# Chapter 5

# **Discrete Random Variables**

Random variables and their associated probability distributions are a basic component of statistical analyses. A statistician will examine the experiment or study and determine the type of observations or data it produces (continuous, discrete, or categorical) and then select a random variable and its distribution to model these data. We examine here three discrete random variables, the binomial, Poisson, and negative binomial, and their probability distributions. There are other discrete random variables but these three are the most commonly encountered in practice. These variables only take integer values and are typically used to model discrete or count data. We will also see how to calculate the mean and variance for a discrete random variable, using its probability distribution and a quantity called the **expected value**.

The basic concept of a **random variable** is to map the outcome of some random event into a number. For example, consider the dice cube example from Chapter 4. Define a number Y that is the number of spots showing on the dice – Y is a random variable. The sample space for Y would be S = 1, 2, 3, 4, 5, 6 and the events any combination of these values. One requirement for Y to be a random variable is that events of the form  $Y \leq y$ for any real number y are events in the probability space (Mood et al. 1974). For example, suppose that y = 3.5 for the dice cube example. The set defined by  $Y \leq 3.5$  corresponds to the event  $A = \{1, 2, 3\}$  and so is a member of the probability space for this example. This requirement is necessary in order to calculate probabilities for the random variable, and there is always a probability distribution associated with a particular random variable.

## 5.1 Binomial distribution

Binomial random variables are commonly used to model categorical observations or data that have two outcomes or states. For example, suppose we are sampling animals and classifying them into two age classes, say either adult (an event A) or juvenile (J). If we sample a single individual and classify it, the sample space would be  $S = \{A, J\}$ . We could then define a probability distribution such that  $P[\{A\}] = p$  and  $P[\{J\}] = 1 - p$ , where p is the probability of observing an adult. Then, a random variable Y equal to the **number** of adults would be a binomial random variable. The random variable Y would have a sample space  $S = \{0, 1\}$  corresponding to the number of adults. We could write the probability distribution for these two events as

$$P[Y = y] = p^{y}(1-p)^{1-y},$$
(5.1)

where y = 0 or 1. To see how this formula works, suppose we want the probability for Y = 1, so that y = 1. Inserting y = 1 in the above formula, we obtain

$$P[Y=1] = p^{1}(1-p)^{1-1} = p^{1}(1-p)^{0} = p.$$
(5.2)

To find the probability for Y = 0, we insert y = 0 in the formula to find

$$P[Y=0] = p^{0}(1-p)^{1-0} = p^{0}(1-p)^{1} = 1-p.$$
(5.3)

Suppose that we now sample two animals and let Y again be the number of adults. The sample space for Y would now be  $S = \{0, 1, 2\}$ . What would be the probability distribution for this random variable? Assuming the two animals sampled are independent events, the probability of seeing two adults (Y = 2) in a row would be  $p \times p = p^2$ , while two juveniles (Y = 0) would be  $(1-p) \times (1-p) = (1-p)^2$ . There are two ways of having one adult and one juvenile, a adult first and a juvenile second, or vice versa. The probability for each is  $p \times (1-p)$ , so the probability of seeing one adult would be twice that, or 2p(1-p). A general formula describing the probability distribution for this variable would be

$$P[Y = y] = {\binom{2}{y}} p^{y} (1-p)^{2-y}.$$
(5.4)

where

$$\binom{2}{y} = \frac{2!}{y!(2-y)!}.$$
(5.5)

#### 5.1. BINOMIAL DISTRIBUTION

The quantity  $\binom{2}{y}$ , known as a binomial coefficient, provides a way of calculating the number of ways y adults can occur among 2 sampled animals. It is often read as '2 choose y'. It makes use of factorials, which are defined for an integer j as the product  $j \times (j-1) \times (j-2) \dots \times 1$ . For example,  $4! = 4 \times 3 \times 2 \times 1$ . By convention, 0! = 1.

To see how this distribution works, we will calculate the probability for different values of y. We have

$$P[Y=0] = {\binom{2}{0}} p^0 (1-p)^{2-0} = \frac{2!}{0!(2-0)!} (1-p)^2$$
(5.6)

$$=\frac{2\times 1}{1(2\times 1)}(1-p)^2$$
(5.7)

$$=\frac{2}{2}(1-p)^2 = (1-p)^2$$
(5.8)

and

$$P[Y=1] = {\binom{2}{1}} p^1 (1-p)^{2-1} = \frac{2!}{1!(2-1)!} p(1-p)$$
(5.9)

$$=\frac{2\times 1}{1(1)}p(1-p)$$
(5.10)

$$=\frac{2}{1}p(1-p) = 2p(1-p).$$
(5.11)

Finally, we have

$$P[Y=2] = {\binom{2}{2}} p^2 (1-p)^{2-2} = \frac{2!}{2!(2-2)!} p^2$$
(5.12)

$$=\frac{2\times 1}{(2\times 1)1}p^2$$
(5.13)

$$=\frac{2}{2}p^2 = p^2.$$
 (5.14)

Do these probabilities sum to 1, satisfying this requirement for a probability distribution? We have  $(1-p)^2+2p(1-p)+p^2 = (1-p)(1-p)+2p-2p^2+p^2 = 1-2p+p^2+2p-2p^2+p^2 = 1$ .

Suppose that we continue to sample l different animals, and let Y be the number of adults. The sample space for this binomial random variable would be  $S = \{0, 1, 2, ..., l\}$ . The probability distribution for this random variable

is called the **binomial distribution**, and can be written using the formula

$$P[Y = y] = f(y) = {l \choose y} p^{y} (1-p)^{l-y}$$
(5.15)

where y = 0, 1, 2, ..., l (Mood et al. 1974). The notation f(y) is often used to denote a probability distribution, which is a function of y given the parameter values.

### 5.1.1 Binomial distribution - SAS demo

The SAS program below calculates and plots the binomial probabilities for different values of y using the SAS function pdf, given the values of the binomial parameters l and p. The probabilities are plotted for three different values of p, with l = 10. We see that for p = 0.5 the probability distribution has a peak at y = 5 (Fig. 5.1), indicating that five adults is the most likely outcome in 10 sampled animals. For p = 0.25 an adult occurs only 25% of the time, and so the probability distribution shifts to the left, with y = 2 having the highest probability (Fig. 5.2). For an adult almost certain, p = 0.9, then the probability distribution is shifted to the right with the peak at y = 9(Fig. 5.3).

```
____ SAS Program -
```

```
* binom_plot.sas;
options pageno=1 linesize=80;
goptions reset=all;
title "Plot probabilities for the binomial distribution";
title2 "k = 10, p = 0.5";
data binom_plot;
    * Binomial parameters here;
    1 = 10;
    p = 0.5;
    do y=0 to 1;
        * Binomial distribution function;
        proby = pdf('binomial',y,p,l);
        * Output y and proby to SAS data file;
        output;
    end;
run;
* Print data;
proc print data=binom_plot;
```

```
run;
* Plot probabilities;
proc gplot data=binom_plot;
    plot proby*y=1 / vref=0 wvref=3 vaxis=axis1 haxis=axis1;
    symbol1 i=needle v=dot c=red width=3 height=2;
    axis1 label=(height=2) value=(height=2) width=3 major=(width=2) minor=none;
run;
quit;
```

