

Vignette: Poisson Statistics

1. Poisson Statistics describes distributions of discrete and small numbers. It is often useful in biology and in thinking about the set-up and interpretation of experiments. You probably will see a problem similar to these on Midterm I.
2. P = probability of event: Q = probability of non-event.

$$P+Q = 1$$

A. Remember also that probabilities multiply:

$$P(x,y) = P(x) \cdot P(y)$$

3. For small (discrete) numbers:

$$P(x) = \frac{m^x}{x!} e^{-m}$$

Where: x = number of events
 m = mean number of events
 $P(x)$ = probability of x events

4. This expression is useful for all sorts of experimental considerations and problems. e.g.:

Problem 1: You are interested in determining the "burst size" of a bacteriophage growing on *E. coli*. In order to do that, you mix suspensions of phage and *E. coli* as follows:

1 ml *E. coli* at 5×10^8 / ml.

20 μ l phage at 2.5×10^{10} / ml

- What portion of the bacteria are infected with exactly one phage?
Two phage? Three phage?
- What portion of the bacteria are uninfected? (I.e., what is $P(0)$. Remember here that, by convention, $n^0 = 1$ and $0! = 1$.), so

$$P(0) = e^{-m}$$

Continuing with the experiment, you dilute the infected culture so that when you put 100 μ l into different wells of a multiple-well plate you are putting, on average (m), one bacterial cell per well; you will incubate to lysis and then assay the number of phage in the individual wells to extract burst-size. In the distribution into wells:

- What fraction of the wells gets exactly one cell?
- What fraction of the wells gets one infected cell?

Problem 2. What would the phage/bacterium ratio have to be so that exactly 95% of the bacteria are infected?

Problem 3. You are engaged in a genome project, and are dancing between costs and maximum coverage. You choose a random approach. If you sequence single-pass (one sequence run on a one-strand genome equivalent) on your library, what would be the coverage (fraction nucleotides sequenced single-pass)?

- How many passes for 95% double-strand coverage (a common goal)?

Problem 4: Make-up and solve a problem using Poisson statistics.

Poisson Problems Solved

1. One ml Eco at $5 \times 10^8/\text{ml} = 5 \times 10^8$ cells
 20 μ page at $2.5 \times 10^{10}/\text{ml} = 5 \times 10^8$ phage

$$m = 1 \text{ phage per cell}$$

- a. Cells with exactly one phage?

$$P(1) = [1^1/1!]e^{-m} = e^{-1} = 0.37 \quad (\text{remember that } e^{-1} = \sim 0.37)$$

- b. Cells with exactly two phage?

$$P(2) = [1^2/2!]e^{-1} = e^{-1}/2 = 0.18$$

- c. Cells that are uninfected?

$$P(0) = [1^0/0!]e^{-1} = e^{-1} = 0.37 \quad (\text{remember: } N^0 = 1, 0! = 1)$$

- d. Distribute into wells: $m=1$

$$P(\text{one cell/well}) = 0.37$$

$$P(\text{one infected cell}) = 0.37 \times 0.37 = \sim 0.14$$

2. $m=?$ for $P(\text{infection}) = 0.95$

$$\text{Flip the question: } P(0) = 0.05 = e^{-m}, \quad m \sim 3$$

3. Single-pass means $m=1$

a. $P(\text{sequence for any nt}) = e^{-m} = 0.37$

- b. Double vs. single-strand coverage hinges only on the number of nucleotides covered.

$$\text{For 95\% coverage: as above, } P(0) = 0.05 = e^{-m} : m = \sim 3$$

$$\text{For 95\% double strand coverage: } m = \text{twice single pass} = \sim 6$$