

(Hidden) assumptions of simple compartmental ODE models

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MMED 2019

Goals

- Review the main uses of applied epidemiological modelling
- Introduce our conceptual framework for applied modelling
- Review commonly overlooked assumptions that are inherent in the structure of simple compartmental ODE models
- Discuss when these assumptions might be problematic, and when they may be desirable
- Begin to explore some alternative model structures that relax the assumptions



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For application to the real world

The use of simplification to represent the key components of something you're trying to understand more clearly

Related to the distribution and determinants of healthrelated states and events





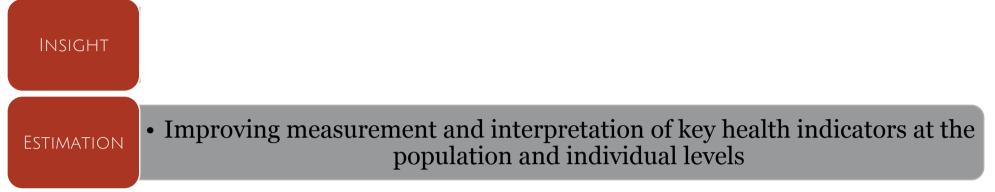


INSIGHT

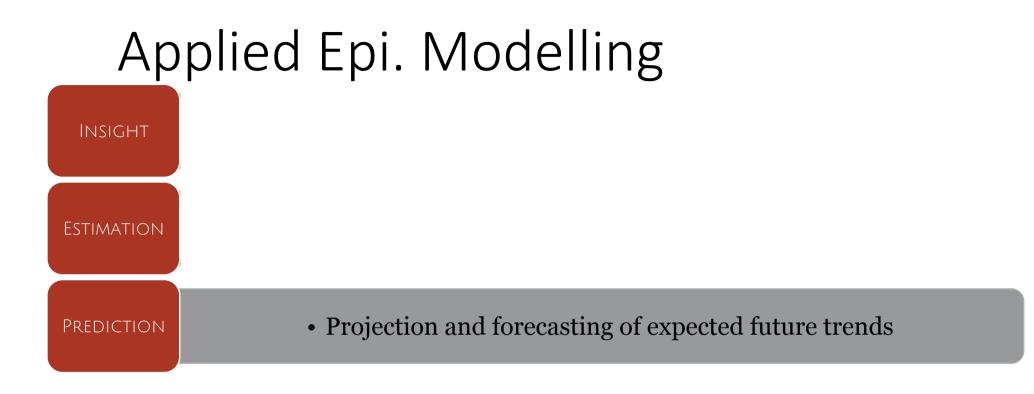
• Improving understanding of the dynamics of health and disease

