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# Processes on A Network: Diffusion

IV124

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# Processes on Networks: Diffusion

Similar principles in different contexts

- Technical networks: cascading failures
- Biological networks: epidemics
- Social networks: opinion formation, information spreading

Today: models of these processes.

# Diffusion: Basic Concepts and Principles

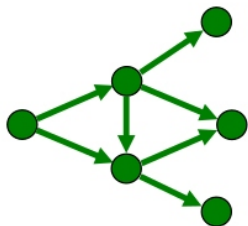
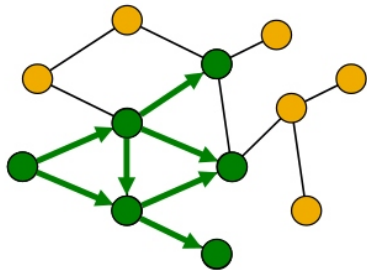
## Components of the model

- **What is spreading:** infection, information, choice, etc.
- **Time-point of spread:** change of choice, infection, failure, etc.
- **Outcome:** epidemic, group-decision, excluded nodes, etc.

We consider a **discrete time domain** – the model evolves in iterative steps. We model a dynamic process on a **static** network.

## Cascade on a Network

One run of the model on a network forms a directed graph – a cascade.



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<sup>1</sup>Leskovec CS224W: Machine Learning with Graphs  
<http://web.stanford.edu/class/cs224w/>

# Forms of Diffusion/Infection

## Simple spreading

- Each node infects its neighbors with a certain probability at each step

## Complex spreading

- Spreading occurs only if a certain fraction of neighboring nodes are *infected*

# Coordination Game on a Network

## Task:

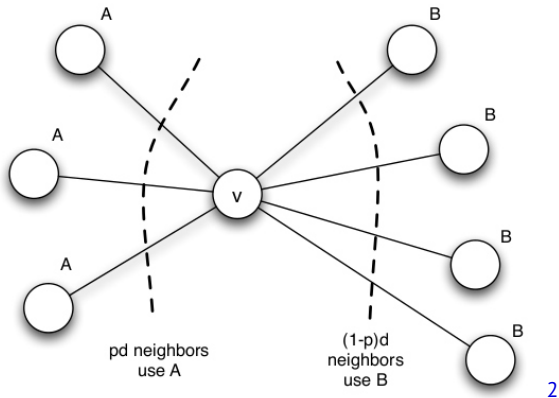
- Choose between A and B (e.g., VHS vs. Betamax, iPhone vs. Samsung)

## Rewards for neighboring nodes $u$ and $v$ :

- Both A: payoff  $a > 0$
- Both B: payoff  $b > 0$
- Disagreement: no payoff

Each node plays on its own. Spreading is monotonic (nodes do not take their decision back).

# Threshold for Changing Decision



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<sup>2</sup>Kleinberg, ch. 19.

# Threshold for Changing Decision

A is a better choice if

$$pda \geq (1 - p)db$$

Thus:

$$p \geq \frac{b}{a + b} = q$$



# Coordination Game - Properties

Equilibrium states:

- Everyone chooses A
- Everyone chooses B
- Incomplete cascade

The initiation of a cascade depends on the **topology of the network**, **initial conditions**, and the value of  $q$ .

## Cascades vs. Clusters

Clusters represent an obstacle for cascades

- dense internal connectivity
- small number of edges to the rest of the graph

Density  $\rho$  of cluster  $C \subseteq G$ :

- each node  $u \in C$  has at least fraction  $\rho$  of edges in  $C$

Cascades vs. density:

- cascades cannot propagate to clusters with  $\rho > (1 - q)$
- **conversely**: if the cascade stops, there is a cluster in the graph with  $\rho > (1 - q)$

# Cascades vs. Weak Ties

Reminder: **weak ties** are bridges between communities

## Role in cascades

- key for information diffusion (e.g. awareness of innovation)
- impenetrable to higher threshold phenomena (e.g. actual adoption of innovation)
- e.g. rapid global dynamics of sharing on social networks vs. slow and local dynamics of political mobilization

## Extensions to Coordination Game

### Bilingual nodes

- a node can choose state  $AB$
- reward  $AB-A$ :  $a$
- reward  $AB-B$ :  $b$
- reward  $AB-AB$ :  $\max(a, b)$
- nodes choosing  $AB$  additionally pay fixed cost  $c$

