

# Online Social Networks and Media

Cascading Behavior in Networks  
Epidemic Spread  
Influence Maximization

# Introduction

**Diffusion:** process by which a piece of information is spread and reaches individuals through interactions.

# **CASCADING BEHAVIOR IN NETWORKS**

# Innovation Diffusion in Networks

How new behaviors, practices, opinions and technologies spread from person to person through a social network as people influence their friends to adopt new ideas

Information effect: choices made by others can provide indirect information about what they know

Old studies:

- Adoption of hybrid seed corn among farmers in Iowa
- Adoption of tetracycline by physicians in US

Basic observations:

- Characteristics of **early adopters**
- Decisions made in the context of **social structure**

# Spread of Innovation

**Direct-Benefit Effect:** there are direct payoffs from copying the decisions of others (relative advantage)

Spread of technologies such as the phone, email, etc

Common principles:

- ✓ *Complexity* of people to understand and implement
- ✓ *Observability*, so that people can become aware that others are using it
- ✓ *Trialability*, so that people can mitigate its risks by adopting it gradually and incrementally
- ✓ *Compatibility* with the social system that is entering (homophily?)

# A Direct-Benefit Model

An *individual* level model of *direct-benefit effects* in networks due to S. Morris

The benefits of adopting a new behavior increase as more and more of the social network neighbors adopt it

## A Coordination Game

Two players (nodes),  $u$  and  $w$  linked by an edge

Two possible behaviors (strategies): A and B

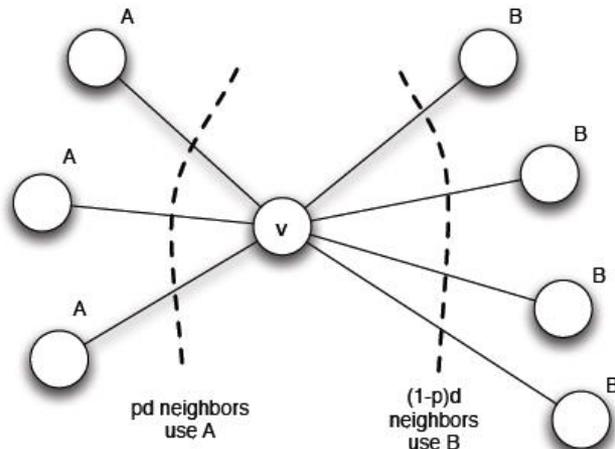
- If both  $u$  and  $w$  adopt A, get payoff  $a > 0$
- If both  $u$  and  $w$  adopt B, get payoff  $b > 0$
- If opposite behaviors, then each get a payoff 0

		$w$	
		A	B
$v$	A	$a, a$	$0, 0$
	B	$0, 0$	$b, b$

# Modeling Diffusion through a Network

$u$  plays a copy of the game with each of its neighbors, its payoff is the *sum* of the payoffs in the games played on each edge

Say some of its neighbors adopt A and some B, what should  $u$  do to maximize its payoff?



Threshold  $q = b/(a+b)$  for preferring A  
(at least  $q$  of the neighbors follow A)

Two obvious equilibria, which ones?

# Modeling Diffusion through a Network: Cascading Behavior

Suppose that initially everyone is using B as a default behavior  
A small set of “initial adopters” decide to use A

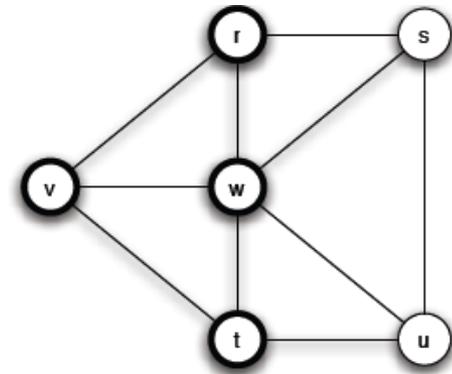
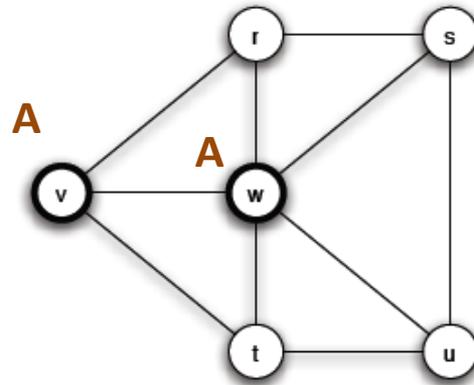
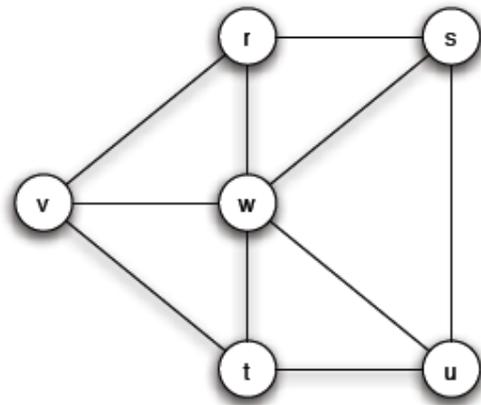
- ✓ When will this result in everyone eventually switching to A?
- ✓ If this does not happen, what causes the spread of A to stop?

Depends on the choice of the *initial adapters* and threshold  $q$

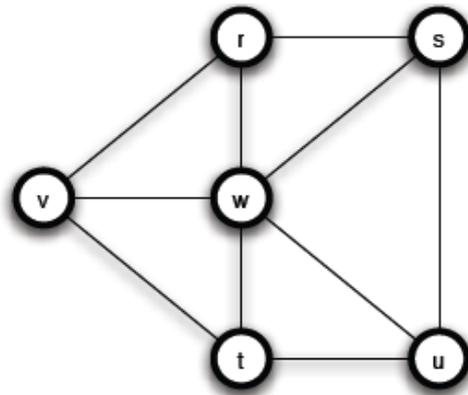
Observation: strictly progressive sequence of switches from B to A

# Modeling Diffusion through a Network: Cascading Behavior

$$a = 3, b = 2, q = 2/5$$



Step 1

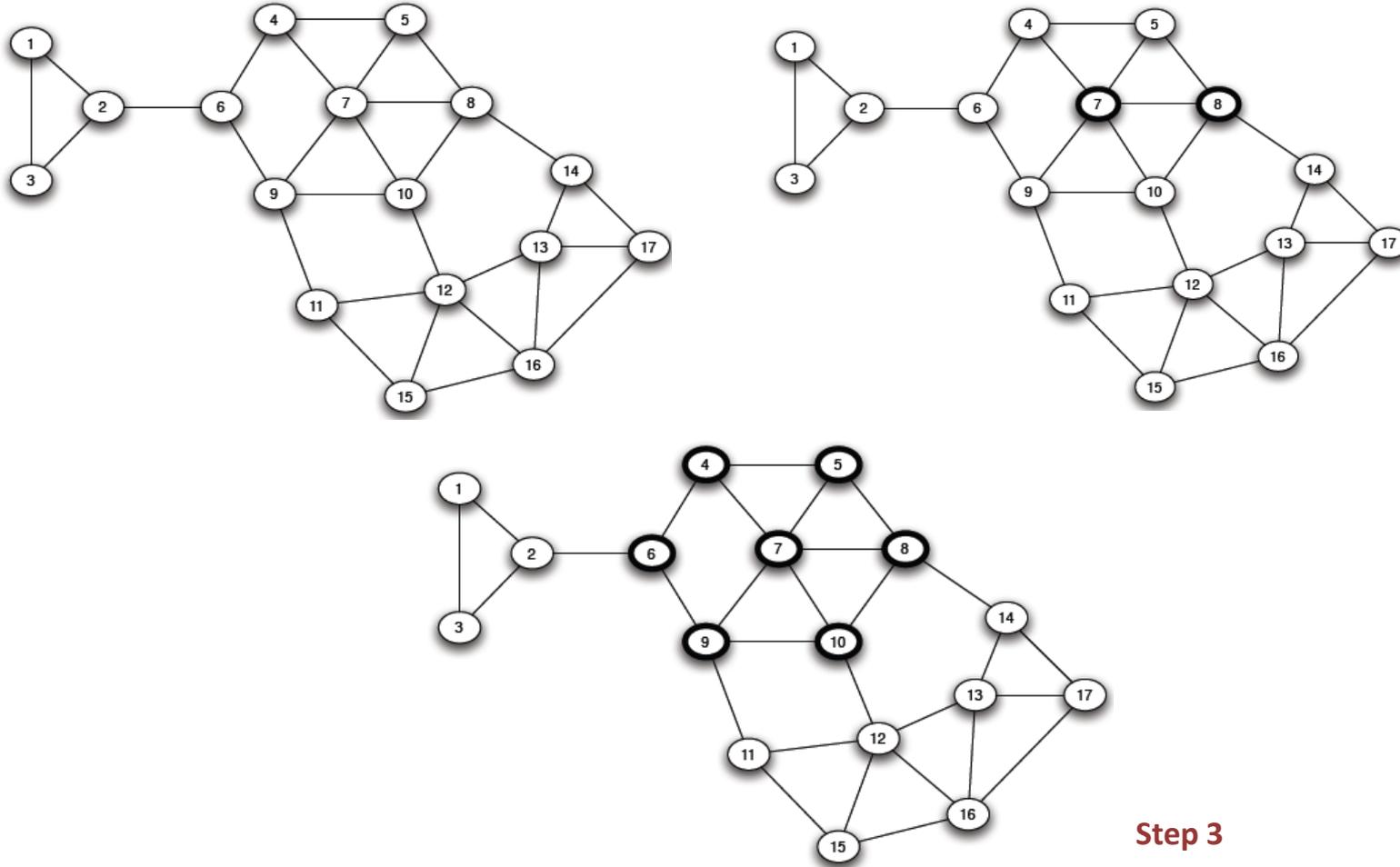


Step 2

Chain reaction of switches to B  $\rightarrow$  A **cascade** of adoptions of A

# Modeling Diffusion through a Network: Cascading Behavior

$a = 3, b = 2, q = 2/5$



# Modeling Diffusion through a Network: Cascading Behavior

1. A set of initial adopters who start with a new behavior  $A$ , while every other node starts with behavior  $B$ .
2. Nodes repeatedly evaluate the decision to switch from  $B$  to  $A$  using a threshold of  $q$ .
3. If the resulting cascade of adoptions of  $A$  eventually causes every node to switch from  $B$  to  $A$ , then we say that the set of initial adopters causes a complete cascade at threshold  $q$ .

# Modeling Diffusion through a Network: Cascading Behavior and “Viral Marketing”

Tightly-knit communities in the network can work to hinder the spread of an innovation

(examples, age groups and life-styles in social networking sites, Mac users, political opinions)

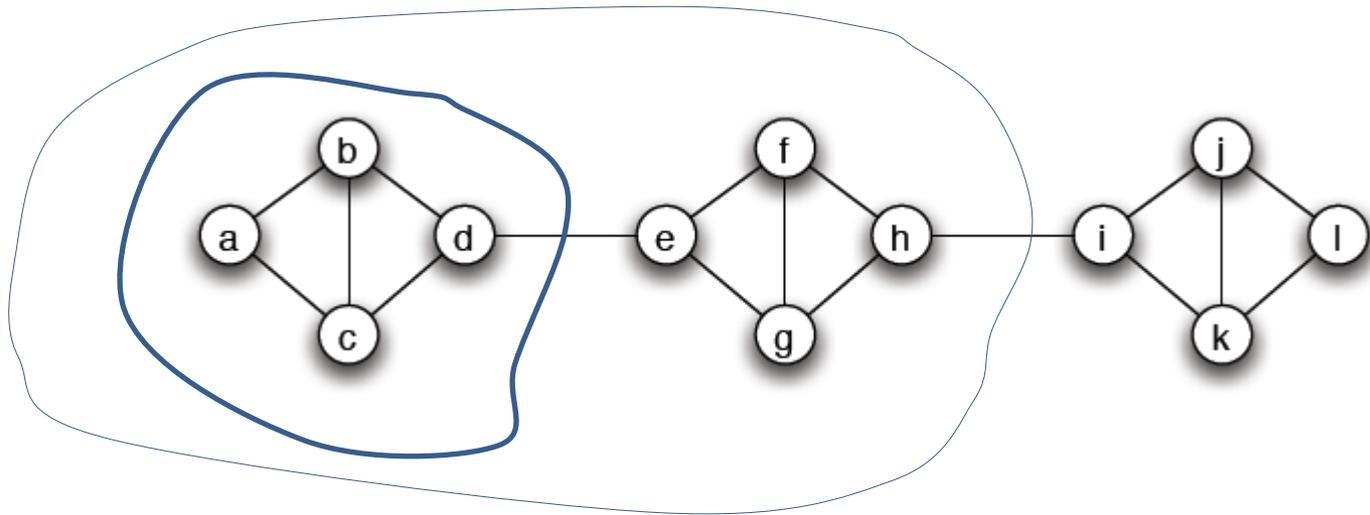
## Strategies

- Improve the quality of A (increase the payoff  $a$ ) (in the example, set  $a = 4$ )
- Convince a small number of *key people* to switch to A

Network-level cascade innovation adoption models vs population-level

# Cascades and Clusters

A **cluster of density  $p$**  is a set of nodes such that each node in the set has *at least* a  $p$  fraction of its neighbors in the set

