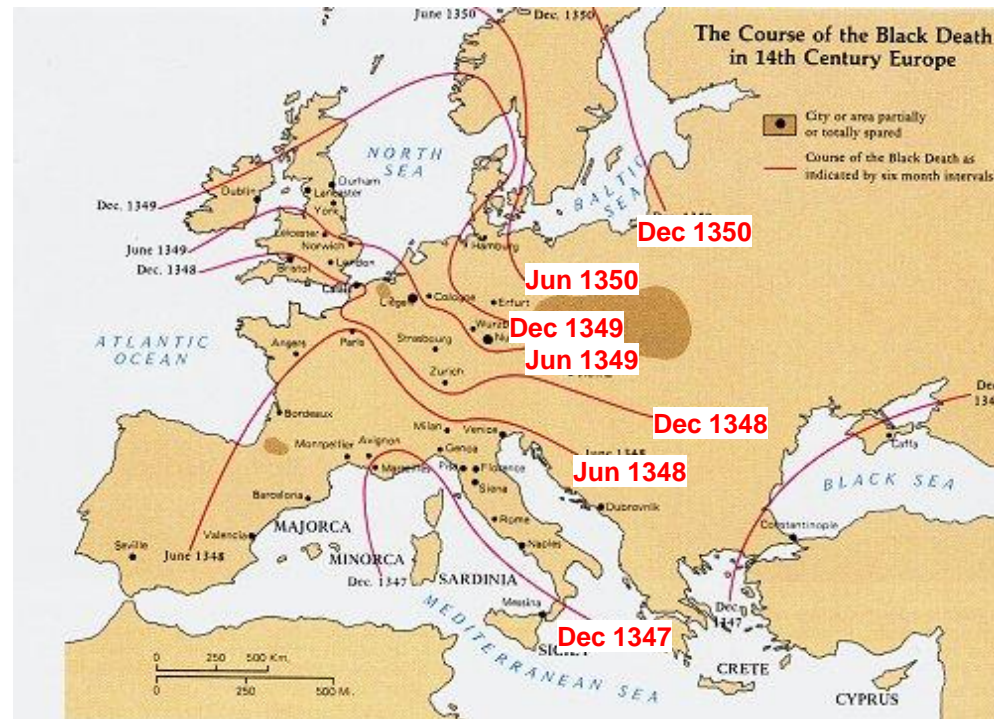


Network Structure

- The Bass model does not take into account network structure
- Let's see some examples

Example: Black Death (Plague)

- Started in 1347 in a village in South Italy from a ship that arrived from China
- Propagated through rats, etc.



Example: Mad-cow disease

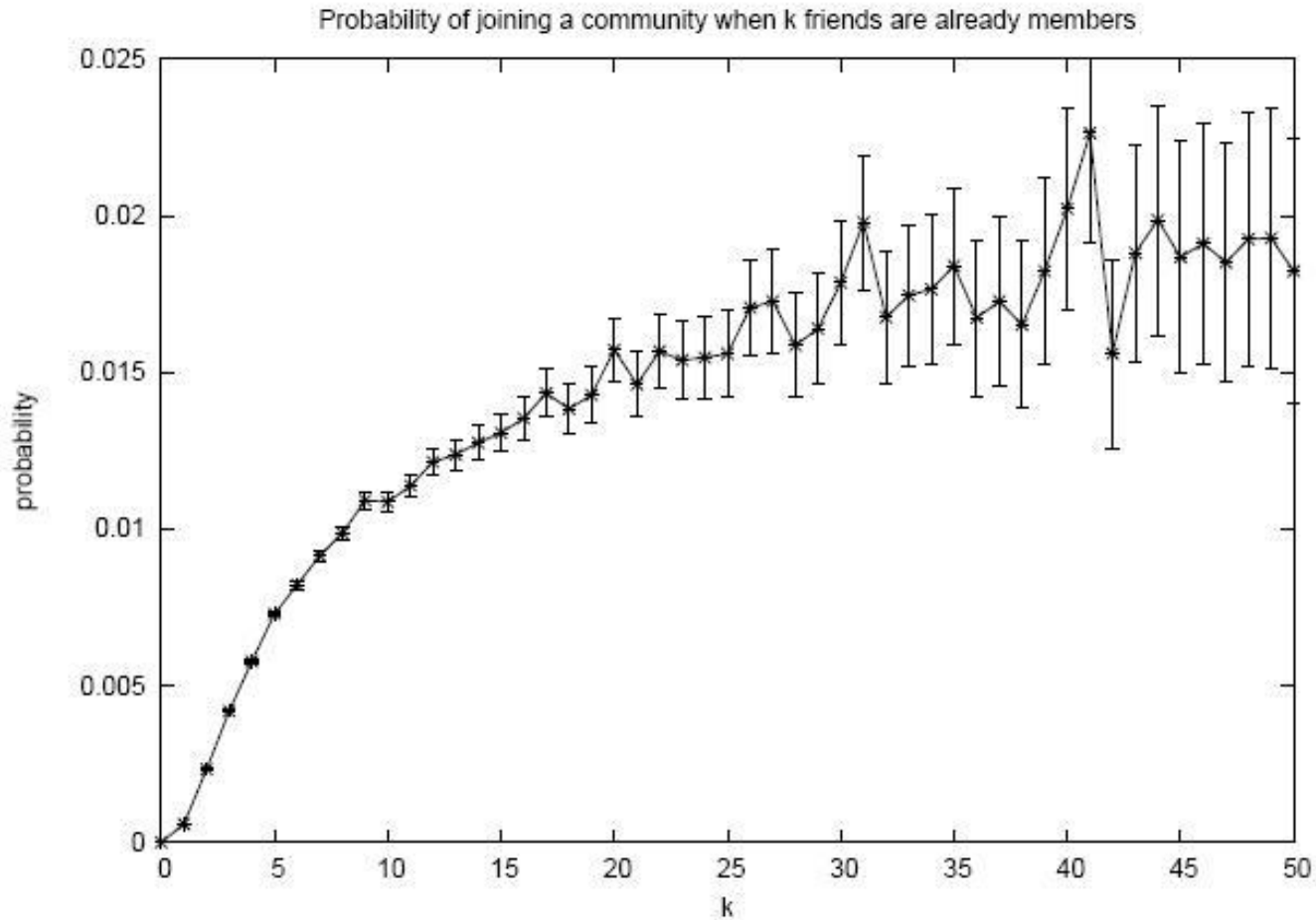
- Jan. 2001: First cases observed in UK
- Feb. 2001: 43 farms infected
- Sep. 2001: 9000 farms infected