### Heterogeneous thresholds

- allows to incorporate differences in susceptibility

# Netlogo Demo

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# Models of Spreading: Epidemics

So far: Complex spreading

- Spreading via the majority of neighbors
- Sociological applications

Now: **Simple spreading**

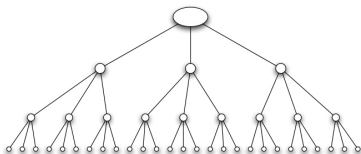
- Stochastic model
- Applications in biological and technical networks

## Note on Stochastic Models

Simple deterministic models can be made more complex (by adding rules)

- Increases the range of possible behaviors
- Analysis becomes more demanding
- At some point, it becomes easier to summarize a large number of real-world events into a single variable

# Branching Processes



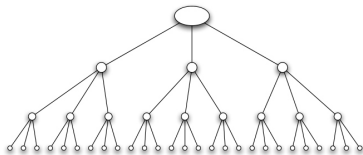
(a) The contact network for a branching process

The simplest tree model

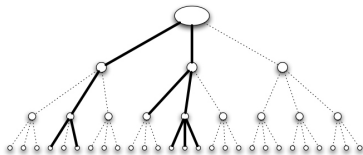
- Patient zero enters population and meets  $k$  individuals
- Probability of transmission upon meeting is  $p$
- $k$  and  $p$  remain constant in every subsequent wave
- Results in a tree of contacts between potentially infected individuals and the subtree of actual infection



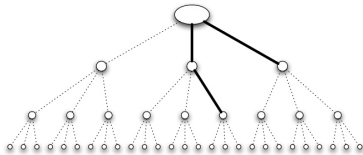
## Simple Spreading



(a) *The contact network for a branching process*



(b) *With high contagion probability, the infection spreads widely*



(c) *With low contagion probability, the infection is likely to die out quickly*

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<sup>3</sup>Easley and Kleinberg 2010

# Branching Processes

## Possible outcomes

- Infection stops after a while (dies out)
- Large epidemic

## Reproduction number $R_0$

- Expected number of new infections caused by a single individual
- Describes the viability and aggressiveness of the infection
- Here,  $R_0 = pk$

# Branching Processes

Development depending on  $R_0$ :

- $R_0 \ll 1$ : rapid end of spread
- $R_0 \gg 1$ : aggressive epidemic
- $R_0 \approx 1$ : the extent of the infection can vary significantly between runs; even small changes in the spread mechanism determine the outbreak of the epidemic

# SIR Model

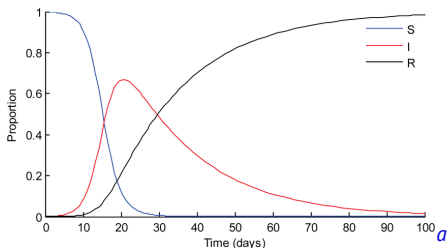
Three consecutive states of a node:

1. **Susceptible**: susceptible to infection from neighbors
2. **Infectious**: infected node spreading the disease for  $t_I$  steps
3. **Removed (Recovered)**: immune/dead node

In each step, nodes in state  $l$  transmit the disease to all their neighbors with probability  $p$ .

# Classic Epidemiological Models: SIR model

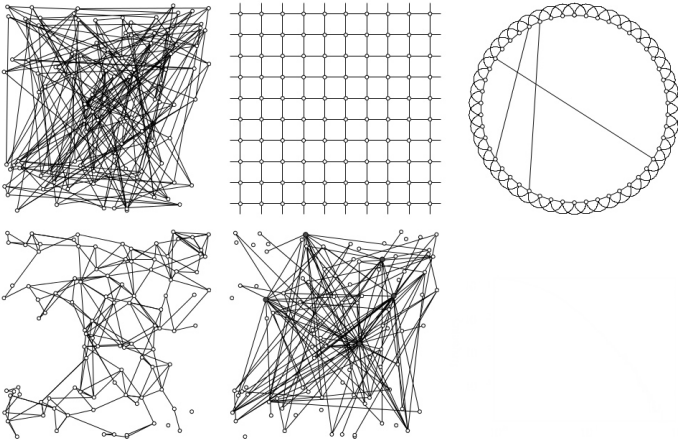
Assumes possible contact between subject and any other population member. Defined by differential equations:



$$\begin{aligned}\frac{dS}{dt} &= -\frac{\beta SI}{N} \\ \frac{dI}{dt} &= \frac{\beta SI}{N} - \gamma I \\ \frac{dR}{dt} &= \gamma I\end{aligned}$$