		SAS Out	put		
Plot prob	abiliti	es for ] = 10.	the bino $p = 0.5$	mial distribution	1 March 17, 2010
		± 10,	p 0.0	obioi weamebady,	nuron 17, 2010
Ob	os l	р	У	proby	
	1 10	0.5	0	0.00098	
	2 10	0.5	1	0.00977	
	3 10	0.5	2	0.04395	
	4 10	0.5	3	0.11719	
	5 10	0.5	4	0.20508	
	6 10	0.5	5	0.24609	
	7 10	0.5	6	0.20508	
	8 10	0.5	7	0.11719	
	9 10	0.5	8	0.04395	
1	.0 10	0.5	9	0.00977	
1	1 10	0.5	10	0.00098	







Figure 5.2: Binomial distribution for l=10, p=0.25 Plot probabilities for the binomial distribution  $_{\rm I=10,\,p=0.25}$ 

Figure 5.3: Binomial distribution for l = 10, p = 0.9Plot probabilities for the binomial distribution I = 10, p = 0.9



# 5.2 Poisson distribution

Poisson random variables are commonly used to model counts of organisms or events in either space or time. For example, a Poisson random variable could be used to model the number of organisms in a sampling quadrat, or the number of flu infections per week in a city. The sample space for a Poisson random variable Y is  $S = \{0, 1, 2, ..., \infty\}$ , implying there is no upper limit on the counts. The Poisson distribution is given by the formula

$$P[Y = y] = f(y) = \frac{e^{-\lambda}\lambda^y}{y!}$$
 (5.16)

where  $y = 0, 1, 2, ..., \infty$ . The parameter  $\lambda$  controls the shape of the distribution and is equal to the mean value of Y. For example, suppose the  $\lambda = 2$ . We have

$$P[Y=0] = f(0) = \frac{e^{-2}2^0}{0!} = \frac{0.13534(1)}{1} = 0.13534,$$
 (5.17)

$$P[Y=1] = f(1) = \frac{e^{-2}2^1}{1!} = \frac{0.13534(2)}{1} = 0.27068,$$
 (5.18)

$$P[Y=2] = f(2) = \frac{e^{-2}2^2}{2!} = \frac{0.13534(4)}{2} = 0.27068,$$
 (5.19)

$$P[Y=3] = f(3) = \frac{e^{-2}2^3}{3!} = \frac{0.13534(8)}{6} = 0.18045,$$
(5.20)

$$P[Y=4] = f(4) = \frac{e^{-2}2^4}{4!} = \frac{0.13534(16)}{24} = 0.09023$$
(5.21)

and so forth.

The Poisson distribution can arise in nature if certain assumptions hold true about the underlying process generating the data or observations (Mood et al. 1974, Snyder & Miller 1991). Suppose that we define an occurrence as a plant being present in a quadrat, or a case of disease occurring in a particular interval of time. For the distribution of occurrences to be Poisson, we first need the probability of more than one occurrence to be small relative to the probability of exactly one occurrence, for a sufficiently small area of space (or short period of time). In other words, two events are unlikely to occur in a small area or period of time. Second, the number of occurrences in different areas of space (or time intervals) should be independent. Another way of obtaining the Poisson distribution is as a limiting case of the binomial distribution. It can be shown that if lp is held constant (by making p small) while  $l \to \infty$ , the binomial distribution approaches a Poisson with  $\lambda = lp$ .

### 5.2.1 Poisson distribution - SAS demo

The following SAS program illustrates how the Poisson distribution varies for different values of  $\lambda$ . It is similar to the binomial distribution program, using the SAS function pdf to again find the probabilities (see below). We see that as  $\lambda$  increases, the Poisson distribution shifts to the right (Fig. 5.4, 5.5).

\_ SAS Program \_

```
* Poisson_plot.sas;
options pageno=1 linesize=80;
goptions reset=all;
title "Plot probabilities for the Poisson distribution";
title2 "lambda = 2";
data poisson_plot;
    * Poisson parameter here;
    lambda = 2;
    * Maximum value of y for plot;
    ymax = 20;
    do y=0 to ymax;
        * Poisson distribution function;
        proby = pdf('poisson',y,lambda);
        * Output y and proby to SAS data file;
        output;
    end;
run;
* Print data;
proc print data=poisson_plot;
run;
* Plot probabilities;
proc gplot data=poisson_plot;
    plot proby*y=1 / vref=0 wvref=3 vaxis=axis1 haxis=axis1;
    symbol1 i=needle v=dot c=red width=3 height=2;
    axis1 label=(height=2) value=(height=2) width=3 major=(width=2) minor=none;
run;
quit;
```

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		SA:	S Output				
Plot	probal	oilities	for the	Poisso	n distribution		
	-		lambda =	2	09:04 Wednesday	, March 17,	201
	Obs	lambda	ymax	У	proby		
	1	2	20	0	0.13534		
	2	2	20	1	0.27067		
	3	2	20	2	0.27067		
	4	2	20	3	0.18045		
	5	2	20	4	0.09022		
	6	2	20	5	0.03609		
	7	2	20	6	0.01203		
	8	2	20	7	0.00344		
	9	2	20	8	0.00086		
	10	2	20	9	0.00019		
	11	2	20	10	0.00004		
	12	2	20	11	0.00001		
	13	2	20	12	0.00000		
	14	2	20	13	0.00000		
	15	2	20	14	0.00000		
	16	2	20	15	0.00000		
	17	2	20	16	0.00000		
	18	2	20	17	0.00000		
	19	2	20	18	0.00000		
	20	2	20	19	0.00000		
	21	2	20	20	0.00000		









# 5.3 Negative binomial distribution

Another useful tool for modeling count data is the negative binomial distribution. It can be thought of as a mixture of Poisson distributions, each with a different value of  $\lambda$ . For example, suppose that we are sampling insects in a forest across a number of locations. At the *ith* location the distribution of insects might be Poisson with parameter  $\lambda_i$ , but  $\lambda_i$  also differs among locations. Then the distribution of insects, considered across all locations, may have a negative binomial distribution. Because the density of most organisms typically varies in space, the negative binomial distribution often provides a better description of count data than the Poisson. The sample space for a negative binomial random variable Y is  $S = \{0, 1, 2, ..., \infty\}$ , the same as the Poisson. The probability distribution for the negative binomial is given by the formula

$$P[Y = y] = f(y) = \frac{\Gamma(k+y)}{\Gamma(y+1)\Gamma(k)} \frac{(m/(k+m))^y}{(1+m/k)^k}$$
(5.22)

where  $y = 0, 1, 2, ..., \infty$ . The  $\Gamma$  symbol stands for the gamma function, which behaves like the factorial function but can be applied to non-integer quantities. The negative binomial distribution has two parameters, m and k, with m the mean of the distribution and k controlling its shape. For large values of k the negative binomial distribution approaches the Poisson distribution, while for small k the distribution becomes increasingly skewed to the right. See Bliss and Fisher (1953) for further information on this distribution.