- Measures to stop: Banned movement, killed millions of animals

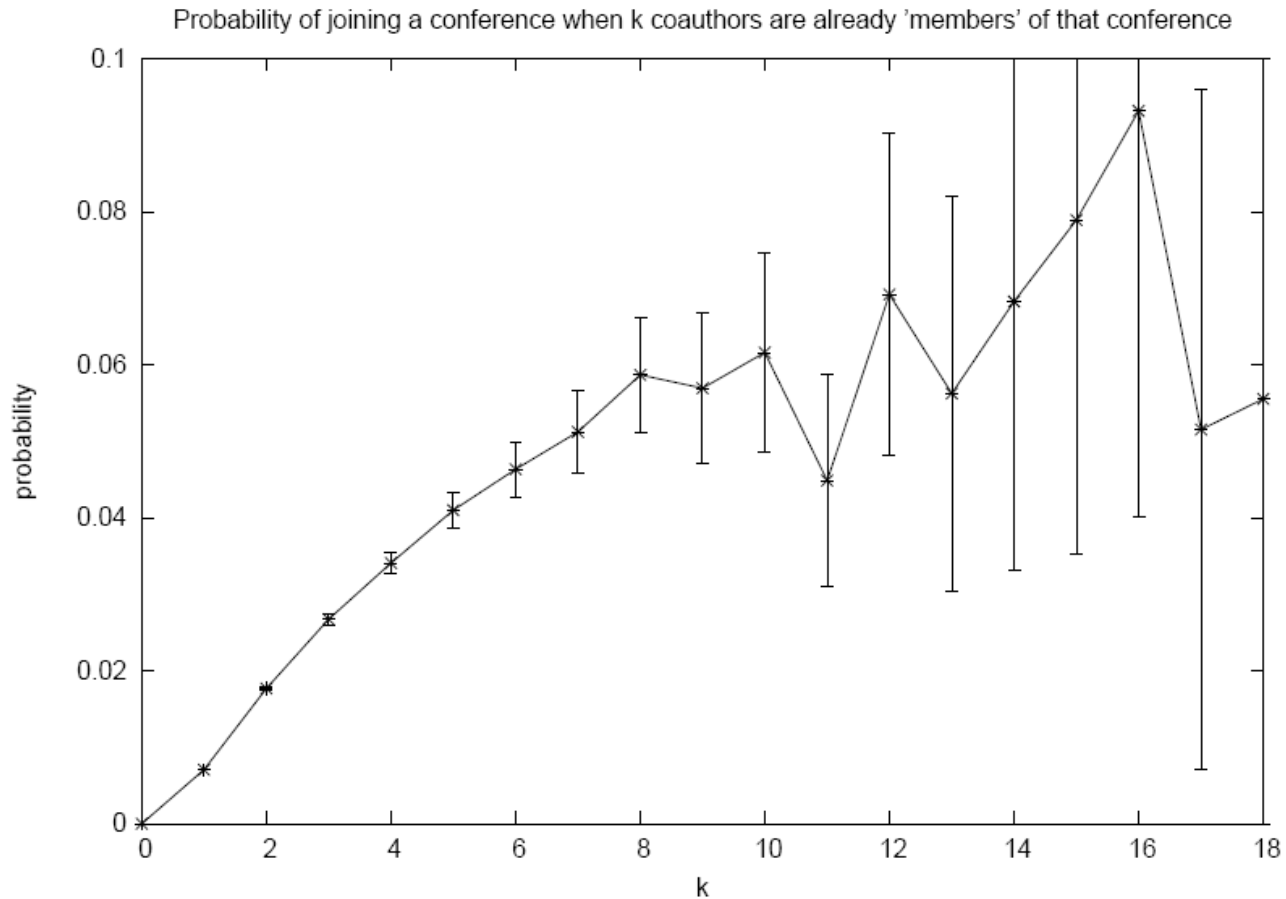
Network Impact

- In the case of the plague it is like moving in a lattice
- In the mad cow we have **weak ties**, so we have a small world
 - Animals being bought and sold
 - Soil from tourists, etc.
- To protect:
 - Make contagion harder
 - Remove weak ties (e.g., mad cows, HIV)

