

# Week 2: Compartmental Models & Expanded

Dr. Rachel Sippy University of Cambridge

#### Week 2 Overview

- Monday, August 2:
  - Relating SIR models to epidemic parameters
  - Estimating parameters in R
- Tuesday, August 3:
  - Guest lecture by Caroline Trotter
  - Modeling meningitis
  - Guided practice in R
- Thursday, August 5:
  - Using serological data for modeling
  - Guided practice in R

## Objectives

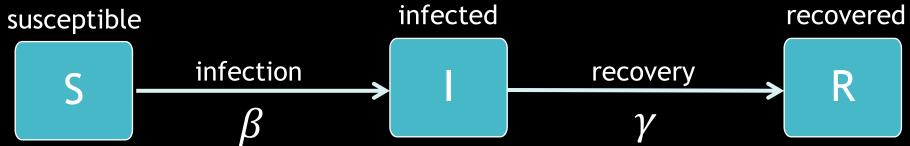
- Learn how to include births and deaths in an SIR model
- Modify model structure to create an SEIR model

# Post Questions in the Chat!

(or ask over microphone)

# Workshop Schedule

| Time         | Topics                             |
|--------------|------------------------------------|
| 2:00-2:10 pm | Greetings                          |
| 2:10-2:50 pm | Other Compartmental Models         |
| 2:50-3:00 pm | Break                              |
| 3:00-3:50 pm | Guest Lecture: Modeling meningitis |
| 3:50-4:00 pm | Break                              |
| 4:00-5:00 pm | R Session                          |



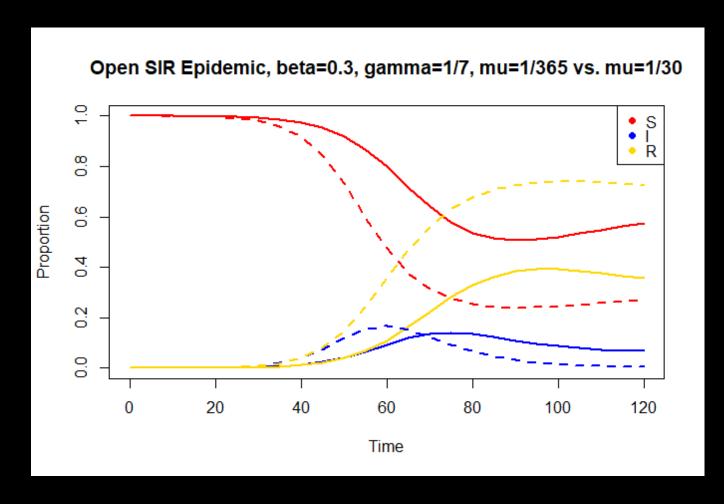
- this is a closed model
  - B is a transmission coefficient
  - γ is the recovery rate

$$\frac{dS(t)}{dt} = -\beta S(t)I(t)$$

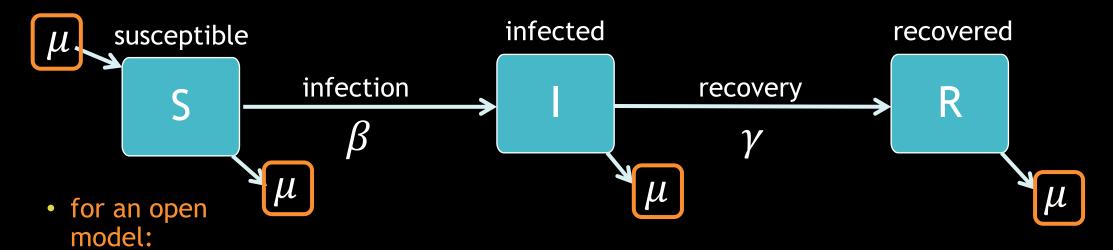
$$\frac{dI(t)}{dt} = \beta S(t)I(t) - \gamma I(t)$$

$$\frac{dR(t)}{dt} = \gamma I(t)$$

### SIR Model Output

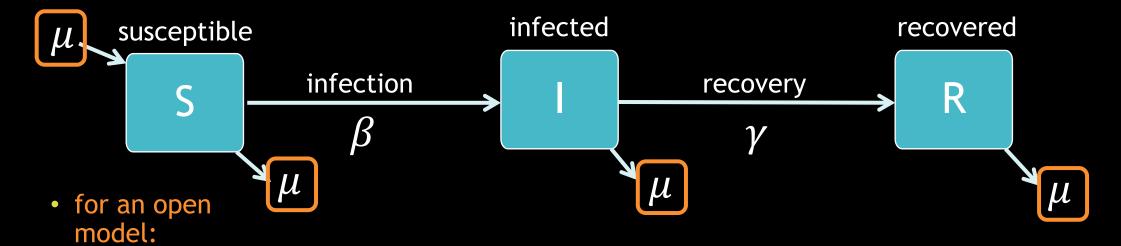


 Having an open population can greatly impact the epidemic dynamics!

