



NATIONAL RESEARCH
UNIVERSITY

Spreading phenomena in networks

Social Network Analysis. MAgOLEGO course.

Lecture 7

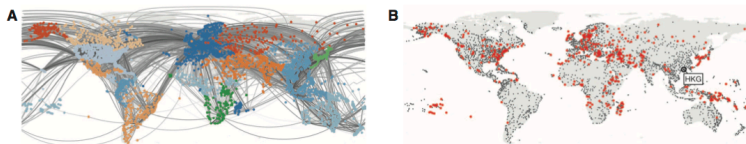
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www.leonidzhukov.net/hse/2019/sna

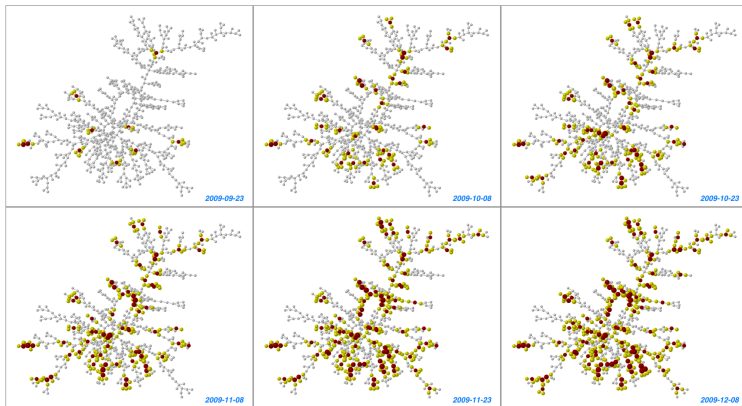
National Research University Higher School of Economics
School of Data Analysis and Artificial Intelligence, Department of Computer Science

Outbreak of SARS in 2003, > 8000 cases, 10% fatality rate, 37 countries



Simulated model:
gray lines - passenger flow, red symbols epidemics location

D. Brockmann, D. Helbing, 2013



Infected - red, friends of infected - yellow

N. Christakis, J. Fowler, 2010

- Given a network \mathbf{G} of potential contacts
- Three states model: susceptible, infected, recovered states
- Probabilistic model (state of a node):
 - $s_i(t)$ - probability that at t node i is susceptible
 - $x_i(t)$ - probability that at t node i is infected
 - $r_i(t)$ - probability that at t node i is recovered
- Model parameters:
 - β - infection rate (probably to get infected on a contact in time δt)
 - γ - recovery rate (probability to recover in a unit time δt)
- connected component - all nodes reachable
- network is undirected (matrix \mathbf{A} is symmetric)
- if graph complete - fully mixing model
- Based upon models from mathematical epidemiology, W.O. Kermack and McKendrick, 1927

Two processes:

- Node infection:



$$P_{inf} \approx \beta s_i(t) \sum_{j \in \mathcal{N}(i)} x_j(t) \delta t$$

- Node recovery:



$$P_{rec} = \gamma x_i(t) \delta t$$

- SI Model

$$S \longrightarrow I$$

- Probabilities that node i : $s_i(t)$ - susceptible, $x_i(t)$ - infected at t

$$x_i(t) + s_i(t) = 1$$

- β - infection rate, probability to get infected in a unit time

$$x_i(t + \delta t) = x_i(t) + \beta s_i(t) \sum_j A_{ij} x_j(t) \delta t$$

- infection equations

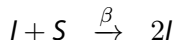
$$\begin{aligned} \frac{dx_i(t)}{dt} &= \beta s_i(t) \sum_j A_{ij} x_j(t) \\ x_i(t) + s_i(t) &= 1 \end{aligned}$$

SI Model

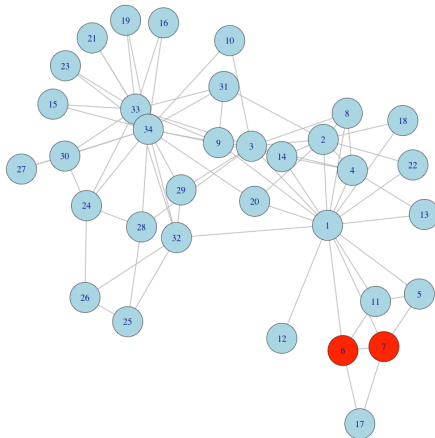
$$S \longrightarrow I$$

1. Every node at any time step is in one state $\{S, I\}$
2. Initialize c nodes in state I
3. On each time step each I node has a probability β to infect its nearest neighbors (NN), $S \rightarrow I$