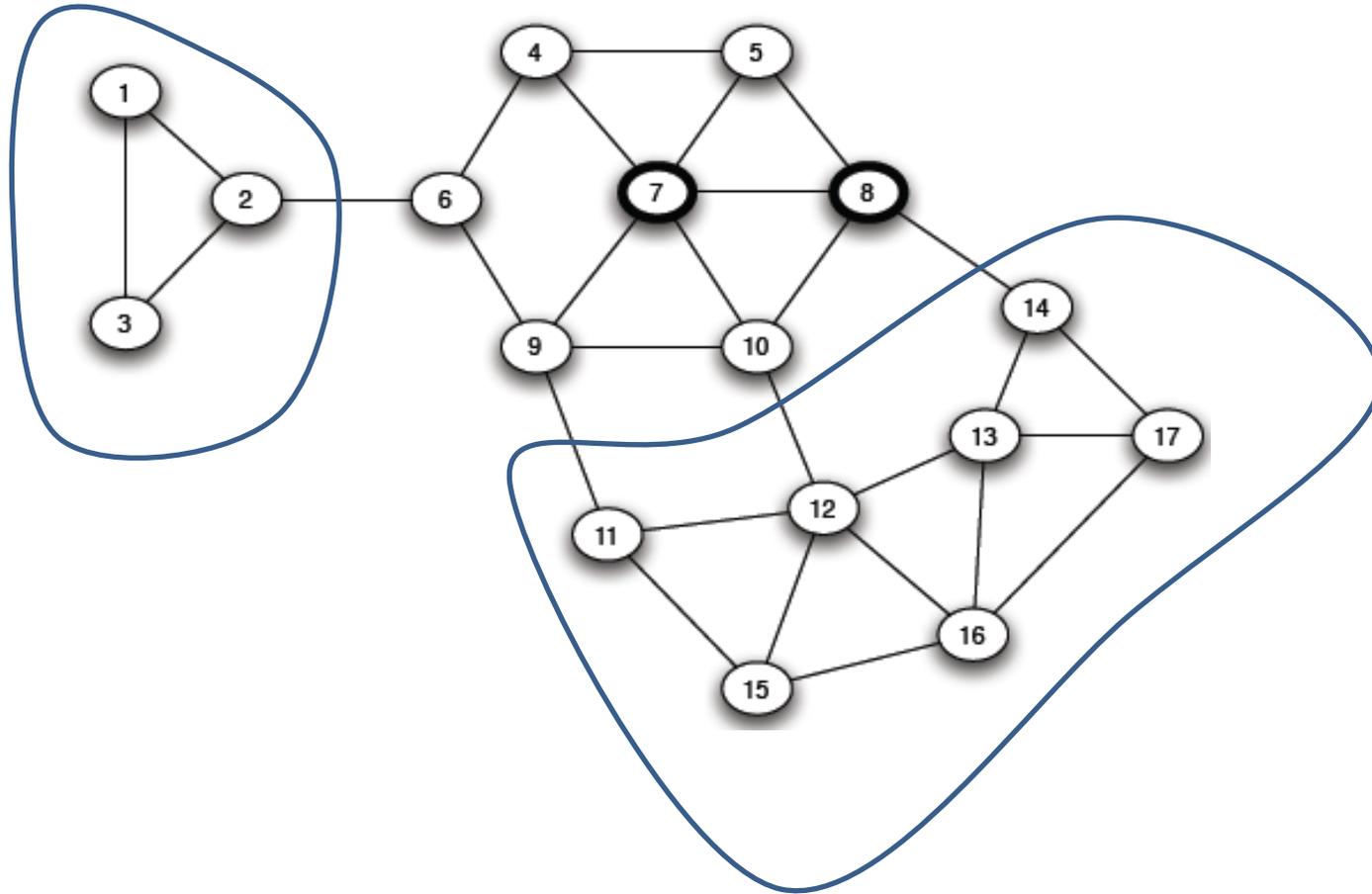


However,

Does not imply that any two nodes in the same cluster necessarily have much in common (what is the density of a cluster with all nodes?)

The union of any two cluster of density  $p$  is also a cluster of density at least  $p$

# Cascades and Clusters



# Cascades and Clusters

**Claim:** Consider a set of initial adopters of behavior  $A$ , with a threshold of  $q$  for nodes in the remaining network to adopt behavior  $A$ .

## (i) (clusters as obstacles to cascades)

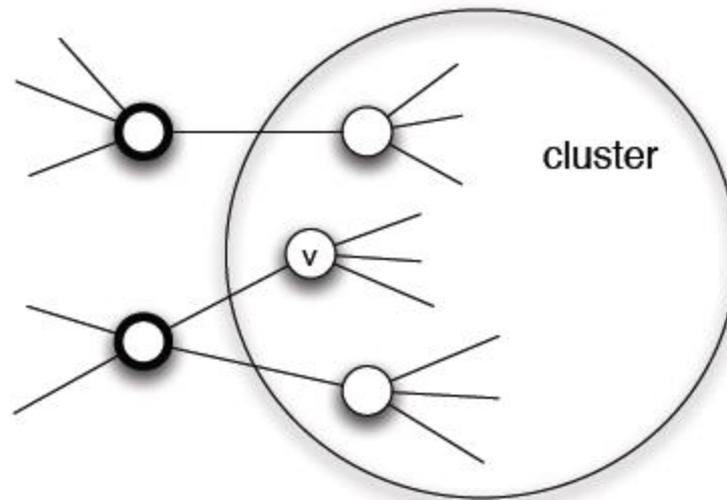
If the remaining network contains a cluster of density greater than  $1 - q$ , then the set of initial adopters will not cause a complete cascade.

## (ii) (clusters are the only obstacles to cascades)

Whenever a set of initial adopters does not cause a complete cascade with threshold  $q$ , the remaining network must contain a cluster of density greater than  $1 - q$ .

# Cascades and Clusters

**Proof of (i)** (clusters as obstacles to cascades)

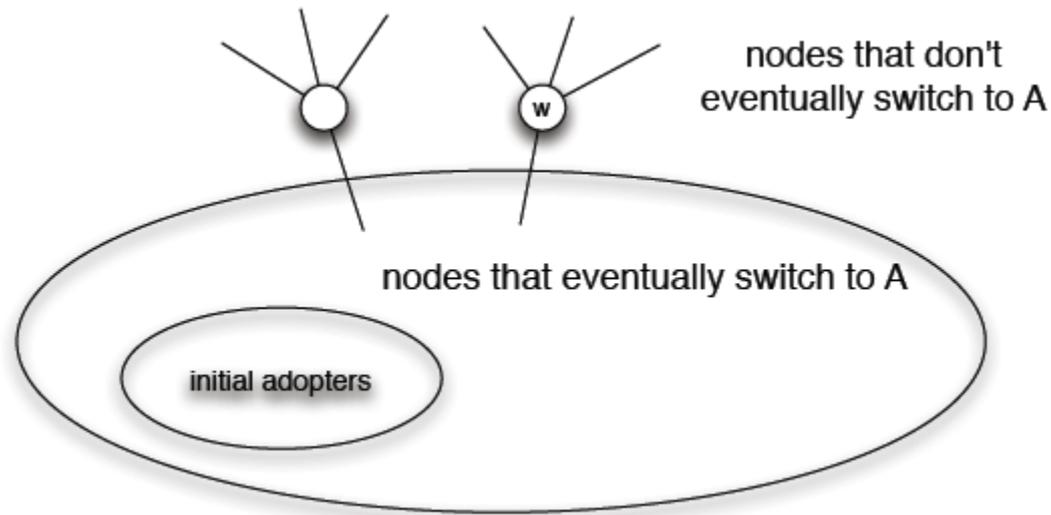


*Proof by contradiction*

Let  $v$  be the first node in the cluster that adopts  $A$

# Cascades and Clusters

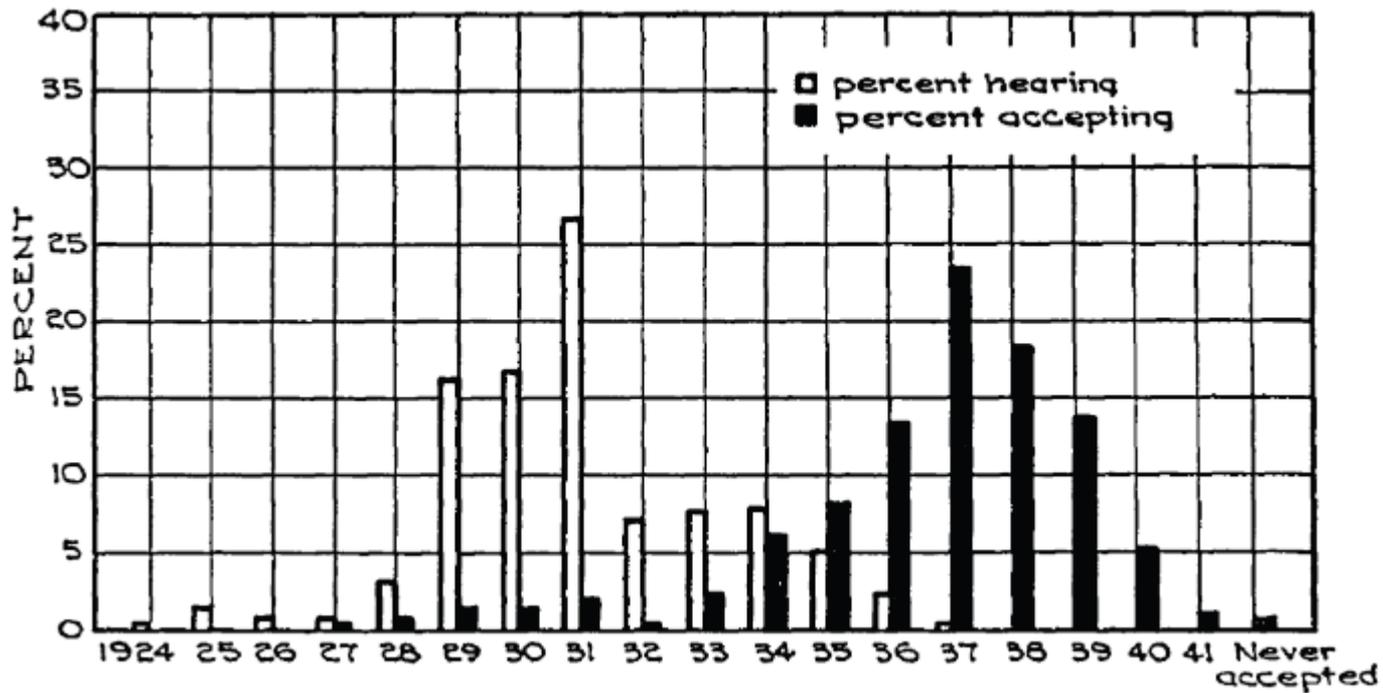
**Proof of (ii) (clusters are the only obstacles to cascades)**



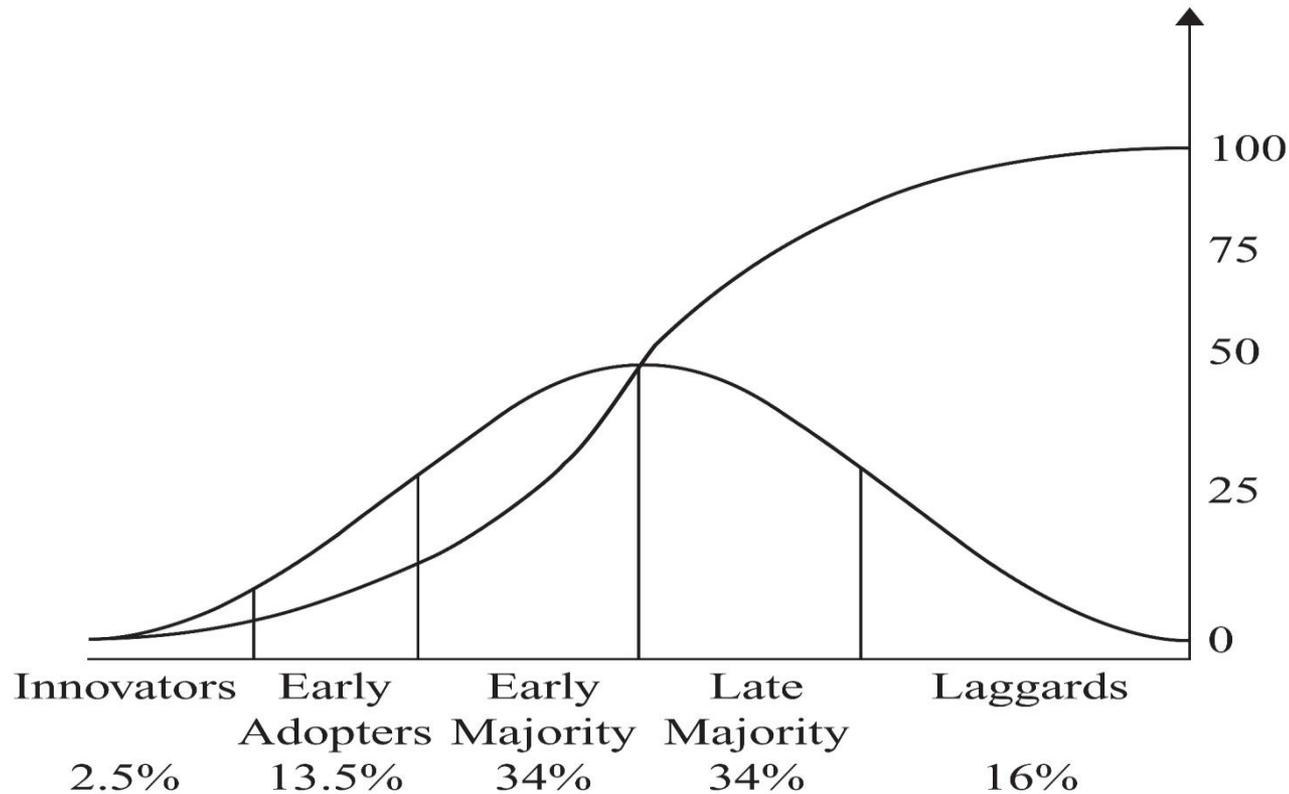
Let  $S$  be the set of all nodes using B at the end of the process  
Show that  $S$  is a cluster of density  $> 1 - q$