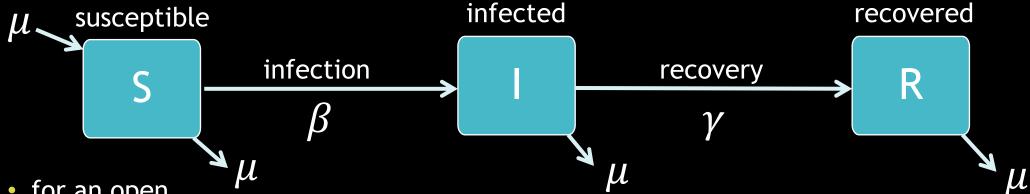


• µ is our birth/death rate

What are our assumptions?



- μ is our birth/death rate
- births are all susceptible
- anyone can die

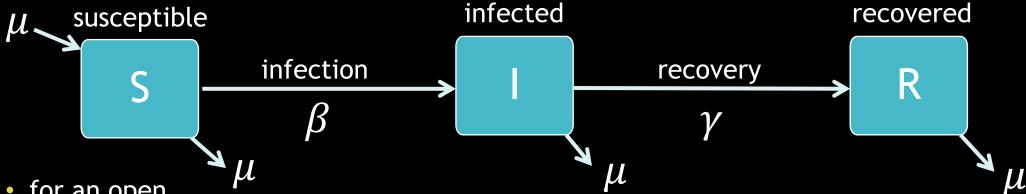


- for an open model:
  - μ is our birth/death rate
  - births are all susceptible but anyone can give birth
  - anyone can die

$$\frac{dS(t)}{dt} = \mu(S(t) + I(t) + R(t)) - \mu S(t) - \beta S(t)I(t)$$

$$\frac{dI(t)}{dt} = \beta S(t)I(t) - \gamma I(t) - \mu I(t)$$

$$\frac{dR(t)}{dt} = \gamma I(t) - \mu R(t)$$



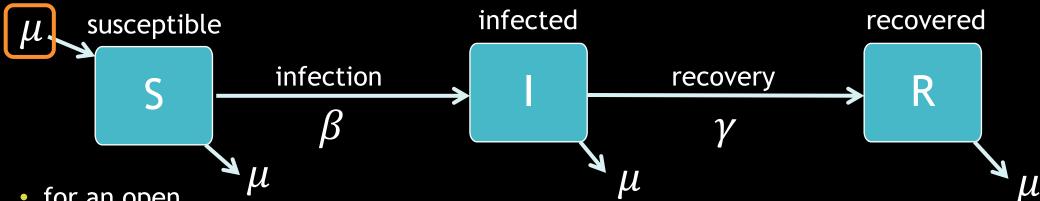
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$$\frac{dR(t)}{dt} = \underline{\gamma I(t)} - \mu R(t)$$

 we still have the same parts from the closed model

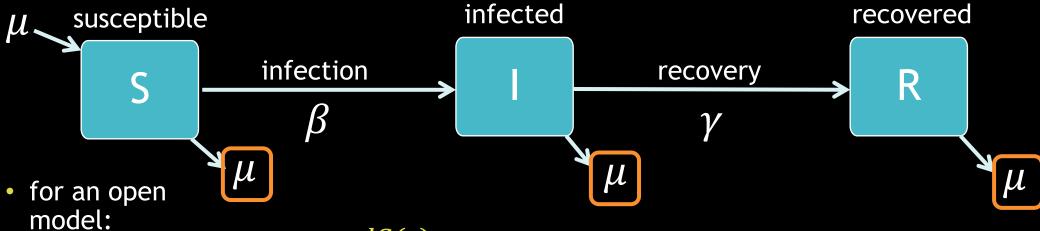


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#### R<sub>0</sub> & Model Structure

- The calculation of R0 depends on the structure of the model being used
- for our closed SIR model:
  - $R_0 = \frac{\beta}{\gamma}$

• for an open SIR model:

• 
$$R_0 = \frac{\beta}{\gamma + \mu}$$

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- for duration of infection:
  - $\frac{1}{\gamma}$

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- for duration of infection:
  - $\frac{1}{\gamma}$
- But usually  $\gamma \gg \mu$  so:

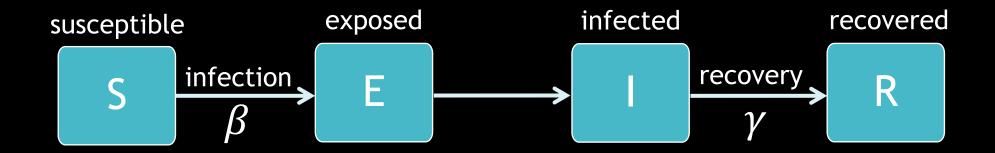
• 
$$R_0 = \frac{\beta}{\gamma} \approx R_0 = \frac{\beta}{\gamma + \mu}$$

