# **Epidemics**

#### Leonid E. Zhukov

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

#### Structural Analysis and Visualization of Networks



#### Lecture outline

- Epidemic models
  - SI model
  - SIS model
  - SIR model

- 2 Branching process
  - Galton-Watson process

# Epidemic dynamics models

- Mathematical epidimiology
- W. O. Kermack and A. G. McKendrick, 1927
- Deterministic compartamental model (population classes)  $\{S, I, T\}$
- S(t) succeptable, number of individuals not yet infected with the disease at time t
- I(t) infected, number of individuals who have been infected with the disease and are capable of spreading the disease.
- R(t) recoverd, number of individuals who have been infected and then recovered from the disease, can't be infected again or to transmit the infection to others.
- Fully-mixing model
- Closed population (no birth, death, migration)
- Models: SI, SIS, SIR, SIRS,...

• S(t) -susceptible , I(t) - infected

$$S \longrightarrow I$$

$$S(t) + I(t) = N$$

- $\bullet$   $\beta$  infection/contact rate, number of contacts per unit time
- Infection equation:

$$I(t + \delta t) = I(t) + \beta \frac{S(t)}{N} I(t) \delta t$$
 
$$\frac{dI(t)}{dt} = \beta \frac{S(t)}{N} I(t)$$

- Fractions: i(t) = I(t)/N, s(t) = S(t)/N
- Equations

$$\frac{di(t)}{dt} = \beta s(t)i(t)$$

$$\frac{ds(t)}{dt} = -\beta s(t)i(t)$$

$$s(t) + i(t) = 1$$

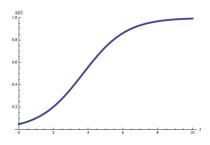
• Differential equation,  $i(t = 0) = i_0$ 

$$\frac{di(t)}{dt} = \beta(1 - i(t))i(t)$$

# Logistic growth function

Solution:

$$i(t) = \frac{i_0}{i_0 + (1 - i_0)e^{-\beta t}}$$



• Limit  $t \to \infty$ 

$$i(t) 
ightarrow 1 \ s(t) 
ightarrow 0$$

#### SIS model

• S(t) -susceptable , I(t) - infected,

$$S \longrightarrow I \longrightarrow S$$
  
 $S(t) + I(t) = N$ 

- ullet  $\beta$  infection rate (on contact),  $\gamma$  recovery rate
- Infection equations:

$$\frac{ds}{dt} = -\beta si + \gamma i$$

$$\frac{di}{dt} = \beta si - \gamma i$$

$$s + i = 1$$

• Differential equation,  $i(t = 0) = i_0$ 

$$\frac{di}{dt} = (\beta - \gamma - i)i$$

# SIS model

Solution

$$i(t) = (1 - \frac{\gamma}{\beta}) \frac{C}{C + e^{-(\beta - \gamma)t}}$$

where

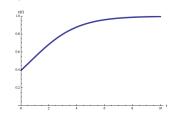
$$C = \frac{\beta i_0}{\beta - \gamma - \beta i_0}$$

• Limit  $t \to \infty$ 

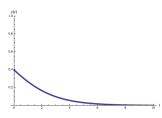
$$eta > \gamma \quad , \quad i(t) o (1 - rac{\gamma}{eta})$$
  $eta < \gamma \quad , \quad i(t) = i_0 e^{(eta - \gamma)t} o 0$ 

# Logistic function

• 
$$\beta > \gamma$$
,  $i(t) \rightarrow (1 - \frac{\gamma}{\beta})$ 



• 
$$\beta < \gamma$$
,  $i(t) = i_0 e^{(\beta - \gamma)t} \rightarrow 0$ 



• S(t) -susceptable , I(t) - infected, R(t) - recovered

$$S \longrightarrow I \longrightarrow R$$

$$S(t) + I(t) + R(t) = N$$

- ullet  $\beta$  infection rate,  $\gamma$  recovery rate
- Infection equation:

$$\frac{ds}{dt} = -\beta si$$

$$\frac{di}{dt} = \beta si - \gamma i$$

$$\frac{dr}{dt} = \gamma i$$

$$s + i + r = 1$$

Equation

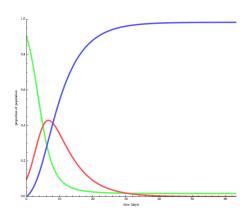
$$\frac{ds}{dt} = -\beta s \frac{dr}{dt} \frac{1}{\gamma}$$

$$s = s_0 e^{-\frac{\beta}{\gamma}r}$$

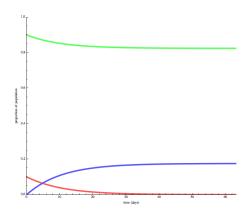
$$\frac{dr}{dt} = \gamma (1 - r - s_0 e^{-\frac{\beta}{\gamma}r})$$

Solution

$$t = \frac{1}{\gamma} \int_0^r \frac{dr}{1 - r - s_0 e^{-\frac{\beta}{\gamma}r}}$$