• Translation of results into decision-making and communication tools

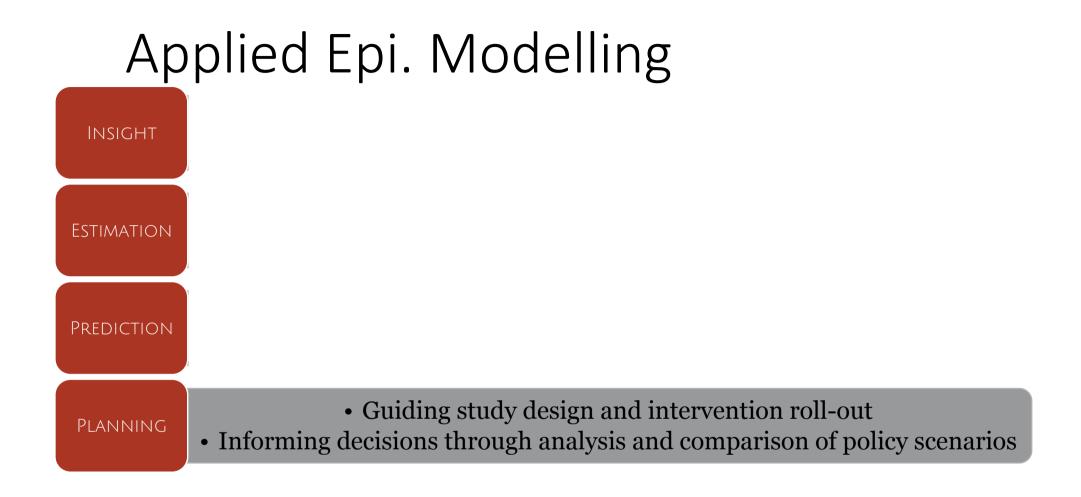




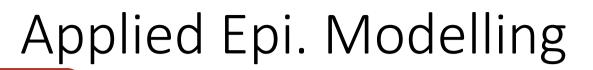


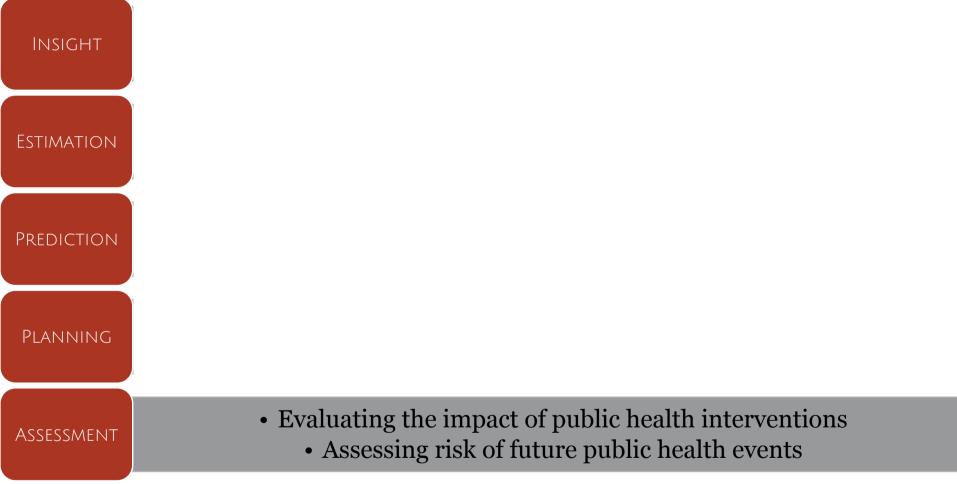














Insight	 Improving understanding of the dynamics of health and disease Translation of results into decision-making and communication tools
Estimation	• Improving measurement and interpretation of key health indicators at the population and individual levels
Prediction	• Projection and forecasting of expected future trends
Planning	 Guiding study design and intervention roll-out Informing decisions through analysis and comparison of policy scenarios
Assessment	 Evaluating the impact of public health interventions Assessing risk of future public health events

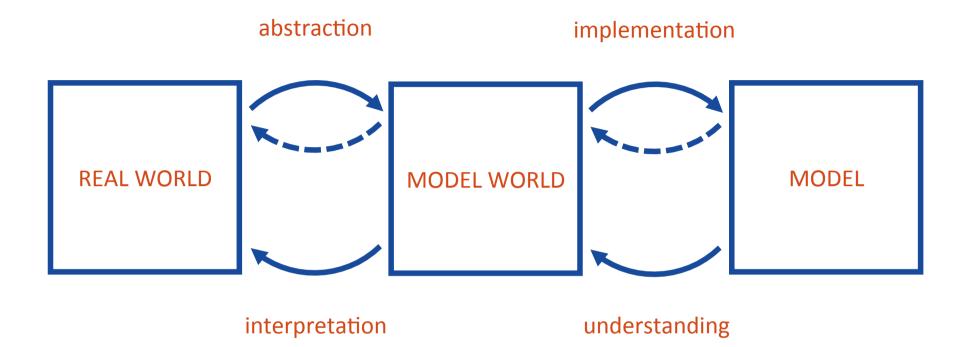


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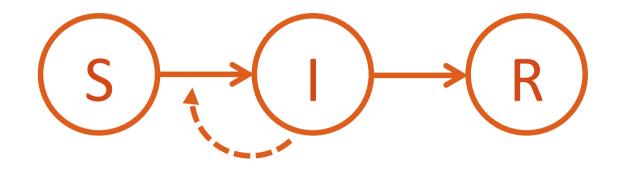


