

①

Lecture 9 a : Spreading Processes on Networks.

- Project proposal / presentation / writeup reminder.
- PSS, 6 smaller

- dynamics on a network (assume G is fixed relative to timescale of dynamics)
- what things spread across a network?
 » (ask for examples)

- state variable $x_i \in \mathcal{P}$
- x_i state variable on each node i ;
 \mathcal{T} various states possible

update function

$$x_i^{t+1} = f(G) = f(k_{nb(i)}^t)$$

instead of ...

{ key idea: Network edges are the mechanism of transmission.
 (representation choice !)

- what are the nodes ?
 » (ask for examples)
- what are edges ?
 » (ask for examples)
- Three kinds of spreading processes on networks. (non-exhaustive)
 - (this wk) » Epidemic models (by pathogens)
 - (next wk) » Social adoption models (preferences)
 - » Biological computation models (neural cascades, boolean networks)

3

{ Two dichotomies

- continuous v.s. discrete time

$$\emptyset \xrightarrow{f} \emptyset \xrightarrow{f} \dots$$

G_t G_{t+1}

- simple v.s. complex contagions
- bio pathogen
- costly social behavior. (peer pressure) *modelling*
- low-cost social behavior

{ Epidemic models of spreading

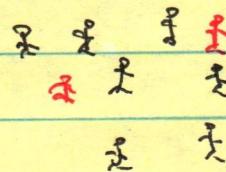
epidemiology : A study of all diseases.

infectious disease ep: : diseases that spread between people

network api : diseases that spread on a social network .

populations

- individuals
- internal systems
- tissues
- cells



molecules

(3)

} ID ep: warm up : the compartmental model

Kermack-McKendrick
(from 1927!)

- let's first ignore the network

assume all-to-all connections,
also called "well-mixed"

- each node has a state variable

$$\left. \begin{array}{l} x_i \in \{S, I, R\} \\ x_i \in \{S, I\} \end{array} \right\}$$

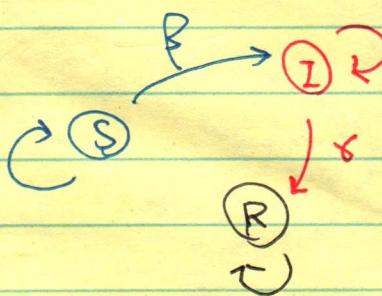
S: susceptible — not infected + can be infected.

I: infected / infectious

R: recovered / removed — not infected + cannot be infected again.

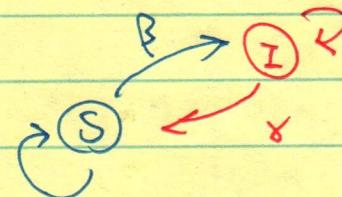
- define state transition rules to update each x_i
these rules are a model of the disease.

} state diagrams $S \xrightarrow{\beta} I_1 \xrightarrow{\gamma} I_2 \xrightarrow{\gamma} I_3 \xrightarrow{\gamma} R$



SIR

- chicken pox
- measles



SIS

influenza
≈ covid common cold.

* anything you can get a childhood vaccine for

* anything for which infection does not confer long-standing immunity

(4)

} Specifying SIR / SIS. (see figure on previous page)

β : transmission rate of infection

higher β : spreads faster / faster rate of $S \rightarrow I$.

lower β : slower spread / S stays larger for longer.

r : recovery rate from infection

higher r : faster recovery

lower r : more time in I .

} SIS dynamics

- let N be total population (constant)

- an infected person transmits disease to a susceptible ($S \rightarrow I$)

- person w/ prob. $\bar{\beta}$ per unit time

- an infected person recovers ($I \rightarrow S$) w/ prob r per unit time

- let $S(t)$ and $I(t)$ denote # S and # I ;

» Note :

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

$$\text{at } t=0 \quad S(0) = N-1, \quad I(0) = 1.$$

- how $S(t)$ evolves

$$S(t + \Delta t) = S(t) - \underbrace{[S(t) + \bar{\beta} \Delta t I(t)]}_{S \rightarrow I} + \underbrace{[I(t) \times r \Delta t]}_{I \rightarrow S}$$

» Then we convert difference eqn to a differential eqn.

(5)

» Variables

$$S = \frac{S}{N}$$

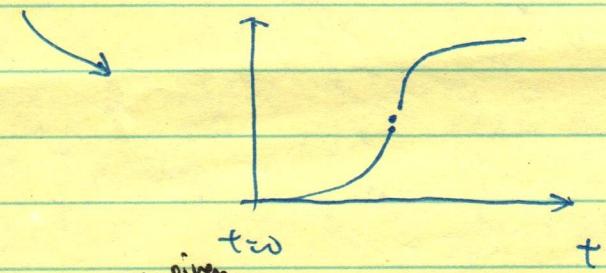
$$I = \frac{I}{N} \quad R = \overline{R} N$$

» rate equation

divide by Δt .

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

solution $I(t)$



Question: Will an epidemic spread? (given small seed)

- let i be very small (relative to N), so approx. $(1-i) \approx 1$.

$$\frac{di}{dt} \approx \beta i - ri = i(\beta - r)$$

- simple ODE w/ logistic function as solution

$$\frac{\beta}{r} : \begin{cases} > 1 & \text{exponential spread} \\ = 1 & \text{high var.} \\ < 1 & \text{does not spread.} \end{cases}$$

Highlights:

$$R_0 = \frac{\beta}{r} \quad (\text{exponential rate when } i \ll 1)$$

(6)

} See Lecture Notes.

$$\left\{ \begin{array}{l} \frac{di}{dt} = 0 \\ i(\infty) = 1 - \beta/r \end{array} \right.$$

Page 7 :

(2.2.2) left \rightarrow Infected eq. at $i = 1 - \beta/r$
(SIS)
right
(SIR)

SIR in the wild :

ideal cond. for SIR are rare (what are they?)

conditions.

100% naive pop.

closed pop.

"homogeneous" mixing.

but, 1978 influenza outbreak @ English boarding school w/
 $n = 763$ boys.

≈ 2.1 days infectious before
isolation.

$$\hat{\beta} = 0.00234$$

$$\gamma = 0.476.$$

Challenging models:
- compartmental

Covid edition : SEIR model w/ death and vaccinations.

see Pabari et al., Science (2021)

- Vaccinated / unvaccinated / no vacc. (refused).
- vaccination rate / infected fatality rate (IFR), etc.

exponential growth : covid-19 $I(t) \propto R_0^t$.

See page 9

Lecture 9 b :

March 14, 2024

"Snow day"

- See the Jupyter Notebook for basic epidemic simulations (on Canvas)