# Innovation Adoption Characteristics

A crucial difference between learning a new idea and actually deciding to accept it



# Innovation Adoption Characteristics

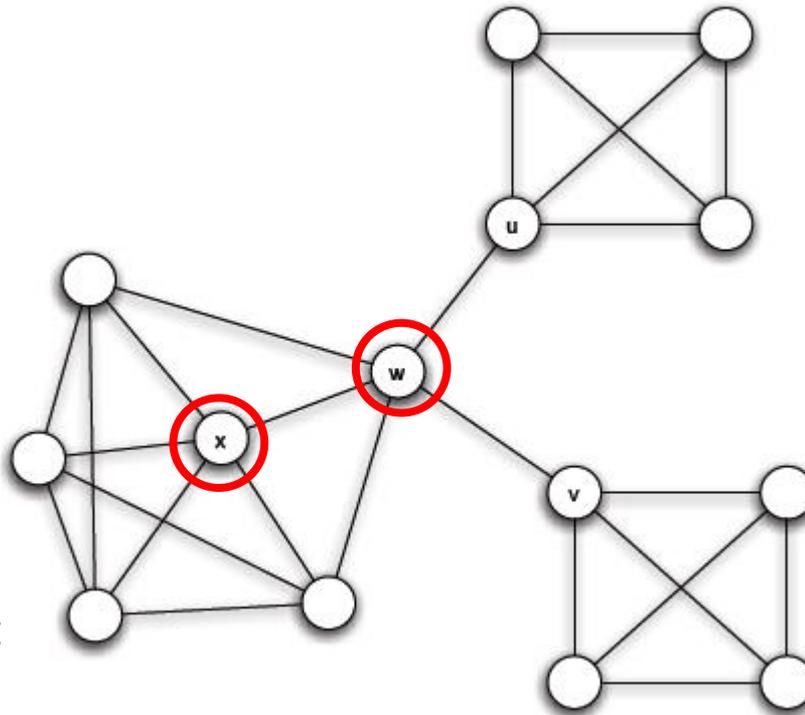


Category of Adopters in the corn study

# Diffusion, Thresholds and the Role of Weak Ties

Relation to weak ties and local bridges

$$q = 1/2$$



Bridges convey awareness  
but are weak at  
transmitting costly to  
adopt behaviors

# Extensions of the Basic Cascade Model: Heterogeneous Thresholds

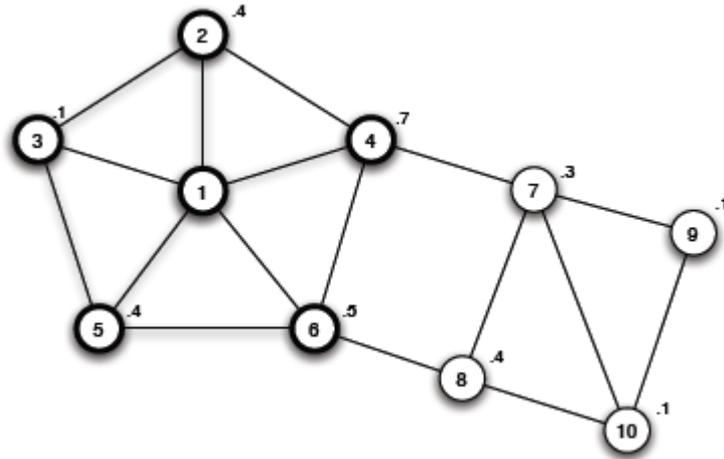
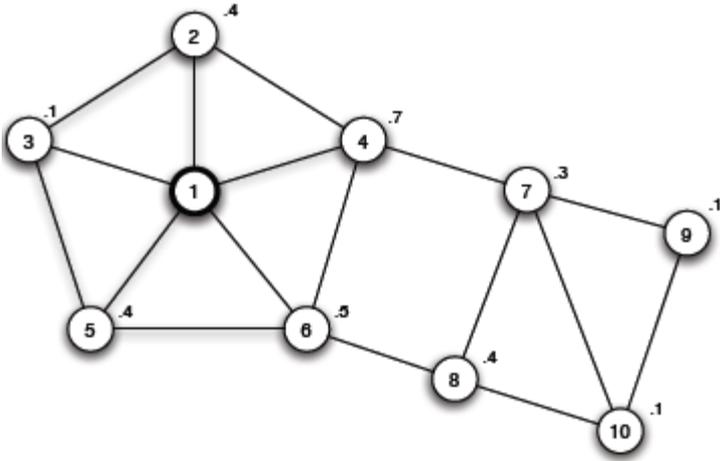
Each person values behaviors A and B differently:

- If both  $u$  and  $w$  adapt A,  $u$  gets a payoff  $a_u > 0$  and  $w$  a payoff  $a_w > 0$
- If both  $u$  and  $w$  adapt B,  $u$  gets a payoff  $b_u > 0$  and  $w$  a payoff  $b_w > 0$
- If opposite behaviors, then each gets a payoff 0

		$w$	
		$A$	$B$
$v$	$A$	$a_u, a_w$	$0, 0$
	$B$	$0, 0$	$b_u, b_w$

Each node  $u$  has its own personal threshold  $q_u \geq b_u / (a_u + b_u)$

# Extensions of the Basic Cascade Model: Heterogeneous Thresholds



- ✓ Not just the power of influential people, but also the extent to which they have access to **easily influenceable people**
- ✓ What about the role of clusters?  
A **blocking cluster** in the network is a set of nodes for which each node  $u$  has more than  $1 - q_u$  fraction of its friends also in the set.

# Knowledge, Thresholds and Collective Action:

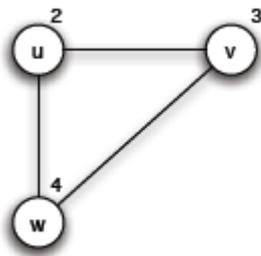
## Collective Action and Pluralistic Ignorance

A *collective action problem*: an activity produces benefits only if enough people participate (population level effect)

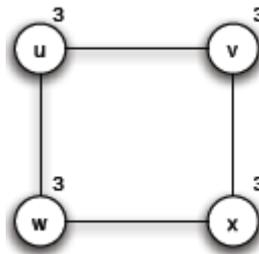
*Pluralistic ignorance*: a situation in which people have wildly erroneous estimates about the prevalence of certain opinions in the population at large (lack of knowledge)

# Knowledge, Thresholds and Collective Action: A model for the effect of knowledge on collective actions

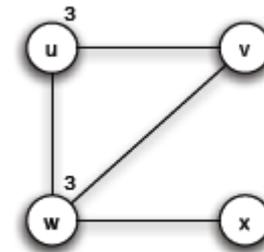
- Each person has a personal threshold which encodes her willingness to participate
- A **threshold of  $k$**  means that she will participate if at least  $k$  people in total (including herself) will participate
- Each person in the network **knows the thresholds of her neighbors** in the network



- w will never join, since there are only 3 people
- v
- u



- Is it safe for u to join?



- Is it safe for u to join?  
(common knowledge)

# Knowledge, Thresholds and Collective Action: Common Knowledge and Social Institutions

- Not just transmit a message, but also make the listeners or readers *aware that many others have gotten the message* as well
- Social networks do not simply allow for interaction and flow of information, but these processes in turn allow individuals to base decisions *on what other knows* and *on how they expect others to behave as a result*

# The Cascade Capacity

Given a network, what is the *largest threshold* at which *any* “*small*” set of initial adopters can cause a *complete cascade*?

Called *cascade capacity* of the network

- Infinite network in which each node has a finite number of neighbors
- Small means finite set of nodes

# The Cascade Capacity: Cascades on Infinite Networks

Same model as before:

- Initially, *a finite set*  $S$  of nodes has behavior A and all others adopt B
- Time runs forwards in steps,  $t = 1, 2, 3, \dots$
- In each step  $t$ , each node other than those in  $S$  uses the decision rule with *threshold*  $q$  to decide whether to adopt behavior A or B
- The set  $S$  causes *a complete cascade* if, starting from  $S$  as the early adopters of A, every node in the network eventually switched permanently to A.

The *cascade capacity* of the network is *the largest value of the threshold*  $q$  for which some finite set of early adopters can cause *a complete cascade*.

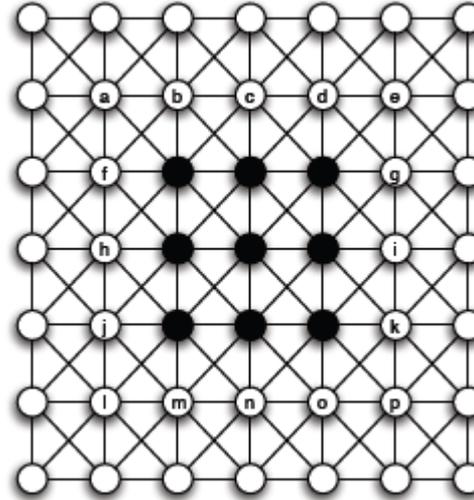
# The Cascade Capacity: Cascades on Infinite Networks

An infinite path



*Spreads if  $\leq 1/2$*

An infinite grid



*Spreads if  $\leq 3/8$*

- ✓ An intrinsic property of the network
- ✓ Even if A better, for  $q$  strictly between  $3/8$  and  $1/2$ , A cannot win

# The Cascade Capacity: Cascades on Infinite Networks

How large can a cascade capacity be?

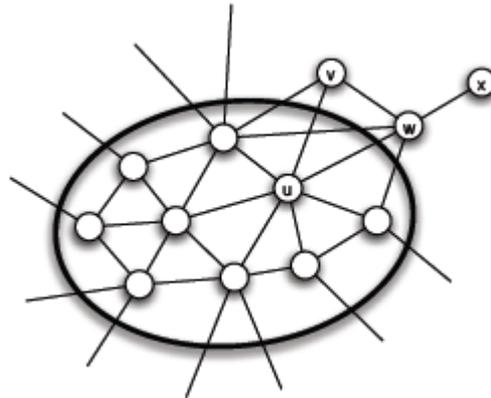
- At least  $1/2$
- *Is there any network with a higher cascade capacity?*
- This will mean that *an inferior technology* can displace a superior one, even when the inferior technology starts at only a small set of initial adopters.

# The Cascade Capacity: Cascades on Infinite Networks

**Claim:** There is no network in which the cascade capacity exceeds  $1/2$

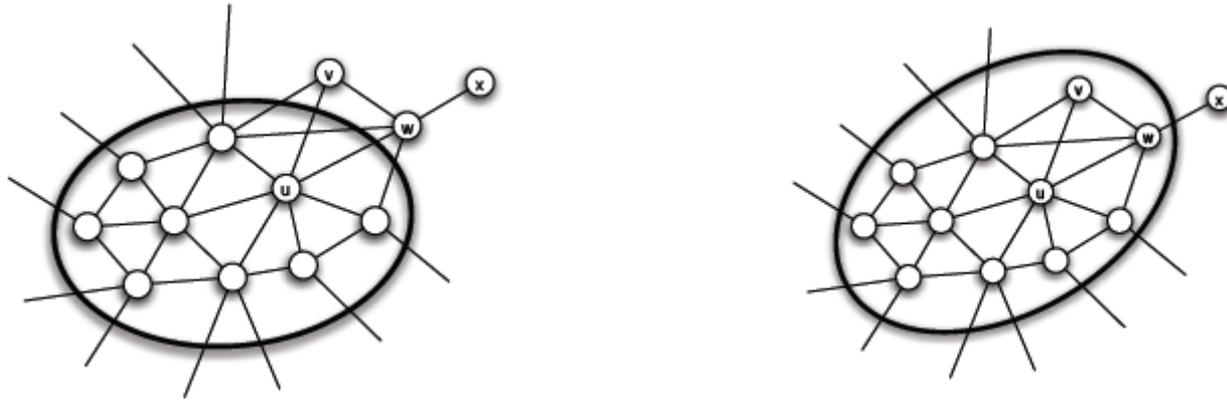
# The Cascade Capacity: Cascades on Infinite Networks

Interface: the set of A-B edges



Prove that in each step the size of the interface strictly decreases  
Why is this enough?