### 5.3.1 Negative binomial distribution - SAS demo

The SAS program below shows how the shape of the negative binomial distribution varies with the parameter k. The program directly calculates the probabilities using the formula above, rather than the SAS pdf function, because we are using a different parameterization of the distribution than SAS. We see that distribution becomes more skewed to the right as k decreases (Fig. 5.6, 5.7).

```
_____ SAS Program _____
```

```
* negbin_plot.sas;
options pageno=1 linesize=80;
goptions reset=all;
title "Plot probabilities for the negative binomial distribution";
title2 "m = 5, k = 5";
data negbin_plot;
    * negative binomial parameters here;
   m = 5; k = 5;
    * Maximum value of y for plot;
   ymax = 20;
    do y=0 to ymax;
        * Negative binomial distribution function;
        proby = (gamma(k+y)/(gamma(y+1)*gamma(k)))*((m/(k+m))**y/(1+m/k)**k);
        * Output y and proby to SAS data file;
        output;
    end;
run;
* Print data;
proc print data=negbin_plot;
run;
* Plot probabilities;
proc gplot data=negbin_plot;
    plot proby*y=1 / vref=0 wvref=3 vaxis=axis1 haxis=axis1;
    symbol1 i=needle v=dot c=red width=3 height=2;
    axis1 label=(height=2) value=(height=2) width=3 major=(width=2) minor=none;
run;
quit;
```

## CHAPTER 5. DISCRETE RANDOM VARIABLES

			SAS 0 <sup>.</sup>	utput _		
Plo	ot probabilit	ies	for the	negati	ve bind	omial distribution
			m =	5, k =	5	15:36 Sunday, March 21, 2
	Obs	m	k	ymax	у	proby
	1	5	5	20	0	0.03125
	2	5	5	20	1	0.07813
	3	5	5	20	2	0.11719
	4	5	5	20	3	0.13672
	5	5	5	20	4	0.13672
	6	5	5	20	5	0.12305
	7	5	5	20	6	0.10254
	8	5	5	20	7	0.08057
	9	5	5	20	8	0.06042
	10	5	5	20	9	0.04364
	11	5	5	20	10	0.03055
	12	5	5	20	11	0.02083
	13	5	5	20	12	0.01389
	14	5	5	20	13	0.00908
	15	5	5	20	14	0.00584
	16	5	5	20	15	0.00370
	17	5	5	20	16	0.00231
	18	5	5	20	17	0.00143
	19	5	5	20	18	0.00087
	20	5	5	20	19	0.00053
	21	5	5	20	20	0.00032





Figure 5.7: Negative binomial distribution for m=5, k=1 Plot probabilities for the negative binomial distribution  $_{\rm m=5,\,k=1}^{\rm m=5,\,k=1}$ 



# 5.4 Expected values for discrete distributions

We have already seen how to calculate the mean, variance, and standard deviation for a set of observations (see Chapter 3). It is possible to calculate analogous quantities for probability distributions, such as the binomial, using the concept of an **expected value**.

Let Y be a random variable with some discrete probability distribution, such as the binomial, Poisson, or other distribution. The expected value or theoretical mean of Y, denoted by the expression E[Y], is defined by the equation

$$E[Y] = \sum_{y} yP[Y = y] = \sum_{y} yf(y).$$
 (5.23)

Here the summation is taken over all possible values of y for the probability distribution. The expected value is a weighted average of each possible value of y, with the weights being the probability associated with each y. It is a measure of the central location of the distribution of Y, in analogy to the sample mean  $\overline{Y}$  for a data set. The expected value of Y can also be thought of as the sample mean  $\overline{Y}$  of an infinitely large number of observations of Y.

For example, let Y have a binomial distribution with l = 5 and p = 0.2. We will first calculate some probabilities for the binomial distribution, then use them to calculate the expected value of Y, or E[Y]. We have

$$P[Y=0] = f(0) = {\binom{5}{0}} 0.2^0 (1-0.2)^{5-0}$$
(5.24)

$$=\frac{5!}{0!(5-0)!}1(0.8^5) \tag{5.25}$$

$$=\frac{120}{1(120)}0.32768\tag{5.26}$$

$$= 0.32768.$$
 (5.27)

$$P[Y=1] = f(1) = {\binom{5}{1}} 0.2^1 (1-0.2)^{5-1}$$
(5.28)

$$=\frac{5!}{1!(5-1)!}0.2(0.8^4) \tag{5.29}$$

$$=\frac{120}{1(24)}0.08192\tag{5.30}$$

$$= 0.40960.$$
 (5.31)

$$P[Y=2] = f(2) = {\binom{5}{2}} 0.2^2 (1-0.2)^{5-2}$$
(5.32)

$$=\frac{5!}{2!(5-2)!}0.04(0.8^3) \tag{5.33}$$

$$=\frac{120}{2(6)}0.02048\tag{5.34}$$

$$= 0.20480.$$
 (5.35)

$$P[Y=3] = f(3) = {\binom{5}{3}} 0.2^3 (1-0.2)^{5-3}$$
(5.36)

$$=\frac{5!}{2!(5-2)!}0.008(0.8^2) \tag{5.37}$$

$$=\frac{120}{2(6)}0.00512\tag{5.38}$$

$$= 0.05120. \tag{5.39}$$

$$P[Y=4] = f(4) = {\binom{5}{4}} 0.2^4 (1-0.2)^{5-4}$$
(5.40)

$$=\frac{5!}{4!(5-4)!}0.0016(0.8^{1}) \tag{5.41}$$

$$=\frac{120}{24(1)}0.00128\tag{5.42}$$

$$= 0.00640.$$
 (5.43)

$$P[Y=5] = f(5) = {\binom{5}{5}} 0.2^5 (1-0.2)^{5-5}$$
(5.44)

$$=\frac{5!}{5!(5-5)!}0.00032(0.8^{0}) \tag{5.45}$$

$$=\frac{120}{120(1)}0.00032\tag{5.46}$$

$$= 0.00032.$$
 (5.47)

These probabilities sum to 1, indicating our calculations are correct. Alternately, we could use the SAS program binom\_plot.sas to find these probabilities.

We will now calculate E[Y] using these probabilities and the formula for E[Y] given above. We have

$$E[Y] = \sum_{y} yf(y) = 0(0.32768) + 1(0.40960) + 2(0.20480)$$
(5.48)

$$+3(0.05120) + 4(0.00640) + 5(0.00032)$$
(5.49)

$$= 0 + 0.40960 + 0.40960 \tag{5.50}$$

$$+0.15360 + 0.02560 + 0.00160 \tag{5.51}$$

$$= 1.00000$$
 (5.52)

So, E[Y] = 1 for the binomial distribution with l = 5 and p = 0.2.