Model Worlds

- A model world is an abstraction of the world that is simple and fully specified, which we construct to help us understand particular aspects of the real world
- A mathematical model is formal description of the assumptions that define a model world
 - We know exactly what assumptions we've made, and we can follow those assumptions to their logical conclusions to address research questions



The SIR Model World





SIR: ODE Model

$$\int \lambda \longrightarrow R \qquad \lambda = \frac{\beta I}{N}$$

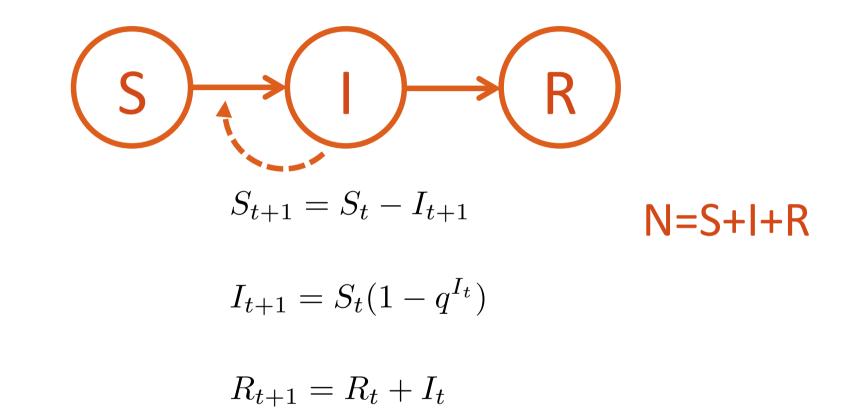
$$\frac{dS}{dt} = -\frac{\beta SI}{N}$$

$$\frac{dI}{dt} = \frac{\beta SI}{N} - \gamma I \qquad \qquad \frac{dR}{dt} = \gamma I$$



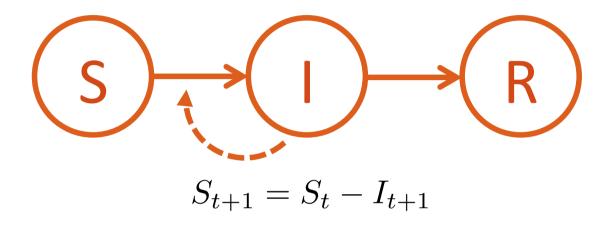
N=S+I+R

SIR: Reed-Frost Model





SIR: Stochastic Reed-Frost



$$\mathbb{P}(I_{t+1} = x) = \binom{S_t}{x} (1 - q^{I_t})^x (q^{I_t})^{S_t - x}$$

$$R_{t+1} = R_t + I_t$$



SIR: Chain Binomial Model

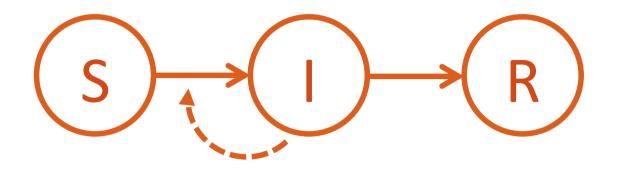
$$S \rightarrow (I) \rightarrow (R)$$

$$S_{t+\Delta t} = S_t - X$$
$$I_{t+\Delta t} = I_t + X - Y$$
$$R_{t+\Delta t} = R_t + Y$$

$$\mathbb{P}(X = x) = \binom{S_t}{x} p^x (1-p)^{S_t - x}$$
$$\mathbb{P}(Y = y) = \binom{I_t}{y} r^y (1-r)^{I_t - y}$$