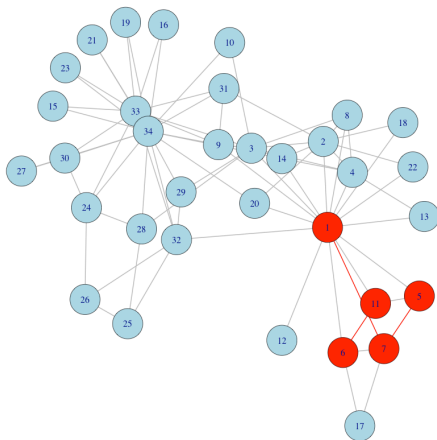
Model dynamics:



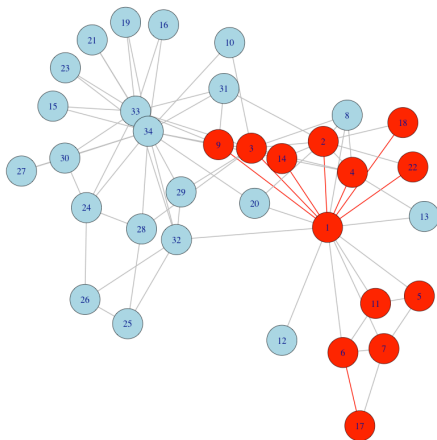
$$\beta = 0.5$$



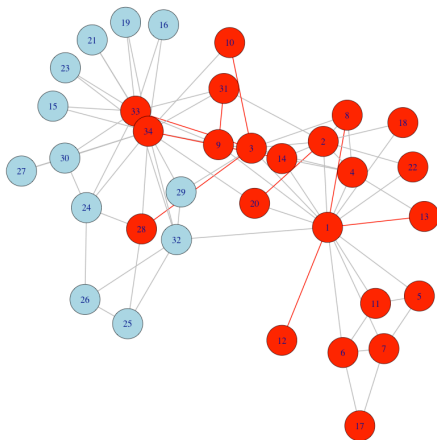
$$\beta = 0.5$$



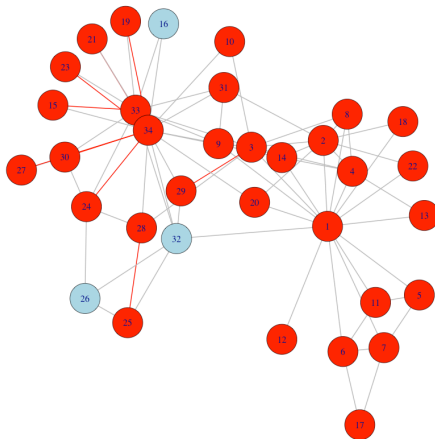
$$\beta = 0.5$$



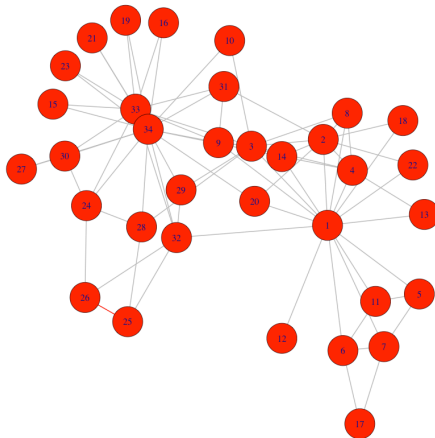
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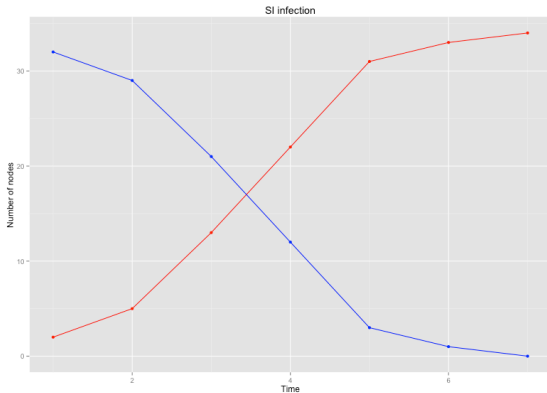


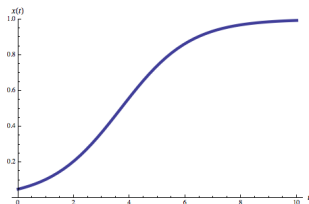
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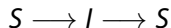




1. growth rate of infections depends on λ_1
2. All nodes in connected component get infected $t \rightarrow \infty$
 $x_i(t) \rightarrow 1$

