## MATH 6397 Homework 3

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1. We will derive the Fokker-Planck equation for the membrane potential distribution under excitatory input. Suppose the input rate of pre-synaptic neuron k at time t is  $v_k(t)$  with weight  $w_k$ . We start with the transition probability moving from state u' to u in time t to  $t + \Delta t$ ,

$$P^{trans}\left(u,t+\Delta t\,\middle|\,u',t\,\right) = \left[1-\Delta t\sum_{k}v_{k}(t)\right]\delta\left(u-u'e^{-\Delta t/\tau_{m}}\right) + \Delta t\sum_{k}v_{k}(t)\delta\left(u-u'e^{-\Delta t/\tau_{m}}-w_{k}\right).$$

The Kolmogorov forward equation gives  $P(u,t+\Delta t) = \int P^{trans}(u,t+\Delta t|u',t)P(u',t)du'$ . Putting this into context, we have

$$\begin{split} P\left(u,t+\Delta t\right) &= \int \left\{ \left[1-\Delta t \sum_{k} v_{k}(t)\right] \delta\left(u-u'e^{-\Delta t/\tau_{m}}\right) + \Delta t \sum_{k} v_{k}(t) \delta\left(u-u'e^{-\Delta t/\tau_{m}}-w_{k}\right) \right\} P\left(u',t\right) du' \\ &= \left[1-\Delta t \sum_{k} v_{k}(t)\right] \int \delta\left(u-u'e^{-\Delta t/\tau_{m}}\right) P\left(u',t\right) du' + \Delta t \sum_{k} v_{k}(t) \int \delta\left(u-u'e^{-\Delta t/\tau_{m}}-w_{k}\right) P\left(u',t\right) du' \end{split}$$

We have that

$$\int \delta \left( u - u' e^{-\Delta t/\tau_m} \right) P\left( u', t \right) du' = \int \delta \left[ -e^{-\Delta t/\tau_m} \left( u' - u e^{\Delta t/\tau_m} \right) \right] P\left( u', t \right) du'$$

$$= \left| -e^{\Delta t/\tau_m} \right| \int \delta \left( u' - u e^{\Delta t/\tau_m} \right) P\left( u', t \right) du'$$

$$= e^{\Delta t/\tau_m} P\left( u e^{\Delta t/\tau_m}, t \right)$$

and

$$\int \delta \left( u - u' e^{-\Delta t/\tau_m} - w_k \right) P\left( u', t \right) du' = \int \delta \left[ -e^{-\Delta t/\tau_m} \left( u' - (u - w_k) e^{\Delta t/\tau_m} \right) \right] P\left( u', t \right) du'$$

$$= \left| -e^{\Delta t/\tau_m} \right| \int \delta \left( u' - (u - w_k) e^{\Delta t/\tau_m} \right) P\left( u', t \right) du'$$

$$= e^{\Delta t/\tau_m} P\left( e^{\Delta t/\tau_m} \left( u - w_k \right), t \right)$$

Applying these simplifications to  $P(u, t + \Delta t)$ , we have

$$\begin{split} P\left(u,t+\Delta t\right) &= \left[1-\Delta t \sum_{k} v_{k}(t)\right] e^{\Delta t/\tau_{m}} P\left(e^{\Delta t/\tau_{m}} u,t\right) + \Delta t \sum_{k} v_{k}(t) e^{\Delta t/\tau_{m}} P\left(e^{\Delta t/\tau_{m}} \left(u-w_{k}\right),t\right) \\ &= e^{\Delta t/\tau_{m}} \left\{P\left(e^{\Delta t/\tau_{m}} u,t\right) + \Delta t \sum_{k} v_{k}(t) \left[-P\left(e^{\Delta t/\tau_{m}} u,t\right) + P\left(e^{\Delta t/\tau_{m}} \left(u-w_{k}\right),t\right)\right]\right\} \end{split}$$

We now expand this expression about  $\Delta t = 0$  and truncate all terms with powers of  $\Delta t$  higher than 1. We use the following truncated expansions.

