

**Suggested solutions for
MVE550 Stochastic Processes and Bayesian Inference
Trial exam autumn 2018**

1. (a) As state 2 is absorbing while the other states are transient, the limiting distribution is $(0, 1, 0)$.

- (b) The transition matrix is $P = \begin{bmatrix} 0 & 0.3 & 0.7 \\ 0 & 1 & 0 \\ 0.6 & 0.4 & 0 \end{bmatrix}$. The part concerning the transient states is $Q = \begin{bmatrix} 0 & 0.7 \\ 0.6 & 0 \end{bmatrix}$ and thus the fundamental matrix is

$$F = (I - Q)^{-1} = \begin{bmatrix} 1 & -0.7 \\ -0.6 & 1 \end{bmatrix}^{-1} = \frac{1}{1 - 0.6 \cdot 0.7} \begin{bmatrix} 1 & 0.7 \\ 0.6 & 1 \end{bmatrix} = \begin{bmatrix} 1.7241 & 1.2069 \\ 1.0345 & 1.7241 \end{bmatrix}.$$

- (c) This can be read of the fundamental matrix: The answer is 1.2069.
2. (a) As $n \rightarrow \infty$ the distribution of each chain will approach the stationary distribution v . Let us write s for the number of states in the state space. We can compute the probability p that the k chains have the same state by summing over the possible states and computing the probability that each chain has this state. Letting $n \rightarrow \infty$, we get

$$p = \sum_{i=1}^s v_i^k.$$

- (b) One may use “coupling”: Pick an ordering of the states in the state space and name the states $1, \dots, s$. Define a function f sending pairs (i, u) , where u is a real number in $[0, 1]$ and i is a state, to a new state, as follows:

$$f(i, u) = \max \left\{ j : \sum_{r=1}^{j-1} P_{ir} \leq u \right\}.$$

One may see that $P(f(i, U) = j) = P_{ij}$ when U is a random variable with a uniform distribution on $[0, 1]$. Thus, if a chain is at state i , one may choose its state in the next step according to $f(i, U)$ where $U \sim \text{Uniform}(0, 1)$.

The values of the k chains are now constructed by using, for each step m , the same random number U_m for all the chains. Each chain will then be a Markov chain with the given transition matrix, while the chains will not be independent, they will be “coupled”.

It should be clear, that, if two chains have the same state at a certain step, they will continue to have the same state at all following steps. Furthermore, although we do not prove this rigorously, it should be clear that unless all rows of the transition matrix are equal, there exist some pairs of states with a non-zero probability that two chains in these two states at one step will have the same state at the following step. Using ergodicity, it may be shown that as $n \rightarrow \infty$, all chains will end up in the same state.

(Note that our coupling will not work if all the rows of the transition matrix are equal. However, in this case, all the rows are equal to v , all chains converge immediately to the limiting distribution, and all questions about convergence are rather uninteresting).

(Note: A fairly short explanation would give students full points for this subquestion).

- (c) Perfect sampling uses the simulation method of (b). In Perfect sampling, one decides on a certain number of steps n , and then simulates s chains, all with different starting states, for n steps, using coupling as above. If all the chains are in the same state at the final step, one knows that this step is a sample from the stationary distribution. For this to happen with a reasonable probability, one needs the effect of (b), i.e., that the chains tend to end up in the same state.

3. We have

$$\begin{aligned} G(s) &= E[s^X] \\ G'(s) &= E[Xs^{X-1}] \\ G''(s) &= E[X(X-1)s^{X-2}] \end{aligned}$$

so that $G'(1) = E[X]$ and $G''(1) = E[X(X-1)] = E[X^2] - E[X]$. Thus

$$\begin{aligned} \text{Var}[X] &= E[X^2] - E[X]^2 \\ &= E[X^2] - E[X] + E[X] - E[X]^2 \\ &= G''(1) + G'(1) - (G'(1))^2. \end{aligned}$$

4. The joint probability mass function for $N_t^{(M)}$ and $N_t^{(F)}$ can be derived as

$$\begin{aligned} &P(N_t^{(M)} = i, N_t^{(F)} = j) \\ &= P(N_t^{(M)} = i, N_t = i + j) \\ &= P(N_t^{(M)} = i \mid N_t = i + j)P(N_t = i + j) \\ &= \frac{(i+j)!}{i! \cdot j!} p^i (1-p)^j e^{-\lambda} \frac{\lambda^{i+j}}{(i+j)!} \\ &= e^{-\lambda} \frac{1}{i!} (p\lambda)^i \frac{1}{j!} ((1-p)\lambda)^j \\ &= \left[e^{-\lambda p} \frac{1}{i!} (p\lambda)^i \right] \left[e^{-\lambda(1-p)} \frac{1}{j!} ((1-p)\lambda)^j \right] \\ &= \text{Poisson}(i; \lambda p) \cdot \text{Poisson}(j; \lambda(1-p)) \end{aligned}$$

From this it follows that the marginal distribution for $M_i^{(M)}$ is $\text{Poisson}(\lambda p)$ and that the marginal distribution for $M_i^{(F)}$ is $\text{Poisson}(\lambda(1-p))$, and that the two random variables are independent.

