

Foundations of dynamic modeling: The SIR Model Family

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DAIDD 2019

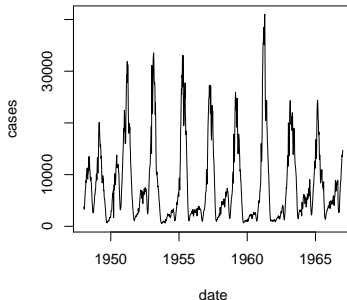
Goals

- ▶ This lecture will:
 - ▶ introduce the idea of dynamical modeling
 - ▶ explain why dynamical modeling is a key tool for understanding infectious disease
 - ▶ discuss and demonstrate simple dynamical models from the SIR model family
 - ▶ investigate some insights that can be gained from these models

Dynamical modeling connects scales



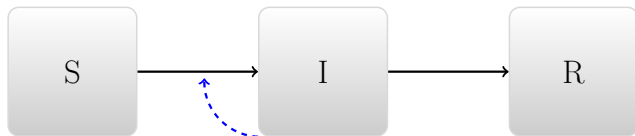
Measles reports from England and Wales



- ▶ Start with rules about how things change in short time steps
 - ▶ Usually based on *individuals*
- ▶ Calculate results over longer time periods
 - ▶ Usually about *populations*

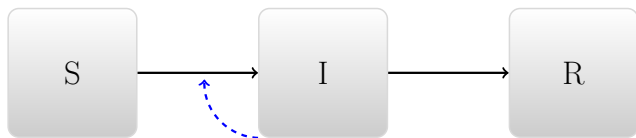
Compartmental models

Divide people into categories:



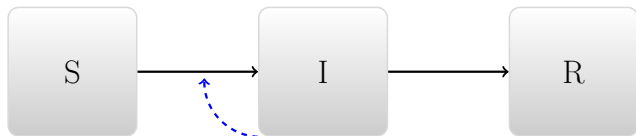
► Susceptible \rightarrow Infectious \rightarrow Recovered

What determines transition rates?



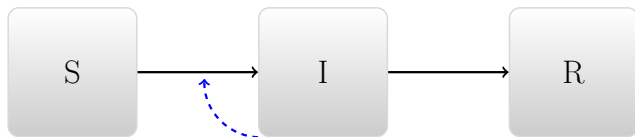
- ▶ People get better independently
- ▶ People get infected by infectious people

Conceptual modeling



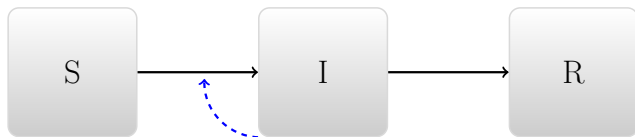
- ▶ What is the final result?
- ▶ When does disease increase, decrease?

Dynamic implementation



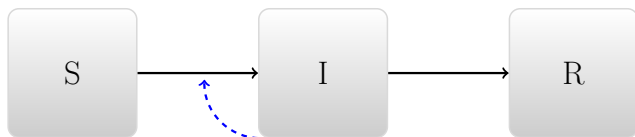
- ▶ Requires assumptions about recovery and transmission
- ▶ The *conceptually simplest* implementation uses Ordinary Differential Equations (ODEs)
 - ▶ Other options may be more realistic
 - ▶ Or simpler in practice

Recovery



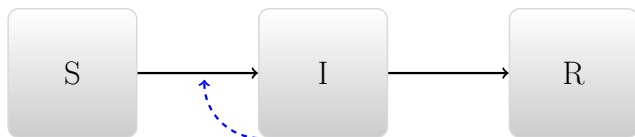
- ▶ Infectious people recover at *per capita* rate γ
 - ▶ Total recovery rate is γI
 - ▶ Mean time infectious is $D = 1/\gamma$

Transmission



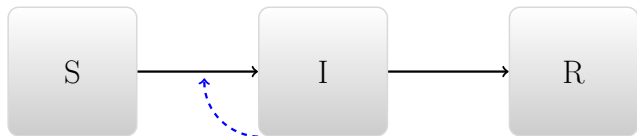
- ▶ Susceptible people get infected by:
 - ▶ Going around and contacting people (rate c)
 - ▶ Some of these people are infectious (proportion I/N)
 - ▶ Some of these contacts are effective (proportion p)
- ▶ Per capita rate of becoming infected is $cpI/N \equiv \beta I/N$
- ▶ Population-level transmission rate is $\mathcal{T} = \beta SI/N$

Another perspective on transmission



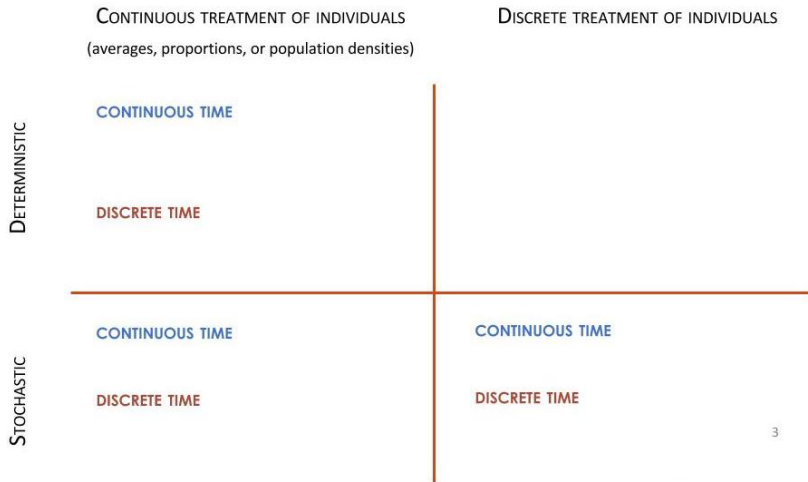
- ▶ Infectious people infect others by:
 - ▶ Going around and contacting people (rate c)
 - ▶ Some of these people are susceptible (proportion S/N)
 - ▶ Some of these contacts are effective (proportion p)
- ▶ Per capita rate of infecting others is $cpS/N \equiv \beta S/N$
- ▶ Population-level transmission rate is $\mathcal{T} = \beta SI/N$

