

Stochastic simulation and chemical inference modules

1. The Gillespie algorithm for stochastic simulation.

Download a copy of the original paper by Gillespie (*Exact Stochastic Simulation of Coupled Chemical Reactions*, D. T. Gillespie, J. Phys. Chem. **81** 2340 (1977)). Locate the file `GillespieMatlab.m`. Write down a description of the function of each variable in the M-file, and the line numbers corresponding to the steps in boxes 1-3 in Fig. 2 of the original paper.

2. A simple stochastic chemical system.

We will first consider the following simple chemical system:



where k_i are the rates of each reaction.

(a) Write down the stoichiometry matrix for this system.

(b) Use `GillespieMatlab.m` to simulate this system for $k_1 = k_2$, $k_1 > k_2$ and $k_1 < k_2$. Plot the resulting stochastic trajectories to verify that the system behaves as expected.

3. Gillespie simulation of the Monod model.

We will now look at a more complicated chemical system, used to model co-operative binding, which we shall see is useful in biological inference. Here, we have a protein which can exist in binding (B) or non-binding (N) states. The protein has n sites, to which sugar molecules (S) may bind. However, the protein may only swap between B and N states when no sugars are bound to it. Sugar binding hence acts to “lock in” a particular state of the protein. This system is described by:



(a) Write down the stoichiometry matrix for this system. Don't forget that sugar molecules are reactants too! Why are some rates a function of n ?

(b) Use `GillespieSwain.m` to simulate this system for $K_R = 100$, $K_T = 10$, $L = 2$, $n = 1$. Explore the occupancy of the B state as S varies over orders of magnitude.

(c) Simulate the system for $n = 2$. What do you notice about the behaviour at different sugar concentrations as n changes?