• 
$$e^{\Delta t/\tau_m} \approx 1 + \frac{\Delta t}{\tau_m}$$

• 
$$P\left(e^{\Delta t/\tau_m}u,t\right) \approx P\left(u,t\right) + \frac{u\Delta t}{\tau_m} \frac{\partial P\left(u,t\right)}{\partial u}$$

• 
$$P\left(e^{\Delta t/\tau_m}\left(u-w_k\right),t\right) \approx P\left(u-w_k,t\right) + \frac{\left(u-w_k\right)\Delta t}{\tau_m} \frac{\partial P\left(u,t\right)}{\partial u}$$

We put things together, and again truncate powers of  $\Delta t$  higher than 1 to have

$$\begin{split} P\left(u,t+\Delta t\right) &\approx \left(1+\frac{\Delta t}{\tau_{m}}\right) \left\{P\left(u,t\right) + \frac{u\Delta t}{\tau_{m}}\frac{\partial P\left(u,t\right)}{\partial u} \right. \\ &+ \Delta t \sum_{k} v_{k}(t) \left[-P\left(u,t\right) - \frac{u\Delta t}{\tau_{m}}\frac{\partial P\left(u,t\right)}{\partial u} + P\left(u-w_{k},t\right) + \frac{\left(u-w_{k}\right)\Delta t}{\tau_{m}}\frac{\partial P\left(u,t\right)}{\partial u}\right]\right\} \\ &\approx & P\left(u,t\right) + \frac{\Delta t}{\tau_{m}} P\left(u,t\right) + \frac{u\Delta t}{\tau_{m}}\frac{\partial P\left(u,t\right)}{\partial u} + \Delta t \sum_{k} v_{k}(t) \left[P\left(u-w_{k},t\right) - P\left(u,t\right)\right] \end{split}$$

We now have

$$P(u,t+\Delta t) - P(u,t) = \frac{\Delta t}{\tau_m} P(u,t) + \frac{u\Delta t}{\tau_m} \frac{\partial P(u,t)}{\partial u} + \Delta t \sum_k v_k(t) \left[ P(u-w_k,t) - P(u,t) \right]$$
$$\frac{P(u,t+\Delta t) - P(u,t)}{\Delta t} = \frac{1}{\tau_m} P(u,t) + \frac{u}{\tau_m} \frac{\partial P(u,t)}{\partial u} + \sum_k v_k(t) \left[ P(u-w_k,t) - P(u,t) \right]$$

Again, we do an expansion but this time in  $w_k$ . For this we need the truncated expansion

$$P(u-w_k,t) \approx P(u,t) - w_k \frac{\partial P(u,t)}{\partial u} + \frac{w_k^2}{2} \frac{\partial^2 P(u,t)}{\partial u^2}.$$

Substitution yields

$$\frac{P(u,t+\Delta t) - P(u,t)}{\Delta t} = \frac{1}{\tau_m} P(u,t) + \frac{u}{\tau_m} \frac{\partial P(u,t)}{\partial u} + \sum_k v_k(t) \left[ -w_k \frac{\partial P(u,t)}{\partial u} + \frac{w_k^2}{2} \frac{\partial^2 P(u,t)}{\partial u^2} \right] 
= \frac{1}{\tau_m} P(u,t) + \frac{\partial P(u,t)}{\partial u} \left( \frac{u}{\tau_m} - \sum_k v_k(t) w_k \right) + \frac{1}{2} \sum_k v_k(t) w_k^2 \frac{\partial^2 P(u,t)}{\partial u^2}.$$

Taking  $\Delta t \rightarrow 0$ , we therefore have

$$\tau_{m} \frac{\partial P(u,t)}{\partial t} = P(u,t) + \frac{\partial P(u,t)}{\partial u} \left( u - \tau_{m} \sum_{k} v_{k}(t) w_{k} \right) + \frac{\tau_{m}}{2} \sum_{k} v_{k}(t) w_{k}^{2} \frac{\partial^{2} P(u,t)}{\partial u^{2}}$$

$$= -\frac{\partial}{\partial u} \left[ \left( -u + \tau_{m} \sum_{k} v_{k}(t) w_{k} \right) P(u,t) \right] + \frac{\tau_{m}}{2} \sum_{k} v_{k}(t) w_{k}^{2} \frac{\partial^{2} P(u,t)}{\partial u^{2}}$$

as desired.