For the binomial distribution in general, it can be shown that

$$E[Y] = lp \tag{5.53}$$

for any value of l and p. Thus, the expected value or theoretical mean for the binomial distribution can be easily calculated given the parameters of this distribution. Plugging l = 5 and p = 0.2 into this equation, we obtain  $E[Y] = 5 \times 0.2 = 1.0$ , the same value as obtained using the expected value formula.

Other probability distributions would have a different formula for the expected value or theoretical mean, but the formula always involves the parameters of the distribution. For the Poisson distribution it can be shown that  $E[Y] = \lambda$ , while for the negative binomial distribution E[Y] = m.

### 5.4.1 Variance for discrete distributions

We can also define the theoretical variance for a random variable Y using expected values. This variance measures the dispersion of Y, and can also be

thought of as the sample variance  $s^2$  of an infinite number of observations. The variance of a discrete random variable Y, denoted by Var[Y], is defined as

$$Var[Y] = \sum_{y} (y - E[Y])^2 P[Y = y] = \sum_{y} (y - E[Y])^2 f(y).$$
(5.54)

Note that this formula makes use of E[Y], so it must be calculated first. As an example, let Y have the same binomial distribution as before, with l = 5and p = 0.2, for which E[Y] = 1. Using the probabilities calculated above, we have

$$Var[Y] = \sum_{y} (y - E[Y])^2 f(y)$$
(5.55)  
=  $(0 - 1)^2 (0.32768) + (1 - 1)^2 (0.40960) + (2 - 1)^2 (0.20480)$ (5.56)  
+  $(3 - 1)^2 (0.05120) + (4 - 1)^2 (0.00640) + (5 - 1)^2 (0.00032)$ (5.57)  
=  $1(0.32768) + 0(0.40960) + (1)0.20480$ (5.58)  
+  $4(0.05120) + 9(0.00640) + (16)0.00032$ (5.59)  
=  $0.32768 + 0 + 0.20480 + 0.20480 + 0.05760 + 0.00512$ (5.60)  
=  $0.8$ .(5.61)

For the binomial distribution, it can be mathematically shown that for any value of l and p, we have

$$Var[Y] = lp(1-p).$$
 (5.62)

Thus, the theoretical variance for the binomial distribution can also be calculated using the parameters of this distribution. Plugging l = 5 and p = 0.2 into this equation, we obtain Var[Y] = 5(0.2)(1 - 0.2) = 0.8, the same value as obtained using the variance formula.

Other probability distributions would have a different formula for the theoretical variance. For the Poisson distribution it can be shown that  $Var[Y] = \lambda$ . Because  $E[Y] = \lambda$  for the Poisson, this implies the mean and variance of a Poisson random variable are equal. For the negative binomial distribution,  $Var[Y] = m + m^2/k$ , while E[Y] = m. This implies the variance of the negative binomial is always greater than its mean. The theoretical standard deviation is simply  $\sqrt{Var[Y]}$ .

# 5.5 Discrete random variables and samples

Discrete random variables like the binomial and Poisson are used to model real observations that are counts. But how well do these mathematical quantities match the behavior of the observations? We will now develop a graphical method of comparing the observed data with the pattern expected for discrete random variables, in particular the Poisson and negative binomial distributions. There are also statistical procedures called goodness-of-fit tests that are used for this purpose, but we defer this to Chapter 20.

### 5.5.1 Parasitic wasps - SAS demo

Small insects are often sampled using sticky-traps, which are small cards covered with a substance called Tanglefoot ( $\mathbb{R}$ ) (The Tanglefoot Company, Grand Rapids, MI). For example, Reeve & Cronin (2010) used this method to sample populations of the parasitic wasp *Anagrus columbi*, which attacks eggs of the planthopper *Prokelisia crocea*. Suppose n = 100 traps are deployed for some period of time, then the traps collected and the wasps counted. If individual wasps are randomly and independently distributed across the field, we would expect the number of wasps per trap to have a Poisson distribution. We can then compare the observed distribution with the expected one for the Poisson distribution, to see if they resemble one another. If so, we can say the Poisson distribution provides an adequate description of these observations.

The first step in this procedure is simply to tabulate the number of traps with 0, 1, 2, 3, ... wasps, which is the observed frequency distribution. We can use proc freq in SAS to accomplish this task as in the following program. The numbers listed as data here are the number of wasps for each of the n = 100sticky-traps. The statement tables y tells proc freq to count the number of observations for each value of y in the data set. The output generated is a table of these frequencies. \_\_\_\_\_ SAS Program \_\_\_\_\_

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	SAS Out	tput _						
	Tabulate Poisson data							1
				10:40	Friday,	March	26,	2010
	Obs	з у						
	1	L 4						
		2 6						
	3	з з						
	4	1 5						
	Ę	5 3						
etc.								
	98	5 3						
	96	6 0						
	97	7 1						
	98	31						
	99	9 0						
	100	) 2						

Tabulate Poisson data

10:40 Friday, March 26, 2010

3

#### The FREQ Procedure

			Cumulative	Cumulative
у	Frequency	Percent	Frequency	Percent
0	6	6.00	6	6.00
1	15	15.00	21	21.00
2	17	17.00	38	38.00
3	29	29.00	67	67.00
4	20	20.00	87	87.00
5	6	6.00	93	93.00
6	5	5.00	98	98.00
7	1	1.00	99	99.00
8	1	1.00	100	100.00

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We now want to compare these observed frequencies with those expected for the Poisson distribution. We first need to estimate the Poisson parameter  $\lambda$  from the observed data using  $\bar{Y}$  (see Chapter 8 for a justification). We then calculate the Poisson probabilities for  $\lambda = \bar{Y}$ , obtaining P[Y = y] for values of y that spans or better exceeds the range of y values in the data set. Because P[Y = y] is the probability or proportion of observations that take the value y, the expected frequency with n observations is therefore equal to  $n \times P[Y = y]$ . We can then compare the observed frequencies with the expected ones generated using the Poisson distribution. These calculations can be automated using the SAS program listed below. The program first uses proc univariate to find  $n, \bar{Y}$ , and the sample variance  $s^2$  for the observed frequencies. We let proc univariate know that the data are in the form of frequencies (the variable obsfreq), rather than individual observations, by adding the command freq obsfreq.