# The Cascade Capacity: Cascades on Infinite Networks



At some step, a number of nodes decide to switch from B to A

*General Remark: In this simple model, a worse technology cannot displace a better and wide-spread one*

# Compatibility and its Role in Cascades

An extension where a single individual can sometimes choose a combination of two available behaviors -> three strategies A, B and AB

## Coordination game with a bilingual option

- Two bilingual nodes can interact using the better of the two behaviors
- A bilingual and a monolingual node can only interact using the behavior of the monolingual node

		<i>w</i>	
		<i>A</i>	<i>B</i>
		<i>AB</i>	
<i>v</i>	<i>A</i>	<i>a, a</i>	<i>0, 0</i>
	<i>B</i>	<i>0, 0</i>	<i>b, b</i>
	<i>AB</i>	<i>a, a</i>	<i>b, b</i>
			<i>(a, b)<sup>+</sup>, (a, b)<sup>+</sup></i>

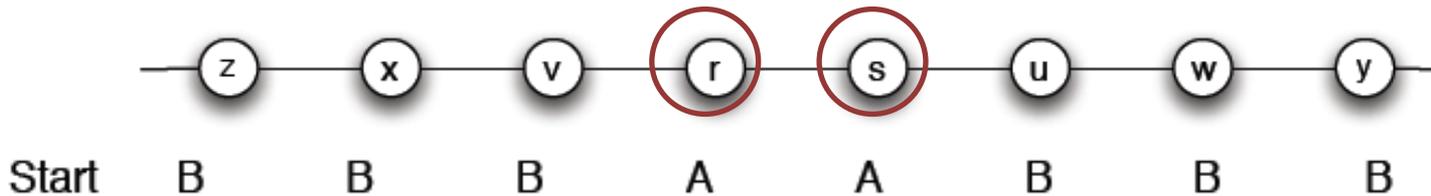
AB is a dominant strategy?

- ✓ **Cost *c*** associated with the AB strategy

# Compatibility and its Role in Cascades

Example ( $a = 2, b = 3, c = 1$ )

	$A$	$B$	$AB$
$A$	$a, a$	$0, 0$	$a, a$
$B$	$0, 0$	$b, b$	$b, b$
$AB$	$a, a$	$b, b$	$(a, b)^+, (a, b)^+$



B:  $0 + b = 3$   
 A:  $0 + a = 2$   
 AB:  $b + a - c = 4 \checkmark$

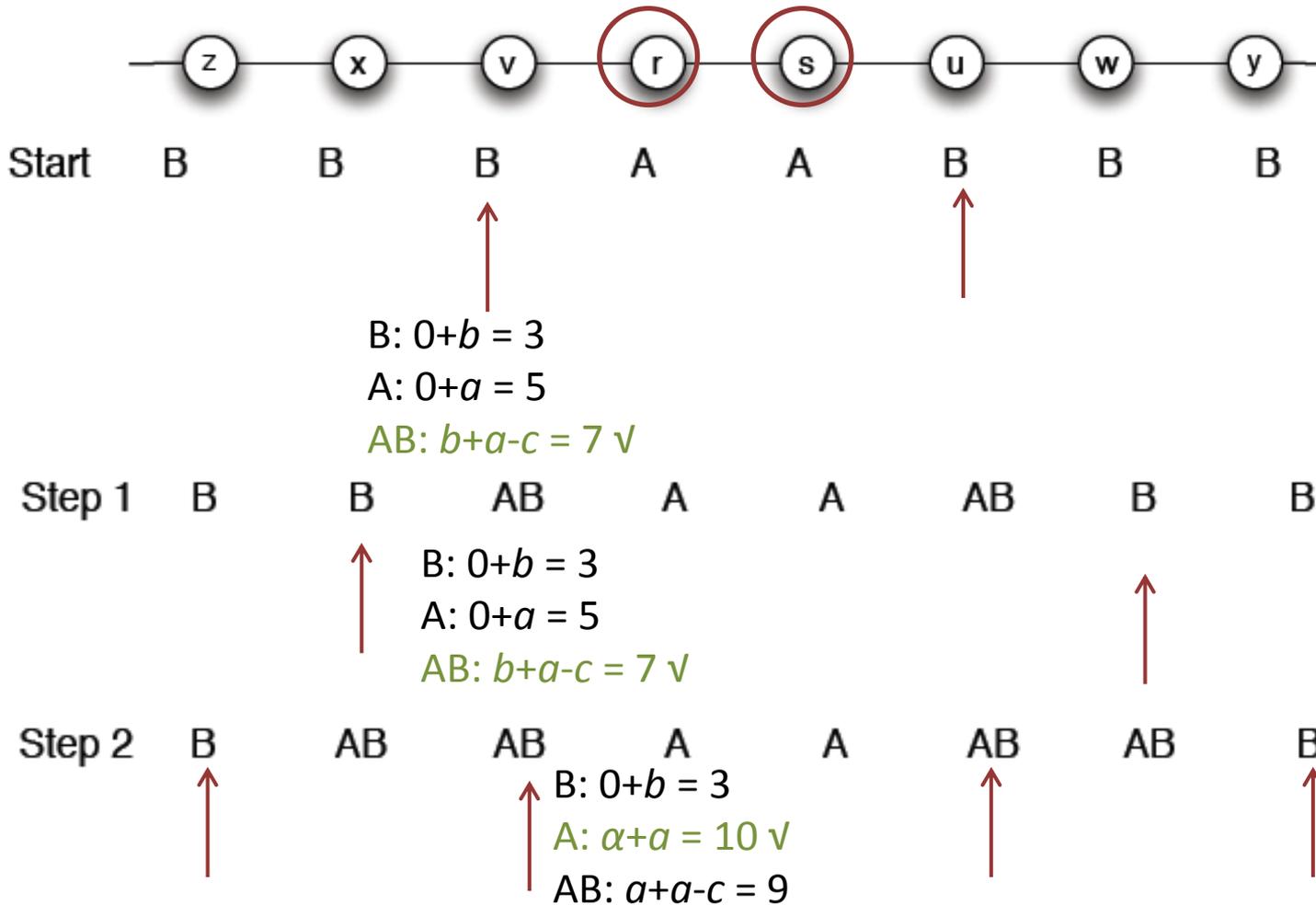


B:  $b + b = 6 \checkmark$   
 A:  $0 + a = 2$   
 AB:  $b + b - c = 5$

# Compatibility and its Role in Cascades

Example ( $a = 5, b = 3, c = 1$ )

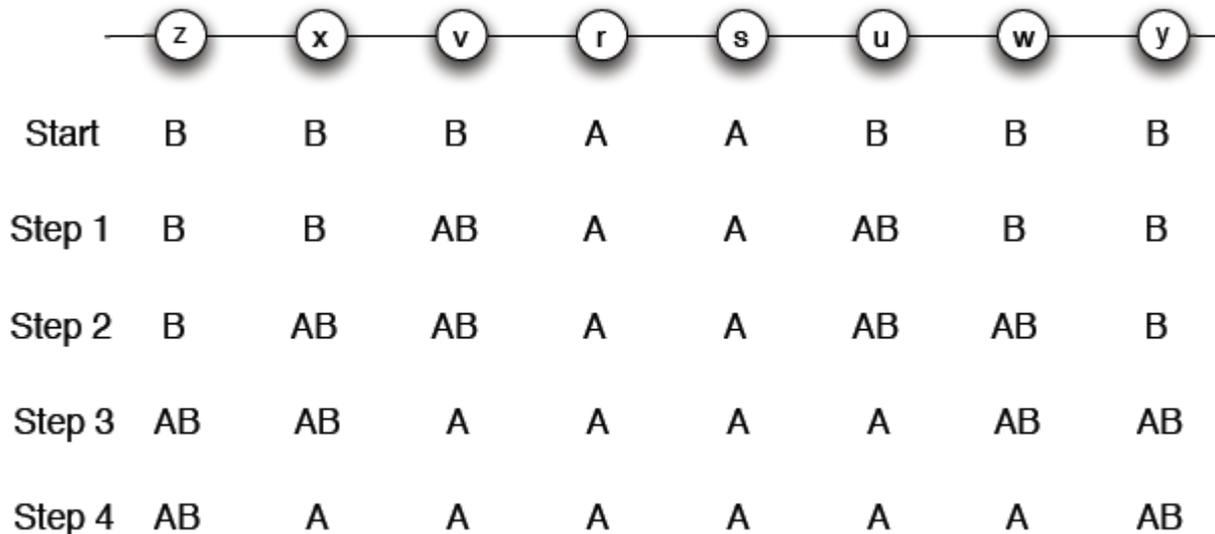
	$A$	$B$	$AB$
$A$	$a, a$	$0, 0$	$a, a$
$B$	$0, 0$	$b, b$	$b, b$
$AB$	$a, a$	$b, b$	$(a, b)^+, (a, b)^+$



# Compatibility and its Role in Cascades

Example ( $a = 5, b = 3, c = 1$ )

		$w$		
		$A$	$B$	$AB$
$v$	$A$	$a, a$	$0, 0$	$a, a$
	$B$	$0, 0$	$b, b$	$b, b$
	$AB$	$a, a$	$b, b$	$(a, b)^+, (a, b)^+$



✓ First, strategy AB spreads, then behind it, nodes switch permanently from AB to A

Strategy B becomes *vestigial*

# Compatibility and its Role in Cascades

Given an infinite graph, for which payoff values of  $a$ ,  $b$  and  $c$ , is it possible for a finite set of nodes to cause a complete cascade of adoptions of  $A$ ?

Fixing  $b = 1$  (default technology)

Given an infinite graph, for which payoff values of  $a$  (how much better the new behavior  $A$ ) and  $c$  (how compatible should it be with  $B$ ), is it possible for a finite set of nodes to cause a complete cascade of adoptions of  $A$ ?

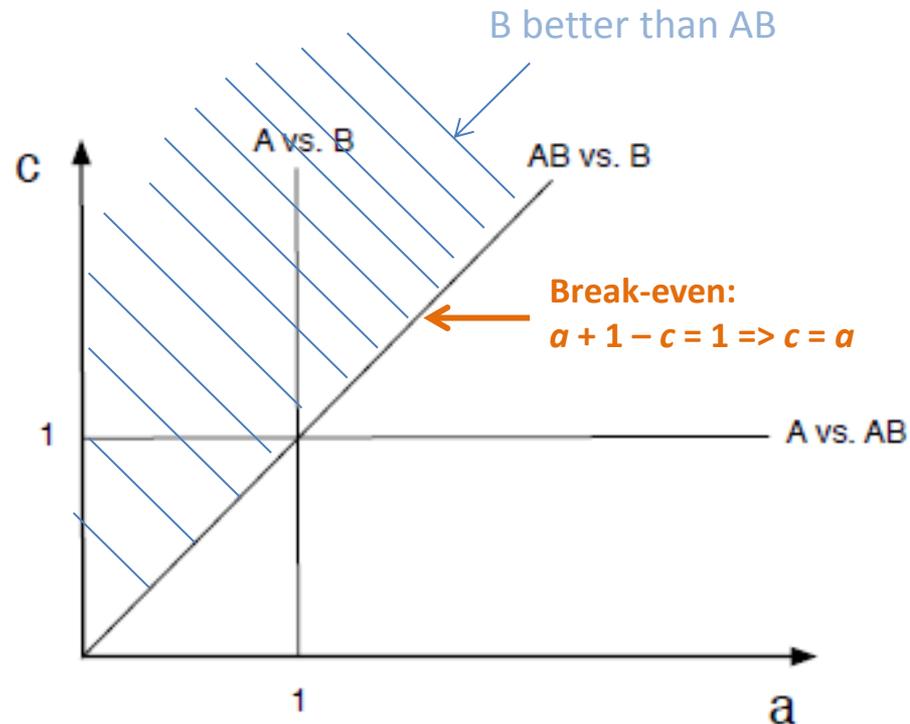
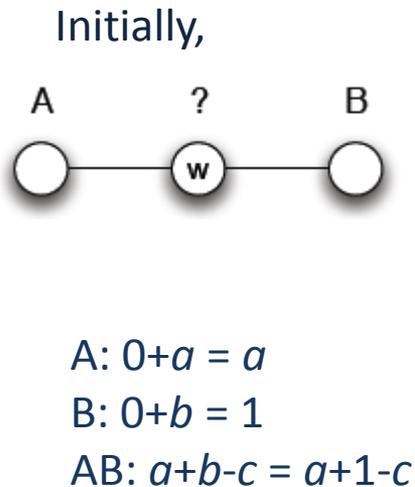
$A$  does better when it has a higher payoff, but in general it has a particularly hard time cascading when the level of compatibility is “intermediate” – when the value of  $c$  is neither too high nor too low

# Compatibility and its Role in Cascades

## Example: Infinite path

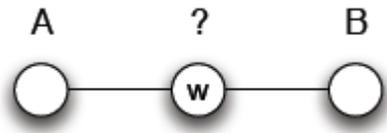
- Spreads when  $q \leq 1/2$ ,  $a \geq b$  (a better technology always spreads)

Assume that the set of initial adopters forms a contiguous interval of nodes on the path  
Because of the symmetry, how strategy changes occur to the right of the initial adopters