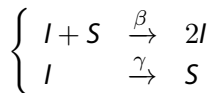
image from M. Newman, 2010

SIS Model

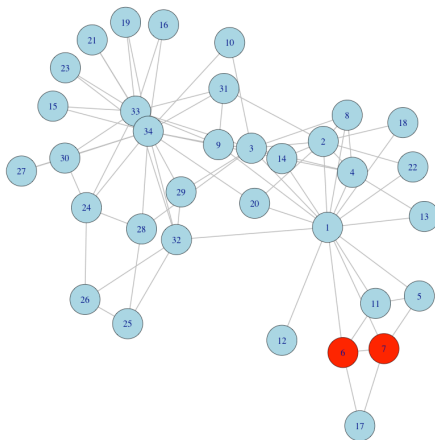


1. Every node at any time step is in one state $\{S, I\}$
2. Initialize c nodes in state I
3. Each node stays infected $\tau_\gamma = 1/\gamma$ time steps
4. On each time step each I node has a probability β to infect its nearest neighbors (NN), $S \rightarrow I$
5. After τ_γ time steps node recovers, $I \rightarrow S$

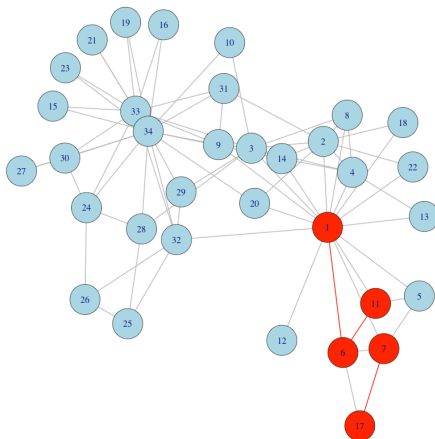
Model dynamics:



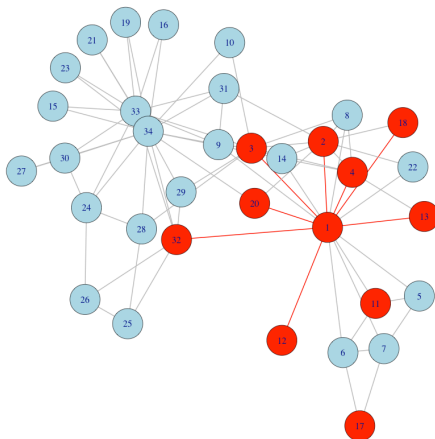
$$\beta = 0.5, \tau = 2$$



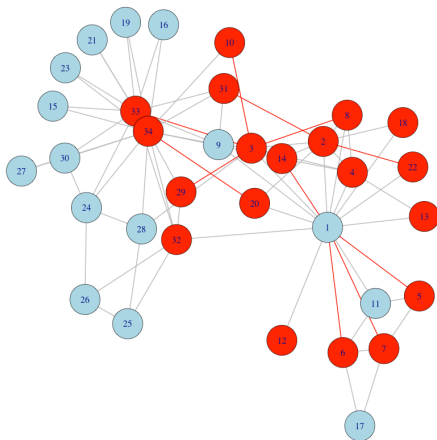
$$\beta = 0.5, \tau = 2$$



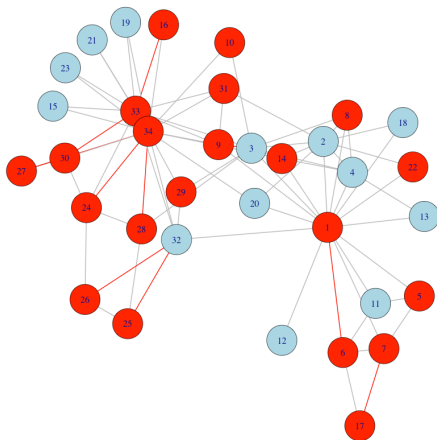
$$\beta = 0.5, \tau = 2$$



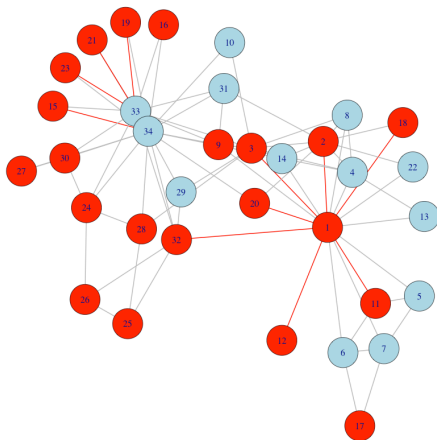
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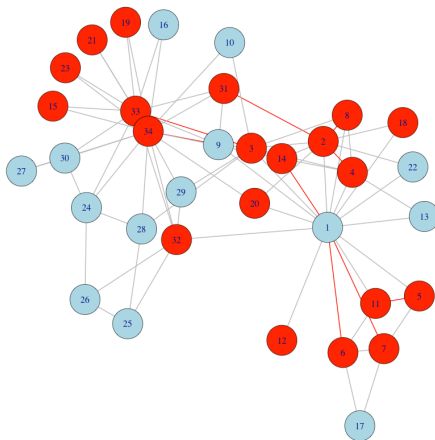
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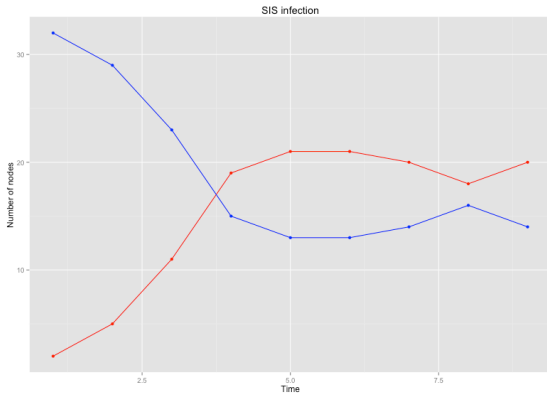


