

# SF2955 Homework 2

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## 1 Introduction

In this report a basic epidemic model whose unknown model parameters will be estimated by Bayesian inference. The posterior will be a complex probability distribution, which will be sampled from using a Markov chain Monte Carlo (MCMC) algorithm.

## 2 Model description

The epidemic is assumed to be spreading in a population of  $P$  individuals. Using a SIR model, a discrete-time stochastic process  $(S_t, I_t, R_t)_{t \in N}$ , where  $S_t$  is individuals who were not infected before day  $t$ ,  $I_t$  is the number of individuals that are infected at the beginning of day  $t$ ,  $R_t$  is the number of individuals that are removed at the beginning of day  $t$ , either recovered, become immune, or died before day  $t$ . The population is assumed to be isolated, each day  $t$ ,  $P = S_t + I_t + R_t$ , so that the process can be described by two states only.

The outbreak starts with an initial number of  $I_0$  cases, so the initial number that are not infected  $S_0 = P - I_0$ . During day  $t$ , an additional number  $\Delta I$  of individuals become infected, which means the number of  $S_t$  decreases the same amount. An additional number  $\Delta R$  either recover or die. So the number of infected individuals increase by  $\Delta I$  and decrease with  $\Delta R$ .

$$\begin{cases} S_{t+1} = S_t - \Delta_t^I, \\ I_{t+1} = I_t + \Delta_t^I - \Delta_t^R, \\ R_{t+1} = R_t + \Delta_t^R, \end{cases} \quad t \in N \quad (1)$$

With a constant probability of  $p$ , the distribution of the flow of  $R_t$  is:

$$\Delta_t^R \sim \text{Bin}(I_t, p^{i \rightarrow r}) \quad (2)$$

The probability that someone becomes infected on day  $t$  is:

$$p_t^{s \rightarrow i} = 1 - \exp(-\lambda(t) \frac{I_t}{P}) \quad (3)$$

The variable  $\lambda(t) > 0$  is dependent on the average number of interactions per individual on day  $t$ .

$$\lambda_t^I \sim \text{NegBin}(\kappa, \varphi) \quad (4)$$

With parameters  $\varphi \in (0, 1)$  and;

$$\kappa = (\frac{1}{\varphi} - 1) S_t p_t^{s \rightarrow i} \quad (5)$$

Throughout this assignment,  $\varphi = 0.995$ . The time frame is divided into  $d$  intervals, where  $\lambda(t)$  is constant.

$$0 = t_0 < t_1 \dots < t_d = T \quad (6)$$

$$\lambda(t) = \sum_{i=1}^{d-1} \lambda_i 1_{t_{i-1}, t_i}(t) + \lambda_d 1_{t_{d-1}, d_i}(t) \quad (7)$$

### 3 Problem 1

Given a parameter vector  $\theta = (\lambda, t, p^{i \rightarrow r})$ , the Markov chain  $(S_t, I_t)_{t \in N}$  transition probabilities are determined:

$$q_0(s_t, i_t; s_{t+1}, i_{t+1}) = P_0(S_{t+1} = s_{t+1}, I_{t+1} = i_{t+1} \mid S_t = s_t, I_t = i_t) \quad (8)$$

Using equation (1) it can be rewritten as follows:

$$P_0(s_t - \Delta_t^I = s_{t+1}, i_t + \Delta_t^I - \Delta_t^R = i_{t+1}) \quad (9)$$

$$P_0(\Delta_t^I = s_t - s_{t+1}) P(\Delta_t^I - \Delta_t^R = i_{t+1} - i_t \mid \Delta_t^I = s_t - s_{t+1}) \quad (10)$$

$$P_0(\Delta_t^I = s_t - s_{t+1}) P(\Delta_t^R = i_t - i_{t+1} + s_t - s_{t+1}) \quad (11)$$