# Compatibility and its Role in Cascades

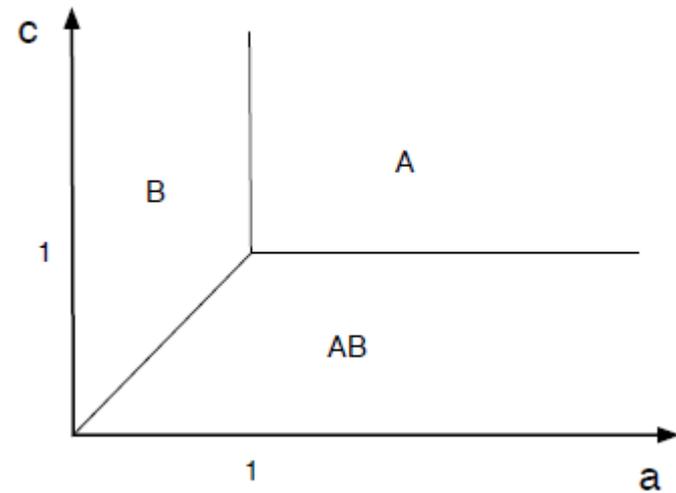
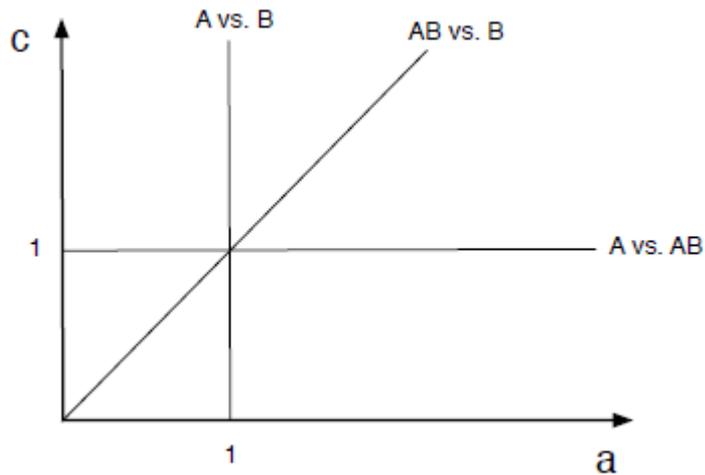
Initially,



$$A: 0+a = a$$

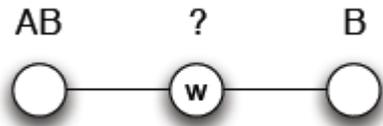
$$B: 0+b = 1$$

$$AB: a+b-c = a+1-c$$



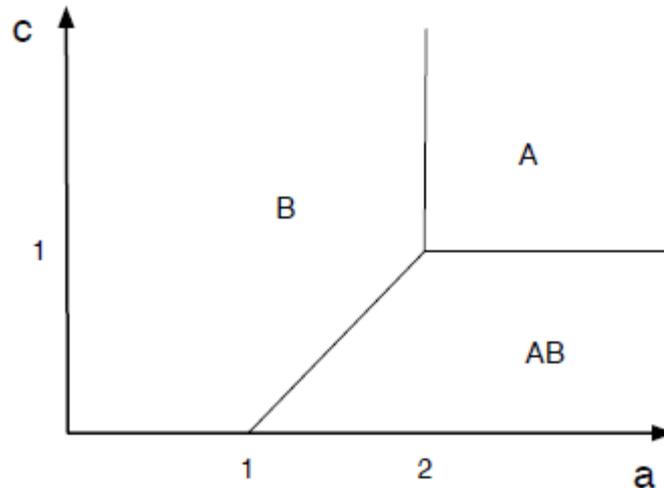
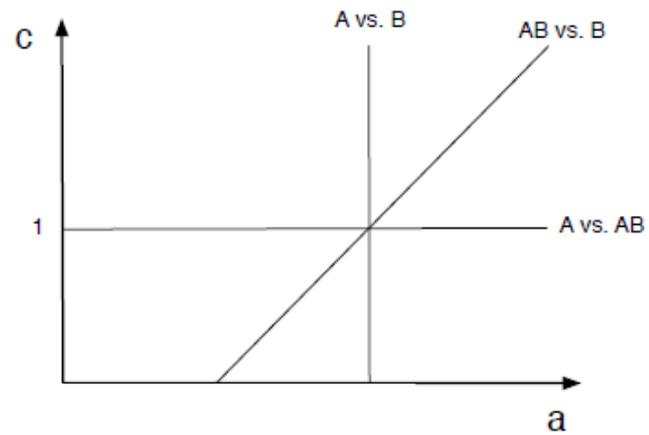
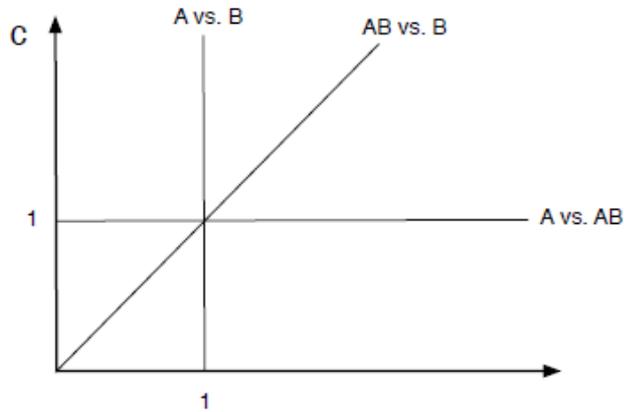
# Compatibility and its Role in Cascades

Then,

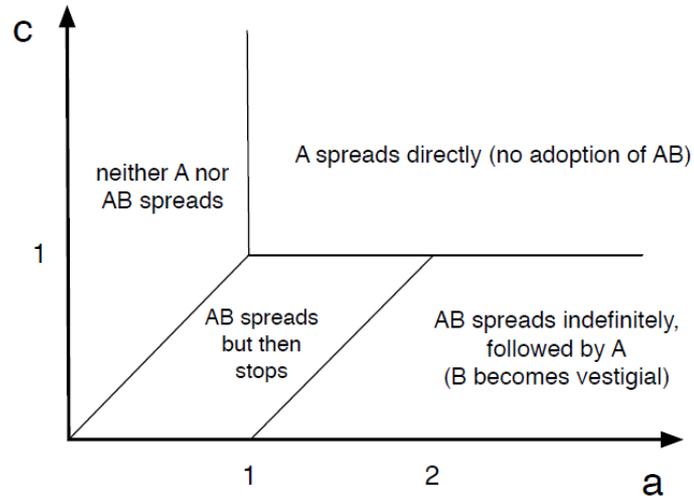
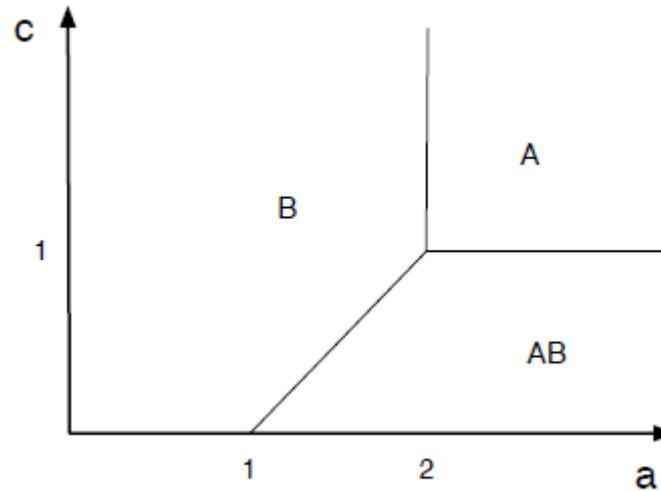
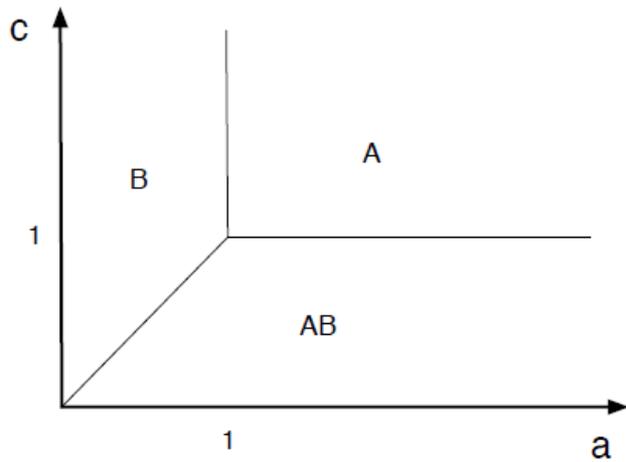


$a < 1$ ,  
 A:  $0+a = a$   
 B:  $b+b = 2 \checkmark$   
 AB:  $b+b-c = 2-c$

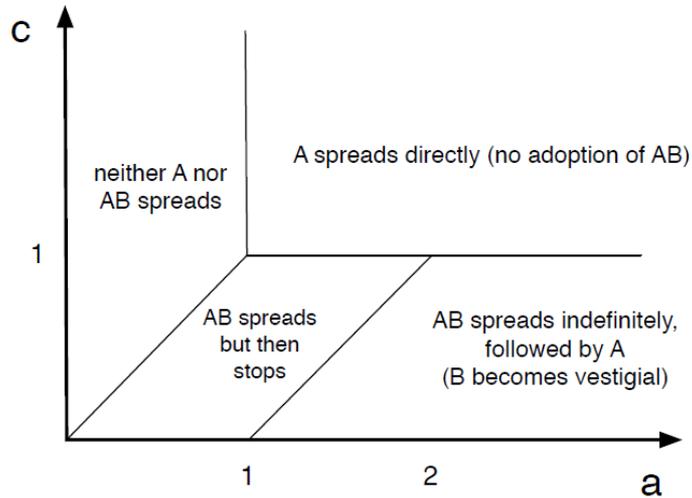
$a \geq 1$   
 A:  $a$   
 B:  $2$   
 AB:  $a+1-c$



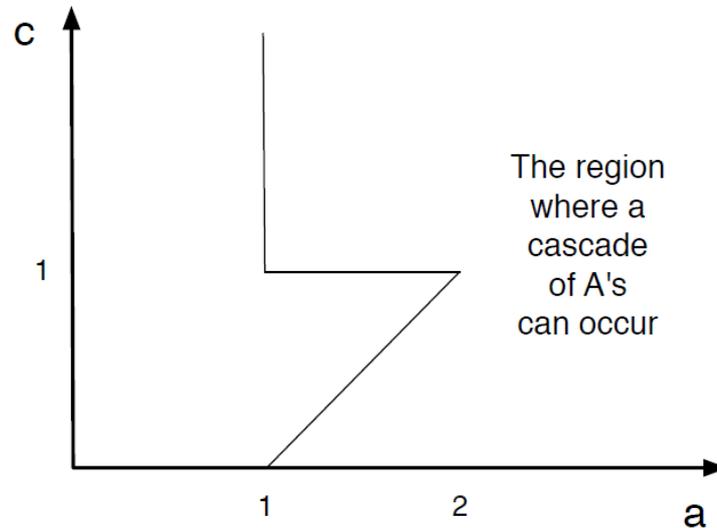
# Compatibility and its Role in Cascades



# Compatibility and its Role in Cascades



What does the triangular cut-out mean?



# Reference

Networks, Crowds, and Markets (Chapter 19)

# **EPIDEMIC SPREAD**

# Epidemics

Understanding the spread of viruses and epidemics is of great interest to

- Health officials
- Sociologists
- Mathematicians
- Hollywood

The underlying **contact network** clearly affects the spread of an epidemic

Model epidemic spread as a **random process** on the graph and study its properties

- Main question: will the epidemic take over most of the network?

**Diffusion of ideas** and the **spread of influence** can also be modeled as epidemics

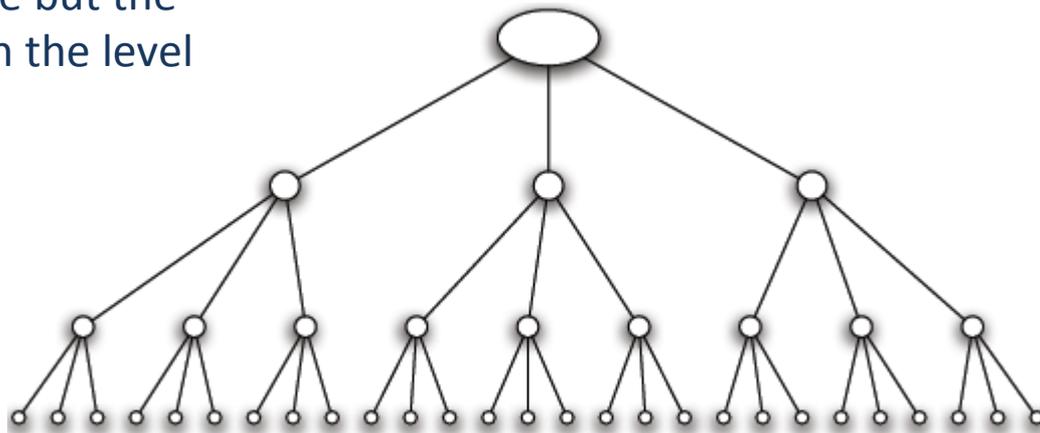


# Branching Processes

- A person transmits the disease to each people she meets *independently with a probability  $p$*
  - Meets  *$k$  people* while she is contagious
1. A person carrying a new disease enters a population, first *wave* of  $k$  people
  2. Second wave of  $k^2$  people
  3. Subsequent waves