5. (a) We get

$$\begin{aligned}
 \pi(a \mid x_0, x_1) &\propto_a \pi(x_0, x_1 \mid a)\pi(a) \\
 &= \pi(x_1 \mid a, x_0)\pi(a) \\
 &= \text{Normal}(x_1; ax_0, 1) \cdot \text{Normal}(a; 1, 1) \\
 &\propto_a \exp\left(-\frac{1}{2}(x_1 - ax_0)^2\right) \exp\left(-\frac{1}{2}(a - 1)^2\right) \\
 &= \exp\left(-\frac{1}{2}\left[a^2x_0^2 - 2ax_0x_1 + x_1^2 + a^2 - 2a + 1\right]\right) \\
 &\propto_a \exp\left(-\frac{1}{2}\left[(x_0^2 + 1)a^2 - 2a(x_0x_1 + 1)\right]\right) \\
 &\propto_a \exp\left(-\frac{1}{2}(x_0^2 + 1)\left(a - \frac{x_0x_1 + 1}{x_0^2 + 1}\right)^2\right)
 \end{aligned}$$

so $a \sim \text{Normal}\left(\frac{x_0x_1+1}{x_0^2+1}, \frac{1}{x_0^2+1}\right)$.

(b) In the above computation a had the prior $\text{Normal}(1, 1)$, but it should be clear that an entirely similar computation would result in a normal posterior also for any other normal prior for a . Furthermore, the posterior for a given a sequence x_0, x_1, \dots, x_k of observations for the variables X_0, X_1, \dots, X_k can be computed by sequentially using each observation as data, and the posterior for a from previous computations as prior for a . The posterior is then at each stage normal, so, by induction, the final posterior, given all observations, is also normal.

6. (a) The chain is ergodic as there are non-zero transition rates for example from 1 to 2 to 4 to 3 and back to 1, and thus only one communication class.

(b)

$$Q = \begin{bmatrix} -4 & 3 & 1 & 0 \\ 0 & -2 & 0 & 2 \\ 3 & 1 & -9 & 5 \\ 0 & 0 & 2 & -2 \end{bmatrix}.$$

(c) We must have $\sum_{i=1}^4 v_i = 1$ and $vQ = 0$. The four columns of Q give rise to 4 dependent equations. Selecting 3 of these, we get, for example,

$$\begin{aligned}
 1 &= v_1 + v_2 + v_3 + v_4 \\
 0 &= -4v_1 + 3v_3 \\
 0 &= 3v_1 - 2v_2 + v_3 \\
 0 &= v_1 - 9v_3 + 2v_4
 \end{aligned}$$

(d)

$$P = \begin{bmatrix} 0 & \frac{3}{4} & \frac{1}{4} & 0 \\ 0 & 0 & 0 & 1 \\ \frac{3}{9} & \frac{1}{9} & 0 & \frac{5}{9} \\ 0 & 0 & 1 & 0 \end{bmatrix}.$$

(e) We must have $\sum_{i=1}^4 w_i = 1$ and $w(I - P) = 0$. The four columns of $I - P$ give rise to 4 dependent equations. Selecting 3 of these, we get, for example,

$$\begin{aligned} 1 &= w_1 + w_2 + w_3 + w_4 \\ 0 &= w_1 - \frac{3}{9}w_3 \\ 0 &= -\frac{3}{4}w_1 + w_2 - \frac{1}{9}w_3 \\ 0 &= -w_2 - \frac{5}{9}w_3 + w_4 \end{aligned}$$

7. (a) We can write

$$B_1 + B_2 + B_3 = B_1 + 2B_2 + B_3 - B_2 = 3B_1 + 2(B_2 - B_1) + B_3 - B_2.$$

where each of B_1 , $B_2 - B_1$, and $B_3 - B_2$ are independent and normally distributed with expectation zero. Thus the sum is normally distributed with expectation zero, and variance

$$\begin{aligned} \text{Var}[B_1 + B_2 + B_3] &= \text{Var}[3B_1 + 2(B_2 - B_1) + B_3 - B_2] \\ &= \text{Var}[B_1] + \text{Var}[2(B_2 - B_1)] + \text{Var}[B_3 - B_2] \\ &= 9 \text{Var}[B_1] + 4 \text{Var}[B_2 - B_1] + \text{Var}[B_3 - B_2] \\ &= 9 \text{Var}[B_1] + 4 \text{Var}[B_1] + \text{Var}[B_1] \\ &= 9 + 4 + 1 = 14. \end{aligned}$$

Thus

$$B_1 + B_2 + B_3 \sim \text{Normal}(0, 14).$$

(b) A Gaussian process $\{X_t\}_{t \geq 0}$ is a continuous-time stochastic process with the property that for all $n = 1, 2, \dots$ and $0 \leq t_1 < t_2 < \dots < t_n$, the random variables $X_{t_1}, X_{t_2}, \dots, X_{t_n}$ have a multivariate Normal distribution.

(c) Note that T_1 is the first hitting time for a . This is a stopping time, and thus the stochastic process $\{X_t\}_{t \geq 0}$ defined by

$$X_t = B_{T_1+t} - B_{T_1}$$

is Brownian motion. This means that it has infinitely many zeroes in the interval $(0, \epsilon)$ for any $\epsilon > 0$. Thus, for any $n > 0$ there is infinitely many sequences $T_1 < t_1 < t_2 < \dots < t_n < T_1 + \epsilon$ such that $B_{t_i} = a$ for $i = 1, \dots, n$. This shows that $T_{100} = T_1$, i.e., its distribution is concentrated in the value 0.