## 4 Prior distributions

In order to provide a Bayesian model, a prior distribution of  $\pi(\theta)$  for  $\theta$  needs to be specified. The parameters are assumed to be priori independent,  $\pi(\theta) = \pi(\lambda)\pi(t)\pi(p^{i \rightarrow r})$ . First  $t$  is assigned a flat prior over the set of breakpoints, the components of  $\lambda$  are priori independent with  $\Gamma(\alpha_i, \beta_i)$ ,

$$\pi(\lambda) = \prod_{i=1}^d \frac{\beta_i^{\alpha_i}}{\Gamma(\alpha_i)} \lambda_i^{\alpha_i-1} e^{-\beta_i \lambda_i} \quad (12)$$

Throughout the assignment,  $\alpha_i = 2$ . Finally, Beta is assigned prior:

$$\pi(p^{i \rightarrow r}) = \frac{1}{B(a,b)} (p^{i \rightarrow r})^{a-1} (1 - p^{i \rightarrow r})^{b-1} \quad (13)$$

## 5 Data

In order to calibrate the model, COVID-19 data from Germany and Iran will be used. The parameters  $\theta$  will be estimated on the basis of two realizations,  $(i_t, r_t)_{t=0}^T$  and  $(I_t, R_t)_{t=0}^T$

## 6 Problem 2

Determining the likelihood, where  $P(S_0 = s_0, I_0 = i_0 \mid \theta) = 1$ :

$$f(y \mid \theta) = P(S_0 = s_0, I_0 = i_0, \dots, S_T = s_T, I_T = i_T \mid \theta) = \quad (14)$$

Using chain rule:

$$P(S_0, I_0)P(S_1, I_1 \mid S_0, I_0)(S_2, I_2 \mid S_0, I_0, S_1, I_1) \dots P(S_{n+1}, I_{n+1} \mid S_n, I_n, S_{n-1}, I_{n-1}, \dots, S_0, I_0) = \quad (15)$$

Simplifying the equation:

$$\prod_{i=0}^n P(S_{n+1}, I_{n+1} \mid S_n, I_n) \quad (16)$$

From problem 1, the expression can be written as:

$$\prod_{i=0}^n P_0(\Delta_t^I = s_t - s_{t+1})P(\Delta_t^R = i_t - i_{t+1} + s_t - s_{t+1}) \quad (17)$$

## 7 Problem 3

Now computing the normalising constant, the full conditionals  $\pi(\lambda \mid y, t, p^{i \rightarrow r})$ ,  $\pi(t \mid y, p^{i \rightarrow r}, \lambda)$ ,  $\pi(p^{i \rightarrow r} \mid y, \lambda, t)$ . Assuming they are priori independent:

$$\pi(\theta \mid y) \propto (y \mid \theta) \pi(\lambda) \pi(t) \pi(p^{i \rightarrow r}) \quad (18)$$

Starting with  $\pi(\lambda \mid y, t, p^{i \rightarrow r})$  and its distributions:

$$\begin{aligned} \pi(\lambda \mid y, t, p^{i \rightarrow r}) &\propto f(y \mid \theta) \pi(\lambda) \propto \\ \prod_{i=1}^d \frac{\beta_i^{\alpha_i}}{\Gamma(\alpha_i)} \lambda_i^{\alpha_i-1} e^{-\beta_i \lambda_i} \prod_{i=0}^T P(\Delta_t^I = s_t - s_{t+1}) P(\Delta_t^R = i_t - i_{t+1} + s_t - s_{t+1}) \end{aligned} \quad (19)$$

Now moving to  $\pi(t \mid y, p^{i \rightarrow r})$

$$\propto \prod_{i=0}^T P(\Delta_t^I = s_t - s_{t+1}) P(\Delta_t^R = i_t - i_{t+1} + s_t - s_{t+1}) 1_{0 < t_1 \dots < t_{d-1} < T}(t) \quad (21)$$

Finally,  $\pi(p^{i \rightarrow r} \mid y, \lambda, t) \propto$

$$\frac{1}{B(a, b)} (p^{i \rightarrow r})^{a-1} (1 - p^{i \rightarrow r})^{b-1} \prod_{i=0}^T P(\Delta_t^I = s_t - s_{t+1}) P(\Delta_t^R = i_t - i_{t+1} + s_t - s_{t+1}) \quad (22)$$

$$\propto (p^{i \rightarrow r})^{a-1} (1 - p^{i \rightarrow r})^{b-1} \prod_{i=0}^T P(\Delta_t^R = i_t - i_{t+1} + s_t - s_{t+1}) \quad (23)$$

