Assignment #5 — Due Friday, October 14 by 4:00 P.M.

Turn in homework to your TA's mailbox using this sheet as the cover page.

Fill in your name and also circle the *lecture section in which you are registered* and circle the *discussion section* you expect to attend to pick up this assignment.

Name:

 Lecture 1 (Hanlon).
 311: Tu 1:00 - 2:15pm
 312: Th 8:00 - 9:15am
 313: We 1:00 - 2:15pm

 Lecture 2 (Larget).
 321: Tu 1:00 - 2:15pm
 322: We 2:30 - 3:45pm
 323: We 1:00 - 2:15pm

The following information will help you with problems in this assignment.

• p-values for the χ^2 or G-test. To compute p-values based on a χ^2 distribution, use the R function pchisq(), which computes a the probability that a χ^2 random variable is less than a given value. As p-values here are the probability that a χ^2 random variable is more than the test statistic, you would use a command as in the following example. The p-value for the G-Test with observed test statistics G=8 and df=2, is given by

```
> 1-pchisq(8,df=2)
```

• Graphing contigency table data. Download the file mosaic.R from the course page. This file contains a function mosaic() that modifies the built-in barchart() from the lattice package in R to create a mosaic plot as described in the text. Load this function into R. (There are many ways to do this, just as you did for gbinom.R last assignment: use source("mosaic.R"), or source(file.choose()), or Run script... or Source File... from the File menu.)

To enter the matrix of counts, type something similar to this example which uses the data on page 214. The function matrix() creates a matrix from the data in the first argument using the second and third arguments to specify the number of rows and columns. The functions rownames() and columns() add names to the rows and columns of the matrix which makes the plot more informative. Note that the data is specified column by column, not row by row, in one long array collected with c(). You can verify that you set the matrix up correctly by typing its name and seeing it.

```
> fish = matrix(c(1,49,10,35,37,9),nrow=2,ncol=3)
> rownames(fish) = c("Eaten","Not eaten")
> colnames(fish) = c("Uninfected","Lightly infected","Highly infected")
> fish
> mosaic(fish)
```

In the example, the plot shows the relative frequencies within each column. If you want to do the same for rows, use t() to transform the matrix.

```
> mosaic(t(fish))
```

Please answer the following questions.

- 1. The following questions ask you to compute a point estimate and confidence interval for a difference in proportions, $p_1 p_2$. We let n_1 denote the number of trials and x_1 the number of observed successes in the first sample; and n_2 denote the number of trials and x_2 the number of observed successes in the second sample.
 - (a) $n_1 = 10$, $x_1 = 7$, $n_2 = 50$, $x_2 = 5$. Compute a point estimate and 95% confidence interval for $p_1 p_2$.
 - (b) $n_1 = 50$, $x_1 = 24$, $n_2 = 55$, $x_2 = 27$. Compute a point estimate and 95% confidence interval for $p_1 p_2$.
 - (c) $n_1 = 1000$, $x_1 = 635$, $n_2 = 556$, $x_2 = 435$. Compute a point estimate and 95% confidence interval for $p_1 p_2$.
- 2. The following questions ask you to compute a point estimate and confidence interval for the odds ratio. We let n_1 denote the number of trials and x_1 the number of observed successes in the first sample; and n_2 denote the number of trials and x_2 the number of observed successes in the second sample.
 - (a) $n_1 = 10$, $x_1 = 7$, $n_2 = 50$, $x_2 = 5$. Compute a point estimate and 95% confidence interval for the odds ratio.
 - (b) $n_1 = 50$, $x_1 = 24$, $n_2 = 55$, $x_2 = 27$. Compute a point estimate and 95% confidence interval for the odds ratio.
 - (c) $n_1 = 1000$, $x_1 = 635$, $n_2 = 556$, $x_2 = 435$. Compute a point estimate and 95% confidence interval for the odds ratio.
- 3. Consider the problem from the notes regarding infection status and the probability of being eaten (also described in the textbook p.213-214). We conduct a second experiment and collect the following data.

	Uninfected	Lightly Infected	Highly Infected	Total
Eaten	2	8	50	60
Not eaten	45	30	5	80
Total	47	38	55	140

For this problem, the null hypothesis and alternative hypotheses are

 H_0 : Parasite infection and being eaten are independent.

 H_A : Parasite infection and being eaten are not independent.

Complete this problem without the use of software, that is, do the calculations by hand.

- (a) Under the null hypothesis, compute the table of expected counts.
- (b) Compute the test statistic X^2 for the χ^2 test of independence.
- (c) Compute the test statistic G for the G-test.
- 4. Use the data set from p.227, Problem 11. But answer these questions.
 - (a) Examine a mosaic plot that compares the estimated conditional probabilities of success of wart removal given the treatment. Include this plot with your solution.
 - (b) Find a point estimate and a 95% confidence interval for the difference in the success probabilities of the two therapies. Interpret the result.
 - (c) Find a point estimate and a 95% confidence interval for the odds ratio of the two therapies. Interpret the result.

- 5. Use the data set from p.229-230, Problem 20. But answer these questions. When conducting a hypothesis test, make sure to include the following: statements of the null and alternative hypotheses, the observed value of the test statistic, the p-value, and an interpretation of the result.
 - (a) Display the data with a mosaic plot to highlight proportions of people in each smoking class within each diet class.
 - (b) Perform the χ^2 test of independence on the data set.
 - (c) Perform the G-test on the data set.
 - (d) Relate the results of these hypothesis tests to what you saw in the mosaic plot. What features apparent in the plot do you think had the largest influence on the results of the tests?
 - (e) The test statistics for the two tests involve summing over all cells in the table. Which cells contributed the most to these sums? How does this relate to your answer of the previous part of the question?
- 6. Extracorporeal membrane oxygenation (ECMO) is a potentially life-saving procedure that is used to treat newborn babies who suffer from respiratory failure. An experiment was conducted to test whether ECMO leads to improved survival relative to conventional medical therapy (CMT). In the experiment 29 babies were treated with ECMO and 10 babies were treated with CMT. The data are displayed below. Perform Fisher's exact test on the observed data.

	CMT	ECMO	Total
Die	4	1	5
Live	6	28	34
Total	10	29	39

When conducting a hypothesis test, make sure to include the following: statements of the null and alternative hypotheses, the observed value of the test statistic, the p-value, and an interpretation of the result.

- 7. Textbook p.125, Problem 22
- 8. Textbook p.125, Problem 23