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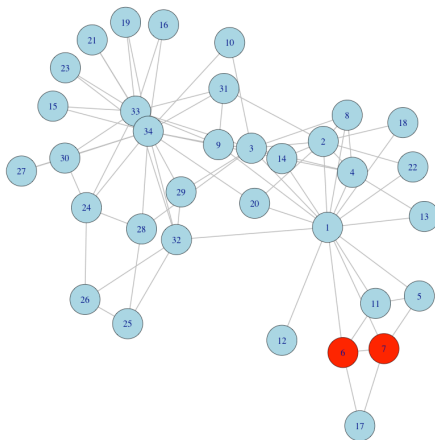


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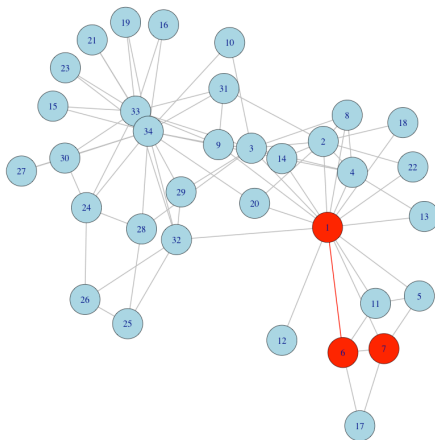