The program then passes these results to a data step where the Poisson probabilities and expected frequencies are calculated, which are then plotted across a range of y values using proc gplot. See SAS output and graph below. We first see that sample mean and variance are similar in magnitude  $(\bar{Y} = 2.910 \text{ vs. } s^2 = 2.628)$ , suggesting these data are close to Poisson (recall that  $E[Y] = Var[Y] = \lambda$  for this distribution). In addition, the observed and expected frequencies are quite similar, again implying an adequate fit by the Poisson distribution. There are some small differences in the observed and expected frequencies, which is to be expected because the observed ones are random quantities.

```
_ SAS Program _
```

```
* Poisson_fit.sas;
options pageno=1 linesize=80;
goptions reset=all;
title 'Fitting the Poisson to frequency data';
data poisson;
    input y obsfreq;
    * Generate offset y values for plot;
    yexp = y - 0.1; yobs = y + 0.1;
    datalines:
0
    6
1
  15
2
  17
3
   29
4
   20
```

```
6
5
6
   5
7
   1
  1
8
9
   0
10 0
;
run;
* Print data set;
proc print data=poisson;
run;
* Descriptive statistics, save ybar, n, and var to data file;
proc univariate data=poisson;
   var y;
   histogram y / vscale=count;
   freq obsfreq;
   output out=stats mean=ybar n=n var=var;
run;
* Print output data file;
proc print data=stats;
run;
* Calculate expected frequencies using ybar;
data poisfit;
   if _n_ = 1 then set stats;
    set poisson;
   poisprob = pdf('poisson',y,ybar);
    expfreq = n*poisprob;
run;
* Print observed and expected frequencies;
proc print data=poisfit;
run;
* Plot observed and expected frequencies;
proc gplot data=poisfit;
    plot expfreq*yexp=1 obsfreq*yobs=2 / overlay legend=legend1 vref=0 wvref=3
    vaxis=axis1 haxis=axis1;
    symbol1 i=needle v=circle c=red width=3 height=2;
    symbol2 i=needle v=square c=blue width=3 height=2;
    axis1 label=(height=2) value=(height=2) width=3 major=(width=2) minor=none;
    legend1 label=(height=2) value=(height=2);
run;
quit;
```

	Š	SAS Outp	ut					
ting	g the	Poisson	to f	requen	cy data 10:40 H	'riday,	March	26
s	У	obsfr	eq	yexp	yobs			
1	0	6		-0.1	0.1			
2	1	15		0.9	1.1			
3	2	17		1.9	2.1			
4	3	29		2.9	3.1			
5	4	20		3.9	4.1			
6	5	6		4.9	5.1			
7	6	5		5.9	6.1			
8	7	1		6.9	7.1			
9	8	- 1		7.9	8.1			
0	9	0		8.9	9 1			
1	10	0		9.9	10.1			
	The U	JNIVARIA Variab	TE Pr le:	ocedur	10:40 H	riday,	March	26
	The (	JNIVARIA Variab Freq:	TE Pr le: obsfr	ocedur y eq	10:40 H	Friday,	March	26
	The U	JNIVARIA Variab Freq: Mome	TE Pr le: obsfr nts	ocedur y eq	10:40 H	Friday,	March	26
	The U	JNIVARIA Variab Freq: Mome 100	TE Pr le: obsfr nts Sum	ocedur y eq Weight	10:40 H e s	Friday,	March	26
	The (	JNIVARIA Variab Freq: Mome 100 2.91	TE Pr le: obsfr nts Sum Sum	ocedur y eq Weight Observ	10:40 H e s ations	Friday,	March 100 291	26
1	The (	JNIVARIA Variab Freq: Mome 100 2.91 L6681	TE Pr le: obsfr nts Sum Sum Vari	ocedur y eq Weight Observ ance	10:40 H e s ations	Friday, 2.62	March 100 291 818182	26
1	The U L.6211	JNIVARIA Variab Freq: Mome 100 2.91 L6681 D9921	TE Pr le: obsfr nts Sum Sum Vari Kurt	ocedur y eq Weight Observ ance osis	10:40 H e s ations	Friday, 2.62 0.31	March 100 291 818182 136421	26
10	The ( 1.6211 ).3950	JNIVARIA Variab Freq: Mome 100 2.91 16681 99921 1107	TE Pr le: obsfr nts Sum Sum Vari Kurt Corr	ocedur y eq Weight Observ ance osis ected	10:40 H e s ations SS	Friday, 2.62 0.31	March 100 291 818182 136421 260.19	26
1 0 5	The U L.6211 D.3950 55.710	JNIVARIA Variab Freq: Mome 100 2.91 16681 09921 1107 01996	TE Pr le: obsfr nts Sum Sum Vari Kurt Corr Std	ocedur y eq Weight Observ ance osis ected Error	10:40 H e ations SS Mean	2.62 0.31 0.16	March 100 291 818182 136421 260.19 211668	26
1 0 5 B	The [ 1.6211 ).3950 55.710 Basic	JNIVARIA Variab Freq: Mome 100 2.91 16681 09921 1107 01996 Statist	TE Pr le: obsfr nts Sum Sum Vari Kurt Corr Std ical	ocedur y eq Weight Observ ance osis ected Error Measur	10:40 H e s ations SS Mean es	2.62 0.31 0.16	March 100 291 818182 136421 260.19 211668	26
1 0 5 8	The U L.6211 D.3950 55.710 Basic	JNIVARIA Variab Freq: Mome 100 2.91 16681 09921 1107 01996 Statist	TE Pr le: obsfr nts Sum Sum Vari Kurt Corr Std ical	ocedur y eq Weight Observ ance osis ected Error Measur Variab	10:40 H e sations SS Mean es ility	2.62 0.31 0.16	100 291 818182 136421 260.19 211668	26
1 0 5 B n	The U L.6211 D.3950 55.710 Basic	JNIVARIA Variab Freq: Mome 100 2.91 16681 09921 1107 01996 Statist	TE Pr le: obsfr nts Sum Vari Kurt Corr Std ical	ocedur y eq Weight Observ ance osis ected Error Measur Variab	10:40 H e s ations SS Mean es ility	2.62 0.31 0.16	March 100 291 818182 136421 260.19 211668	26
1 0 5 B 9100	The U L.6211 D.3950 55.710 Basic	JNIVARIA Variab Freq: Mome 100 2.91 16681 09921 1107 01996 Statist Stat D	TE Pr le: obsfr nts Sum Vari Kurt Corr Std ical	ocedur y eq Weight Observ ance osis ected Error Measur Variab ion	10:40 H e s ations SS Mean es ility	2.62 0.31 0.16 1.621	March 100 291 818182 136421 260.19 211668	26

Interquartile Range 2.00000

Tests for Location: MuO=0

Test	-S	tatistic-	p Valu	1e
Student's t	t	17.95003	Pr >  t	<.0001
Sign	М	47	Pr >=  M	<.0001
Signed Rank	S	2232.5	Pr >=  S	<.0001

Quantiles (Definition 5)

Quantile	Estimate
100% Max 99% 95% 90% 75% Q3 50% Median 25% 01	8.0 7.5 6.0 5.0 4.0 3.0
25% Q1 10% 5% 1%	2.0 1.0 0.0 0.0
0% Min	0.0

