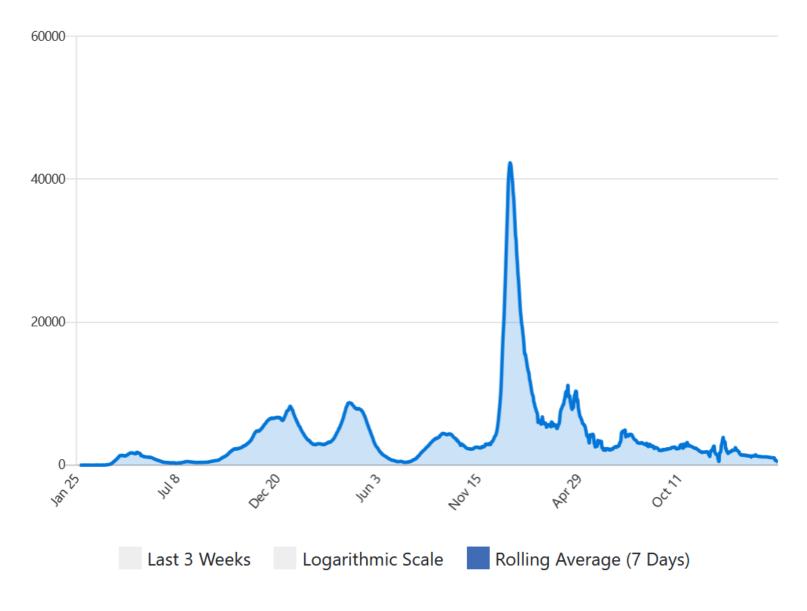
# Epidemic modelling basics (1-variable models) Lecture 12a: 2023-04-03

MAT A35 – Winter 2023 – UTSC Prof. Yun William Yu

Image credit: https://commons.wikimedia.org/wiki/File:Coronavirus\_SARS-CoV-2.jpg

#### Epidemic curves – Covid19 in Canada



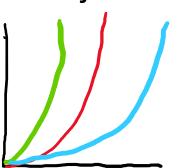
https://covid19tracker.ca/, 7-day rolling average, new cases

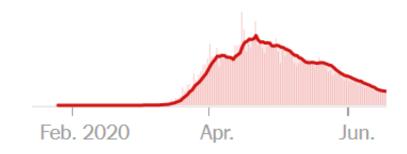
## Infection rates

• Assumption 1: each infected individual infects other individuals at a constant positive rate  $\beta$ .

## Exponential model: $I(t) = I(0)e^{\beta t}$

- Let's focus on the initial months of the pandemic.
- Can use regression to find the good values for  $\beta$ , or even just trial and error.





 Model doesn't take into account finite population size.

What's wrong with the model? A: Model is too simple

- B: Cannot determine good  $\beta$
- C: Doesn't reflect the data
- D: All of the above
- E: None of the above

## Compartmental models

• Assumption 2: there is a total fixed population size N = I(t) + S(t), where S is the number of Susceptible individuals.

• Does this fix the problems from the previous slide?

A: Yes B: No C: Maybe D: ???

## SI Model of Epidemics

 Modified assumption 1: The infection rate is proportional to the average number of times an infected individual encounters a susceptible individual in the population, assuming random encounters.

## Solving SI model qualitatively

$$N = S(t) + I(t)$$
$$\dot{S} = -\frac{\beta SI}{N}$$
$$\dot{I} = \frac{\beta SI}{N}$$

## Solving the SI model exactly

• 
$$\dot{I} = \frac{\beta I}{N} (N - I)$$

What methods should we use?