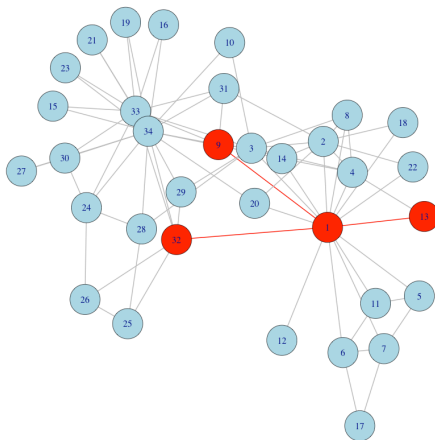
$$\beta = 0.2, \tau = 2$$



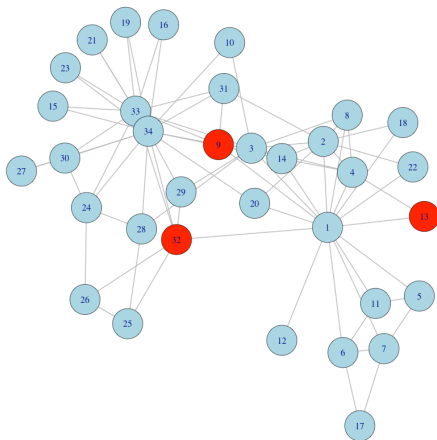
$$\beta = 0.2, \tau = 2$$



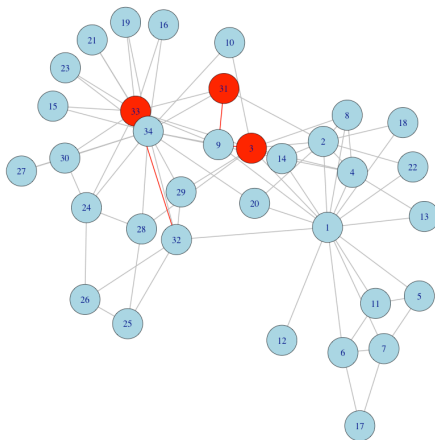
$$\beta = 0.2, \tau = 2$$



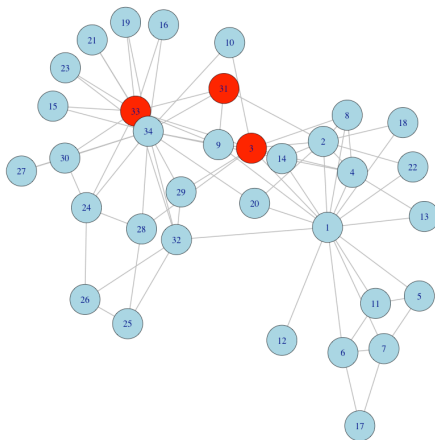
$$\beta = 0.2, \tau = 2$$



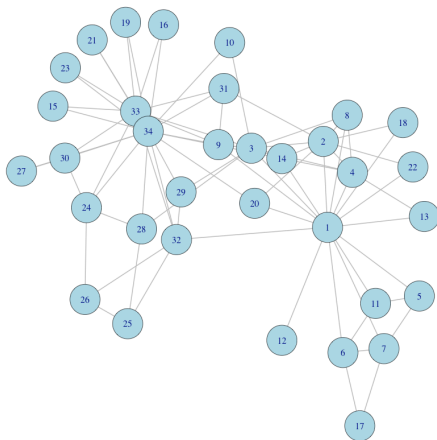
$$\beta = 0.2, \tau = 2$$

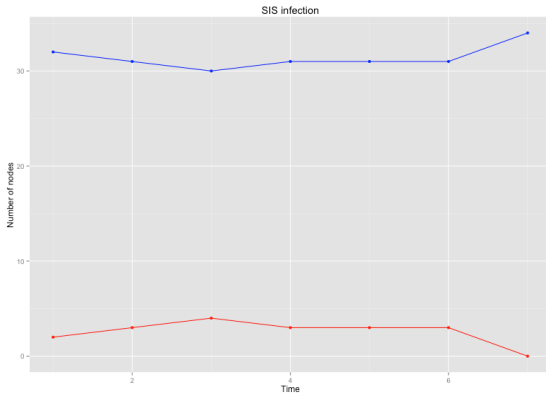


$$\beta = 0.2, \tau = 2$$



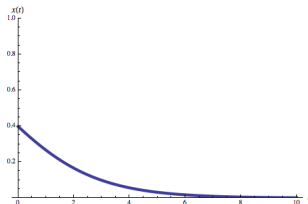
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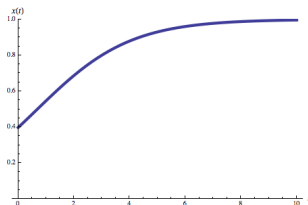


Epidemic threshold R_0 :

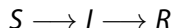
- if $\frac{\beta}{\gamma} < R_0$ - infection dies over time



- if $\frac{\beta}{\gamma} > R_0$ - infection survives and becomes epidemic



SIR Model

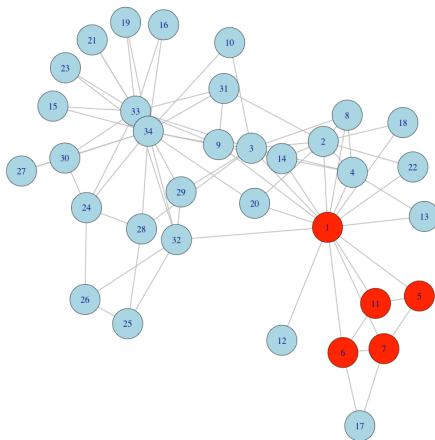


1. Every node at any time step is in one state $\{S, I, R\}$
2. Initialize c nodes in state I
3. Each node stays infected $\tau_\gamma = 1/\gamma$ time steps
4. On each time step each I node has a probability β to infect its nearest neighbours (NN), $S \rightarrow I$
5. After τ_γ time steps node recovers, $I \rightarrow R$
6. Nodes R do not participate in further infection propagation

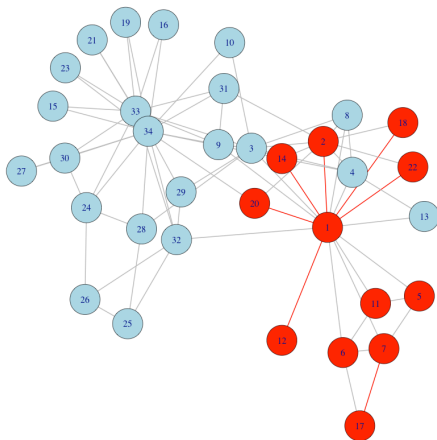
Model dynamics:

$$\left\{ \begin{array}{l} I + S \xrightarrow{\beta} 2I \\ I \xrightarrow{\gamma} R \end{array} \right.$$