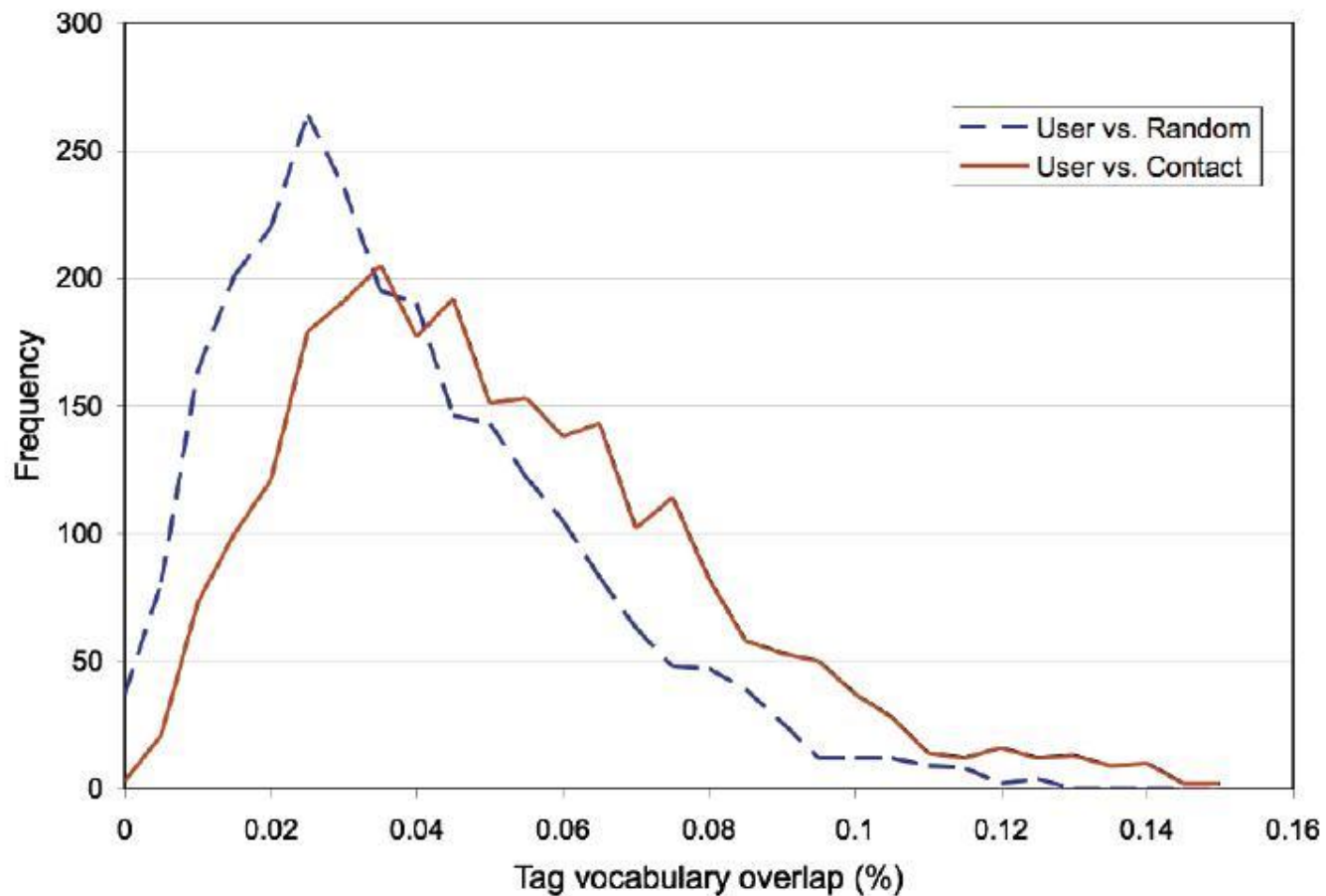
Example: Join an online group



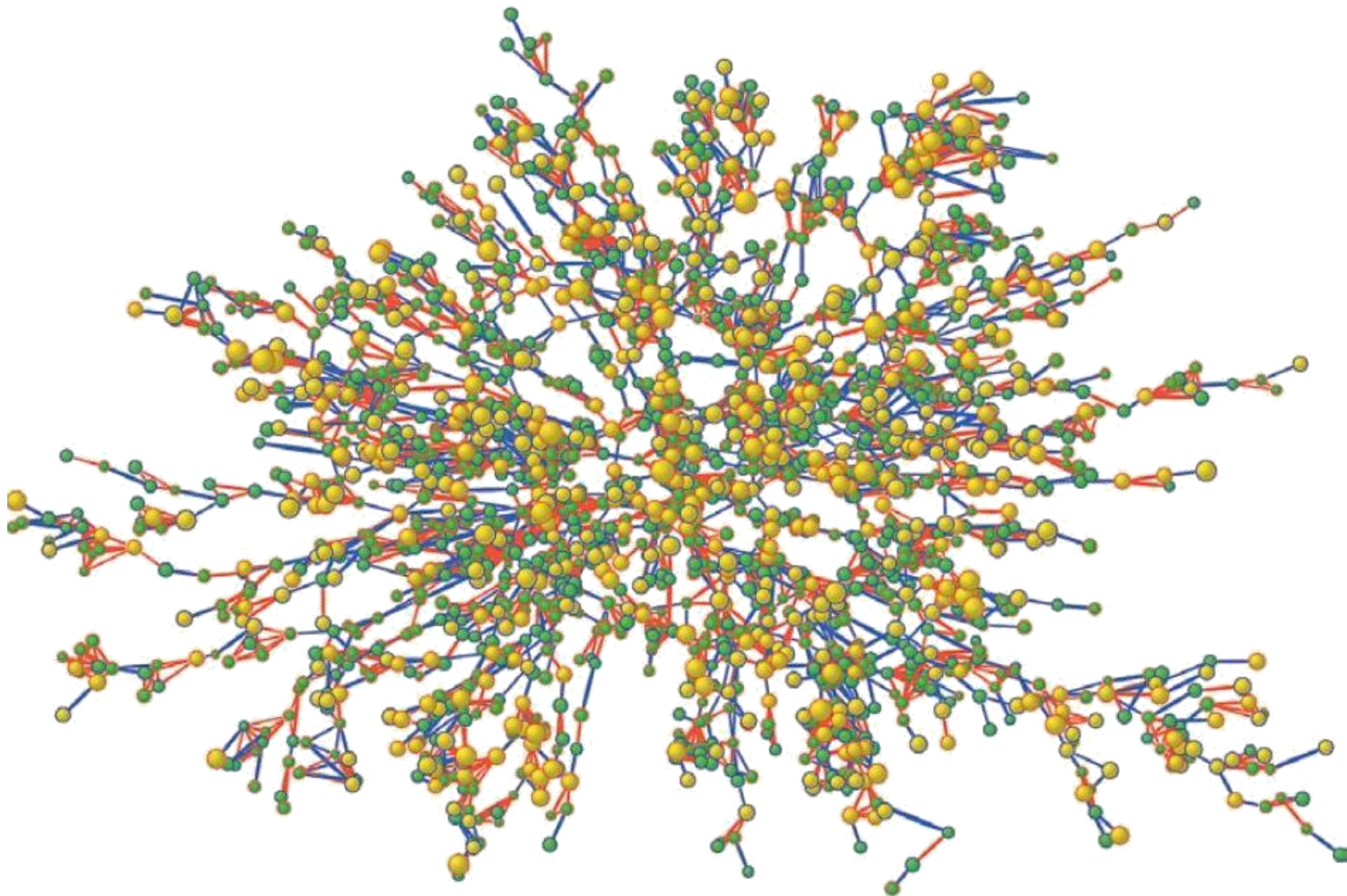
Example: Publish in a conference



Example: Use the same tag



Obesity study



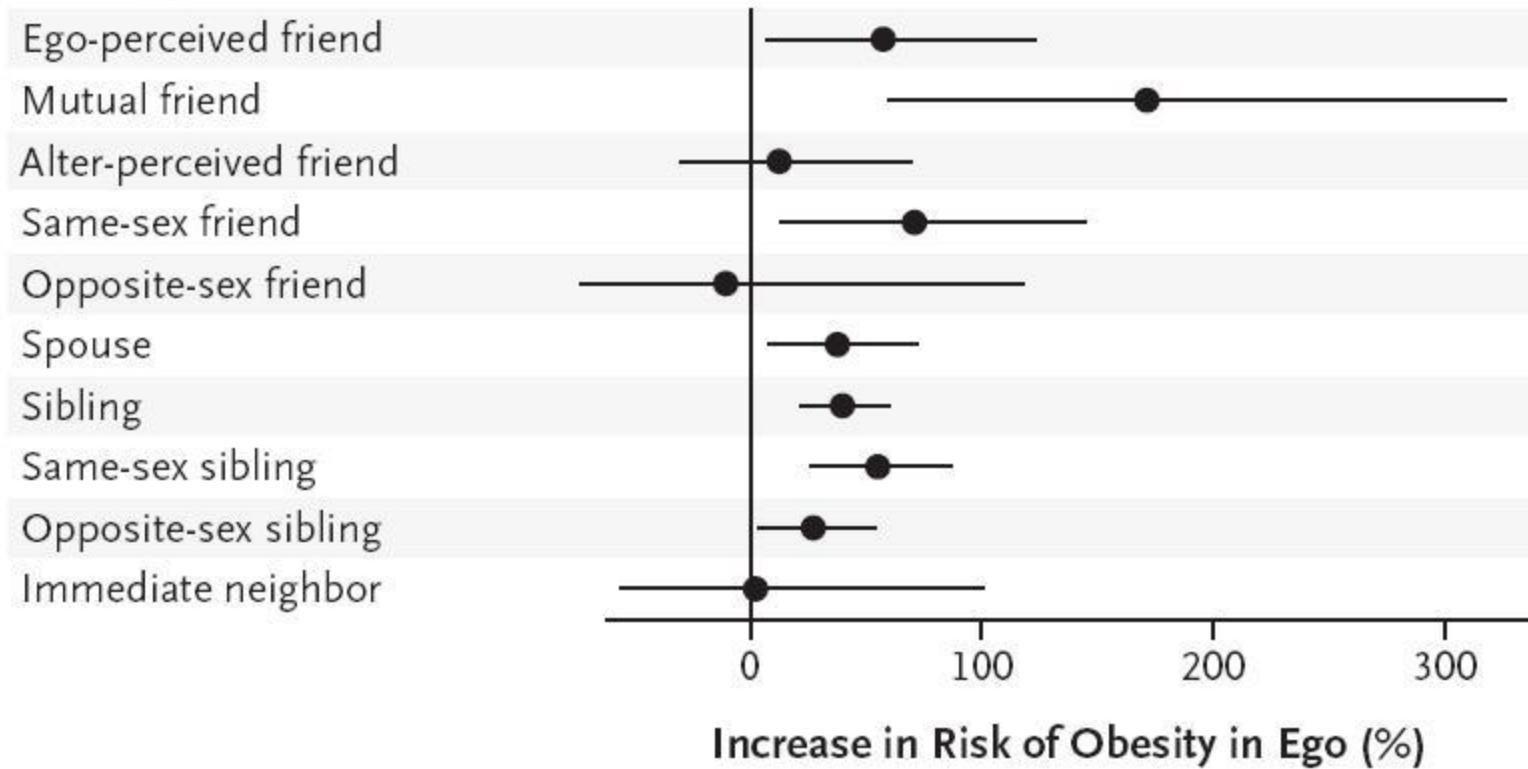
Example: obesity study

Christakis and Fowler, “The Spread of Obesity in a Large Social Network over 32 Years”, New England Journal of Medicine, 2007.

- Data set of 12,067 people from 1971 to 2003 as part of Framingham Heart Study
- Results
 - Having an obese friend increases chance of obesity by 57%.
 - obese sibling → 40%, obese spouse → 37%

Obesity study

Alter Type



Models of Influence

- We saw that often decision is correlated with the number/fraction of friends
- This suggests that there might be influence: the more the number of friends, the higher the influence
- Models to capture that behavior:
 - Linear threshold model
 - Independent cascade model

Linear Threshold Model

- A node v has threshold $\theta_v \sim U[0, 1]$
- A node v is influenced by each neighbor w according to a *weight* b_{vw} such that