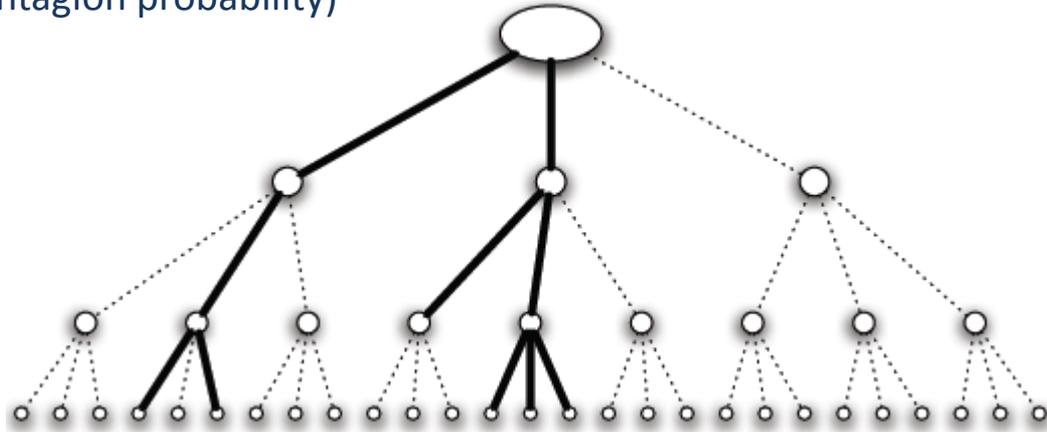
A contact network with  $k = 3$

Tree (root, each node but the root, a single node in the level above it)



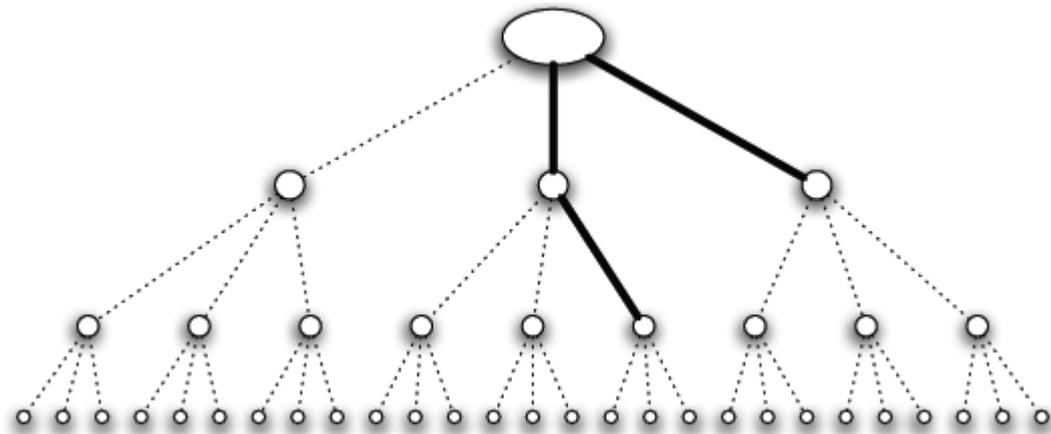
# Branching Processes

Aggressive epidemic (high contagion probability)



Mild epidemic (low contagion probability)

- If it ever reaches a wave where it infects no one, then it dies out
- Or, it continues to infect people in every wave infinitely



# Branching Processes: Basic Reproductive Number

**Basic Reproductive Number** ( $R_0$ ): the expected number of new cases of the disease caused by a single individual

Claim: (a) If  $R_0 < 1$ , then with probability 1, the disease dies out after a finite number of waves. (b) If  $R_0 > 1$ , then with probability greater than 0 the disease persists by infecting at least one person in each wave.

$$R_0 = pk$$

- (a)  $R_0 < 1$  -- Each infected person produces less than one new case in expectation  
Outbreak constantly trends downwards
- (b)  $R_0 > 1$  -- trends upwards, and the disease persists with positive probability  
(when  $p < 1$ , the disease can get unlucky!)

A “knife-edge” quality around the critical value of  $R_0 = 1$

# Branching process

- Assumes no network structure, no triangles or shared neighbors

# The SIR model

- Each node may be in the following states
  - **Susceptible**: healthy but not immune
  - **Infected**: has the virus and can actively propagate it
  - **Removed**: (Immune or Dead) had the virus but it is no longer active
- probability of an Infected node to infect a Susceptible neighbor

# The SIR process

- Initially all nodes are in state S(usceptible), except for a few nodes in state I(nfected).
- An infected node stays infected for  $t_I$  steps.
  - Simplest case:  $t_I = 1$
- At each of the  $t_I$  steps the infected node has probability  $p$  of infecting any of its susceptible neighbors
  - $p$ : Infection probability
- After  $t_I$  steps the node is Removed

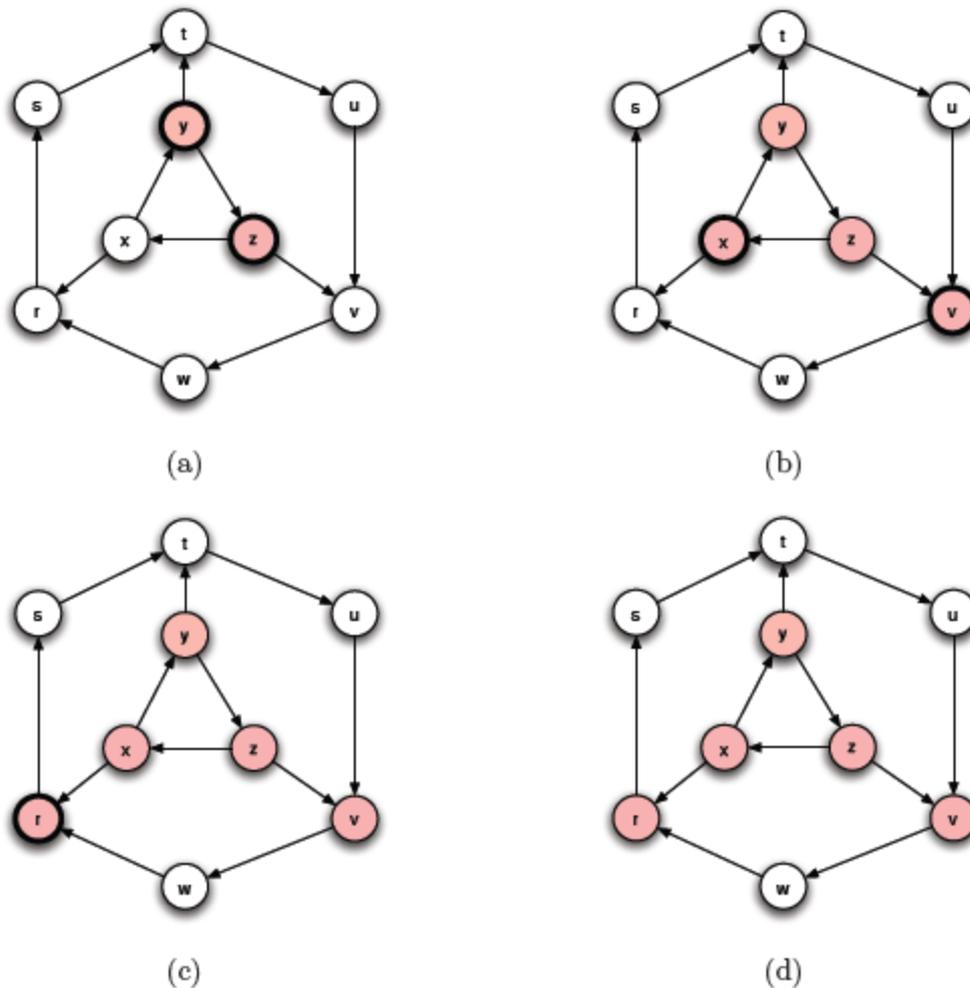


Figure 21.2: The course of an SIR epidemic in which each node remains infectious for a number of steps equal to  $t_I = 1$ . Starting with nodes  $y$  and  $z$  initially infected, the epidemic spreads to some but not all of the remaining nodes. In each step, shaded nodes with dark borders are in the Infectious ( $I$ ) state and shaded nodes with thin borders are in the Removed ( $R$ ) state.

# SIR and the Branching process

- The branching process is a special case where the graph is a tree (and the infected node is the root)
- The basic reproductive number is not necessarily informative in the general case

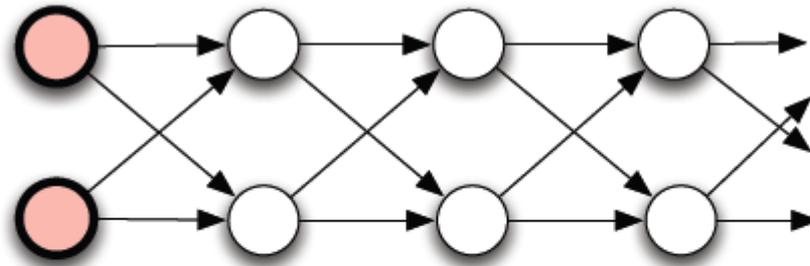


Figure 21.3: In this network, the epidemic is forced to pass through a narrow “channel” of nodes. In such a structure, even a highly contagious disease will tend to die out relatively quickly.

# Percolation

- **Percolation**: we have a network of “pipes” which can carry liquids, and they can be either **open** with probability  $p$ , or **close** with probability  $(1-p)$ 
  - The pipes can be pathways within a material
- If liquid enters the network from some nodes, does it **reach** most of the network?
  - The network **percolates**

# SIR and Percolation

- There is a connection between SIR model and percolation
- When a virus is transmitted from  $u$  to  $v$ , the edge  $(u,v)$  is activated with probability  $p$
- We can assume that all edge activations have happened **in advance**, and the input graph has **only** the **active edges**.
- Which nodes will be infected?
  - The nodes **reachable** from the initial infected nodes
- In this way we transformed the **dynamic SIR process** into a **static** one.

# Example

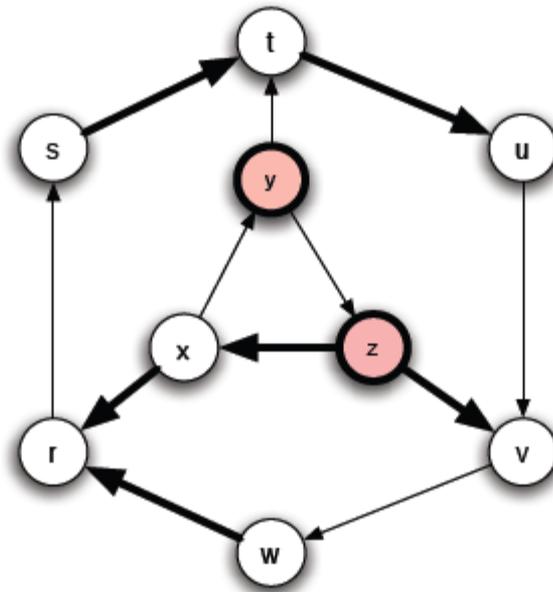


Figure 21.4: An equivalent way to view an SIR epidemic is in terms of *percolation*, where we decide in advance which edges will transmit infection (should the opportunity arise) and which will not.

# The SIS model

- **Susceptible-Infected-Susceptible**
  - Susceptible: healthy but not immune
  - Infected: has the virus and can actively propagate it
- An **Infected** node infects a **Susceptible** neighbor with probability  $p$
- An **Infected** node becomes **Susceptible** again with probability  $q$  (or after  $t_I$  steps)
  - In a **simplified** version of the model  $q = 1$
- Nodes **alternate** between **Susceptible** and **Infected** status

# Example

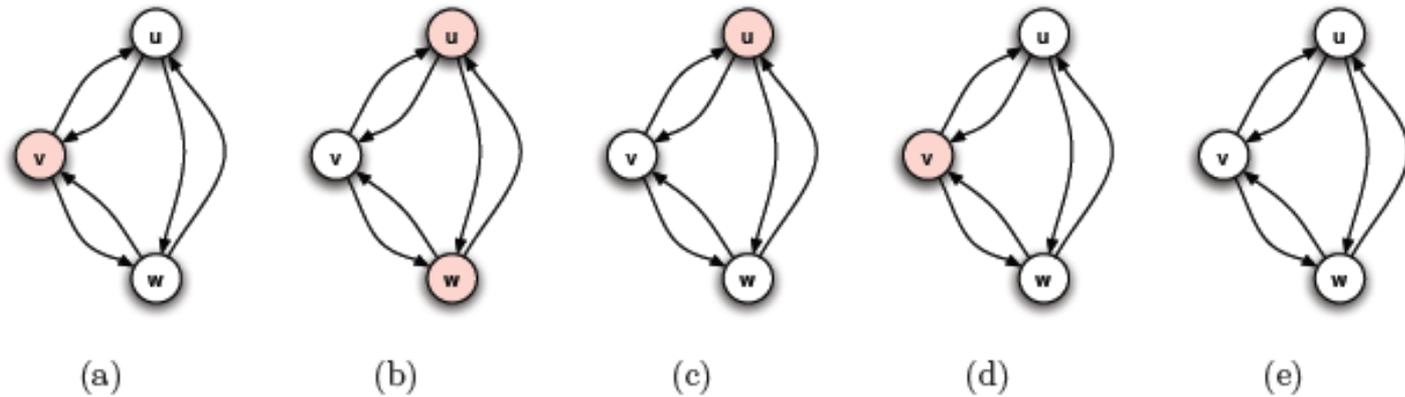


Figure 21.5: In an SIS epidemic, nodes can be infected, recover, and then be infected again. In each step, the nodes in the Infectious state are shaded.

- When no **Infected** nodes, virus dies out
- Question: will the virus die out?

