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Models of network formation

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Part VI

Lecture 6 Epidemic models



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- The SI model is the one of susceptible and infected.
- Each person meets with β persons per unit of time.
- If there are X infected to start with then the average rate of new infections per unit of time is $\beta SX/n$. Hence we have

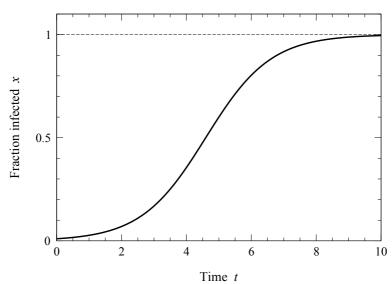
$$\frac{dX}{dt} = \beta \frac{SX}{n}$$
$$\frac{dS}{dt} = -\beta \frac{SX}{n}$$

• By using s = S/n and x = X/n, we get the solution $x(t) = \frac{x_0 e^{\beta t}}{1 - x_0 + x_0 e^{\beta t}}$ which is a logistic function.



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From [Newman, 2010, Fig. 17.1].



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- The SIR model is the one of susceptible, infected and recovered (or dead).
- Each person meets with β persons per unit of time and infected recover (or die) at a constant average rate of γ.
- If there are X infected to start with then the average rate of new infections per unit of time is $\beta SX/n$.



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Yannis Haralambous (IMT Atlantique • How long does an infected take to start recovering (and hence stop infecting others)? The probability of not recovering at δt is $1-y\delta t$ and hence the probability of not recovering in a total time t is $\lim_{\delta t \to 0} (1-y\delta t)^{t/\delta t} = e^{-yt}$. If we add the condition "recovering at time t" we get ye^{-yt} . Hence we have

$$\frac{dX}{dt} = \beta \frac{SX}{n}$$
$$\frac{dS}{dt} = -\beta \frac{SX}{n}$$

• By using s = S/n and x = X/n, we get the solution $x(t) = \frac{x_0 e^{\beta t}}{1 - x_0 + x_0 e^{\beta t}}$ which is a logistic function.



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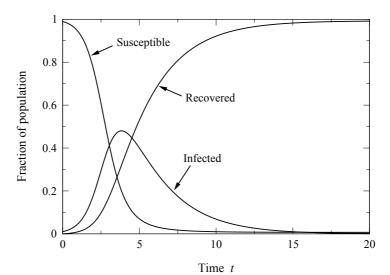
• The differential equations become

$$\begin{aligned} \frac{ds}{dt} &= -\beta sx \\ \frac{dx}{dt} &= \beta sx - \gamma x \\ \frac{dr}{dt} &= \gamma x. \end{aligned}$$



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From [Newman, 2010, Fig. 17.2], here $\beta = 1$, $\gamma = 0.4$, $s_0 = 0.99$, $x_0 = 0.01$, $r_0 = 0$.



The basic reproduction number

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- The basic reproduction number R₀ is the average number of people to which an infected passes the disease before recovering.
- If R < 1 the disease will die out. The value R₀ = 1 is called epidemic threshold.
- In the SIR model,

$$R_0 = \beta \gamma \int_0^\infty t e^{-\gamma t} dt = \frac{\beta}{\gamma}.$$



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- The SIS model is the one of susceptible, infected which can return to susceptible state, and then get reinfected, etc.
- Each person meets with β persons per unit of time and infected recoverat a constant average rate of γ.
- The differential equations are

$$\frac{ds}{dt} = \gamma x - \beta s x$$

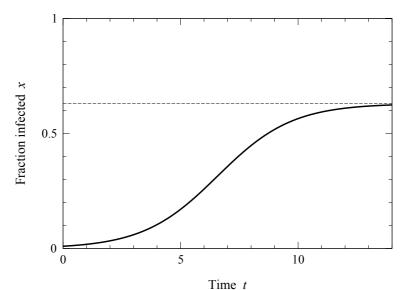
$$\frac{dx}{dt} = \beta sx - \gamma x$$
.

• We get the solution $x(t) = x_0 \frac{(\beta - \gamma)e^{(\beta - \gamma)t}}{\beta - \gamma + \beta x_0 e^{(\beta - \gamma)t}}$



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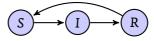
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From [Newman, 2010, Fig. 17.3].



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- The SIRS model is the one of susceptible, infected, recovered which can return to susceptible state, and then get reinfected, etc.
- Each person meets with β persons per unit of time and infected recoverat a constant average rate of y. Let δ the average rate at which people lose immunity.
- The differential equations are

$$\frac{ds}{dt} = \delta r - \beta s x$$
$$\frac{dx}{dt} = \beta s x - \gamma x$$
$$\frac{dr}{dt} = \gamma x - \delta r.$$

• There is no analytic solution for this model.



Epidemic models on networks

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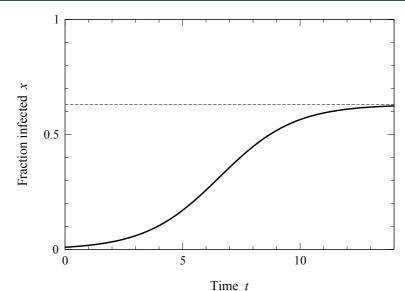
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- The epidemic models we described assumed full mixing of population.
- On a network we can consider that an infected individual can infect only its susceptible neighbors, with probability β per unit of time.
- M.J. Newman says: It is difficult to solve a model such as this
 for a general network, and in many cases the best we can do is to
 simulate it on a computer.
- We can study:
 - 1 late-time properties: what happens when $t \to \infty$?
 - 2 the relation of epidemics with giant components: if there is an infected individual in a giant component the result will be different than if it is in a small component;
 - 3 we can introduce probabilities of pairs of vertices of having given pairs of states;
 - 4 the time-related properties of a model for different degree distributions.
- For further reading, see [House, 2011].



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From [Newman, 2010, Fig. 17.3].



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