Computational Molecular Medicine

Assignment 5

Due April 18

Problem 1: Consider a simple Galton-Watson branching process for a population of (normal) stem cells with the following parameters:

- The population starts from one cell : N(0) = 1.
- At each generation, each stem cell can either die with probability $\frac{1}{2}$ or divide into two stem cells with probability $\frac{1}{2}$.

Simulate this process from generation t = 0 to generation t = 1000.

- Compare the analytical expression of the expected value of the size of population at each generation t with the value estimated with simulations.
- Do the same thing for the probability of extinction of the population by considering t = 1000 as $t = \infty$.

Using the recursive equation between the MGF F(t+1,z) of N(t+1) and the MGF F(t,z) of N(t):

- Derive a method to compute P(N(t) > 0).
- Compare the computed probability with the values estimated from simulations for t = 15.

Problem 2: Consider the same setting as Problem 1 with the difference that the probability of death is now $\frac{20}{41}$.

- Answer the same questions posed in Problem 1.
- Compare the approximation of E(N(t)|N(t) > 0) derived in class for large values of t with its estimated values from your simulations.

Problem 3: In this problem we will take into account mutations of normal stem cells and simultaneously consider both normal and mutant populations. More specifically, we will consider the process $N(t) = (N_0(t), N_1(t))$ where $N_0(t)$ (respectively, $N_1(t)$) is the number of normal (respectively mutant) stem cells at generation t. The definitions and dynamics are:

- Each normal stem cell can either die with probability $\frac{100}{201}$ or divide into two cells. When a normal stem cell divides, there is a possibility that exactly one of the two daughter cells acquires a mutation and becomes a mutant cell. This event has probability $\frac{1}{5000}$ and in this case a cell is added to the mutant population and hence only one normal cell is generated.
- Mutant cells have a fitness advantage: they can either die with probability $\frac{20}{41}$ or divide into two mutant cells. We will not consider the possibility that mutant cells acquire another mutation.
- Every mutant cell whose parent is a normal cell is the founder cell of a mutant "clone" consisting of all the descendants of the mutant founder cell.
- We define the time T of the appearance of cancer as first time t at which there exists a mutant clone of size at least 10^6 . (Note: $T = \infty$ if no such t exists.)

Starting from 100 normal cells and no mutant cells, simulate this process for 500 generations, i.e., t = 1 to t = 500.

• Plot the "cancer incidence" curve $t \to P(T \le t)$, where each generation represents two months. The overall cancer incidence is $P(T \le 500)$.