# An eigenvalue point of view

- If  $A$  is the adjacency matrix of the network, then the virus dies out if

$$\lambda_1(A) \leq \frac{q}{p}$$

- Where  $\lambda_1$  is the first eigenvalue of  $A$

# Multiple copies model

- Each node may have **multiple copies** of the same virus
  - $\mathbf{v}$ : state vector :  $v_i$  : number of virus copies at node  $i$
- At time  $t = 0$ , the state vector is initialized to  $\mathbf{v}^0$
- At time  $t$ ,
  - For each node  $i$ 
    - For each of the  $v_i^t$  virus copies at node  $i$ 
      - the copy is copied to a neighbor  $j$  with prob  $p$
      - the copy dies with probability  $q$

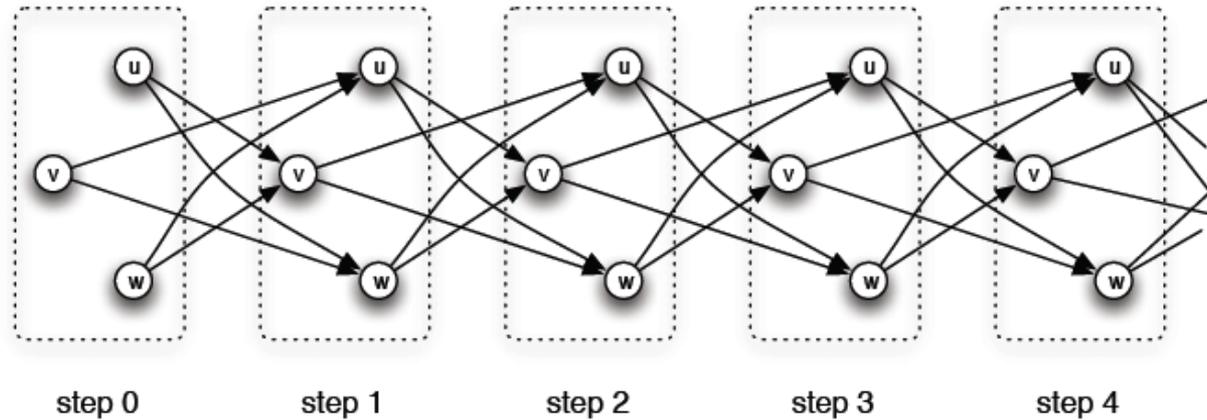
# Analysis

- The expected state of the system at time  $t$  is given by

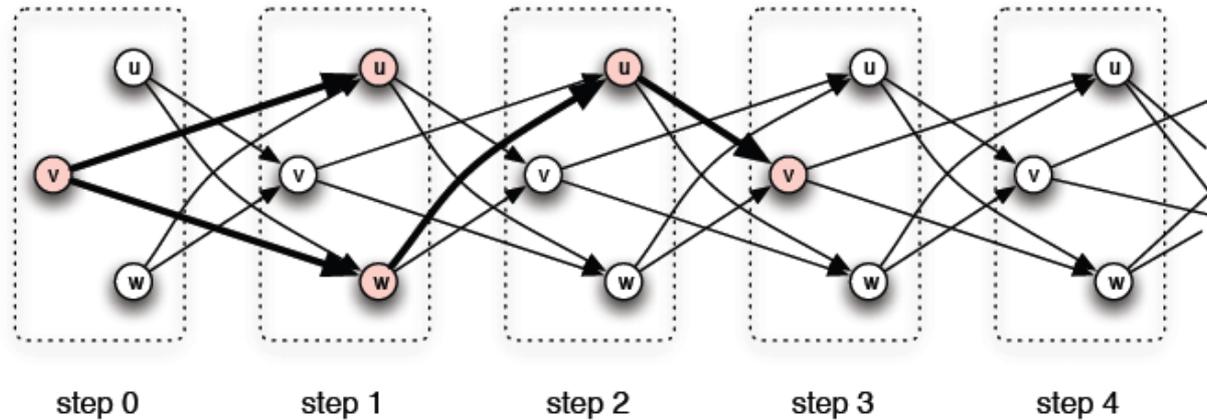
$$\bar{\mathbf{v}}^t = (\mathbf{p}\mathbf{A} + (1-q)\mathbf{I})\bar{\mathbf{v}}^{t-1}$$

- As  $t \rightarrow \infty$ 
  - if  $\lambda_1(\mathbf{p}\mathbf{A} + (1-q)\mathbf{I}) < 1 \Leftrightarrow \lambda_1(\mathbf{A}) < q/p$  then  $\bar{\mathbf{v}}^t \rightarrow \mathbf{0}$ 
    - the probability that all copies die converges to 1
  - if  $\lambda_1(\mathbf{p}\mathbf{A} + (1-q)\mathbf{I}) = 1 \Leftrightarrow \lambda_1(\mathbf{A}) = q/p$  then  $\bar{\mathbf{v}}^t \rightarrow \mathbf{c}$ 
    - the probability that all copies die converges to 1
  - if  $\lambda_1(\mathbf{p}\mathbf{A} + (1-q)\mathbf{I}) > 1 \Leftrightarrow \lambda_1(\mathbf{A}) > q/p$  then  $\bar{\mathbf{v}}^t \rightarrow \infty$ 
    - the probability that all copies die converges to a constant  $< 1$

# SIS and SIR



(a) To represent the SIS epidemic using the SIR model, we use a “time-expanded” contact network

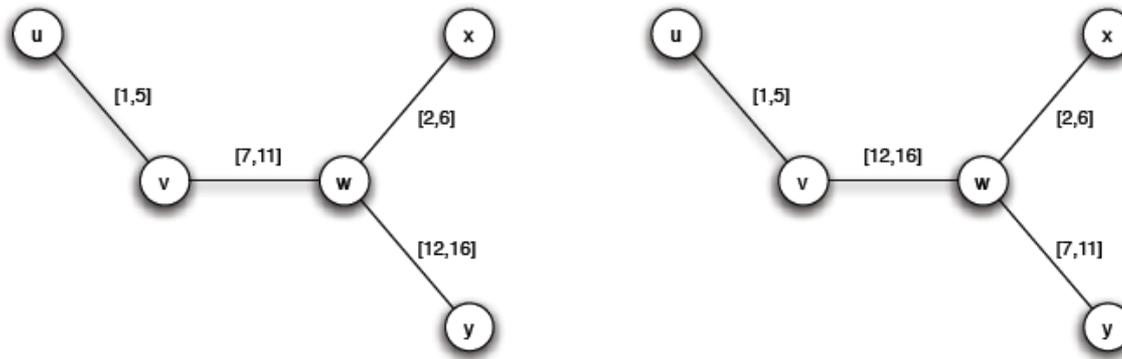


(b) The SIS epidemic can then be represented as an SIR epidemic on this time-expanded network.

Figure 21.6: An SIS epidemic can be represented in the SIR model by creating a separate copy of the contact network for each time step: a node at time  $t$  can infect its contact neighbors at time  $t + 1$ .

# Including time

- Infection can only happen within the **active window**



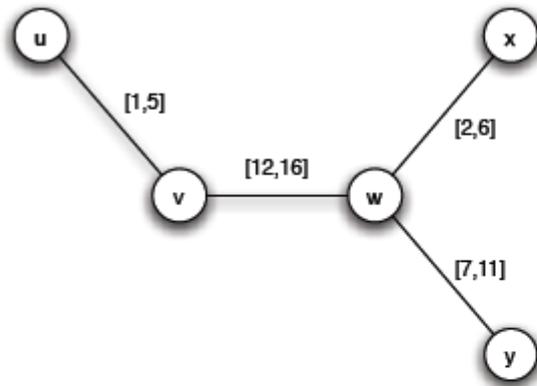
(a) In a contact network, we can annotate the edges with time windows during which they existed.

(b) The same network as in (a), except that the timing of the  $w$ - $v$  and  $w$ - $y$  partnerships have been reversed.

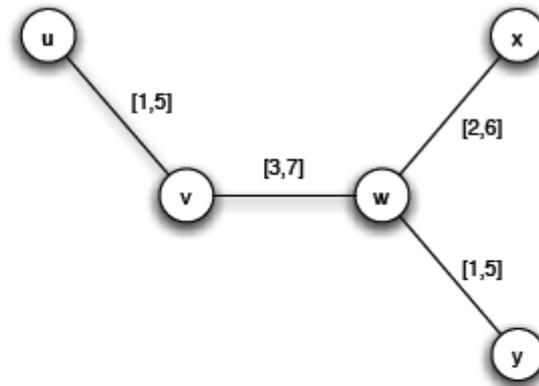
Figure 21.8: Different timings for the edges in a contact network can affect the potential for a disease to spread among individuals. For example, in (a) the disease can potentially pass all the way from  $u$  to  $y$ , while in (b) it cannot.

# Concurrency

- Importance of concurrency – enables branching



(a) *No node is involved in any concurrent partnerships*



(b) *All partnerships overlap in time*

Figure 21.10: In larger networks, the effects of concurrency on disease spreading can become particularly pronounced.

# **INFLUENCE MAXIMIZATION**

# Maximizing spread

- Suppose that instead of a virus we have an **item** (product, idea, video) that propagates through **contact**
  - **Word of mouth propagation.**
- An advertiser is interested in **maximizing the spread** of the item in the network
  - The holy grail of “**viral marketing**”
- Question: which nodes should we “**infect**” so that we maximize the spread? [KKT2003]

# Independent cascade model

- Each node may be **active** (has the item) or **inactive** (does not have the item)
- Time proceeds at discrete time-steps. At time **t**, every node **v** that became active in time **t-1** activates a non-active neighbor **w** with probability  $p_{uw}$ . If it fails, it does not try again
- The same as the simple **SIR model**

# Influence maximization

- **Influence function**: for a set of nodes  $A$  (target set) the influence  $s(A)$  is the expected number of active nodes at the end of the diffusion process if the item is originally placed in the nodes in  $A$ .
- **Influence maximization problem** [KKT03]: Given an network, a diffusion model, and a value  $k$ , identify a set  $A$  of  $k$  nodes in the network that maximizes  $s(A)$ .
- The problem is NP-hard

# A Greedy algorithm

- What is a simple algorithm for selecting the set  $A$ ?

## Greedy algorithm

Start with an empty set  $A$

Proceed in  $k$  steps

At each step add the node  $u$  to the set  $A$  that **maximizes** the **increase** in function  $s(A)$

- The node that activates the most additional nodes

- Computing  $s(A)$ : perform multiple **simulations** of the process and take the average.
- How good is the solution of this algorithm compared to the optimal solution?

# Approximation Algorithms

- Suppose we have a (combinatorial) optimization problem, and  $X$  is an instance of the problem,  $OPT(X)$  is the value of the optimal solution for  $X$ , and  $ALG(X)$  is the value of the solution of an algorithm  $ALG$  for  $X$ 
  - In our case:  $X = (G, k)$  is the input instance,  $OPT(X)$  is the spread  $S(A^*)$  of the optimal solution,  $GREEDY(X)$  is the spread  $S(A)$  of the solution of the Greedy algorithm
- $ALG$  is a good approximation algorithm if the ratio of  $OPT$  and  $ALG$  is **bounded**.

# Approximation Ratio

- For a **maximization** problem, the algorithm **ALG** is an  **$\alpha$ -approximation algorithm**, for  **$\alpha < 1$** , if for all input instances  **$X$** ,  
$$ALG(X) \geq \alpha OPT(X)$$
- The solution of  **$ALG(X)$**  has value **at least  $\alpha\%$**  that of the optimal
- **$\alpha$**  is the **approximation ratio** of the algorithm
  - Ideally we would like  **$\alpha$**  to be a **constant close to 1**

# Approximation Ratio for Influence Maximization

- The **GREEDY** algorithm has approximation ratio  $\alpha = 1 - \frac{1}{e}$

$$GREEDY(X) \geq \left(1 - \frac{1}{e}\right) OPT(X), \text{ for all } X$$

# Proof of approximation ratio

- The spread function  $s$  has two properties:

- $S$  is **monotone**:

$$S(A) \leq S(B) \text{ if } A \subseteq B$$

- $S$  is **submodular**:

$$S(A \cup \{x\}) - S(A) \geq S(B \cup \{x\}) - S(B) \text{ if } A \subseteq B$$

- The addition of node  $x$  to a set of nodes has **greater** effect (more activations) for a **smaller** set.
  - The **diminishing returns** property

# Optimizing submodular functions

- **Theorem:** A **greedy** algorithm that optimizes a **monotone** and **submodular** function  $S$ , each time adding to the solution  $A$ , the node  $x$  that maximizes the gain  $S(A \cup \{x\}) - s(A)$  has approximation ratio  $\alpha = \left(1 - \frac{1}{e}\right)$
- The spread of the Greedy solution is **at least 63%** that of the optimal

# Submodularity of influence

- Why is  $S(A)$  submodular?
  - How do we deal with the fact that influence is defined as an **expectation**?
- We will use the fact that **probabilistic propagation** on a **fixed graph** can be viewed as **deterministic propagation** over a **randomized graph**
  - Express  $S(A)$  as an expectation over the **input graph** rather than the choices of the algorithm

# Independent cascade model

- Each edge  $(u,v)$  is considered only **once**, and it is “activated” with probability  $p_{uv}$ .
- We can assume that all random choices have been made in advance
  - generate a **sample subgraph** of the input graph where edge  $(u,v)$  is included with probability  $p_{uv}$
  - propagate the item **deterministically** on the input graph
  - the active nodes at the end of the process are the nodes **reachable** from the target set  $A$
- The influence function is obviously(?) submodular when propagation is deterministic
- The **linear combination** of submodular functions is also a submodular function

# Linear threshold model

- Again, each node may be **active** or **inactive**
- Every **directed** edge  $(v,u)$  in the graph has a weight  $b_{vu}$ , such that

$$\sum_{v \text{ is a neighbor of } u} b_{vu} \leq 1$$

- Each node  $u$  has a **randomly generated** threshold value  $T_u$
- Time proceeds in discrete time-steps. At time  $t$  an **inactive** node  $u$  becomes **active** if

$$\sum_{v \text{ is an active neighbor of } u} b_{vu} \geq T_u$$

- Related to the game-theoretic model of adoption.

