# TMA4265 Stochastic Modelling – Fall 2023 Project 1

# **Background** information

- The deadline for the project is Sunday October 9 at 23:59.
- This project must be passed to be admitted to the final exam, and a reasonable attempt must be made for each problem to pass this project (E-level effort is not enough).
- The project should be done in groups of **two** or **three** people. You must sign up as a group in Blackboard before submitting your report and code.
- ullet The project report should preferably be prepared using IATEXor R-markdown and must:
  - be a pdf-file.
  - include necessary equations and explanation to justify the answers in each problem.
  - include the required plots in the pdf and reference them in the text.

You do not need to repeat the question text in the report you submit.

- The computer code must be written in R and must be submitted as a separate file **not** as an appendix in the report. If you include any code in the report it should be as small excerpts which serve as natural parts of your answer to the question. Make sure your code runs. We may test it.
- There is a **7 page limit** for the project report. If you submit a longer report, we may not read it. The computer code is submitted as a separate file and may be as long as necessary.
- Make your computer code readable and add comments that describe what the code is doing.
- We will provide help with the project in the exercise classes in weeks 39 and 40, and in the lectures in week 40.
- If you have questions outside the aforementioned times, please post your questions on Discourse (https://mattelab2023h.math.ntnu.no/c/tma4265/19) if possible. If your question is impossible to ask without giving away important parts of the solution, you may email your question to the teaching assistants.
- The pdf-file with the report and the files with computer code must be submitted through our Blackboard pages under "Projects". You need to sign up as a group before you can submit your answer.

# Problem 1: Modelling an outbreak of measles

Throughout this problem you may assume that one year has exactly 365 days and that we are considering a population consisting of a fixed number N > 0 individuals. We use a simplified model where each individual only has three possible states: susceptible (S), infected (I), and recovered and immune (R). We model on a daily scale and let  $n = 0, 1, \ldots$  denote time measured in days. We treat measles as an infectious disease, and we assume that each day

- 1) a susceptible individual can become infected or remain susceptible,
- 2) an infected individual can become recovered or remain infected,
- 3) a recovered individual can lose immunity and become susceptible, or remain recovered.

In the first stage of modelling, we assume that the individuals in the population are independent, and assume that each day, any susceptible individual has a probability  $0 < \beta < 1$  of becoming infected tomorrow, any infected individual has a probability  $0 < \gamma < 1$  of becoming recovered tomorrow, and any recovered individual has a probability  $0 < \alpha < 1$  of losing immunity tomorrow.

a) Consider one specific individual, and let  $X_n$  denote the state of that individual at time n. Let the states 0, 1 and 2 correspond to S, I and R, respectively, and assume that  $X_0 = 0$ . Justify that  $\{X_n : n = 0, 1, \ldots\}$  is a Markov chain and explain why the transition probability matrix is given by

$$\mathbf{P} = \begin{bmatrix} 1 - \beta & \beta & 0 \\ 0 & 1 - \gamma & \gamma \\ \alpha & 0 & 1 - \alpha \end{bmatrix}.$$

State what each parameter represents in your explanation.

b) Assume that  $\beta = 0.01$ ,  $\gamma = 0.10$  and  $\alpha = 0.005$ . Explain why you can be certain that this Markov chain has a limiting distribution, and calculate (by hand) the long-run mean number of days per year spent in each state.

- c) Assume that the individual is susceptible at time 0, and complete the following tasks:
  - (In R) Simulate the Markov chain for 20 years (7300 time steps).
  - (Computer code) For each run of this code, use the last 10 years (of the 20 years) to get an estimate of each of the quantities in b).
  - (In R) Run the code 30 times and compute an approximate 95% confidence interval (CI) for each of the quantities in b) (using the 30 independent estimates).
  - (Report) Explain briefly how you computed the CIs, provide the computed CIs, and discuss whether the computed CIs are compatible with your exact calculations in **b**)?