The SIR Model Family



A mathematical model is formal description of the assumptions that define a model world





Taxonomy of **compartmental** models

CONTINUOUS TREATMENT OF INDIVIDUALS

(averages, proportions, or population densities)

CONTINUOUS TIME

- Ordinary differential equations
- Partial differential equations

DISCRETE TIME

Difference equations

(eg, Reed-Frost type models)

CONTINUOUS TIME

• Stochastic differential equations

DISCRETE TIME

Stochastic difference equations

CONTINUOUS TIME

Gillespie algorithm

DISCRETE TIME

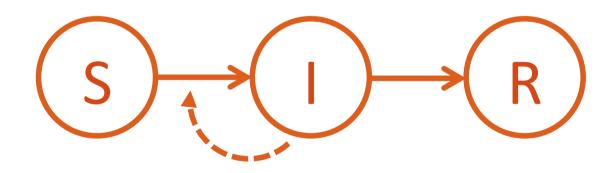
 Chain binomial type models (eg, Stochastic Reed-Frost models)

DISCRETE TREATMENT OF INDIVIDUALS



Taxonomy of **compartmental** models

What is a **compartmental** model?



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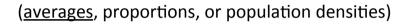


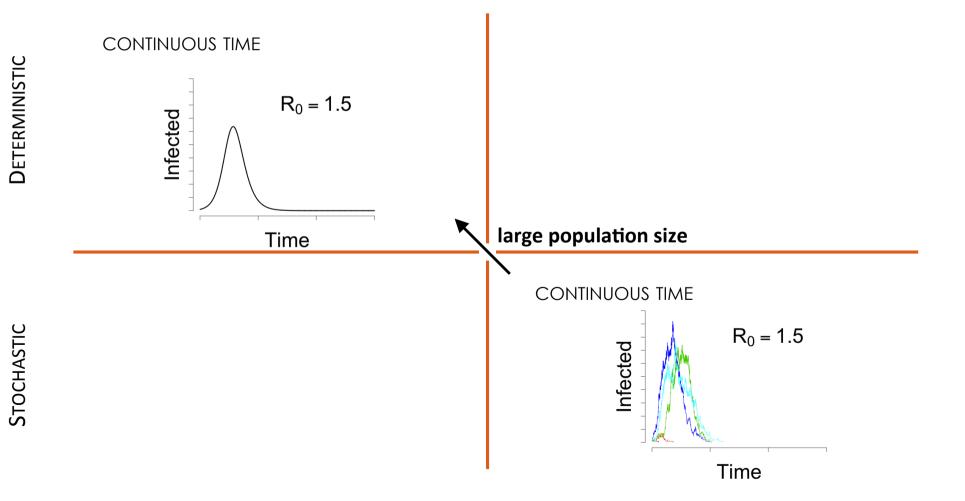


Taxonomy of **compartmental** models

CONTINUOUS TREATMENT OF INDIVIDUALS

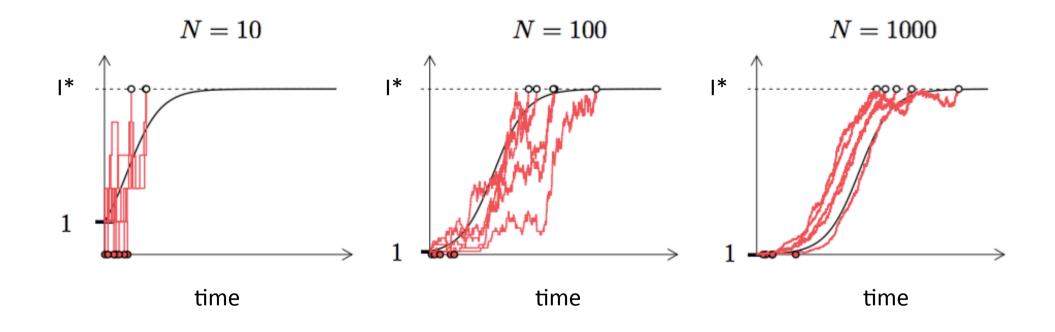
DISCRETE TREATMENT OF INDIVIDUALS







Demographic stochasticity



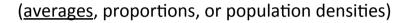
Borchering & McKinley (2018) *Multiscale Modeling and Simulation* DOI: 10.1137/17M1155259

