## CS-E5885 Modeling biological networks

## Exam, February 23, 2023

## Lecturer: Harri Lähdesmäki

You are NOT allowed to use calculators or any other additional equipments/material in the exam. Please write your answers in English. Please write carefully. Use mathematical notation and equations in your answers whenever possible and reasonable. The use of diagrams and drawings is also encouraged.

## Questions:

- 1. Consider random variables  $\mathbf{x} = (x_1, \ldots, x_6)$ .
  - (a) Write the joint probability distribution for x in product-form in two different ways: such that if factorizes according to the undirected graphical model (also known as Markov random field) shown in Figure 1 a), and such that if factorizes according to the directed graphical model (also known as Bayesian network) shown in Figure 1 b). Explain all the mathematical notations used in your answer. (5 points)
  - (b) Convert the static directed graphical model in Figure 1 b) to a dynamic Bayesian network for x(t) = (x<sub>1</sub>(t),...,x<sub>6</sub>(t)), t = 0, 1, 2, ..., by "unrolling" the static network over one time step, from time t-1 to t. Draw the resulting dynamic Bayesian network model as a directed graph for one time step, from time t 1 to t. Also write the joint distribution over a finite length time-series x(0), x(1),...,x(T) in a product form such that the product involves conditional probabilities only for univariate variables. You can assume that the initial state x(0) is fixed, i.e., x(0) does not have any uncertainty. (5 points)

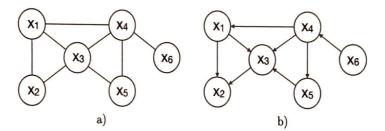


Figure 1: a) Undirected and b) directed graphical model for random variables  $\mathbf{x} = (x_1, \dots, x_6)$ .

2. The reaction equations for the well-known Lotka-Volterra can be written as

$$\begin{array}{c} Y_1 \rightarrow 2Y_1 \\ Y_1 + Y_2 \rightarrow 2Y_2 \\ Y_2 \rightarrow \emptyset. \end{array}$$

Write the ODE version of the Lotka-Volterra model for concentrations  $[Y_1]$  and  $[Y_2]$  assuming the mass-action kinetics with deterministic reaction rates  $k_1$ ,  $k_2$  and  $k_3$  for the three reactions. (5 points)

3. Consider modeling continuous-time dynamics for n variables,  $\mathbf{x}(t) = (x_1(t), \dots, x_n(t))$ , using ODEs, where the model for the *i*th variable can be written in a general form as

$$rac{dx_i(t)}{dt} = f_i(\mathbf{x}(t)| heta_i), \quad i = \{1,\ldots,n\}.$$

Assume you have N + 1 measurements  $\mathbf{y}(t_0), \mathbf{y}(t_1), \dots, \mathbf{y}(t_N)$  from a single time-series trajectory that are measured at arbitrary time points  $t_0, t_1, \dots, t_N$ . Each measurement

 $\mathbf{y}(t) = (y_1(t), \dots, y_n(t))$  contains a measurement for all *n* variables, where  $y_i(t) = x_i(t) + \epsilon_i(t)$ , where  $\epsilon_i(t)$  denotes measurement error. Describe the gradient matching method for learning the parameters  $\theta_i$  separately for each variable  $x_i$ . (5 points)

4. Background: Recall the Poisson timestep method for approximative simulation of coupled chemical reaction networks. Briefly, time is discretized into small time increments  $\Delta t$ . For each time increment we need to simulate a v-dimensional reaction vector r such that  $r_i \sim \text{Po}(h_i(x, c_i)\Delta t)$  and then the updated state x and time t are obtained by x := x + Sr and  $t := t + \Delta t$ , where S is the stoichiometric matrix.

**Question:** Describe how you can obtain the chemical Langevin equation (CLE) as a further approximation of the Poisson timestep method, and write down a general equation for CLE in terms of a stochastic differential equation (SDE) model (or the corresponding Euler-Maruyama simulation algorithm). In your answer, remember to describe the statistical aspects and motivations for deriving the CLE. (10 points total)