- A: Separation of variables
- **B:** Integrating factor
- C: u-substitution
- D: All of the above
- E: None of the above

#### Integrating factor and u-substitution

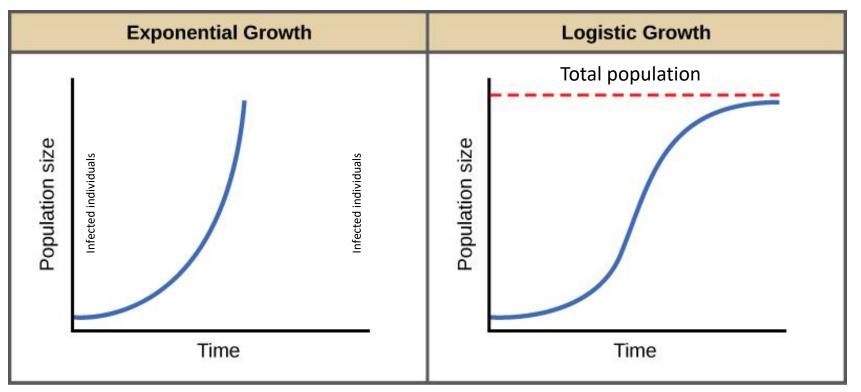
• 
$$\dot{I} = \frac{\beta I}{N} (N - I)$$

## Separation of variables

$$\dot{I} = \frac{\beta I}{N} (N - I)$$

## Logistic growth equation

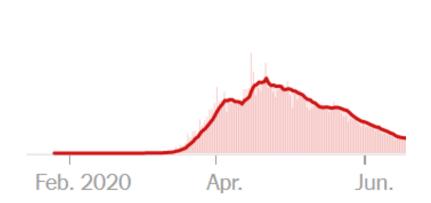
• SI model has logistic growth, which starts out like exponential growth, but levels out as everyone is infected.

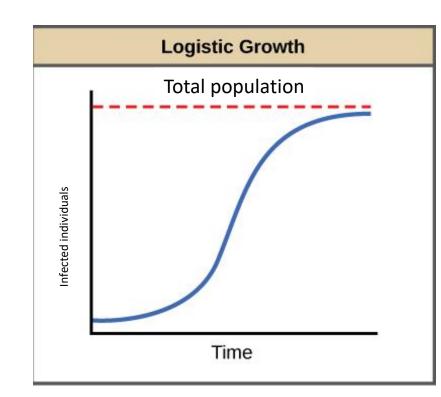


https://commons.wikimedia.org/wiki/File:Figure\_45\_03\_01.jpg

## Further improvements

- We now no longer go off to infinity, which is good.
- But, we still are missing the downward part of the epidemic curve.
- How do we get the number of infected to go back down in our model?
  - A: Add a recovery termB: Add a mortality rateC: Add an immunization rateD: All of the aboveE: None of the above





## SIS Model

• Assumption 3: Individuals recover at rate  $\gamma$ , and become susceptible to re-infection.

## Multiple cases

• 
$$\frac{dI}{dt} = \left(\beta - \frac{\beta I}{N} - \gamma\right) I$$

- Note, we have two parameters,  $\beta$  and  $\gamma$ , so there are three cases to consider.
- What are your guesses for behavior in each of the following?
- Case 1:  $\beta = \gamma$
- Case 2:  $\beta < \gamma$
- Case 3:  $\beta > \gamma$ 
  - A: Disease dies out
  - B: Number of infected goes to nonzero constant
  - C: Number of infected oscillates up and down
  - D: All of the above
  - E: None of the above

Case 1: 
$$\beta = \gamma$$
  
•  $\frac{dI}{dt} = \left(\beta - \frac{\beta I}{N} - \gamma\right) I$ 

Case 2:  $\beta < \gamma$  $\bullet \frac{dI}{dt} = \left(\beta - \frac{\beta I}{N} - \gamma\right)I$ 

Case 3:  $\beta > \gamma$  $\bullet \frac{dI}{dt} = \left(\beta - \frac{\beta I}{N} - \gamma\right) I$ 

#### Basic reproduction number

- Case 1:  $\beta = \gamma$ . Disease dies out.
- Case 2:  $\beta < \gamma$ . Disease dies out.
- Case 3:  $\beta > \gamma$ . Disease persists.
- In the SIS model, we call the ratio  $R_0 = \frac{\beta}{\gamma}$  the *basic* reproduction number of the system because it is the number of secondary infections  $\beta$  caused during the infectious period  $\frac{1}{\gamma}$ .
- When  $R_0 > 1$ , disease persists.
- When  $R_0 \leq 1$ , disease dies out.