



## Introduction to Fitting an SIR model practical

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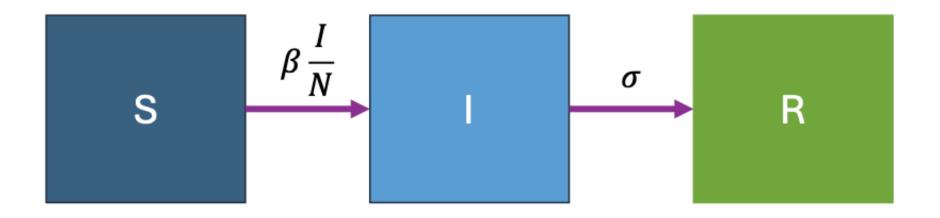
Modern Statistics and Machine Learning for Population Health in Africa

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### Fitting two different models: SIR model



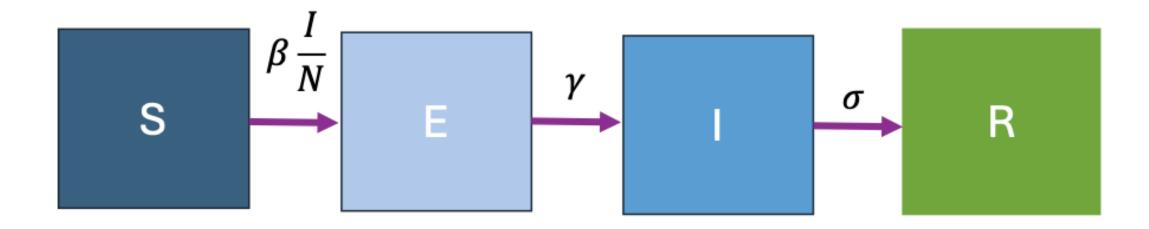
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## Fitting two different models: SEIR model



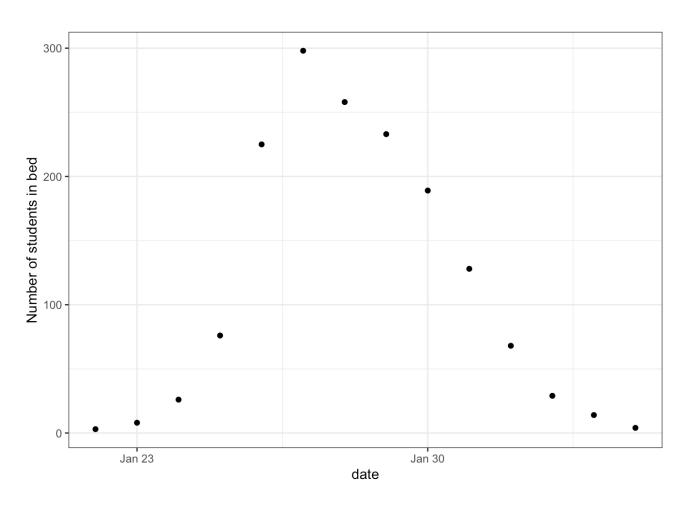
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#### To two types of data: Prevalence



We examine an outbreak of influenza A (H1N1) in 1978 at a British boarding school. The data consists of the daily number of students in bed (prevalence data), spanning over a time interval of 14 days.

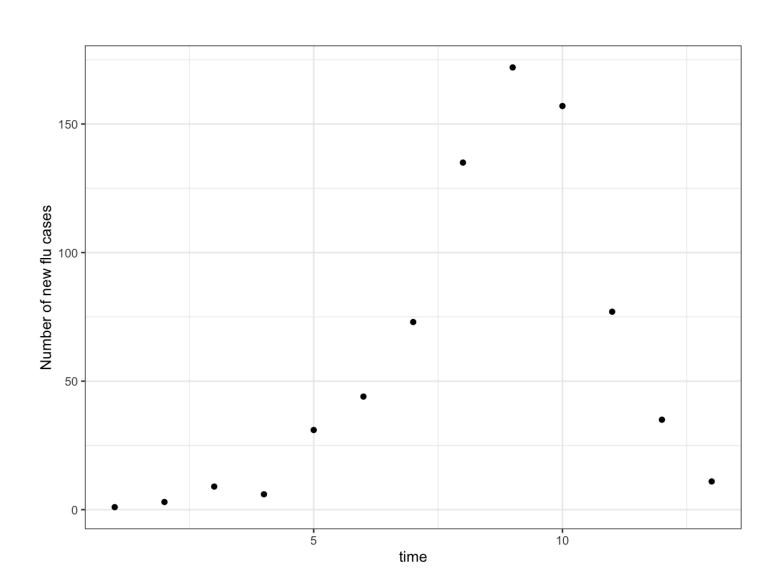


### To two types of data: Incidence

MRC Centre for Global Infectious Disease Analysis

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New cases of flu each day



#### Stan model file

```
functions {
  vector sir(real t,
             vector y,
             real beta,
             real sigma,
             real N) {
      vector[3] dydt;
      real S = y[1];
      real I = y[2];
      real R = y[3];
      dydt[1] = -beta * I * S / N;
      dydt[2] = beta * I * S / N - sigma * I;
      dvdt[3] = sigma * I;
      return dydt;
data {
parameters {
transformed parameters{
model {
generated quantities {
```



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#### Fitting the model in python



```
1 import numpy as np
3 # Set the population size
4 N = 763
6 # Define initial conditions
7 i0 = 1
8 	 s0 = N - i0
9 r0 = 0
10 y0 = [s0, i0, r0]
12 # Make the data struture
13 stan_data = {
"n_days": len(df.in_bed),
15 "y0": y0,
16 "t0": 0,
17 "t": np.arange(1, len(df.in_bed)+1),
18 "N": N,
19 "cases": df.in_bed
20 }
```

Then we compile our stan model.

```
1 sir_model = CmdStanModel(stan_file = 'sir_model.stan')
2
```

Finally we run MCMC. For this problem, it suffices to use Stan's defaults. Note that, as is standard practice, we run 4 Markov chains.