# MATH3290 Mathematical Modeling 2023/2024 <br> Assignment 1 <br> Due Date: 5pm, February 20th 

Note: Submit your assignment via Blackboard. Late submissions are not allowed.

1. Consider the data sets in Table 1

| $x$ | 1 | 2 | 3 | 4 | 5 | 6 | 7 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $y_{1}$ | 7 | 15 | 33 | 61 | 99 | 147 | 205 |
| $y_{2}$ | 4.5 | 20 | 90 | 403 | 1,808 | 8,130 | 36,316 |

Table 1: Data set for Problem 1.
(a) For $\left(x, y_{1}\right)$, construct a divided difference table. What conclusions can you make about $y_{1}$ ? Would you use a low-order polynomial as an empirical model? If so, what order?
(b) For $\left(x, y_{2}\right)$, construct a divided difference table. Would you use a low-order polynomial as an empirical model? If not, give the reason.
2. The following data were obtained for the growth of a sheep population introduced into a new environment on the island of Tasmania.

| $t$ (year) | 1814 | 1824 | 1834 | 1844 | 1854 | 1864 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $p($ population $)$ | 125 | 275 | 830 | 1200 | 1750 | 1650 |

Table 2: Data set for Problem 2.
(a) Plot the change in population versus year. Is there a trend?
(b) Formulate a discrete dynamical system model. Use the least-squares criterion to find the model parameter.
(c) Predict the sheep population in the year 1869 .
3. (Markov process) A certain protein molecule can have three configurations which are denoted as $C_{1}, C_{2}$ and $C_{3}$. Every second, a protein molecule can make a transition from one configuration to another configuration with the following probabilities:

$$
\begin{array}{lll}
P\left(C_{1} \rightarrow C_{1}\right)=0.3 & P\left(C_{1} \rightarrow C_{2}\right)=0.2 & P\left(C_{1} \rightarrow C_{3}\right)=0.5 \\
P\left(C_{2} \rightarrow C_{1}\right)=0.3 & P\left(C_{2} \rightarrow C_{2}\right)=0.5 & P\left(C_{2} \rightarrow C_{3}\right)=0.2 \\
P\left(C_{3} \rightarrow C_{1}\right)=0.4 & P\left(C_{3} \rightarrow C_{2}\right)=0.2 & P\left(C_{3} \rightarrow C_{1}\right)=0.4
\end{array}
$$

The configuration transition are demonstrated in Figure 1. (For example, the molecule will transit from $C_{1}$ to $C_{2}$ with probability 0.2 .)
Consider a living body with a fixed number of protein molecules. We let $C_{i}^{n}(i=1,2,3 ; n=$ $0,1,2, \ldots)$ be the percentage of molecules that are in configuration $C_{i}(i=1,2,3)$ at the end of the $n$-th second.


Figure 1: An illustration of the configuration transition process for Problem 3.
(a) Formulate a model for $C_{i}^{n}$ using a system of difference equations.
(b) Find the equilibrium point, and determine its stability.
(c) Consider the three initial conditions in Table 3. Compute $C_{1}^{5}, C_{2}^{5}, C_{3}^{5}$ for each case. Does the long term behaviour sensitive to the initial condition?

| Percentage | $C_{1}^{0}$ | $C_{2}^{0}$ | $C_{3}^{0}$ |
| :---: | :---: | :---: | :---: |
| Case A | 0 | 0 | 1 |
| Case B | 0 | 0.5 | 0.5 |
| Case C | 0.2 | 0.2 | 0.6 |

Table 3: Data set for Problem 3.