Fitting the Poisson to frequency data 3 10:40 Friday, March 26, 2010

The UNIVARIATE Procedure Variable: y

#### Freq: obsfreq

#### Extreme Observations

	Lowest			Highest	
Value	Freq	Obs	Value	Freq	Obs
0 1	6 15	1 2	4 5	20 6	5 6

2	17	3	6	5	7	
3	29	4	7	1	8	
4	20	5	8	1	9	

Fitting the Poisson to frequency data 4 10:40 Friday, March 26, 2010

Obs	n	ybar	var
1	100	2.91	2.62818

Fitting the Poisson to frequency data 5 10:40 Friday, March 26, 2010

Obs	n	ybar	var	У	obsfreq	yexp	yobs	poisprob	expfreq
1	100	2.91	2.62818	0	6	-0.1	0.1	0.05448	5.4476
2	100	2.91	2.62818	1	15	0.9	1.1	0.15852	15.8524
3	100	2.91	2.62818	2	17	1.9	2.1	0.23065	23.0653
4	100	2.91	2.62818	3	29	2.9	3.1	0.22373	22.3733
5	100	2.91	2.62818	4	20	3.9	4.1	0.16277	16.2766
6	100	2.91	2.62818	5	6	4.9	5.1	0.09473	9.4730
7	100	2.91	2.62818	6	5	5.9	6.1	0.04594	4.5944
8	100	2.91	2.62818	7	1	6.9	7.1	0.01910	1.9100
9	100	2.91	2.62818	8	1	7.9	8.1	0.00695	0.6947
10	100	2.91	2.62818	9	0	8.9	9.1	0.00225	0.2246
11	100	2.91	2.62818	10	0	9.9	10.1	0.00065	0.0654

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Figure 5.8: Observed and expected frequencies - Poisson distribution Fitting the Poisson to frequency data



### 5.5.2 Corn borers - SAS demo

We now examine the spatial distribution of an insect pest, the European corn borer Ostrinia nubilalis, as reported by Bliss and Fisher (1953). The number of borers was recorded for 120 hills in which corn was planted. These data are already tabulated and can be directly inserted in the SAS program poisson\_fit.sas (see below). For this example, we see that the Poisson distribution provides a relatively poor fit (see Fig. 5.9) - there are more zeroes (y = 0) and large values  $(y \ge 7)$  in the observed frequencies than predicted by the Poisson. We also note that the sample variance  $s^2 = 7.770$  is considerably larger than the mean  $\overline{Y} = 3.167$ , while for the Poisson these two quantities should be equal. This finding also suggests that these data are not Poisson in distribution.

\_ SAS Program \_

```
* Poisson_fit2.sas;
options pageno=1 linesize=80;
goptions reset=all;
title 'Fitting the Poisson to frequency data';
data poisson;
        input y obsfreq;
        * Generate offset y values for plot;
        yexp = y - 0.1; yobs = y + 0.1;
        datalines;
   24
0
1
   16
2
   16
3
   18
4
   15
5
    9
    6
6
7
    5
8
    3
9
    4
   3
10
11
   0
12
   1
;
run;
* Print data set;
proc print data=poisson;
run:
* Descriptive statistics, save ybar, n, and var to data file;
```

```
proc univariate data=poisson;
        var y;
        histogram y / vscale=count;
        freq obsfreq;
        output out=stats mean=ybar n=n var=var;
run;
* Print output data file;
proc print data=stats;
run;
* Calculate expected frequencies using ybar;
data poisfit;
        if _n_ = 1 then set stats;
        set poisson;
        poisprob = pdf('poisson',y,ybar);
        expfreq = n*poisprob;
run;
* Print observed and expected frequencies;
proc print data=poisfit;
run;
* Plot observed and expected frequencies;
proc gplot data=poisfit;
        plot expfreq*yexp=1 obsfreq*yobs=2 / overlay legend=legend1 vref=0 wvref=3
        vaxis=axis1 haxis=axis1;
        symbol1 i=needle v=circle c=red width=3 height=2;
        symbol2 i=needle v=square c=blue width=3 height=2;
        axis1 label=(height=2) value=(height=2) width=3 major=(width=2) minor=none;
        legend1 label=(height=2) value=(height=2);
run;
quit;
```

\_ SAS Output \_\_\_

Fitting the Poisson to frequency data 1 16:30 Monday, March 29, 2010 Obs y obsfreq yexp yobs

105	У	oppited	yerb	yous
1	0	24	-0.1	0.1
2	1	16	0.9	1.1
3	2	16	1.9	2.1
4	3	18	2.9	3.1
5	4	15	3.9	4.1
6	5	9	4.9	5.1
7	6	6	5.9	6.1
8	7	5	6.9	7.1

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9	8	3	7.9	8.1
10	9	4	8.9	9.1
11	10	3	9.9	10.1
12	11	0	10.9	11.1
13	12	1	11.9	12.1

Fitting the Poisson to frequency data 2 16:30 Monday, March 29, 2010

#### The UNIVARIATE Procedure Variable: y

#### Freq: obsfreq

#### Moments

N	120	Sum Weights	120
Mean	3.16666667	Sum Observations	380
Std Deviation	2.78752724	Variance	7.77030812
Skewness	0.91183392	Kurtosis	0.32893349
Uncorrected SS	2128	Corrected SS	924.666667
Coeff Variation	88.0271761	Std Error Mean	0.25446526

#### Basic Statistical Measures

Location

Variability

Mean	3.166667	Std Deviation	2.78753
Median	3.000000	Variance	7.77031
Mode	0.000000	Range	12.00000
		Interquartile Range	4.00000

#### Tests for Location: Mu0=0

Test	-St	atistic-	p Value	e
Student's t	t	12.4444	Pr >  t	<.0001
Sign	М	48	Pr >=  M	<.0001
Signed Rank	S	2328	Pr >=  S	<.0001

Quantiles (Definition 5)

Quantile	Estimate
100% Max	12
99%	10
95%	9
90%	7
75% Q3	5
50% Median	3
25% Q1	1
10%	0
5%	0
1%	0
0% Min	0

Fitting the Poisson to frequency data 3 16:30 Monday, March 29, 2010

> The UNIVARIATE Procedure Variable: y

#### Freq: obsfreq

#### Extreme Observations

	Lowest			Highest	
Value	Freq	Obs	Value	Freq	Obs
0	24	1	7	5	8
1	16	2	8	3	9
2	16	3	9	4	10
3	18	4	10	3	11
4	15	5	12	1	13

Fitting the Poisson to frequency data 4 16:30 Monday, March 29, 2010

Obs	n	ybar	var
1	120	3.16667	7.77031

								, , , , , , , , , , , , , , , , , , ,	
Obs	n	ybar	var	У	obsfreq	yexp	yobs	poisprob	expfreq
1	120	3.16667	7.77031	0	24	-0.1	0.1	0.04214	5.0573
2	120	3.16667	7.77031	1	16	0.9	1.1	0.13346	16.0147
3	120	3.16667	7.77031	2	16	1.9	2.1	0.21130	25.3565
4	120	3.16667	7.77031	3	18	2.9	3.1	0.22304	26.7652
5	120	3.16667	7.77031	4	15	3.9	4.1	0.17658	21.1892
6	120	3.16667	7.77031	5	9	4.9	5.1	0.11183	13.4198
7	120	3.16667	7.77031	6	6	5.9	6.1	0.05902	7.0827
8	120	3.16667	7.77031	7	5	6.9	7.1	0.02670	3.2041
9	120	3.16667	7.77031	8	3	7.9	8.1	0.01057	1.2683
10	120	3.16667	7.77031	9	4	8.9	9.1	0.00372	0.4462
11	120	3.16667	7.77031	10	3	9.9	10.1	0.00118	0.1413
12	120	3.16667	7.77031	11	0	10.9	11.1	0.00034	0.0407
13	120	3.16667	7.77031	12	1	11.9	12.1	0.00009	0.0107