First, we verify that for a constant input, at  $t \to \infty$ , the Gaussian density with mean  $RI_0$  and variance  $\sigma/\sqrt{2}$  is a solution of the Fokker-Planck equation. Note that at  $t \to \infty$ , we have  $\frac{\partial P(u,\infty)}{\partial t} = 0$ . Hence, we only have to show that the right-hand side of the equation sums up to zero.

$$-\frac{\partial}{\partial u}\left[\left(-u+\tau_{m}\sum_{k}v_{k}w_{k}\right)P\left(u,\infty\right)\right]=P\left(u,\infty\right)+\frac{\partial P}{\partial u}\left(u-RI_{0}\right)$$

and

$$\frac{\tau_m}{2} \sum_{k} v_k w_k^2 \frac{\partial^2 P}{\partial u^2} = \frac{1}{2} \sigma^2 \frac{\partial}{\partial u} \left[ \frac{1}{\sqrt{\pi} \sigma} \exp\left(-\frac{(u - RI_0)^2}{\sigma^2}\right) \left(-\frac{2(u - RI_0)}{\sigma^2}\right) \right] 
= \frac{1}{2} \sigma^2 \frac{\partial}{\partial u} \left[ P(u, \infty) \left(-\frac{2(u - RI_0)}{\sigma^2}\right) \right] = -\frac{\partial}{\partial u} \left[ P(u, \infty) \left(u - RI_0\right) \right] 
= -\frac{\partial P}{\partial u} (u - RI_0) - P(u, \infty)$$

Combining this two make up he right-hand side of the Fokker-Planck equation which is now clearly zero.

Consider a leaky integrate and fire neuron with Poisson input without threshold. Suppose that there is a single pre-synaptic neuron so that the input rate is v = 5, synaptic weight is w = 1, and membrane time constant is  $\tau_m = 1$ . With this, we have  $RI_0 = \tau_m vw = 5$  and  $\sigma^2 = \tau_m vw^2 = 5$ . A simulation of such a neuron, up to time T = 1000 ms, is found in Figure 1.

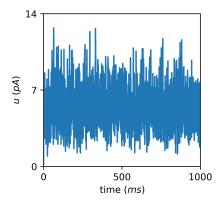


Figure 1: Simulation of a leaky integrate and fire model without threshold.

As given in Eq. 8.46, after a long observation time, the membrane potential follows a Gaussian distribution with mean  $RI_0$  and variance  $\sigma\sqrt{2}$ . To verify this, we use the simulation above and sample the membrane potential trajectory at 100,000 time points and compare the resulting histogram to the analytic prediction. We see in Figure 2 that indeed this is the case.

2. Consider a leaky integrate and fire neuron with Poisson input with threshold  $\theta = 5$ . Suppose that there is a single pre-synaptic neuron so that the input rate is v = 5, synaptic weight is w = 1, and membrane time constant is  $\tau_m = 1$ . With this, we have  $RI_0 = \tau_m vw = 5$  and  $\sigma^2 = \tau_m vw^2 = 5$ . Given that the resting potential is  $u_{rest} = 0$ , the analytic prediction for the expected time to firing, T(x) is the solution of the differential equation  $A(x)\partial_x T(x) + 0.5B(x)\partial_x T(x) = -1$ , with the initial conditions  $T(a \to \infty) = 0$  and T(5) = 0. If we assume that both A(x) and B(x) are constant, and assign  $A(x) = RI_0$  and  $B(x) = \sigma^2$ , the solution of the ODE is given by

$$T(x,a) = -\frac{e^{-2a}(-a+b)}{5(-e^{-2a}+e^{-2b})} + \frac{a}{5} + \frac{e^{-2x}(-a+b)}{5(-e^{-2a}+e^{-2b})} - \frac{t}{5}.$$

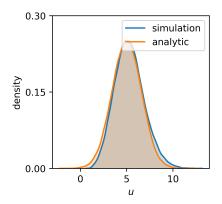


Figure 2: A comparison of the analytic prediction of the distribution of membrane potential and simulated values.

It we start at  $u_{rest}$ , then  $\langle T(u_{rest})\rangle = \lim_{a \to -\infty} T(u_{rest}) = \lim_{a \to -\infty} T(0) = 1$  ms. All these computations were done in MATLAB. Empirical results show that the mean passage time is around 2 ms, as seen in Figure 3.

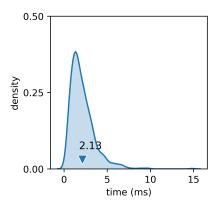


Figure 3: Distribution of passage time at threshold.

3. Suppose inter-arrival times X of spikes follow a gamma distribution  $\Gamma(\alpha,\beta)$ ,  $\alpha$  being the shape parameter, and  $\beta$  the rate. Then  $E[X] = \frac{\alpha}{\beta}$  and  $Var[X] = \frac{\alpha}{\beta^2}$ . It follows that the coefficient of variation is  $CV = \frac{\sqrt{Var[X]}}{E[X]} = \frac{1}{\sqrt{\alpha}}$ .