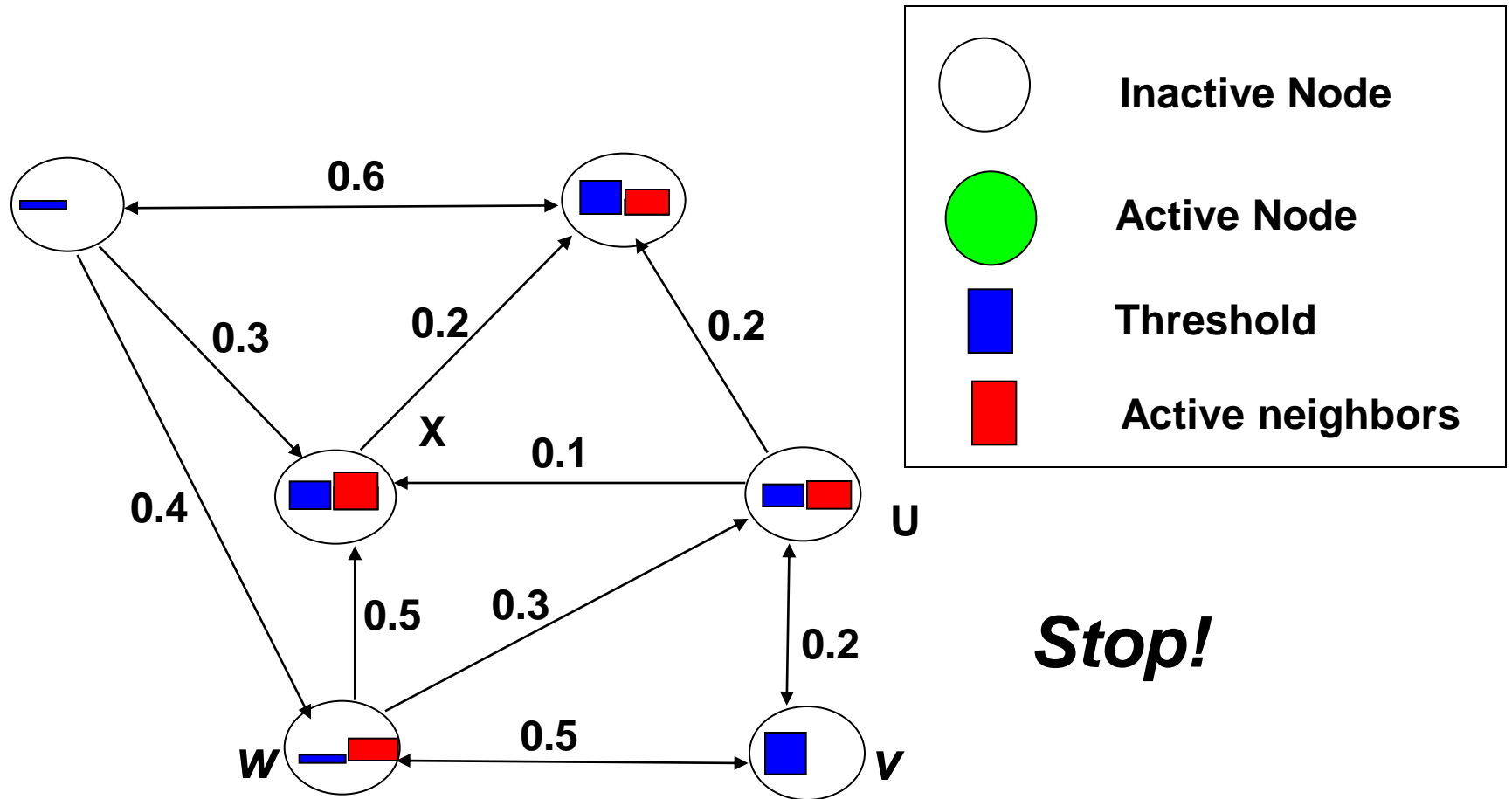
$$\sum_{w \text{ neighbor of } v} b_{v,w} \leq 1$$

- A node v becomes active when at least (weighted) θ_v fraction of its neighbors are active

$$\sum_{w \text{ active neighbor of } v} b_{v,w} \geq \theta_v$$

Examples: riots, mobile phone networks

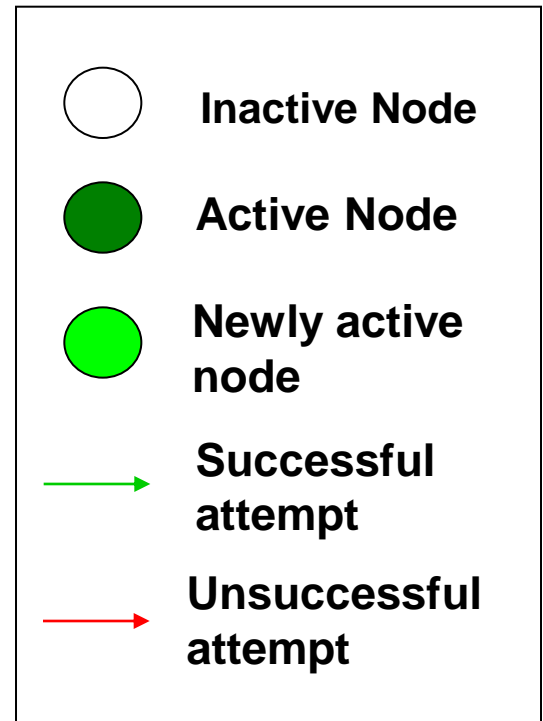
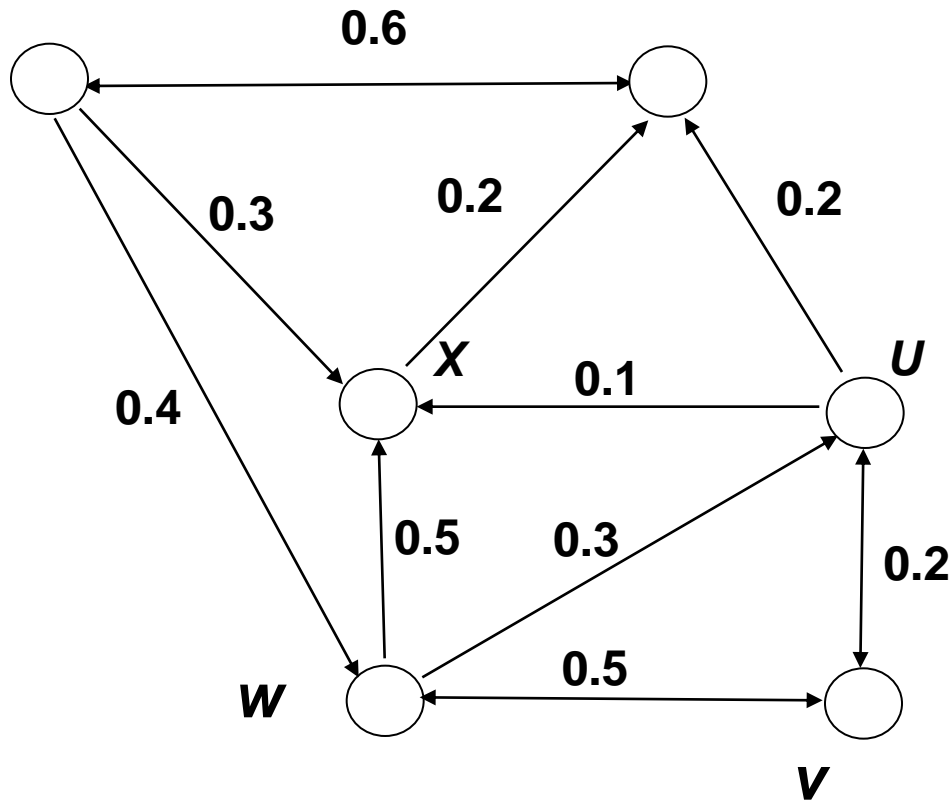
Example



Independent Cascade Model

- When node v becomes active, it has a **single** chance of activating each currently inactive neighbor w .
- The activation attempt succeeds with probability p_{vw} .

