

**Probability distributions and expectation practice problems**

1. **Geometric distribution.** The geometric distribution represents the number of trials  $Y$  until a “success”, where at each trial the probability of success is  $p$ . If we want to find the probability we will succeed after  $y$  trials, we will have  $y - 1$  “failures”, each with probability  $(1 - p)$  and then a success. This gives us the probability mass function (pmf):

$$P_Y(y) = (1 - p)^{y-1}p$$

- (a) Verify that the total probability over all values  $y \in \{1, 2, \dots, \infty\}$  sums to 1. *Hint: what is the sum of an infinite geometric series?*
- (b) Verify that the expected value of the geometric distribution is  $E[Y] = \frac{1}{p}$ . *Hint: differentiate the sum of an infinite geometric series twice.*

2. **Exponential distribution.** The continuous analog of the geometric distribution is the exponential distribution. We can think of an exponential random variable  $X$  as the “waiting time” to success without discrete trials (i.e. time it takes to wait for the bus). The probability density function (pdf) for the exponential distribution with parameter  $\lambda$  is:

$$P_X(x) = \lambda e^{-\lambda x}$$

Instead of summing over all possible values  $x$ , with a continuous probability distribution we need to integrate over all possible  $x \in [0, \infty)$ .

- (a) Verify that the total probability over all  $x \in [0, \infty)$  sums to 1.
- (b) Verify that the expected value of the exponential distribution is  $E[X] = \frac{1}{\lambda}$ . *Hint: use integration by parts.*

**Coalescent practice problems**

1. Let  $\mu$  be the per base, per generation mutation rate. Given that the expected time to coalescence for two lineages is  $2N$  generations, how many differences do we expect between two sequences?

2. The expected value of  $T_i$  (time when there are  $i$  lineages) is:

$$E[T_i] = \frac{1}{\binom{i}{2}} = \frac{2}{i(i-1)}.$$

Let  $T_{\text{total}}$  be the total branch length of the coalescent genealogy (sum of all branch lengths). Making use of  $E[T_i]$ , what is  $E[T_{\text{total}}]$  (your result can include a summation)?

3. Using  $E[T_i]$  again, what is the expected value of  $T_{\text{MRCA}}$ , the time to most recent common ancestor of the entire sample? Let the sample size be  $n$ . Simplify your result so it does not include a summation.