A special case of the gamma distribution is the Erlang distribution where the shape parameter  $\alpha$  is a positive integer. We parametrize the Erlang distribution with the shape parameter k and rate parameter k. Given K following the Erlang distribution, we have that  $E[X] = \frac{k}{\lambda}$  and  $Var[X] = \frac{k}{\lambda^2}$ . It follows that the coefficient of variation is  $CV = \frac{\sqrt{Var[X]}}{E[X]} = \frac{1}{\sqrt{k}}$ .

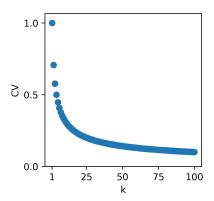


Figure 4: Coefficient of variation of an Erlang distributed random variable given increasing values of the shape parameter k.

As seem in Figure 4, increasing the parameter k decreases the CV. This means that as the number of exponential random variable summands increase, the variation around the resulting mean decreases.

4. We will derive the Fokker-Planck equation corresponding to the haploid Moran model with mutation. The are are two alleles, a and A, with a single locus. The rate of mutation from a to A is given by u, and from A to a is v. Suppose that the total population is N and that  $u, v \in O\left(\frac{1}{N}\right)$ . Given the population n of allele a, we do the substution  $x = \frac{n}{N}$ . First, we setup the following transition probabilities.

$$T\left(x\left|x+\frac{1}{N}\right.\right) = (1-v)\left(x+\frac{1}{N}\right)\left(1-x-\frac{1}{N}\right) + u\left(x+\frac{1}{N}\right)^{2}$$

$$T\left(x\left|x-\frac{1}{N}\right.\right) = (1-u)\left(1-x+\frac{1}{N}\right)\left(x-\frac{1}{N}\right) + v\left(1-x+\frac{1}{N}\right)^{2}$$

$$T\left(x+\frac{1}{N}|x\right) = (1-u)\left(1-x\right)x + v(1-x)^{2}$$

$$T\left(x-\frac{1}{N}|x\right) = (1-v)x(1-x) + ux^{2}$$

We also need the following Taylor expansions of *P*.

$$P\left(x + \frac{1}{N}, t\right) = P(x, t) + \frac{1}{N} \frac{\partial P}{\partial x} + \frac{1}{2N^2} \frac{\partial^2 P}{\partial x^2} + O\left(\frac{1}{N^3}\right)$$
$$P\left(x - \frac{1}{N}, t\right) = P(x, t) - \frac{1}{N} \frac{\partial P}{\partial x} + \frac{1}{2N^2} \frac{\partial^2 P}{\partial x^2} + O\left(\frac{1}{N^3}\right)$$

Putting things together, we have

$$\begin{split} \frac{\partial P\left(x,t\right)}{\partial t} &= T\left(x\left|x+\frac{1}{N}\right.\right) P\left(x+\frac{1}{N},t\right) + T\left(x\left|x-\frac{1}{N}\right.\right) P\left(x-\frac{1}{N},t\right) - P(x,t) \left[T\left(x+\frac{1}{N}|x\right) + T\left(x-\frac{1}{N}|x\right)\right] \\ &= \left\{(1-v)\left(x+\frac{1}{N}\right)\left(1-x-\frac{1}{N}\right) + u\left(x+\frac{1}{N}\right)^2\right\} \left[P(x,t) + \frac{1}{N}\frac{\partial P}{\partial x} + \frac{1}{2N^2}\frac{\partial^2 P}{\partial x^2}\right] \\ &\quad + \left\{(1-u)\left(1-x+\frac{1}{N}\right)\left(x-\frac{1}{N}\right) + v\left(1-x+\frac{1}{N}\right)^2\right\} \left[P(x,t) - \frac{1}{N}\frac{\partial P}{\partial x} + \frac{1}{2N^2}\frac{\partial^2 P}{\partial x^2}\right] \\ &\quad - \left[(1-u)\left(1-x\right)x + v(1-x)^2 + (1-v)x(1-x) + ux^2\right]P(x,t) + O\left(\frac{1}{N^3}\right) \\ &= \frac{1}{N^2}\left\{\frac{\partial P}{\partial x}\left(2-u-3v-4x+4ux-4vx\right) + 2P(u+v-1) + \frac{\partial^2 P}{\partial x^2}\left(\frac{u}{2} + x-x^2 - \frac{ux}{2} - \frac{3vx}{2} + ux^2 + vx^2\right)\right\} \\ &\quad + \frac{1}{N}\left\{\frac{\partial P}{\partial x}\left(ux+vx-v\right) + P(u+v)\right\} + O\left(\frac{1}{N^3}\right) \end{split}$$