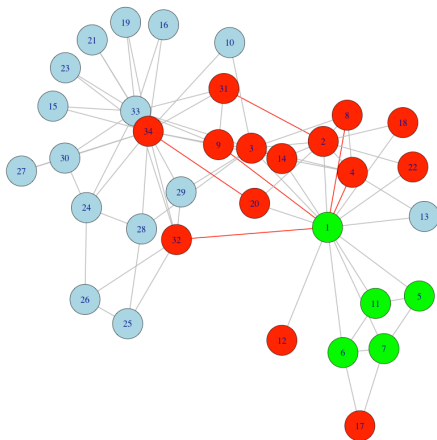
$$\beta = 0.5, \tau = 2$$



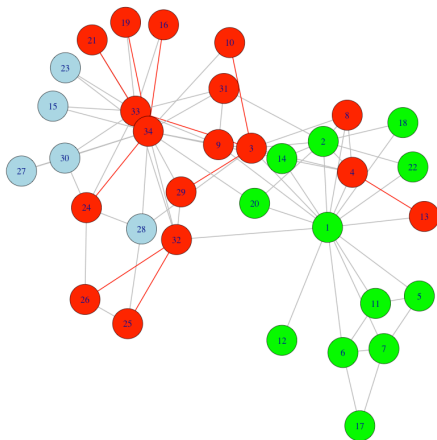
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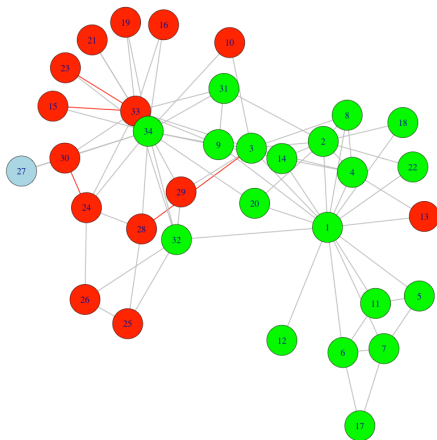
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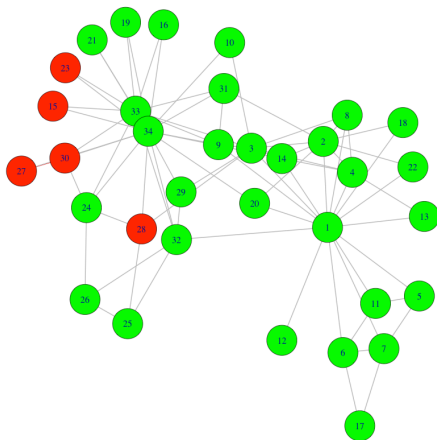
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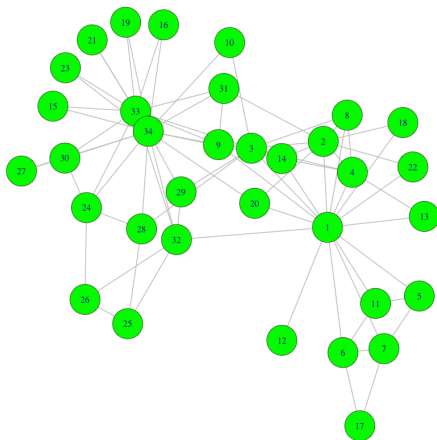
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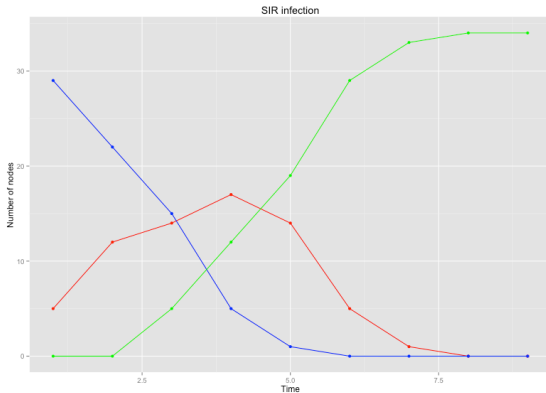


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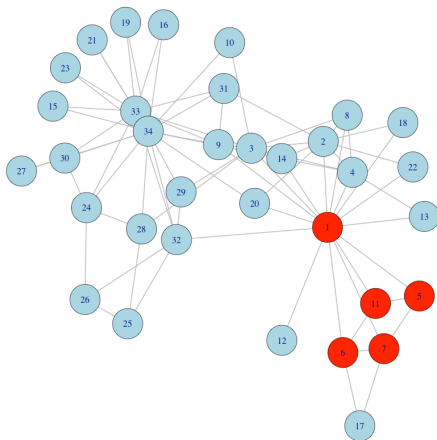


