Due: Monday, May 6, 2019

This is the last homework assignment of the semester. Please hand in a **physical copy** to my office by **6 pm on Monday, May 6**th.

1. As discussed in class, the Luria-Delbrück distribution can be generalized to different (deterministic) growth rates for the sensitive and resistant bacteria. Assume, as in class, the following growth laws:

$$\frac{dn}{dt} = \alpha(n+m,t)n$$
$$\frac{dm}{dt} = c\alpha(n+m,t)m$$

Here n and m denote the number of sensitive and resistant cells, respectively, at time t, and c > 0 is a constant.

- (a) Let Z denote the random variable of the number of mutants when the sensitive population is at size N. Find an approximate expression for the cumulant generating function of Z, ψ_Z(s). *Hint:* As in class, write it as a sum, and approximate the sum via an integral.
- (b) Your expression in (a) should involve the mutation rate ν . Using a Taylor series calculation, find the leading-order (i.e. first order) terms in ν . Recall that this approximation is valid when ν is small, which we assume here.
- (c) Using your approximation in part (b), find the mean and variance of Z.
- 2. Consider the chemical reaction network modeling the production and degradation of mRNA:

$$\emptyset \xrightarrow{\alpha} M \xrightarrow{\beta} \emptyset$$

Recall that the corresponding chemical master equation (CME) is given by

$$\frac{dp_k}{dt} = \alpha p_{k-1} + (k+1)\beta p_{k+1} - \alpha p_k - k\beta p_k,$$

so that the stationary distribution π must satisfy

$$\alpha \pi_{k-1} + (k+1)\beta \pi_{k+1} - \alpha \pi_k - k\beta \pi_k = 0.$$
(1)

Here $\pi = (\pi)_{k=0}^{\infty}$ is a distribution vector (note: it is infinite in length). The corresponding solution is given by

$$\pi_k = e^{-\lambda} \frac{\lambda^k}{k!},\tag{2}$$

where

$$\lambda = \frac{\alpha}{\beta}.$$

Verify this claim. That is, show that π defined by (2) satisfies (1).

3. Consider now a simple model of transcription and translation:

$$\begin{split} \emptyset \xrightarrow{\alpha} M \xrightarrow{\beta} \emptyset \\ M \xrightarrow{\theta} M + P \\ P \xrightarrow{\delta} \emptyset \end{split}$$

Here P represents the protein translated from M.

- (a) Write down the stochiometry matrix Γ for above network.
- (b) Assuming mass-action kinetics, write down the propensity functions (i.e. rates) ρ_j^{σ} , for j = 1, 2, 3, 4. Note that the above system contains 4 reactions.
- (c) Write down the chemical master equation (CME) for the transcription/translation model. Recall here that

$$k = \left(\begin{array}{c} k_1 \\ k_2 \end{array}\right)$$

is now a vector, as their are two chemical species in the network (M and P).