Since  $u, v \in O\left(\frac{1}{N}\right)$ , we have  $\frac{u}{N^2}, \frac{v}{N^2} \in O\left(\frac{1}{N^3}\right)$ . The above equation now simplifies to

$$\frac{\partial P}{\partial t} = \frac{1}{N^2} \left\{ \frac{\partial^2 P}{\partial x^2} \left( x - x^2 \right) + \frac{\partial P}{\partial x} \left( 2 - 4x \right) - 2P \right\} + \frac{1}{N} \left\{ \frac{\partial P}{\partial x} \left( ux + vx - v \right) + P \left( u + v \right) \right\} + O \left( \frac{1}{N^3} \right)$$

After removing the  $O\left(\frac{1}{N^3}\right)$  term, a simple expansion will verify that the above is equivalent to

$$\frac{\partial P}{\partial t} = \frac{1}{N} \frac{\partial}{\partial x} \left[ (ux - v(1 - x)) P \right] + \frac{1}{N^2} \frac{\partial^2}{\partial x^2} \left[ x(1 - x) P \right]$$

as required.

5. Here we simulate a two-step adaptation process and track the number of individuals with genotypes 1, 2, and 3, which we will denote by  $n_1$ ,  $n_2$ , and  $n_3$ . Suppose that at any given time  $n_1 + n_2 + n_3 = N$ . Also, assume that genotype 1 mutates to genotype 2 with rate  $\mu_{12}$  and genotype 2 mutates to genotype 3 with rate  $\mu_{23}$ . Given these mutation rates, the transition probabilities are given by

$$\begin{split} &P(n_1+1,n_2-1,n_3\,|n_1,n_2,n_3) = \left(1-\mu_{12}\right)\left(\frac{n_1}{N}\right)\left(\frac{n_2}{N}\right) \\ &P(n_1+1,n_2,n_3-1\,|n_1,n_2,n_3) = \left(1-\mu_{12}\right)\left(\frac{n_1}{N}\right)\left(\frac{n_3}{N}\right) \\ &P(n_1-1,n_2+1,n_3\,|n_1,n_2,n_3) = \left(1-\mu_{23}\right)\left(\frac{n_2}{N}\right)\left(\frac{n_1}{N}\right) + \mu_{12}\left(\frac{n_1}{N}\right)\left(\frac{n_1}{N}\right) \\ &P(n_1-1,n_2,n_3+1\,|n_1,n_2,n_3) = \left(\frac{n_3}{N}\right)\left(\frac{n_1}{N}\right) \\ &P(n_1,n_2+1,n_3-1\,|n_1,n_2,n_3) = \left(1-\mu_{23}\right)\left(\frac{n_2}{N}\right)\left(\frac{n_3}{N}\right) \\ &P(n_1,n_2-1,n_3+1\,|n_1,n_2,n_3) = \left(\frac{n_3}{N}\right)\left(\frac{n_2}{N}\right) + \mu_{23}\left(\frac{n_2}{N}\right)\left(\frac{n_2}{N}\right) \end{split}$$

Taking N = 1000 and  $\mu_{12} = \mu_{23} = \mu$ , with different mutation rates, we perform 3000 simulations of this process, and record the time when the first individual with genotype 3 arrives. The distribution

of the arrival times is presented in Figure 5 and a comparison of the mean arrival time and expected arrival time when  $\mu_{\ll} \frac{1}{N^2}$  is seen in Table 1. A quick look shows that the actual mean arrival times are far larger than expectation. This may in fact be due to the fact that for all the considered cases,  $\mu$  is not significantly less than  $\frac{1}{N^2} = 0.000001$ .

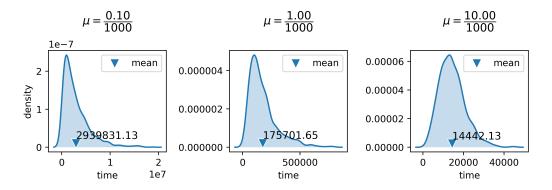


Figure 5: Distribution of time to first arrival of an individual with genotype 3 given different rates  $\mu$ .

	empirical mean	$\frac{1}{\mu}$
$\mu = 0.1/N$	2,939,831	10,000
$\mu = 1/N$	175,701	1,000
$\mu = 10/N$	14,442	100

Table 1: Comparison of the mean time of arrival of the first individual with genotype 3 and the expected time  $\frac{1}{\mu}$ .