$$\beta = 0.5, \tau = 2$$

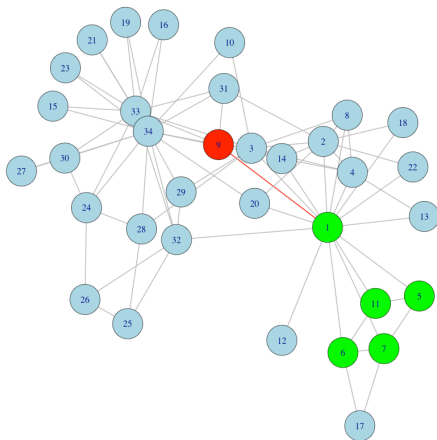




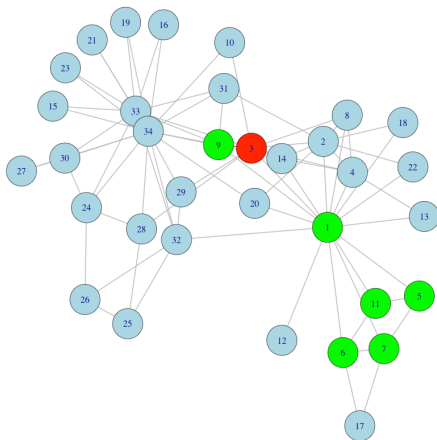
$$\beta = 0.2, \tau = 2$$



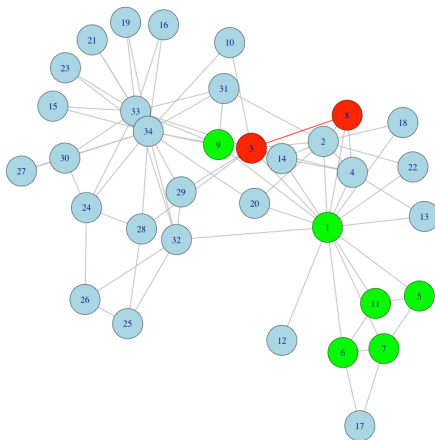
$$\beta = 0.2, \tau = 2$$



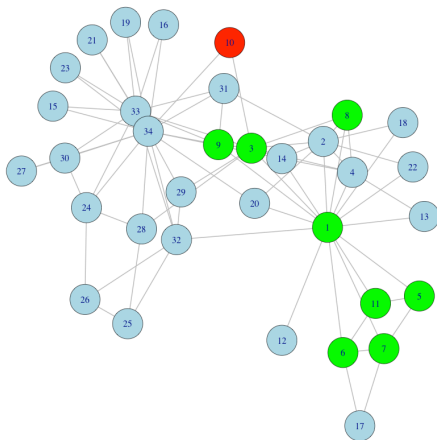
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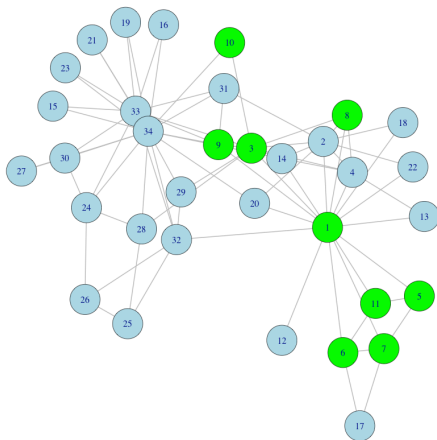
$\beta = 0.2, \tau = 2$

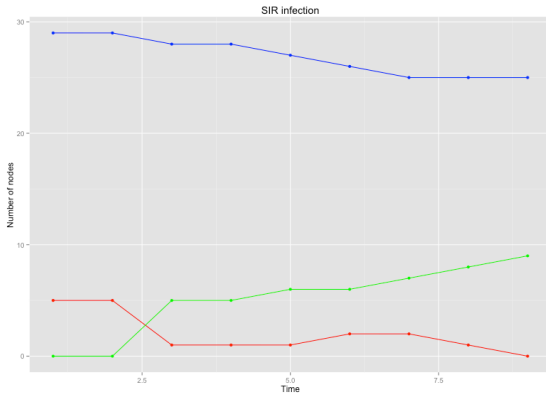


$$\beta = 0.2, \tau = 2$$



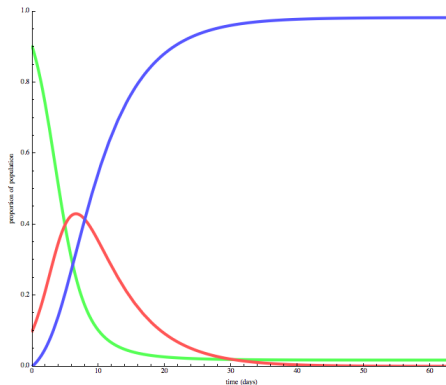
$$\beta = 0.2, \tau = 2$$





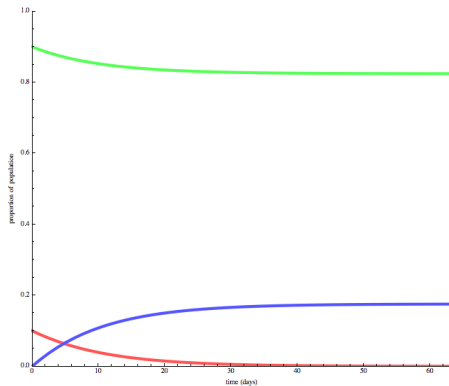
Epidemic threshold R_0 :

