



# Agent-Based Simulation in Complex Networks

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### Miguel Rebollo (@mrebollo)

# Session 5. Dynamics

Complex networks display a surprising degree of robustness: although key components malfunction, local failures rarely lead to the loss of the global information.

We'll study why this effect appears.

#### PERCOLATION THEORY

Impact of node removal

# Percolation theory

removal of one link has limited impact, but remove several can break a network into components. How many nodes we have to delete?



# Percolation

Which is the expected size of the largest cluster?

What is the average cluster size?

When it is formed?



# Percolation

Average cluster size <s>

 $\langle s \rangle \sim |p - p_c|^{-\gamma_p}$ 

Order parameter  $P_{\infty}$ 

 $P_{\infty} \sim (p-p_c)^{\beta_p}$ 

**Correlation length** 

 $\xi \sim |p - p_c|^{-\nu}$ 

γ, β and υ: critical exponents

 $p_c = 0.593$  universal value



### ROBUSTNESS

Robust (from oak -roble-) system that resist errors and failures without degrading

### Robustness

Inverse percolation: node removal until disconnection

Critical exponents are the same

Fragmentation process is abrupt, not gradual



 $P_{\infty} \sim |f - f_c|^{\beta}$ 

### Efficiency

$$E = \frac{1}{N(N-1)} \sum_{i, j \neq i} \frac{1}{d_{ij}}$$

### The bigger the distance, the less efficient the network Networks with short paths are more efficient

### Vulnerability

Measures how failures affect to the efficiency (variation of eff when a link/node disappears)

Dependence on the topology



#### ATTACK TOLERANCE

What happens in the network when s sabotage is intended?

# **Deliberate Attacks**

#### Random



#### Hubs play a significant role in failure tolerance

Scale-free

#### CASCADE FAILURES

Because failures do not arrive isolated

### Cascades





# Three regimen



Subcritical <k> < 1

Supercritical <k> > 1 Critical <k> = 1

### DIFFUSION

Diseases, computer viruses, innovation or memes, all they share the same transmission scheme

### Closing credits. Planet of the Apes



#### **EPIDEMIC MODELING**

Mathematical models the simulates how a disease evolves

### Based on two concetps

**compartmentalization**: classification of individuals in groups attending to their state (SI / SIS / SIR)

**homogenous mixing**: all individuals have same prob. to get infected (no contact network)

# SI model



### susceptible



# SI model

Infection increases until all population infected





# SIS model

#### Two regimen

Endemic state ( $\gamma < \beta$ )

Disease free ( $\gamma > \beta$ )



$$\tau = \frac{1}{\gamma(R_0 - 1)}$$

Characteristic time

Reproductive number

< 1 extinction

 $R_0 = \frac{\beta \langle k \rangle}{\gamma}$ 

> 1 propagation

## SIR model



### susceptible

infected

### removed

# SIR model



#### **EPIDEMIC IN NETWORKS**

Contacts in a network limits the paths a virus takes in a population

# **Contact networks**

Contacts constrained by the network

Real degree instead of average

Differences on random and scale-free topologies



#### IMMUNIZATION

Can we protect? Immunization strategies. Herd immunization

### How to control a pandemic?

- Interventions to reduce transmission (masks, gloves,...)
- Contact-reducing interventions (quarentine)
- Vaccination: remove nodes from network

Who should be first vaccinated?

### Robustness and immunization



## Strategies

- Random immunization
  - Random selection
- Selective immunization
  - Random selection (G0)
  - Neighbors of G0 (G1)
  - illusion of majority deg(G1) > deg(G0)
    - $\rightarrow$  G1 vacinned