## 8 Problem 4

Implementing a hybrid sampler simulating from the joint posterior  $\pi(\theta \mid y)$  for the available data sets, using a standard Gibbs step for  $p^{i \rightarrow r}$ , and the components  $\lambda$  and  $t$  are updated using local Metropolis-Hastings moves. Using a Gaussian random walk proposal for each  $\lambda_i$ , a candidate is generated according to:

$$\lambda_i^* = \lambda_i + \sigma \epsilon \quad (24)$$

Where  $\epsilon$  is normally distributed and  $\sigma > 0$  is an algorithmic parameter. For each breakpoint, a candidate is generated using a random walk proposal:

$$t_i^* = t_i + \epsilon \quad (25)$$

Where  $\epsilon$  has a discrete uniform distribution,  $U(-M, M)$ .

## 9 Problem 5

Investigating the sensitivity of the posteriors and mixing with regard to the hyperparameters  $(\beta_i)$ ,  $a, b$  showed that they did not have any significant impact on the results, only the beta distribution. The algorithmic parameters  $\sigma, M$ , causes higher variance to  $\lambda, t$  with an lower values.

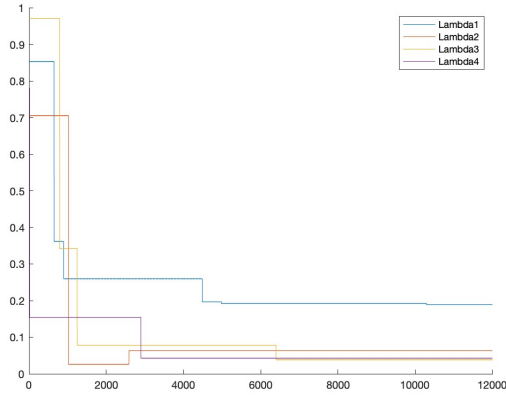


Figure 1: Proposed lambdas

In figure 1,  $\sigma = 100$ . Increasing  $\beta_i$  seems to decrease the variance of the proposed lambdas, especially the first one.

## 10 Problem 6

Starting with the German data, with  $\sigma = 0.05$ , the breakpoint converges to  $t = 34$ , which is the turning point for the infected in Germany.

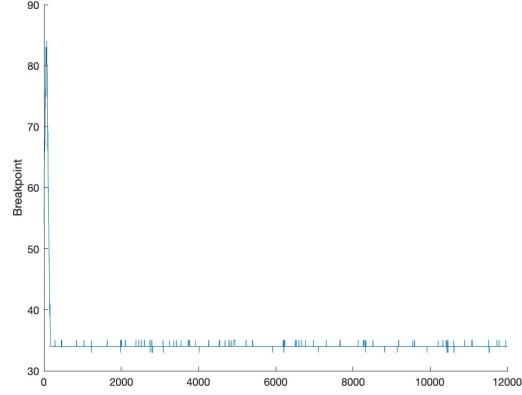


Figure 2: Breakpoint for German data

Having instead 3 breakpoints, they converge to 20, 28, and 34. Which has about the same time between them as the implemented COVID-19 containment laws. The mean of  $p^{i \rightarrow r}$  was computed to be 0.065. The lambdas showed an increase in variance the more they were, but their values were as expected.

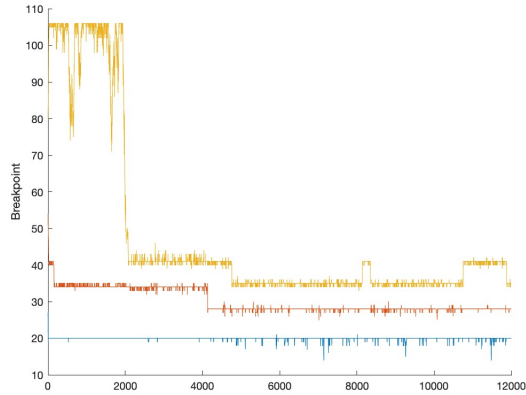


Figure 3: 3 breakpoints for German data

Looking at Iran's data of infected, two breakpoints should be made out easily, at around 35 and 65.