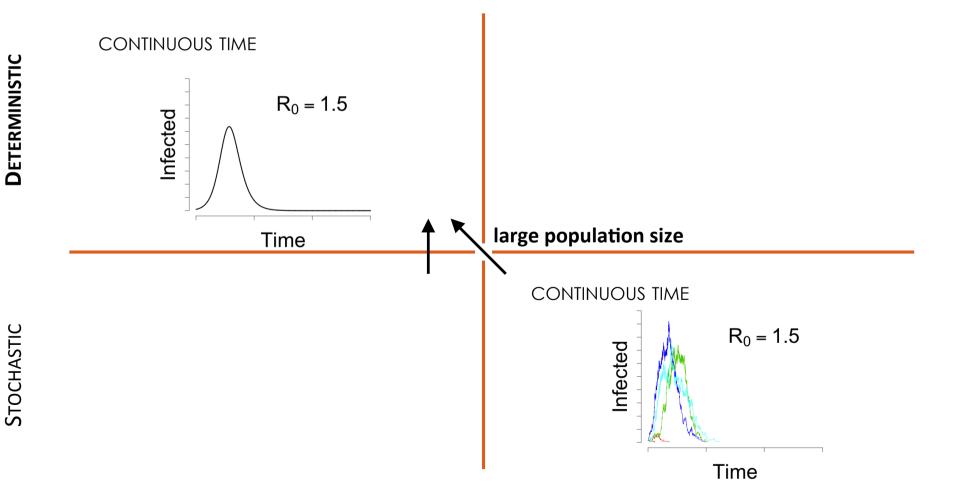


Taxonomy of **compartmental** models

CONTINUOUS TREATMENT OF INDIVIDUALS

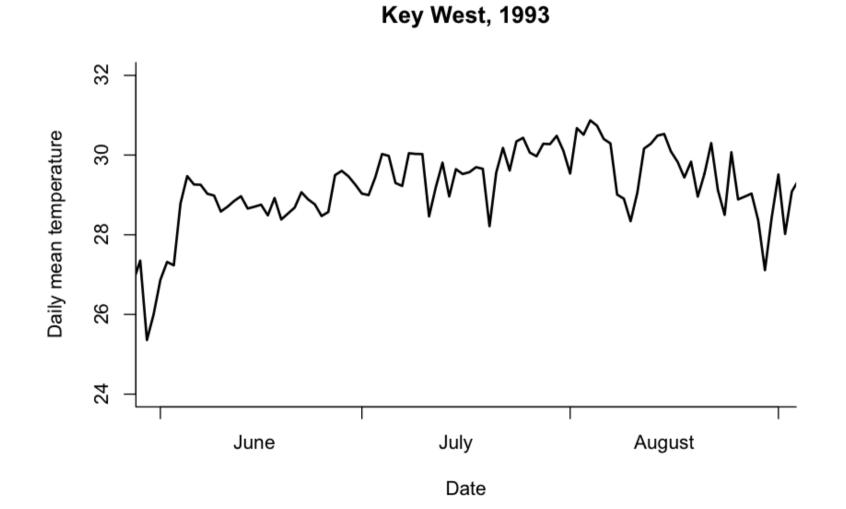
DISCRETE TREATMENT OF INDIVIDUALS







Environmental stochasticity



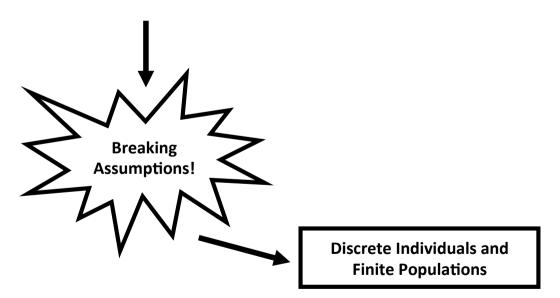
Compartmental ODE models assume

- Large population size
- Deterministic progression
 - for a given set of initial conditions and parameter values, a deterministic model always gives the same outcome
- Quickly assess what model outcomes are possible, and when



