

# Thesis

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## 1 Introduction

Measles is a childhood disease  
TSIR.

$$\begin{aligned} S_{t+1} &= S_t + B_t - I_{t+1} \\ I_{t+1} &= \beta_t I_t^\alpha S_t / N_t \end{aligned} \tag{1}$$

## 2 Susceptible reconstruction

Recall that the susceptible population dynamics can be written as follows:

$$S_{t+1} = S_t + B_t - I_{t+1}. \tag{2}$$

Rewriting incidence as product of observed cases,  $C_{t+1}$ , divided by the reporting rate,  $\rho_{t+1}$ , we obtain the following:

$$S_{t+1} = S_t + B_t - \frac{C_{t+1}}{\rho_{t+1}}. \tag{3}$$

Iteratively solving the equation yields the following relation:

$$S_{t+1} = S_1 + \sum_{i=1}^t B_i - \sum_{i=1}^t \frac{C_{i+1}}{\rho_{i+1}}. \tag{4}$$

Representing susceptible dynamics as a deviation from the mean number of susceptibles  $S_t = \bar{S} + Z_t$ , Finkenstadt and Grenfell (2000) showed dynamics of  $Z_t$  and  $\rho_t$  can be recovered from the data by fitting a smooth curve to cumulative birth as a function of cumulative cumulative cases. Dalziel et al. (2016) then claimed that the result can be generalized by representing susceptible dynamics as a deviation from moving average,  $S_t = \sigma N_t + W_t$ , where  $\sigma$  is proportion susceptible:

$$\sum_{i=1}^t B_i = \sum_{i=1}^t \frac{C_{i+1}}{\rho_{i+1}} + W_{t+1} - W_1 + \sigma(N_{t+1} - N_1). \tag{5}$$

However, they did not provide sufficient theoretical justification; here, we analyze the claimed generalization.

### Constant reporting rate

First, consider the case when the reporting is constant ( $\rho_i = \rho$  for all  $\rho \in \mathbb{N}$ ), equation 5 simplifies to

$$\sum_{i=1}^t B_i = \frac{1}{\rho} \sum_{i=1}^t C_{i+1} + W_{t+1} - W_1 + \sigma(N_{t+1} - N_1). \quad (6)$$

Writing  $X_{t+1} = \sum_{i=1}^t C_{i+1}$  and  $Y_{t+1} = \sum_{i=1}^t B_i$ , we obtain a familiar form:

$$Y_{t+1} = \frac{1}{\rho} X_{t+1} + W_{t+1} - W_1 + \sigma(N_{t+1} - N_1), \quad (7)$$

When population size stays constant ( $N_t = N_1$  for all  $t \in \mathbb{N}$ ), the equation simplifies to

$$Y_{t+1} = \frac{1}{\rho} X_{t+1} + W_{t+1} - W_1, \quad (8)$$

which suggests that linear regression can be used to estimate the reporting rate and susceptible dynamics (Finkenstadt and Grenfell, 2000). In general, one can write

$$N_{t+1} = N_t + B_t - D_t,$$

where  $D_t$  is the number of individuals that died between time  $t$  and  $t+1$ . Then, we have

$$Y_{t+1} = \frac{1}{\rho} X_{t+1} + W_{t+1} - W_1 + \sigma \left( \sum_{i=1}^t B_i - \sum_{i=1}^t D_i \right). \quad (9)$$

Writing  $M_t = \sum_{i=1}^t D_i$ , we get

$$(1 - \sigma)Y_{t+1} = \frac{1}{\rho} X_{t+1} + W_{t+1} - W_1 + \sigma M_t, \quad (10)$$

which suggests that a linear regression is likely to overestimate susceptible dynamics and underestimate reporting rate.

### Non-constant reporting rate

Consider the following equation:

$$\sum_{i=1}^t B_i = \sum_{i=1}^t \frac{C_{i+1}}{\rho_{i+1}} + W_{t+1} - W_1 + \sigma(N_{t+1} - N_1). \quad (11)$$

Define  $\hat{\rho}$  such that  $1/\hat{\rho} = E[1/\rho_t]$ . Then,

$$Y_{t+1} = \frac{1}{\hat{\rho}} X_{t+1} + \sum_{i=1}^t \left( \frac{1}{\rho_{i+1}} - \frac{1}{\hat{\rho}} \right) C_{i+1} + W_{t+1} - W_1 + \sigma(N_{t+1} - N_1). \quad (12)$$

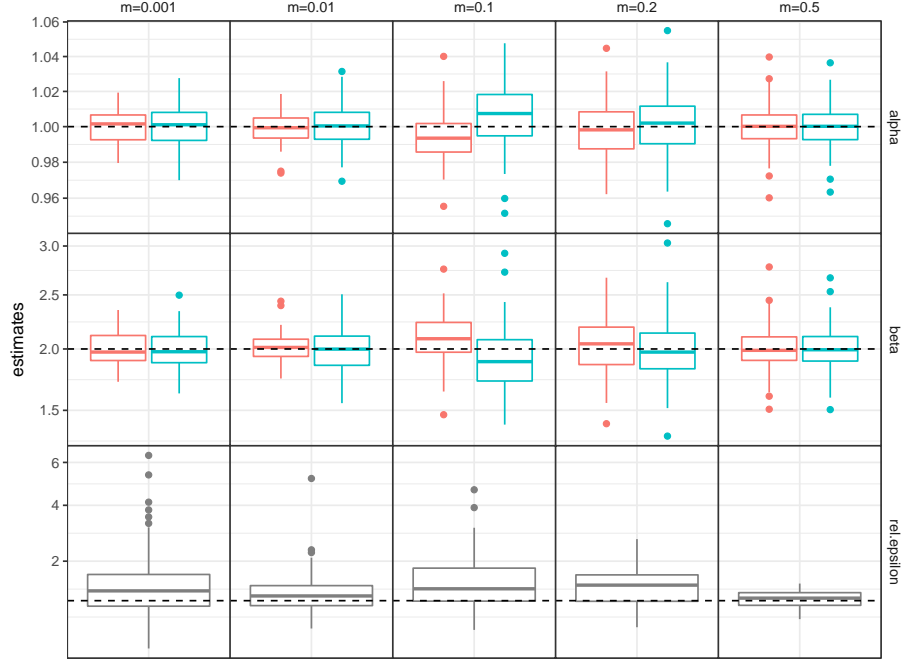


Figure 1: **SIR model** TODO: caption

Let

$$R_{t+1} = \sum_{i=1}^t \left( \frac{1}{\rho_{i+1}} - \frac{1}{\hat{\rho}} \right) C_{i+1}.$$

Then, we obtain an expression that generalizes the result derived by Finkenstadt and Grenfell (2000) by accounting for changes in population sizes.

$$\begin{aligned} Y_{t+1} &= \frac{1}{\hat{\rho}} X_{t+1} + \left( \frac{1}{\rho_{t+1}} - \frac{1}{\hat{\rho}} \right) C_{t+1} + R_t + W_{t+1} - W_1 + \sigma(N_{t+1} - N_1) \\ &= \frac{1}{\rho_{t+1}} X_{t+1} + \left( \frac{1}{\hat{\rho}} - \frac{1}{\rho_{t+1}} \right) X_t + R_t + W_{t+1} - W_1 + \sigma(N_{t+1} - N_1) \end{aligned} \quad (13)$$

### 3 Spatial coupling

### 4 Case study: measles

## References

Dalziel, B. D., O. N. Bjørnstad, W. G. van Panhuis, D. S. Burke, C. J. E. Metcalf, and B. T. Grenfell (2016). Persistent chaos of measles epidemics in the

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