Example



Stop!

Optimization problems

- Given a particular model, there are some natural optimization problems.
 1. How do I select a set of users to give coupons to in order to maximize the total number of users infected?
 2. How do I select a set of people to vaccinate in order to minimize influence/infection?
 3. If I have some sensors, where do I place them to detect an epidemic ASAP?

Influence Maximization Problem

- Influence of node set S : $f(S)$
 - **expected** number of active nodes at the end, if set S is the initial active set
- Problem:
 - Given a parameter k (budget), find a k -node set S to maximize $f(S)$
 - Constrained optimization problem with $f(S)$ as the objective function

$f(S)$: properties (to be demonstrated)

- Non-negative (obviously)
- Monotone: $f(S + v) \geq f(S)$
- Submodular:
 - Let N be a finite set
 - A set function $f : 2^N \mapsto \mathfrak{R}$ is submodular *iff*
$$\forall S \subset T \subset N, \forall v \in N \setminus T,$$
$$f(S + v) - f(S) \geq f(T + v) - f(T)$$

(diminishing returns)

Bad News

- For a submodular function f , if f only takes non-negative values, and is monotone, finding a k -element set S for which $f(S)$ is maximized is an NP-hard optimization problem [GFN77, NWF78].
- It is NP-hard to determine the optimum for influence maximization for both independent cascade model and linear threshold model.

Good News

- We can use Greedy Algorithm!
 - Start with an empty set S
 - For k iterations:
 - Add node v to S that maximizes $f(S + v) - f(S)$.
- How good (bad) it is?
 - Theorem: The greedy algorithm is a $(1 - 1/e)$ approximation.
 - The resulting set S activates at least $(1 - 1/e) > 63\%$ of the number of nodes that any size- k set S could activate.

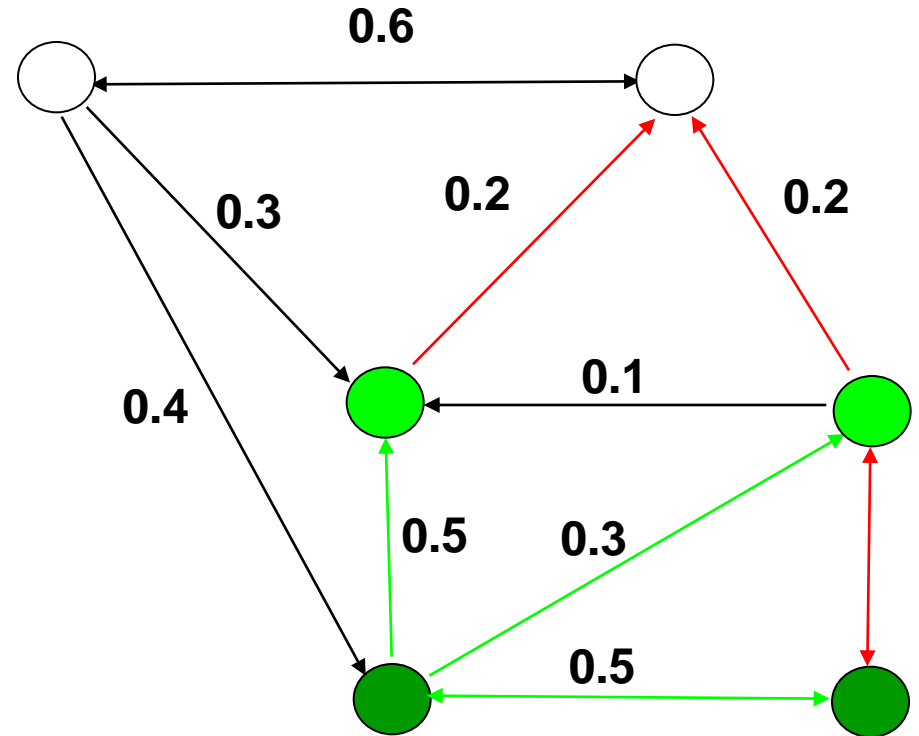
Key 1: Prove submodularity

$$\forall S \subset T \subset N, \forall v \in N \setminus T,$$

$$f(S + v) - f(S) \geq f(T + v) - f(T)$$

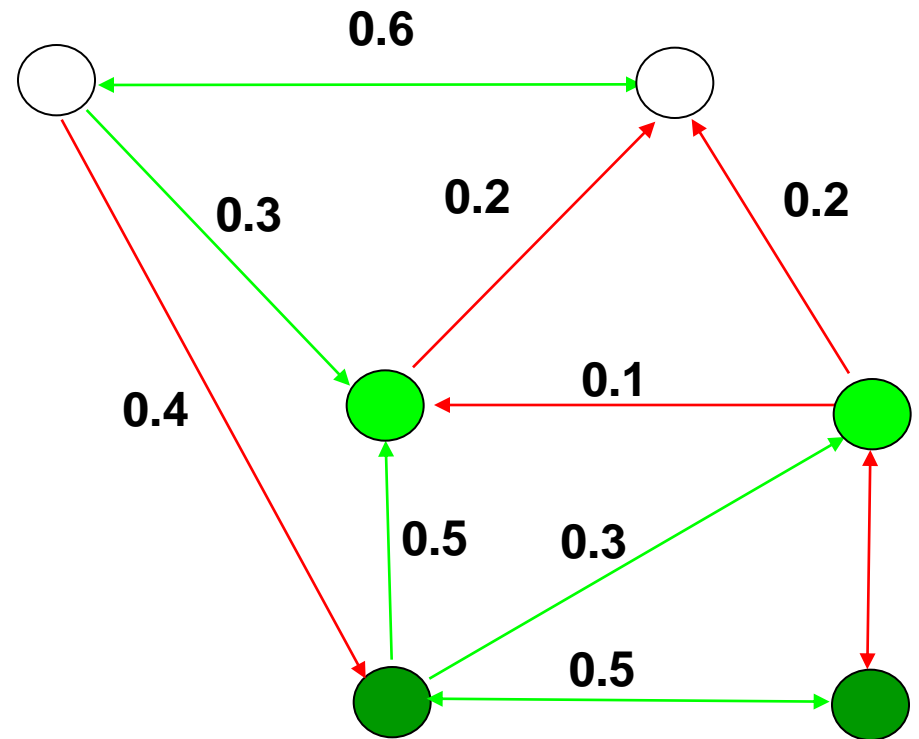
Submodularity for Independent Cascade

- Coins for edges are flipped during activation attempts.