Due to the highly infectious nature of measles, the proportions of susceptible, infected and recovered individuals in the population will change with time. This in turn means that it is highly unrealistic to assume that  $\beta$  does not change with time. Assume that the population size N=1000 is constant through time and at each time step consists of  $S_n$  susceptible individuals,  $I_n$  infected individuals, and  $R_n$  recovered individuals.

Assume that for each time step n, the probability that a susceptible individual becomes infected is  $\beta_n = 0.5 I_n/N$ , the probability that an infected individual recovers is  $\gamma = 0.10$ , and the probability that a recovered individual becomes susceptible is  $\alpha = 0.005$ . Assume that the N = 1000 individuals change states independently of each other at each time step given the values of  $\beta_n$ ,  $\gamma$  and  $\alpha$ . Define the discrete-time stochastic process  $\{Y_n : n = 0, 1, \ldots\}$ , where  $Y_n = (S_n, I_n, R_n)$ .

**d)** Let  $Z_n = (S_n, I_n)$  for  $n = 0, 1, \ldots$  Are  $\{I_n : n = 0, 1, \ldots\}$ ,  $\{Z_n : n = 0, 1, \ldots\}$ , and  $\{Y_n : n = 0, 1, \ldots\}$  Markov chains? Give definite arguments for all three.

Introduce 50 infected individuals in the population at time n = 0 by  $Y_0 = (950, 50, 0)$ .

- e) Complete the following tasks:
  - (In R) Simulate  $\{Y_n : n = 0, 1, 2, ...\}$  until time step n = 300. Hint: At each time step, the number of new susceptible individuals, new infected individuals, and new susceptible individuals are all binomial distributions. After you determine the number of trials and the success probability for each of them, you can simulate them, for example, using rbinom in R.
  - (Report) Choose one realization and show the temporal evolutions of  $S_n$ ,  $I_n$  and  $R_n$  together in one figure. Give a short discussion of why the behaviour of the Markov chain is very different in time intervals 0–50 and 50–300.
- f) A major interest in the modelling of infectious diseases lies in the explosive behaviour during the initial outbreak of the disease. Based on 1000 simulations of the outbreak for time steps  $n = 0, 1, \ldots, 300$ :
  - (In R) Estimate the expected maximum number of infected individuals during the simulated time steps,  $E[\max\{I_0, I_1, \dots, I_{300}\}]$ , and the expected time at which the number of infected individuals first takes its highest value,  $E[\min\{\arg\max_{n\leq 300}\{I_n\}\}]$ .
  - (In R) Compute approximate 95% CIs for the two expected values.
  - (Report) Provide the computed CIs, and discuss how would you use them assess the potential severity of the outbreak?
- g) This question is open to different solutions, but you need to justify and describe the approach you choose in your report.

A strategy to avoid large outbreaks is immunisation programmes. Assume vaccination provides life-long immunity, and consider three cases: 100 individuals are vaccinated, 600 individuals are vaccinated, and 800 individuals are vaccinated.

We want to introduce 50 infected individuals among the unvaccinated individuals, and study the behaviour of the number of infected individuals through time. Modify the above model to accommodate the vaccinated individuals, and use your modified model to generate four realizations of the temporal evolution of the number of infected individuals in the same figure: the three cases described above, and the original case of no vaccinated individuals. Discuss what changes as more and more individuals are vaccinated.

Compute and discuss changes in the estimated expected values from 1f) in the three new cases compared to the original case of no vaccinated individuals.

## Problem 2: Insurance claims

Let X(t) denote the number of claims received by an insurance company in the time interval [0,t]. We will assume that  $\{X(t):t\geq 0\}$  can be modelled as a Poisson process, where t is measured in days since January 1st at 00:00.00. Assume that the rate of the Poisson process is given by  $\lambda(t)=1.5,\,t\geq 0$ .

#### a) Complete the following tasks:

- (Report) Compute the probability that there are more than 100 claims at March 1 at 00:00.00 (59 days).
- (In R) Write code to verify the calculation by simulating 1000 realizations of the Poisson process.
- (Report) Compare the estimated probability with the exact calculation and comment. Further, make a figure that shows 10 realizations of X(t),  $0 \le t \le 59$ , plotted in the same figure.