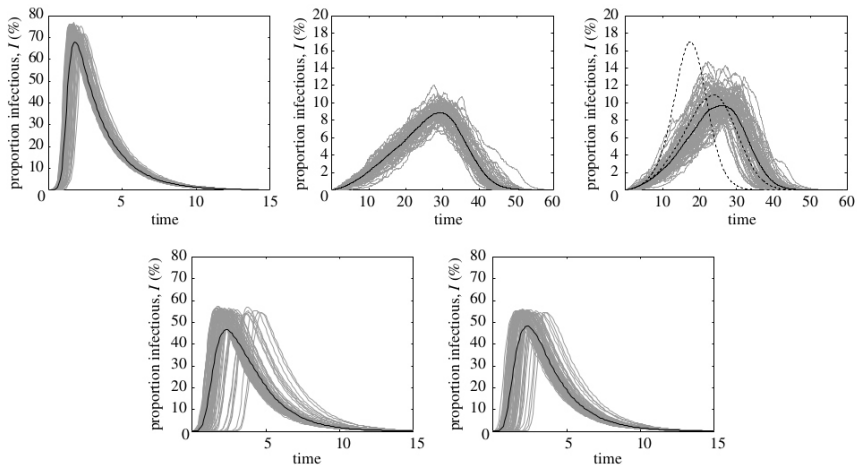
<sup>a</sup>Luz, P., et al (2010)

# SIR vs. Networks<sup>4</sup>



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<sup>4</sup>Keeling & Eames 2005

SIR vs. Networks<sup>5</sup>


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<sup>5</sup>Keeling & Eames 2005

## SIR Model: Extensions

The dynamics are simple (branching process on a network)

Possible extensions:

- Weighted graph: non-homogeneous transmission probability  $p$
- Non-homogeneous  $I_t$
- Division of  $I$  into more detailed categories: infectious incubation, less infectious period with symptoms, ...



# SIS Model

We allow for repeated infection.

1. **Susceptible:** susceptible to infection from neighbors
2. **Infectious:** infected node spreading infection for  $t_i$  steps
3. **Susceptible**

Compared to the SIR model, SIS allows for very long runs on a finite network.

## SIRS Model

In the occurrence of real diseases, we observe significant oscillations, which are not captured even by the SIS model.

We add time-limited immunity.

1. **Susceptible:** susceptible to infection from neighbors
2. **Infectious:** infected node spreading infection for  $t_I$  steps
3. **Recovery:** recovered and immune node for  $t_R$  steps
4. **Susceptible**

## Global vs. Local Oscillations

SIRS exhibits local oscillations on a general network.

Global oscillations

- require homophilic (local) links and long-range shortcuts
- correspond to the characteristic of small worlds
- the specific dynamics is closely related to the network topology

# SIRS and Small Worlds

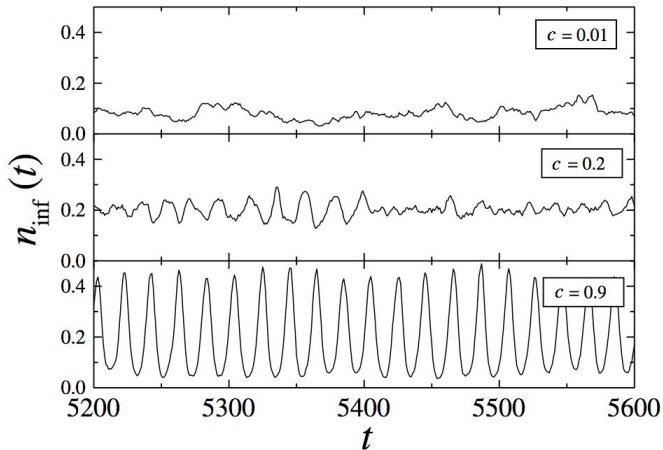
Small-world Watts-Strogatz network model (reminder):

- ring with local connections; with probability  $c$ , the edges are rewired to a random target

SIRS dynamics

- global oscillations (synchronization) depend on the number of *shortcuts* – weak links
- small  $c$  – local infection, large  $c$  – global oscillations

# SIRS and Small Worlds



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<sup>6</sup>Kiperman et al. 2001

# Demos

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