Fitting the Poisson to frequency data 5 16:30 Monday, March 29, 2010

Figure 5.9: Observed and expected frequencies - Poisson distribution Fitting the Poisson to frequency data



As an alternative to the Poisson, we can try fitting the negative binomial distribution using a similar SAS program. This distribution has two parameters, m and k, that must also be estimated before we can fit the distribution. The parameter m can be estimated using  $\bar{Y}$  as with the Poisson, but k is best estimated using a technique called maximum likelihood (see Chapter 8). We will use a SAS procedure that can model count data using the negative binomial distribution, proc genmod, in order to estimate k (SAS Institute Inc. 2014). The output of proc genmod is manipulated in several data steps to combine these estimates with the observed frequency data, and then the negative binomial probabilities and expected frequencies calculated and plotted. See SAS program and output below.

We see that the expected frequencies for the negative binomial distribution provide a better match to the observed ones for this data set (Fig. 5.10). We also note that the variance predicted for the negative binomial distribution is close to the observed variance. From the negative binomial fit, we have m = 3.167 and k = 1.760, and so the estimated variance is  $m + m^2/k = 3.167 + 3.167^2/1.760 = 7.459$ , while the observed variance is  $s^2 = 7.770$ . This further implies the negative binomial provides a better fit to these data than the Poisson distribution.

```
_____ SAS Program _____
```

```
* negbin_fit2.sas;
options pageno=1 linesize=80;
goptions reset=all;
title 'Fitting the negative binomial to frequency data';
data negbin;
        input y obsfreq;
        * Generate offset y values for plot;
        yexp = y - 0.1; yobs = y + 0.1;
        datalines;
0 24
1 16
2 16
3 18
4 15
5
   9
6
   6
7
   5
8
   3
9
   4
10 3
11 0
12 1
;
run;
* Print data set;
proc print data=negbin;
run;
* Descriptive statistics, save ybar, n, and var to data file;
proc univariate data=negbin;
        var y;
        histogram y / vscale=count;
        freq obsfreq;
        output out=stats mean=ybar n=n var=var;
run;
* Print output data file;
proc print data=stats;
run;
* Estimate m and k for the negative binomial distribution;
proc genmod data=negbin;
       model y = / dist=negbin;
        freq obsfreq;
        ods output ParameterEstimates=params;
run;
```

```
* Pick out value of m from genmod output;
data m;
        set params;
        if _n_ = 1;
        m = exp(Estimate);
        keep m;
run;
* Pick out value of k from genmod output;
data k;
        set params;
        if _n_ = 2;
        k = 1/Estimate;
       keep k;
run;
* Put m and k in one data file;
data params;
        merge m k;
run;
* Calculate expected frequencies using m and k;
data nbfit;
        if _n_ = 1 then set stats;
        if _n_ = 1 then set params;
        set negbin;
        nbprob = (gamma(k+y)/(gamma(y+1)*gamma(k)))*((m/(k+m))**y/(1+m/k)**k);
        expfreq = n*nbprob;
run;
* Print observed and expected frequencies;
proc print data=nbfit;
run;
* Plot observed and expected frequencies;
proc gplot data=nbfit;
        plot expfreq*yexp=1 obsfreq*yobs=2 / overlay legend=legend1 vref=0 wvref=3
        vaxis=axis1 haxis=axis1;
        symbol1 i=needle v=circle c=red width=3 height=2;
        symbol2 i=needle v=square c=blue width=3 height=2;
        axis1 label=(height=2) value=(height=2) width=3 major=(width=2) minor=none;
        legend1 label=(height=2) value=(height=2);
run;
quit;
```

```
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```

		SAS Output				
Fitting the	nega	ative binomial	to fr	equency data		1
				16:30 Monday, N	March 29,	2010
Obs	У	obsfreq	yexp	yobs		
1	0	24	-0.1	0.1		
2	1	16	0.9	1.1		
3	2	16	1.9	2.1		
4	3	18	2.9	3.1		
5	4	15	3.9	4.1		
6	5	9	4.9	5.1		
7	6	6	5.9	6.1		
8	7	5	6.9	7.1		
9	8	3	7.9	8.1		
10	9	4	8.9	9.1		
11	10	3	9.9	10.1		
12	11	0	10.9	11.1		
13	12	1	11.9	12.1		

Fitting the negative binomial to frequency data 2 16:30 Monday, March 29, 2010

#### The UNIVARIATE Procedure Variable: y

### Freq: obsfreq

#### Moments

N	120	Sum Weights	120
Mean	3.16666667	Sum Observations	380
Std Deviation	2.78752724	Variance	7.77030812
Skewness	0.91183392	Kurtosis	0.32893349
Uncorrected SS	2128	Corrected SS	924.666667
Coeff Variation	88.0271761	Std Error Mean	0.25446526

#### Basic Statistical Measures

Location		Variability			
Mean	3.166667	Std Deviation	2.78753		

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Median	3.000000	Variance	7.77031
Mode	0.000000	Range	12.00000
		Interquartile Range	4.00000

#### Tests for Location: MuO=0

Test	-Statistic-		p Value		
Student's t	t	12.4444	Pr >  t	<.0001	
Sign	М	48	Pr >=  M	<.0001	
Signed Rank	S	2328	Pr >=  S	<.0001	

#### Quantiles (Definition 5)

Quantile	Estimate
100% Max 99%	12 10
95%	9
90%	7
75% Q3	5
50% Median	3
25% WI	1
10%	0
5% 1%	0
1/0 OV Min	0
0% 1111	0

Fitting the negative binomial to frequency data 3 16:30 Monday, March 29, 2010

The UNIVARIATE Procedure Variable: y

#### Freq: obsfreq

#### Extreme Observations

-----Highest------Value Freq Obs Value Freq Obs

### 5.5. DISCRETE RANDOM VARIABLES AND SAMPLES

0	24	1	7	5	8
1	16	2	8	3	9
2	16	3	9	4	10
3	18	4	10	3	11
4	15	5	12	1	13

Fitting the negative binomial to frequency data 4 16:30 Monday, March 29, 2010

Obs	n	ybar	var
1	120	3.16667	7.77031

Fitting the negative binomial to frequency data 5 16:30 Monday, March 29, 2010

The GENMOD Procedure

#### Model Information

Data Set	WORK.NEGBIN
Distribution	Negative Binomial
Link Function	Log
Dependent Variable	У
Frequency Weight Variable	obsfreq

Number of Observations Rea	ad 13
Number of Observations Use	ed 12
Sum of Frequencies Read	120
Sum of Frequencies Used	120
Missing Values	1

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	119	140.6185	1.1817
Scaled Deviance	119	140.6185	1.1817
Pearson Chi-Square	119	104.3325	0.8767
Scaled Pearson X2	119	104.3325	0.8767
Log Likelihood		92.3889	

Full Log Likelihood	-272.1343
AIC (smaller is better)	548.2685
AICC (smaller is better)	548.3711
BIC (smaller is better)	553.8435

Algorithm converged.

#### Analysis Of Maximum Likelihood Parameter Estimates

			Standard	Wald	95%	Wald	