Compartmental ODE models assume

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 - for a given set of initial conditions and parameter values, a deterministic model always gives the same outcome







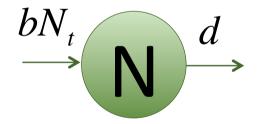
Taxonomy of **compartmental** models

CONTINUOUS TREATMENT OF INDIVIDUALS (averages, proportions, or population densities)

DISCRETE TREATMENT OF INDIVIDUALS

CONTINUOUS TIME

Ordinary differential equations



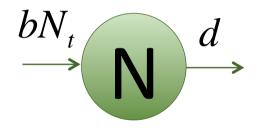
STOCHASTIC

Simple ODE models assume

- Time proceeds in a continuous manner
- Parameter values remain constant

CONTINUOUS TIME

Ordinary differential equations



$$\frac{dN_t}{dt} = bN_t - dN_t$$



Simple ODE models assume

- Time proceeds in a continuous manner
- Parameter values remain constant

CONTINUOUS TIME

Ordinary differential equations

$$N_{t+\Delta t} = N_t e^{r\Delta t}$$





CONTINUOUS TIME

• Ordinary differential equations

$$bN_t \longrightarrow d \qquad r = b - d \qquad \frac{dN}{dt} = rN$$

$$N_{t+\Delta t} = N_t e^{r\Delta t}$$

DISCRETE TIME

• Difference equations

$$\lambda_{\Delta t} = e^{r\Delta t}$$

 $N_{t+\Delta t} = \lambda_{\Delta t} N_t$

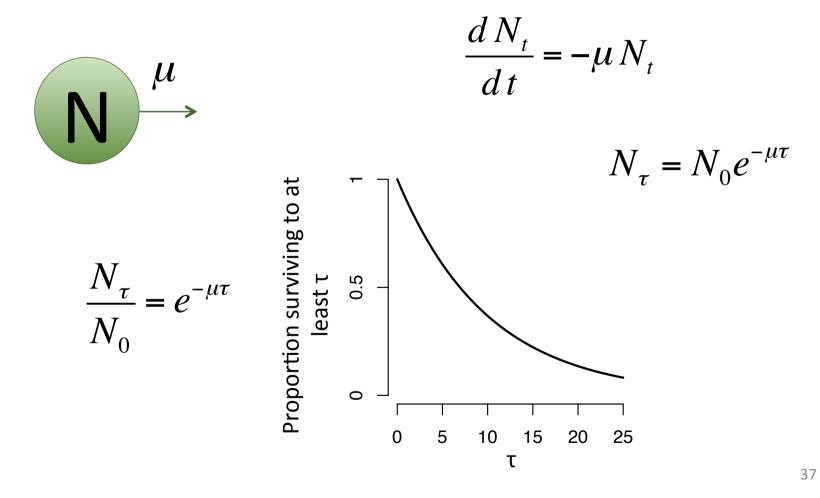
Simple compartmental ODE models assume

- Homogeneity within compartments
- Large population size
- Deterministic progression
- Time proceeds in a continuous manner
- Parameter values remain constant
- Memory-less processes