Submodularity for Independent Cascade

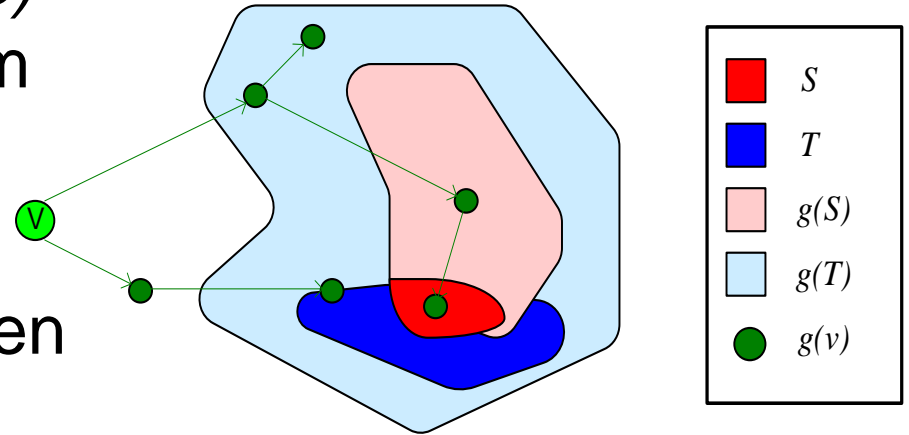
- Coins for edges are flipped during activation attempts.
- Can pre-flip all coins and reveal results immediately.



- Active nodes in the end are reachable via green paths from initially targeted nodes.
- Study reachability in green graphs

Submodularity, Fixed Graph

- Fix “green graph” G . $g(S)$ are nodes reachable from S in G .
- Submodularity: $g(T + v) - g(T) \subseteq g(S + v) - g(S)$ when $S \subseteq T$.



- $g(S + v) - g(S)$: nodes reachable from $S + v$, but not from S .
- From the picture: $g(T + v) - g(T) \subseteq g(S + v) - g(S)$ when $S \subseteq T$ (*indeed!*).

Submodularity of the Function

Fact: A non-negative linear combination of submodular functions is submodular

$$f(S) = \sum_G \text{Prob}(G \text{ is green graph}) \cdot g_G(S)$$

- $g_G(S)$: nodes reachable from S in G .
- Each $g_G(S)$: is submodular (previous slide).
- Probabilities are non-negative.

Submodularity for Linear Threshold

- Use similar “green graph” idea.
- Once a graph is fixed, “reachability” argument is identical.
- How do we fix a green graph now?
- Each node picks at most one incoming edge, with probabilities proportional to edge weights.
- Equivalent to linear threshold model (trickier proof).

Key 2: Evaluating $f(S)$

Evaluating $f(S)$

- How to evaluate $f(S)$?
- Still an open question of how to compute efficiently
- But: very good estimates by simulation
 - repeating the diffusion process often enough (polynomial in n ; $1/\varepsilon$)
 - Achieve $(1 - \varepsilon)$ -approximation to $f(S)$.
- Generalization of Nemhauser/Wolsey proof shows: Greedy algorithm is now a $(1 - 1/e - \varepsilon')$ -approximation.

Experiment Data

- A collaboration graph obtained from co-authorships in papers of the arXiv high-energy physics theory section
- co-authorship networks arguably capture many of the key features of social networks more generally
- Resulting graph: 10748 nodes, 53000 distinct edges