# Influence Maximization

- KKT03 showed that in this case the influence  $S(A)$  is still a **submodular** function, using a similar technique
  - Assumes **uniform random thresholds**
- The **Greedy** algorithm achieves a  $(1-1/e)$  approximation

# Proof idea

- For each node  $u$ , pick **one** of the edges  $(v, u)$  incoming to  $u$  with probability  $b_{vu}$  and make it **live**. With probability  $1 - \sum b_{vu}$  it picks no edge to make live
- Claim: Given a set of seed nodes  $A$ , the following two **distributions** are the **same**:
  - The **distribution over the set of activated nodes** using the Linear Threshold model and seed set  $A$
  - The **distribution over the set of nodes of reachable nodes** from  $A$  using live edges.

# Proof idea

- Consider the special case of a **DAG** (Directed Acyclic Graph)
  - There is a **topological ordering** of the nodes  $v_0, v_1, \dots, v_n$  such that edges go from left to right
- Consider node  $v_i$  in this ordering and assume that  $S_i$  is the set of **neighbors** of  $v_i$  that are **active**.
- What is the probability that node  $v_i$  becomes active in either of the two models?
  - In the **Linear Threshold** model the random threshold  $\theta_i$  must be greater than  $\sum_{u \in S_i} b_{ui} \geq \theta_i$
  - In the **live-edge** model we should pick one of the edges in  $S_i$
- This proof idea generalizes to general graphs
  - Note: if we know the thresholds in advance submodularity does not hold!

# Experiments

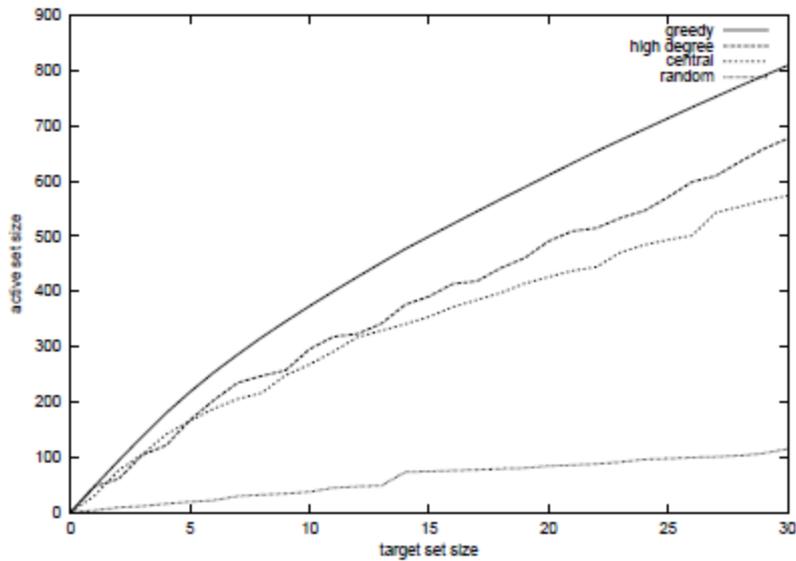


Figure 2: Results for the weighted cascade model

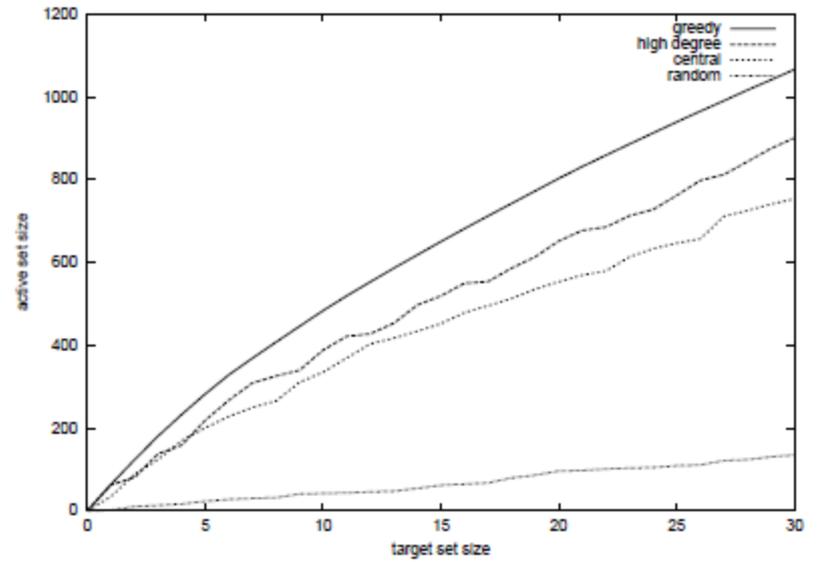
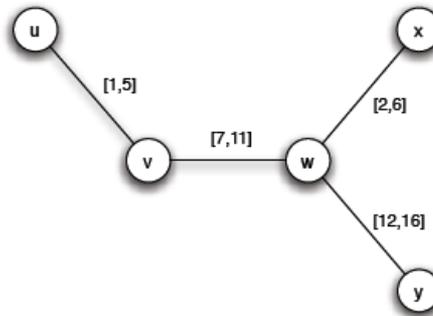
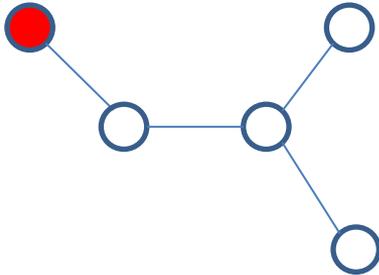


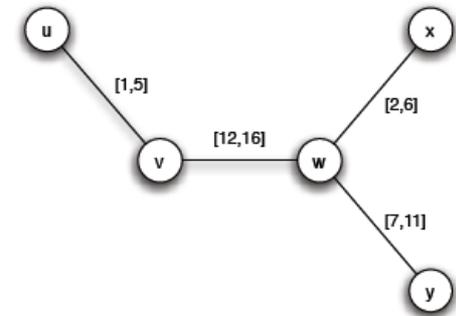
Figure 1: Results for the linear threshold model

# Another example

- What is the spread from the red node?



(a) In a contact network, we can annotate the edges with time windows during which they existed.



(b) The same network as in (a), except that the timing of the  $w-v$  and  $w-y$  partnerships have been reversed.

- Inclusion of **time** changes the problem of influence maximization

- N. Gayraud, E. Pitoura, P. Tsaparas, Diffusion Maximization on Evolving networks, submitted to SDM 2015

# Evolving network

- Consider a network that **changes** over time
  - Edges and nodes can appear and disappear at **discrete time steps**
- Model:
  - The evolving network is a sequence of graphs  $\{G_1, G_2, \dots, G_n\}$  defined over the same set of vertices  $V$ , with different edge sets  $E_1, E_2, \dots, E_n$ 
    - Graph snapshot  $G_i$  is the graph at time-step  $i$ .

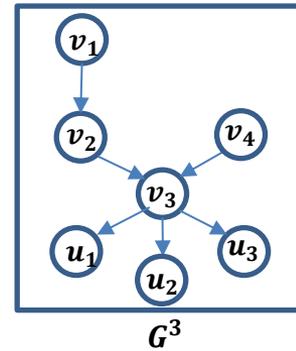
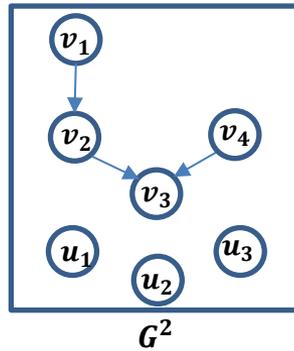
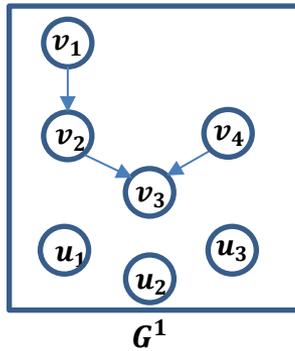
# Time

- How does the evolution of the network **relates** to the evolution of the diffusion?
  - How much physical time does a diffusion step last?
- Assumption: The two processes are **in sync**. One diffusion step happens in on one graph snapshot
- **Evolving IC model**: at time-step  $t$ , the infectious nodes try to infect their neighbors in the graph  $G_t$ .
- **Evolving LT model**: at time-step  $t$  if the weight of the active neighbors of node  $v$  in graph  $G_t$  is greater than the threshold the nodes gets activated.

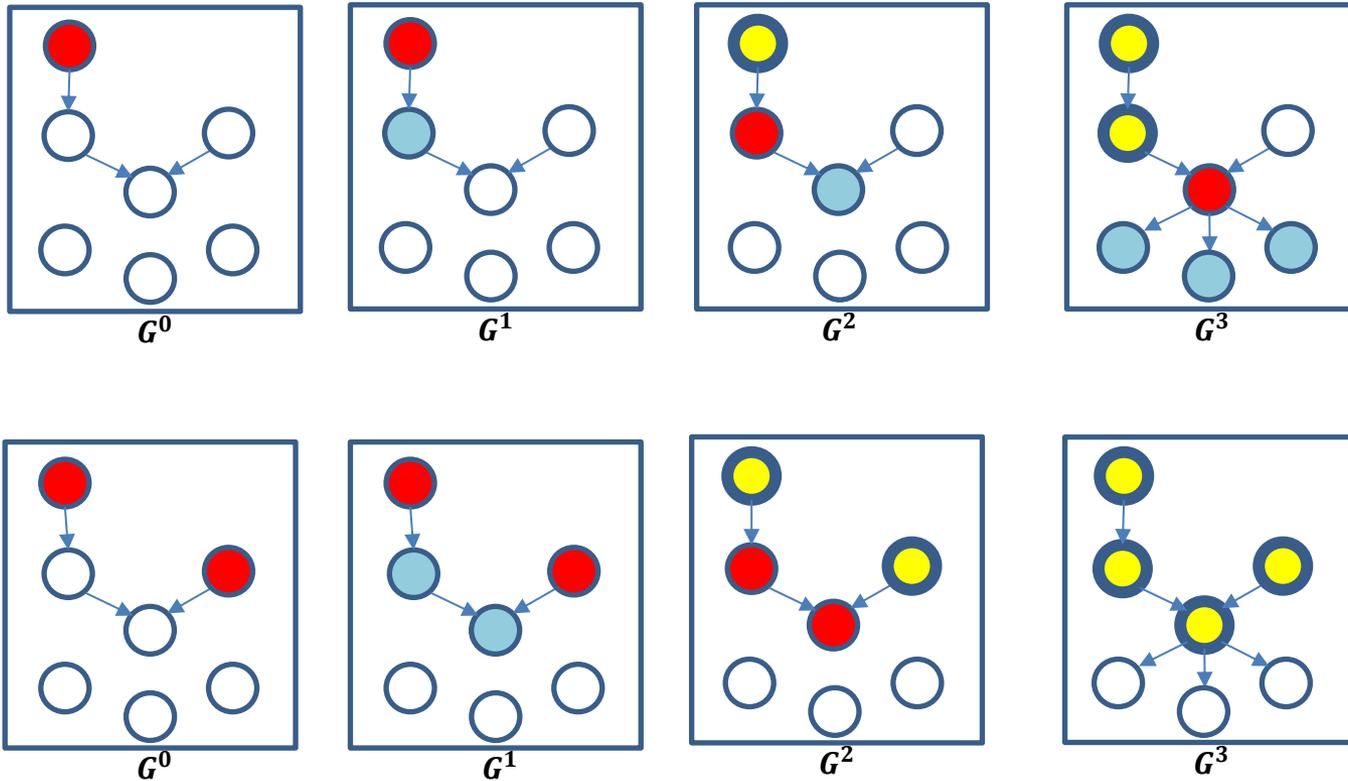
# Submodularity

- Will the spread function remain monotone and submodular?
- No!

# Evolving IC model



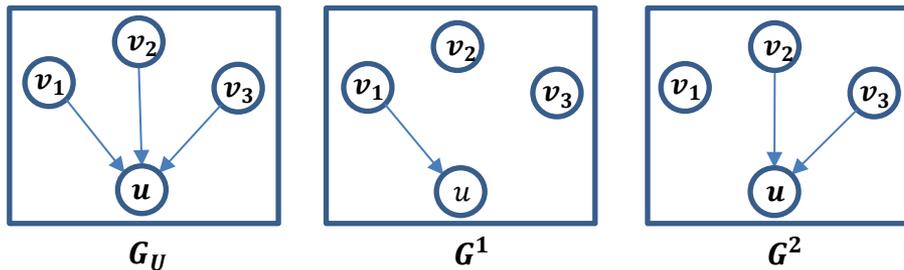
# Evolving IC model



The spread is **not even monotone** in the case of the Evolving IC model

# Evolving LT model

- The evolving LT model is monotone but it is **not submodular**



- Expected Spread:** the probability that  $u$  gets infected
  - Adding node  $v_3$  has a **larger effect** if  $v_2$  is already in the set.