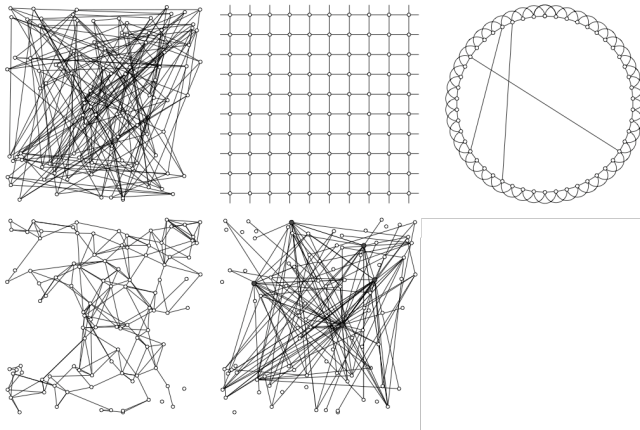
$\frac{\beta}{\gamma} > R_0$ - infection survives and becomes epidemic



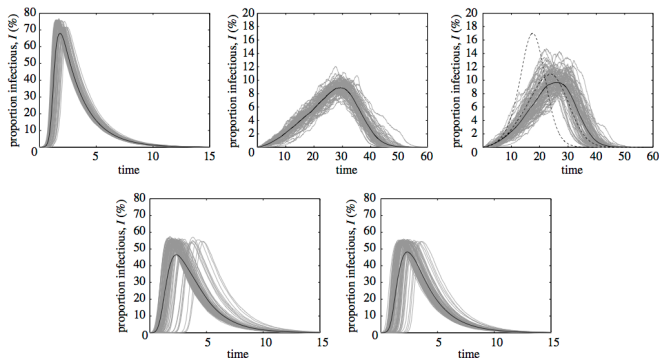
Epidemic threshold R_0 :

$\frac{\beta}{\gamma} < R_0$ - infection dies over time



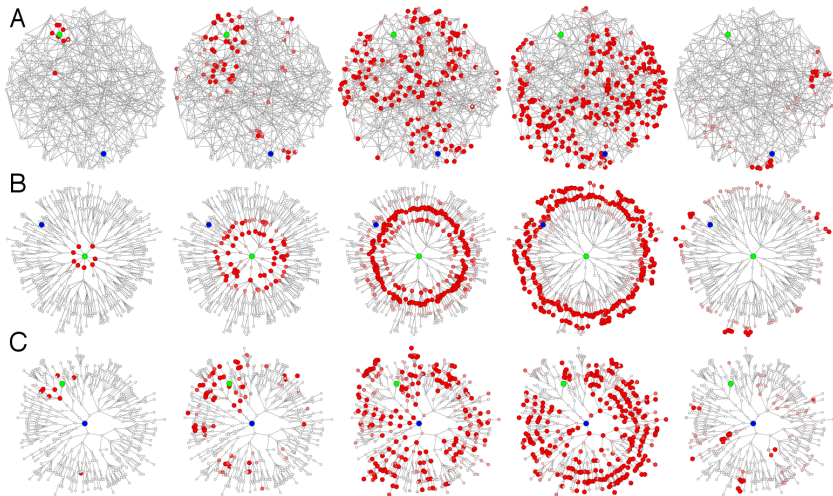


Networks: 1) random, 2) lattice, 3) small world, 4) spatial, 5) scale-free



Networks: 1) random, 2) lattice, 3) small world, 4) spatial, 5) scale-free

Keeling et al, 2005

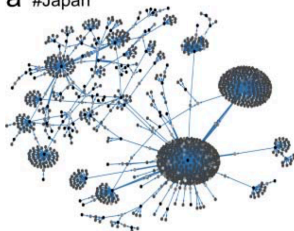


Social contagion phenomena refer to various processes that depend on the individual propensity to adopt and diffuse knowledge, ideas, information.

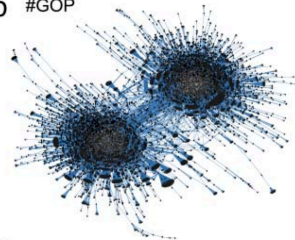
- Similar to epidemiological models:
 - "susceptible" - an individual who has not learned new information
 - "infected" - the spreader of the information
 - "recovered" - aware of information, but no longer spreading it
- Two main questions:
 - if the rumor reaches high number of individuals
 - rate of infection spread

Mem diffusion on Twitter

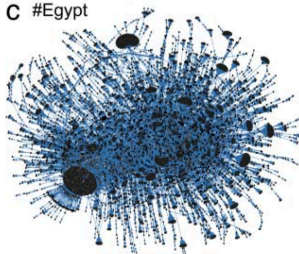
a #Japan



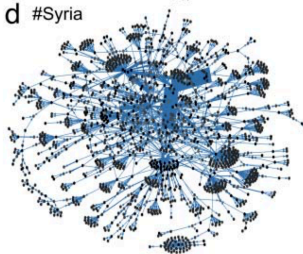
b #GOP



c #Egypt



d #Syria



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