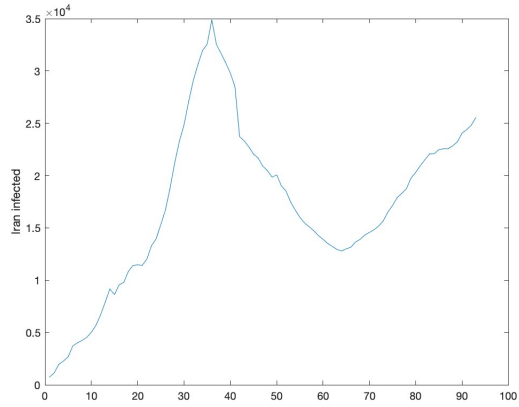
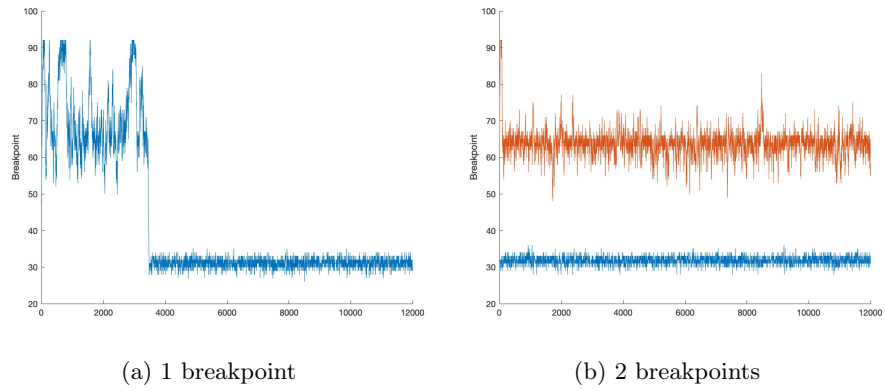


Figure 4: Iran infected

With one breakpoint the logarithm seems indecisive between the two, while two breakpoints converges much more nicely.



(a) 1 breakpoint

(b) 2 breakpoints

Figure 5: Iran breakpoints

Looking at the countries' lambdas, the stages of COVID-19 containment is clearer in Germany's lambdas, while both countries have a bigger lambda with an increase in infection.

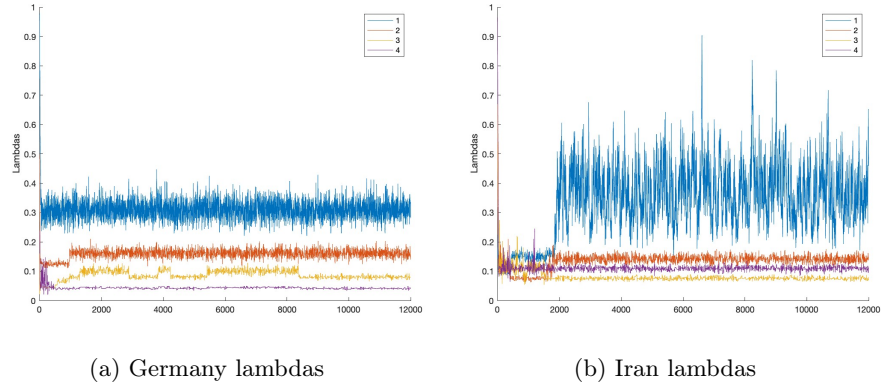


Figure 6: 4 lambdas

## 11 Problem 7

Since  $p^{i \rightarrow r}$  does not require  $\lambda$  or  $t$ , an MCM algorithm is not required. Looking the data given one can instead decide on parameters to use with a Beta distribution.