Parameter	DF	Estimate	Error	Confidenc	ce Limits	Chi-Square	Pr > ChiSq
Intercept	1	1.1527	0.0858	0.9845	1.3209	180.40	<.0001
Dispersion	1	0.5680	0.1298	0.3136	0.8225		

NOTE: The negative binomial dispersion parameter was estimated by maximum likelihood.

Fitting the negative binomial to frequency data 6 16:30 Monday, March 29, 2010

Obs	n	ybar	var	m	k	у	obsfreq	yexp	yobs	nbprob	expfreq
1	120	3.16667	7.77031	3.16667	1.76049	0	24	-0.1	0.1	0.16335	19.6023
2	120	3.16667	7.77031	3.16667	1.76049	1	16	0.9	1.1	0.18483	22.1793
3	120	3.16667	7.77031	3.16667	1.76049	2	16	1.9	2.1	0.16396	19.6748
4	120	3.16667	7.77031	3.16667	1.76049	3	18	2.9	3.1	0.13209	15.8503
5	120	3.16667	7.77031	3.16667	1.76049	4	15	3.9	4.1	0.10103	12.1237
6	120	3.16667	7.77031	3.16667	1.76049	5	9	4.9	5.1	0.07481	8.9770
7	120	3.16667	7.77031	3.16667	1.76049	6	6	5.9	6.1	0.05417	6.5008
8	120	3.16667	7.77031	3.16667	1.76049	7	5	6.9	7.1	0.03860	4.6319
9	120	3.16667	7.77031	3.16667	1.76049	8	3	7.9	8.1	0.02717	3.2599
10	120	3.16667	7.77031	3.16667	1.76049	9	4	8.9	9.1	0.01893	2.2722
11	120	3.16667	7.77031	3.16667	1.76049	10	3	9.9	10.1	0.01309	1.5714
12	120	3.16667	7.77031	3.16667	1.76049	11	0	10.9	11.1	0.00900	1.0797
13	120	3.16667	7.77031	3.16667	1.76049	12	1	11.9	12.1	0.00615	0.7379

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# 5.6 Classifying spatial or temporal patterns

The spatial distribution of organisms, or the temporal occurrence of events like cases of disease, is often compared with the Poisson distribution. This distribution essentially assumes a random, independent distribution of organisms or events, and if the observed distribution differs from the Poisson then this could indicate some interesting biology. For example, if the observed frequencies have a distribution with more extreme values (low or high) than the Poisson, with  $s^2 > \overline{Y}$ , this implies organisms are unevenly distributed in space, or events in time. A pattern like this is often called an overdis**persed** distribution, or alternatively a clumped, aggregated, or contagious distribution (Pielou 1977, Begon et al. 2006). One method of quantifying the level of overdispersion is to fit the negative binomial distribution to the data and use the value of k as an index. Small values of k (say k < 5) imply an overdispersed distribution, while larger ones indicate a distribution close to Poisson. More rarely, an observed distribution may have fewer extreme values than the Poisson, with  $s^2 < \overline{Y}$ , implying the organisms are evenly distributed in space (or events in time). This is called an **underdispersed** distribution, also known as a regular, even, or repulsed distribution.

The figures below provide examples of spatial distributions that are overdispersed, Poisson, or underdispersed. Note the obvious clusters of organisms in the overdispersed example (Fig. 5.11). This might occur because the clusters are offspring from a single parent, the organisms are responding to resources that are clumped in space, or because the organisms are attracted to one another. The Poisson data also show a few clusters (Fig. 5.12), but these are chance occurrences. If we were to divide this graph into quadrats and count the number of organisms per quadrat, we would find the frequency distribution is close to Poisson. In contrast to the other examples, the organisms are spaced apart to some extent in the underdispersed example (Fig. 5.13). This could occur because they are territorial, compete for resources, or otherwise regulate their numbers in some fashion (Ridout & Besbeas 2004).



Figure 5.11: Overdispersed distribution of organisms in space  $${\rm Overdispersed\ distribution}$$ 

Figure 5.12: Poisson distribution of organisms in space Poisson distribution of organisms



 $\begin{array}{c} Figure \ 5.13: \ Underdispersed \ distribution \ of \ organisms \ in \ space \\ \textbf{Underdispersed \ distribution} \end{array}$ 



# 5.7 References

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# 5.8 Problems

- 1. Consider the dice cube example from Chapter 4, and define a random variable Y that is the number of spots showing on the dice cube. Find E[Y] and Var[Y] for this random variable. Show your work.
- 2. Suppose that a random variable Y has a discrete distribution with the following probabilities:
  - $\begin{array}{ll} y & P[Y=y] \\ 0 & 0.5000 \\ 1 & 0.2500 \\ 2 & 0.1250 \\ 3 & 0.0625 \\ 4 & 0.0625 \end{array}$
  - (a) What is the expected value of Y, or E[Y]?
  - (b) What is the variance of Y, or Var[Y]?
- 3. An entomologist studies the spatial distribution of aphids in a field. They randomly select 100 locations within the field and count the number of aphids on the plants at each location. The following observed frequency distribution was obtained:

Aphids $(y)$	Frequency
0	19
1	22
2	16
3	10
4	11
5	11
6	6
7	2
8	1
9	1
10	1
11	0

- (a) Use the SAS program Poisson\_fit.sas to calculate  $\bar{Y}$  and  $s^2$ , and generate a plot of the observed frequencies vs. those expected for the Poisson distribution. Attach your SAS program and output.
- (b) Based on the above results, do the data have a Poisson distribution? Explain your answer using the pattern of observed and expected frequencies, and the values of  $\bar{Y}$  and  $s^2$ . Is the pattern random (Poisson), overdispersed, or underdispersed?
- (c) What are some possible biological explanations for this pattern?
- 4. A field is surveyed for golden mice (*Ochrotomys nuttalli*) using a grid of baited traps. A total of 100 traps were deployed and the number of mice counted in each trap. The following frequency distribution was obtained:

Mice $(y)$	Frequency
0	55
1	21
2	10
3	7
4	4
5	2
6	1
7	0
8	0

- (a) Use the program Poisson\_fit.sas to calculate to calculate  $\bar{Y}$  and  $s^2$ , and generate a plot of the observed frequencies vs. those expected for the Poisson distribution. Attach your program and output.
- (b) Based on the above results, do the data have a Poisson distribution? Explain your answer using the pattern of observed and expected frequencies, and the values of  $\bar{Y}$  and  $s^2$ . Is the pattern random (Poisson), overdispersed, or underdispersed?

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