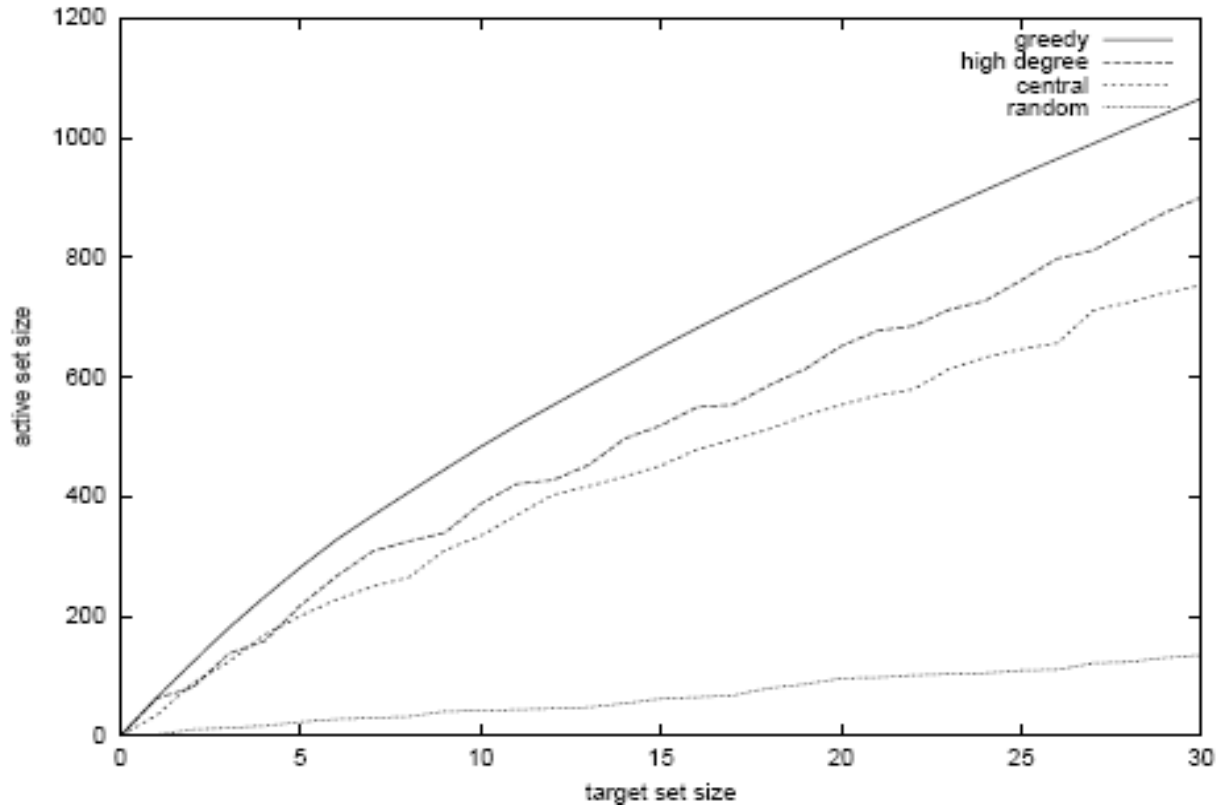
Experiment Settings

- Linear Threshold Model: multiplicity of edges as weights
 - $\text{weight}(v \rightarrow \omega) = C_{vw} / dv$, $\text{weight}(\omega \rightarrow v) = C_{wv} / dw$
- Independent Cascade Model:
 - Case 1: uniform probabilities p on each edge
 - Case 2: edge from v to ω has probability $1/dw$ of activating ω .
- Simulate the process 10000 times for each targeted set, re-choosing thresholds or edge outcomes pseudo-randomly from $[0, 1]$ every time
- Compare with other 3 common heuristics
 - (in)degree centrality, distance centrality, random nodes.

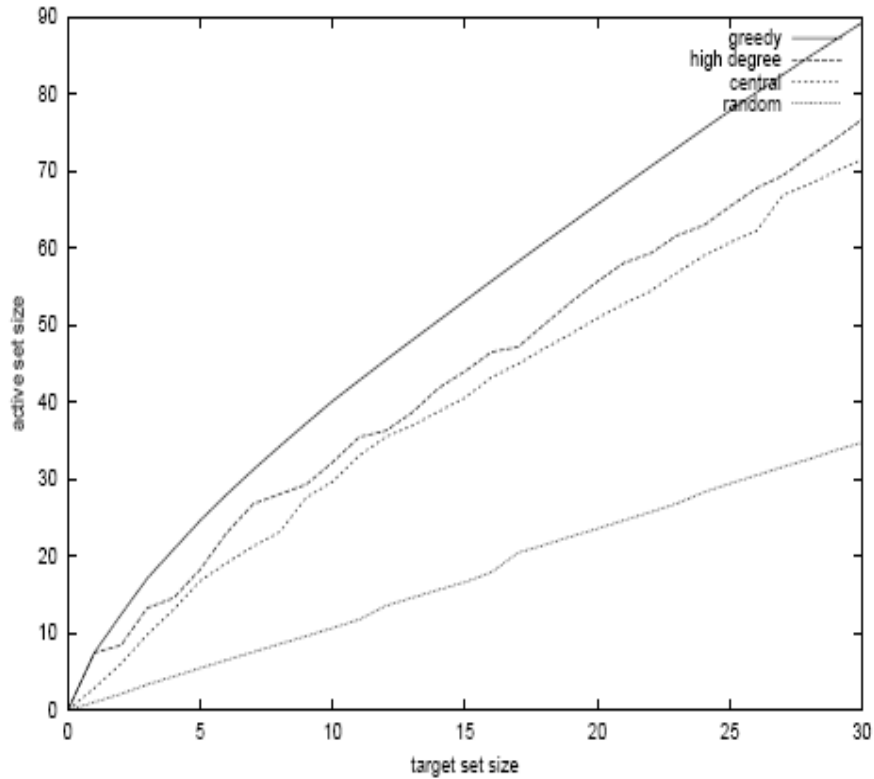
Outline

- Models of influence
 - Linear Threshold
 - Independent Cascade
- Influence maximization problem
 - Algorithm
 - Proof of performance bound
 - Compute objective function
- Experiments
 - Data and setting
 - **Results**

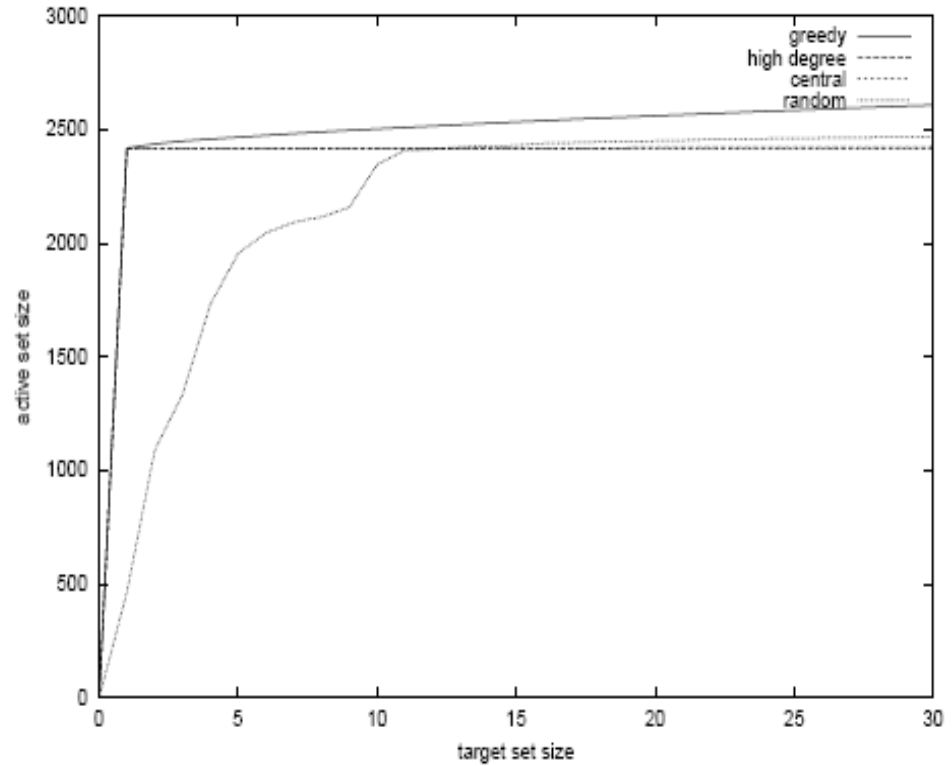
Results: linear threshold model



Independent Cascade Model – Case 1



$P = 1\%$



$P = 10\%$

Independent Cascade Model – Case 2

Reminder: linear
threshold model