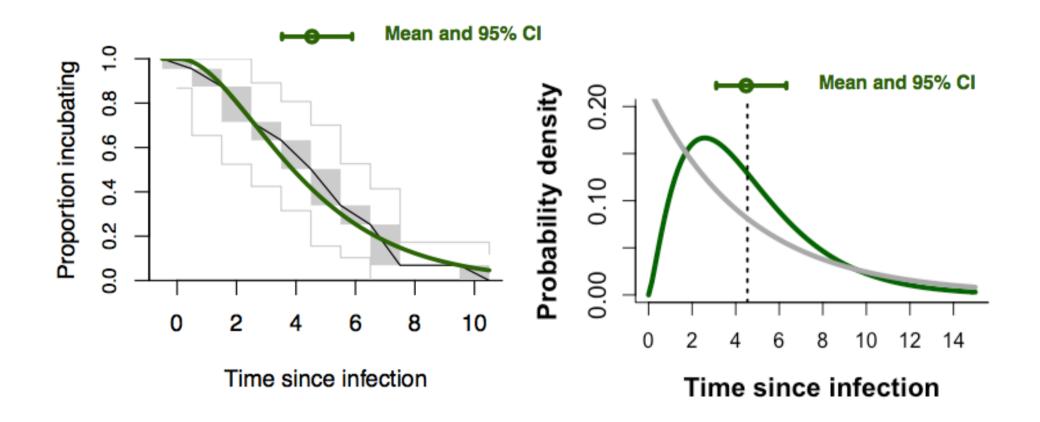
Simple ODE models assume

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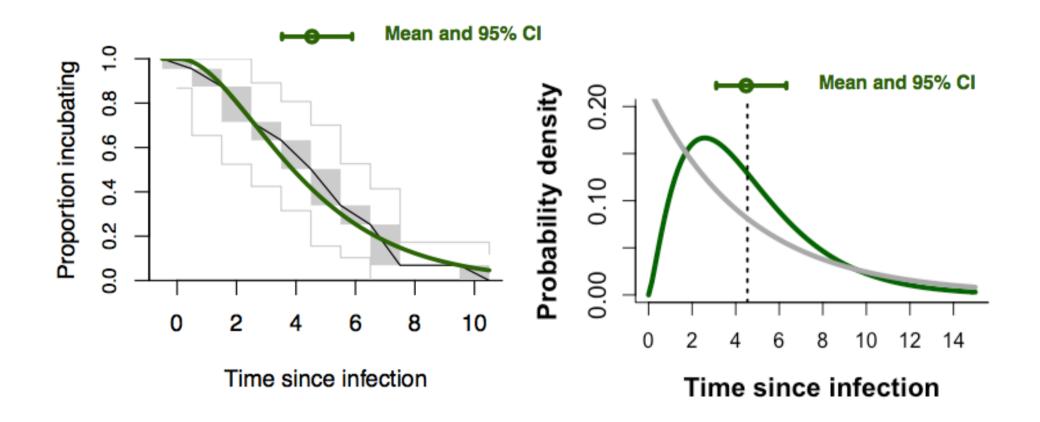
Realistic waiting times



³⁸ Estimate based on data from Joshi *et al.* (2009) *Transactions of the Royal Society of Tropical Medicine and Hygiene*