- $\frac{\beta}{\gamma} = 4$   $i_0 = 0.1$



- $\frac{\beta}{\gamma} = 0.5$   $i_0 = 0.1$

Equation

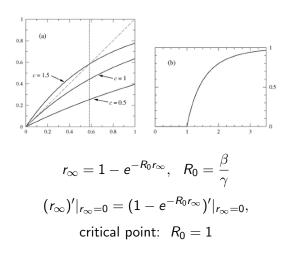
$$\frac{dr}{dt} = \gamma (1 - r - s_0 e^{-\frac{\beta}{\gamma}r})$$

• Limits:  $t \to \infty$ ,  $\frac{dr}{dt} = 0$ ,  $r_{\infty} = const$ ,

$$1 - r_{\infty} = s_0 e^{-\frac{\beta}{\gamma} r_{\infty}}$$

• Initial conditions: r(0) = 0, i(0) = c/N,  $s(0) = 1 - c/N \approx 1$ 

$$1-r_{\infty}=e^{-\frac{\beta}{\gamma}r_{\infty}}$$



- $r_{\infty}$  the total size of the outbreak
- Epidemic threshold

Epidemics: 
$$R_0 > 1$$
,  $\beta > \gamma$  ,  $r_\infty = const > 0$ 

No epidemics: 
$$R_0 < 1, \quad \beta < \gamma \quad , \quad r_\infty \to 0$$

Basic reproduction number

$$R_0 = \frac{\beta}{\gamma}$$

It is average number of people infected by a person before his recovery

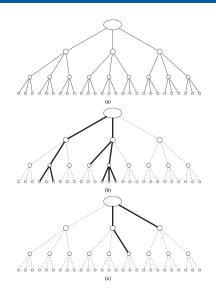
$$R_0 = E[\beta \tau] = \beta \int_0^\infty \gamma \tau e^{-\gamma \tau} d\tau = \frac{\beta}{\gamma}$$

# Model of contagion

Simple model of contagion (decease transmission)

- 1st-wave: first infected person enters the population and transmits to each person he meets with probability *p*. Suppose he meets *k* people while contagious
- 2nd-wave: Each infected person from 1st wave meets k new people and independently transmits infection with probability p
- 3rd-wave: ....

This is Galton-Watson branching stochastic process (Proposed by Francis Galton 1889 as a model for extinction of family names)



#### Random branching process:

- let  $\xi_i^n$  number of transmitted infections by *i*th node on level *n*
- let  $Z_n$  number of infected on level n,  $Z_0 = 1$ . Then:

$$Z_{n+1} = \sum_{i=1}^{Z_n} \xi_i^{(n)}$$

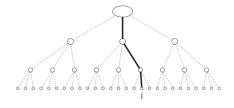
- If each node has k neighbors, transmits infection with probability p, Average number of infected people  $E[\xi_i^n] = pk = R_0$  basic reproductive number
- Recursion

$$E[Z_{n+1}] = E[\sum_{i=1}^{Z_n} \xi_i^{(n)}] = E[\xi_i^{(n)}] \ E[Z_n] = pk \ E[Z_n]$$

$$E[Z_n] = (pk)^n = R_0^n$$

Galton-Watson branching random process:

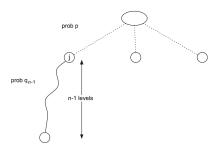
- if  $R_0 = 1$ , the mean of number of infected nodes does not change
- if  $R_0 > 1$ , the mean grows geometrically as  $R_0^n$
- ullet if  $R_0 < 1$ , the mean shrinks geometrically as  $R_0^n$



 $R_0 = 1$  - point of phase transition

#### Extinction probability

- let  $q_n$  probability that infection persists n steps (levels of the tree)
- $pq_{n-1}$  probability that spreads through one first contact and then survives n-1 levels

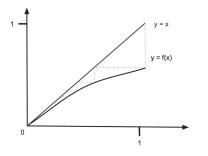


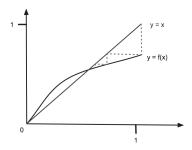
•  $(1 - pq_{n-1})^k$  - probability that will not spread through any of the subtries

$$(1 - pq_{n-1})^k = 1 - q_n$$

• Recurrence  $(q_n$  - probability that infection persists through n steps)

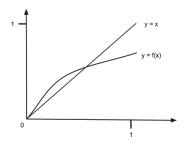
$$q_n = 1 - (1 - pq_{n-1})^k$$





• limiting probability  $q^* = \lim_{n \to \infty} q_n$ 

$$q^* = 1 - (1 - pq^*)^k$$



Slope:

$$pk(1-pq)^{k-1}\big|_{q=0}=1$$

• When  $R_0 = pk > 1$ , there is a non zero probability of infection persists

07.04.2015

#### References

- A Contribution to the Mathematical Theory of Epidemics. , Kermack, W. O. and McKendrick, A. G. , Proc. Roy. Soc. Lond. A 115, 700-721, 1927.
- The Mathematics of Infectious Disease, Herbert W. Hethcote, SIAM Review, Vol. 42, No. 4, p. 599-653, 2000