# Informing Policy via Dynamic Models: Cholera in Haiti

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#### Introduction

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# **Advantages of Statistical Modeling**

- Nonlinear-dynamic statistical models have proven to be a useful tool for modeling infectious disease outbreaks (TODO: CITE)
- Most common examples are SIR models and their various extensions.
- These models enable the modeling of scientifically meaningful states, prediction of the future of the outbreak, and modeling the potential effects of interventions (such as vaccinations) (TODO: CITE)

#### Concerns

- Despite their utility, there exist many cautionary warnings against the use of these types of models (TODO: CITE).
- Concerns include:
  - TODO
  - TODO
  - TODO
- Despite these warnings, there is very little practical advice on how to approach these issues.

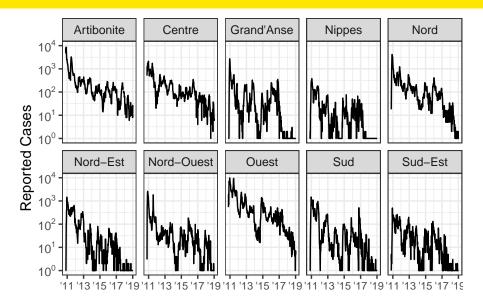
#### Cholera in Haiti

- We consider the 2010-2019 cholera outbreak in Haiti.
- Cholera was introduced to Haiti in 2010 following the devastating earthquake of the same year.
- Although some new cases have been detected, there were no recorded cholera cases in Haiti between February, 2019 and September 2022 (TODO: Cite).

#### Data



#### Data



#### **Models**

• We build on the study by Lee et. al (2020), in which four independent teams built non-linear models to describe cholera dynamics.

	Model 1	Model 2	Model 3	Model 4
Deterministic / Stochastic	Stochastic	Deterministic	Stochastic	Agent Based
Spatial Model	No	Yes	Yes	Yes
Fitting Method	IF2	Trajectory Matching	PIF / IF2	NA

# **Model Fitting**

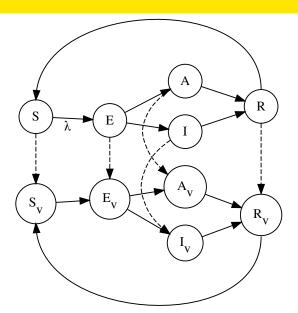
## Computing

- Parameters of nonlinear dynamic models are often fit by finding their posterior distribution or by maximizing some objective measure.
- Because of the non-linear nature of the models, this can be computationally expensive.
- Great care should be taken to determine the necessary amount of computation needed to solve the problem at hand.

#### **Nested Models**

- Non-linear dynamic models make assumptions about the dynamics of the system in question.
- Consider testing scientifically meaningful nested hypothesis.
- For example, we consider adding a linear trend in transmission to
   Model 1 in order to account for the apparent decrease in cholera cases.

## Model 1



## Model 1 (Continued...)

• Individuals move  $S \to E$  at time t with a rate of  $\lambda(t)$ , where:

$$\lambda(t) = (I + \epsilon A)^{\nu} \frac{d\Gamma(t)}{dt} \beta(t)/N,$$

$$\log \beta(t) = \sum_{j=1}^{6} \beta_j s_j(t) + \frac{\xi \bar{t}}{t}$$

•  $\frac{d\Gamma}{dt}$  is multiplicative Gamma white-process noise,  $\epsilon, \nu, \xi, \beta_{1:6}$  are parameters to be estimated,  $s_{1:6}(t)$  are a B-spline basis.

# **Model Diagnostics**

## Don't just rely on simulations

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#### For me

#### **Model Forecasts**

## Reproducibility