Realistic waiting times

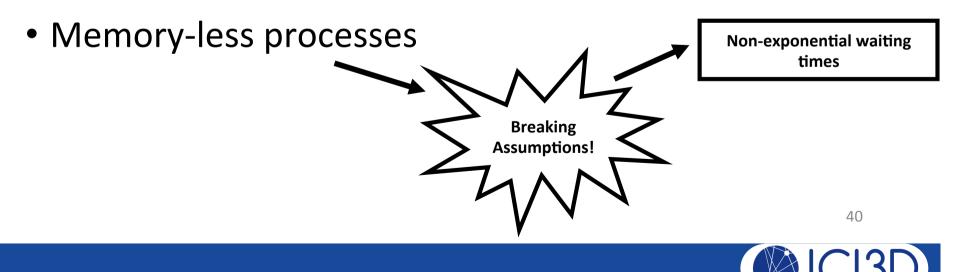


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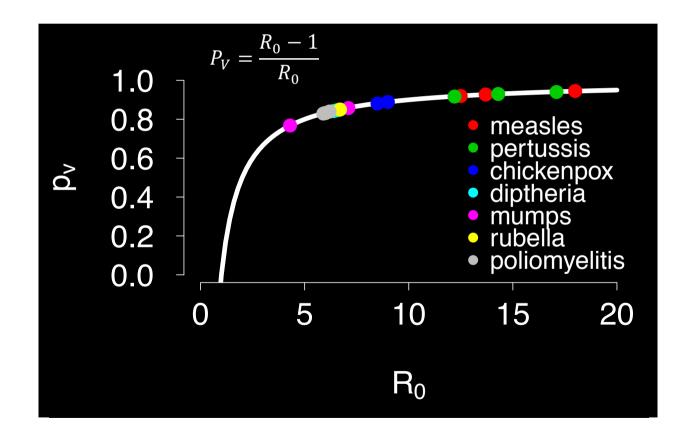
ICI3D

Simple compartmental ODE models assume

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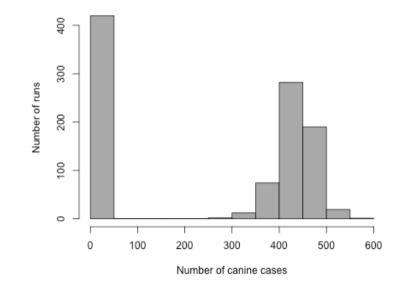


 Simple ODE models are important tools for building understanding



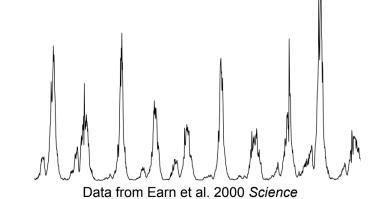


- Simple ODE models are important tools for building understanding
- It's important to recognize the assumptions built into these models
 - When populations are small, average behaviors can be misleading





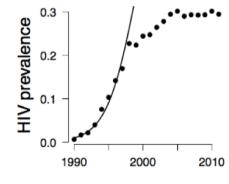
- Simple ODE models are important tools for building understanding
- It's important to recognize the assumptions built into these models
 - When populations are small, average behaviors can be misleading
 - When rates vary, simple ODEs can fail to reproduce important (observed) dynamics



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CISD)

- Simple ODE models are important tools for building understanding
- It's important to recognize the assumptions built into these models
 - When populations are small, average behaviors can be misleading
 - When rates vary, simple ODEs can fail to reproduce important (observed) dynamics



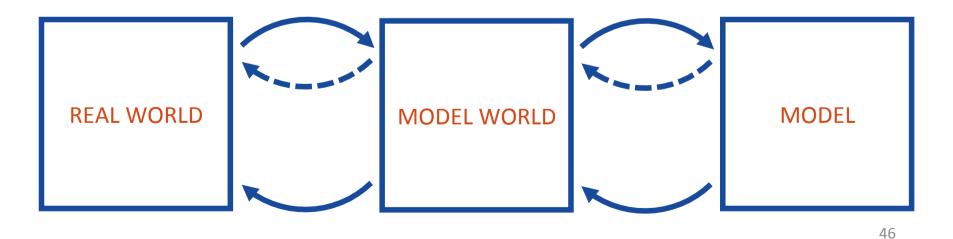
Dushoff lecture on heterogeneity (Wed)



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 - When populations are small, average behaviors can be misleading
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- The applied epidemiological modelling process requires
 - abstraction
 - specification and implementation
 - gaining an understanding of the dynamics
 - interpretation





UNIVERSITY OF GEORGIA

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(Hidden) assumptions of simple compartmental ODE models. DOI: <u>10.6084/m9.figshare.5044606</u>

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https://figshare.com/articles/Mathematical_Assumptions_of_Simple_ODE_models/5044606

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