ODE implementation



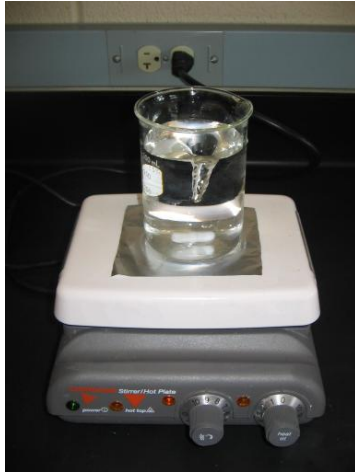
$$\begin{aligned}\frac{dS}{dt} &= -\beta \frac{SI}{N} \\ \frac{dI}{dt} &= \beta \frac{SI}{N} - \gamma I \\ \frac{dR}{dt} &= \gamma I\end{aligned}$$

Model taxonomy



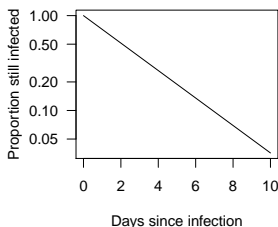
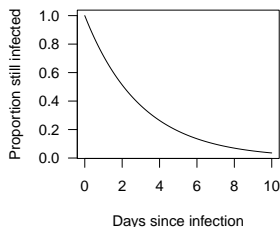
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ODE assumptions



- ▶ Lots and lots of people
- ▶ Perfectly mixed

ODE assumptions



- ▶ Waiting times are exponentially distributed
- ▶ Rarely realistic

More about transmission



- ▶ $\beta = pc$
 - ▶ What is a contact?
 - ▶ What is the probability of transmission?
- ▶ Sometimes this decomposition is clear
- ▶ But usually it's not

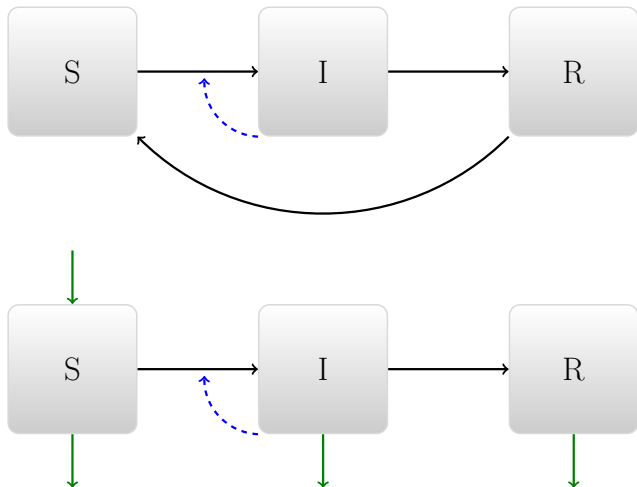
Population sizes

- ▶ How does β change with population size?
- ▶ We can make different assumptions about this
 - ▶ It may increase with population size, or not
- ▶ If population size changes we have to *consider* the question

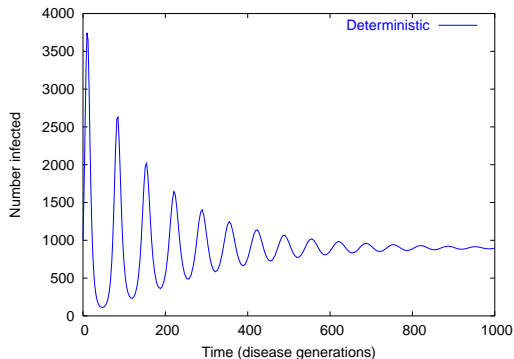
Digression – units

- ▶ $\mathcal{T} = \beta SI/N : [\text{ppl}/\text{time}]$
- ▶ $\beta : [1/\text{time}]$
 - ▶ The true β always has people in the numerator and the denominator
 - ▶ $\beta/\gamma = \beta D : [1]$

Closing the circle

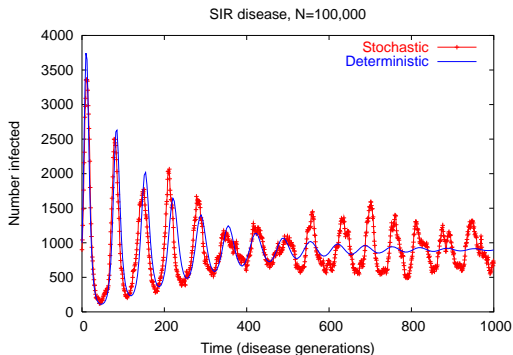


Tendency to oscillate



- ▶ Many susceptibles \rightarrow many infections \rightarrow few susceptibles \rightarrow few infections $\rightarrow \dots$
- ▶ Oscillations in simple models tend to be “damped”

With individuality



- ▶ Treating individuals as individuals can produce substantial oscillations even in large populations
- ▶ Interaction between random effects and the different time scales (of infection and recovery)

Summary

- ▶ Dynamic models are an essential tool because they allow us to link between scales
- ▶ There are many ways to construct and implement dynamic models
- ▶ Very simple models can provide useful insights
 - ▶ Reproductive numbers and thresholds
 - ▶ Tendency for oscillation (and tendency for damping)
- ▶ More complex models can provide more detail, but also require more assumptions, and more choices
- ▶ Understanding simple models can help guide our understanding of more complicated models



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