Assume that the monetary claims are independent, and are independent of the claim arrival times. Each claim amount (in mill. kr.) has an exponential distribution with rate parameter  $\gamma = 10$ . This means that claim  $C_i \sim \text{Exp}(\gamma)$ ,  $i = 1, 2, \ldots$  The total claim amount at time t is defined by  $Z(t) = \sum_{i=1}^{X(t)} C_i$ .

**NB:** Since  $\gamma$  is a **rate** parameter, the exponential distribution is parametrized as  $f(c) = \gamma e^{-\gamma c}$ , c > 0. This may differ from what you have seen in other courses, but we will exclusively use this parametrization of the exponential distribution in this course.

### b) Complete the following tasks:

- (In R) Write code that uses 1000 simulations to estimate the probability that the total claim amount exceeds 8 mill. kr. at March 1 at 00:00.00 (59 days).
- (Report) Provide the estimated probability, and make a figure that shows 10 realizations of Z(t),  $0 \le t \le 59$ , plotted in the same figure.
- c) The policy of the insurance company is to investigate a claim if and only if it exceeds 250000 kr.. For  $t \ge 0$ , let  $Y_t$  denote the number of claims, which need to be investigated, received in the time interval [0,t]. Prove that  $\{Y(t): t \ge 0\}$  is a Poisson process and compute its rate.

# Problem 3: Metropolis algorithm example (OPTIONAL)

This problem is entirely optional. It is intended as a brief application of the Metropolis algorithm (a type of Markov chain Monte Carlo) for any interested.

Define constants ...,  $c_{-1}, c_0, c_1, ...$ , with  $0 < c_i < \infty$  for each  $i \in \mathbb{Z}$ . Let  $X_0 = 0$ , and define  $\{X_n\}$  for n = 1, 2, ... algorithmically in the following way:

- 1. Generate proposal  $X_{n+1}^*$ , with  $X_{n+1}^*$  equal to  $X_n 1$  with probability 0.5 and equal to  $X_n + 1$  otherwise
- 2. Generate  $U \sim \text{Unif}(0,1)$
- 3. Set

$$X_{n+1} = \begin{cases} X_{n+1}^*, & \text{if } U < \min\left\{\frac{c_{X_{n+1}^*}}{c_{X_n}}, 1\right\} \\ X_n, & \text{otherwise.} \end{cases}$$

If we repeat the above process for n = 0, 1, ..., we can generate a realization of  $\{X_n\}$  up to a certain number of time steps.

- a) Is  $\{X_n\}$  a Markov chain? Is it reducible/irreducible and what is its periodicity? What if, for some  $i, c_i = 0$ ?
- **b)** Show that the transition probabilities take the form:

$$P_{ij} = \begin{cases} 0.5 \cdot \min\left\{1, \frac{c_j}{c_i}\right\}, & j = i \pm 1\\ 1 - 0.5\left(\min\left\{1, \frac{c_{i+1}}{c_i}\right\} + \min\left\{1, \frac{c_{i-1}}{c_i}\right\}\right), & j = i\\ 0, & \text{otherwise.} \end{cases}$$

- c) Show that, if  $\{X_n\}$  is positive recurrent, then the long run proportion of time spent by  $\{X_n\}$  in state i is  $\frac{c_i}{\sum_i c_i}$ .
- d) Is the stationary distribution in c) a limiting distribution? Why or why not?

**NB:** by a theorem outside of the scope of this class,  $\{X_n\}$  is positive recurrent if and only if  $\sum_{i=-\infty}^{\infty} c_i < \infty$ .

**NB** 2: We have found a method for sampling distributions on  $\mathbb{Z}$  with probability proportional to  $c_i$ . Importantly, the algorithm outlined in the recursive definition of  $\{X_n\}$  does not depend on  $\sum_i c_i$ . In other words, it does not depend on the normalizing constant of the distribution, only the relative values of subsequent  $c_i$ . This algorithm is a special case of the Metropolis algorithm, which in turn is a special case of Metropolis-Hastings. More information on these algorithms and Markov chain Monte Carlo (MCMC) is given in TMA4300.