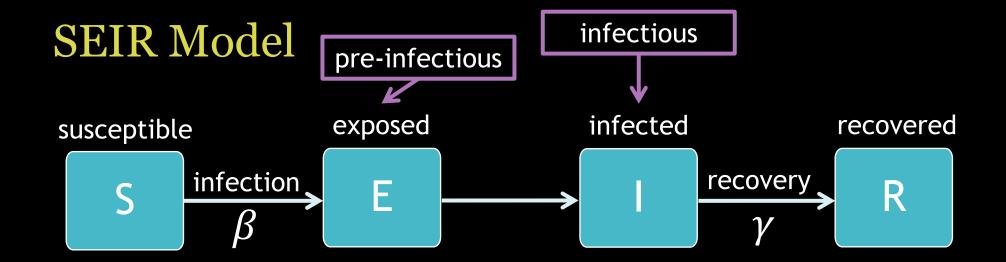
• for an open SIR model:

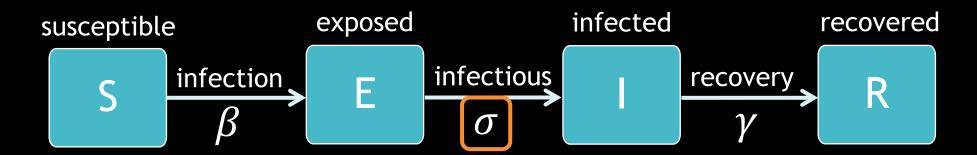
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$$R_0 = \frac{\beta}{\gamma + \mu}$$

for duration of infection:

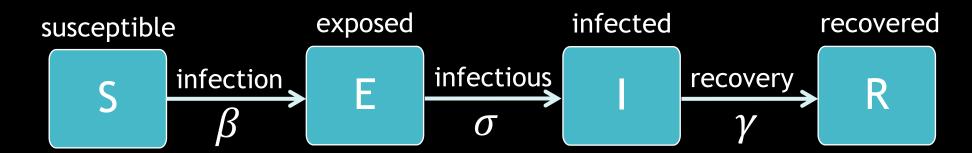
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$$\frac{1}{\gamma + \mu}$$







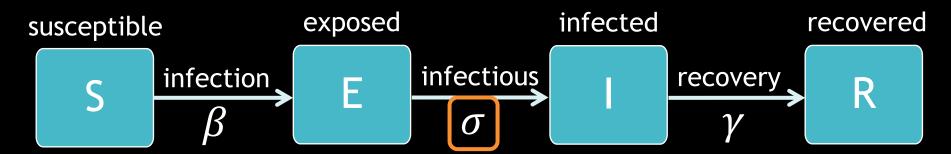
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  - σ is the rate at which people change from exposed to infectious
  - 1/σ is the latent period



- for an SEIR model:
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$$\frac{dS(t)}{dt} = -\beta S(t)I(t) \longleftarrow \bullet$$

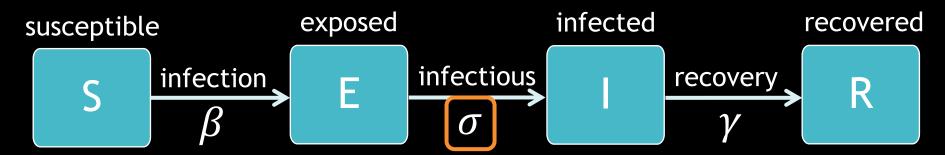
 the first equation is exactly the same



- for an SEIR model:
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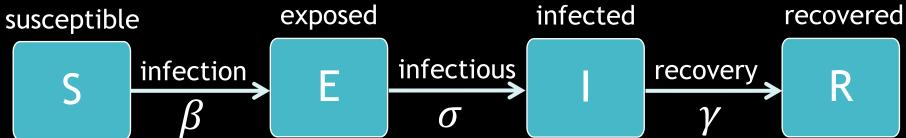


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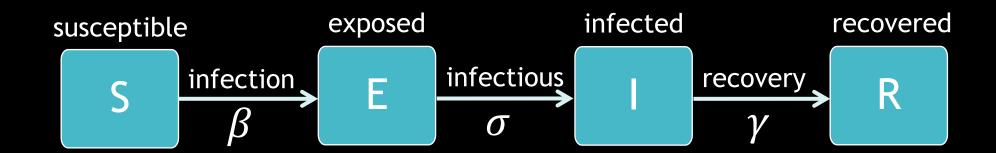
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$$\frac{dI(t)}{dt} = \sigma E(t) - \gamma I(t)$$

$$\frac{dR(t)}{dt} = \gamma I(t)$$
• the last equation is exactly the same



- We could also make the SEIR (or a model with any other structure) be open (include birth/death/migration)
  - we can also make different death rates for each compartment
  - models are extremely flexible!

# Questions?

10 minute break

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