#### Chapter 9 Notes, Part 1 - Inference for Proportion and Count Data

We want to estimate the proportion p of a population that have a specific attribute, like "what percent of houses in Cambridge have a mouse in the house?"

We are given  $X_1, \ldots, X_p$  where  $X_i$ 's are Bernoulli, and  $P(X_i = 1) = p$ .

 $X_i$  is 1 if house *i* has a mouse.

Let  $Y = \sum_{i} X_i$  so  $Y \sim Bin(n, p)$ .

An estimator for p is:

$$\hat{p} = \frac{Y}{n} = \frac{1}{n} \sum_{i} X_i.$$

 $\hat{p}$  is a random variable. For large n (rule of thumb,  $n\hat{p} \ge 10, n(1-\hat{p}) \ge 10$ ) the CLT says that approximately:

$$\hat{p} \sim N\left(p, \frac{pq}{n}\right)$$
 where  $q = 1 - p$ .

Questions: What's up with that rule of thumb? Where did the pq/n come from?

### **Confidence Intervals**

The CI can be computed in 2 ways (here for 2-sided case):

• CI first try:

$$P\left(-z_{\alpha/2} \le \frac{\hat{p}-p}{\sqrt{pq/n}} \le z_{\alpha/2}\right) = 1 - \alpha.$$

We could solve it for p but the expression is quite large...

• CI second try:

$$P\left(-z_{\alpha/2} \le \frac{\hat{p} - p}{\sqrt{\hat{p}\hat{q}/n}} \le z_{\alpha/2}\right) \approx 1 - \alpha$$

which yields

$$\hat{p} - z_{\alpha/2} \sqrt{\frac{\hat{p}\hat{q}}{n}} \le p \le \hat{p} + z_{\alpha/2} \sqrt{\frac{\hat{p}\hat{q}}{n}}.$$

So that's the approximate CI for p.

Sample size calculation for CI

Want a CI of width 2E:

$$\hat{p} - E \le p \le \hat{p} + E$$

 $\mathbf{SO}$ 

$$E = z_{\alpha/2} \sqrt{\frac{\hat{p}\hat{q}}{n}}$$
 which means  $n = \left(\frac{z_{\alpha/2}}{E}\right)^2 \hat{p}\hat{q}$ .

Question: We'll take  $\hat{p}\hat{q}$  to be its largest possible value,  $(1/2) \times (1/2)$ . Why do we do this? Why don't we just use the  $\hat{p}$  and  $\hat{q}$  that we measure from the data?

So, we need:

$$n = \left(\frac{z_{\alpha/2}}{E}\right)^2 \frac{1}{4}$$
 observations.

Hypothesis testing on proportion for large n

$$\begin{array}{rcl} H_0 & : & p = p_0 \\ H_1 & : & p \neq p_0. \end{array}$$

Large *n* and  $H_0$  imply  $\hat{p} \approx N\left(p_0, \frac{p_0 q_0}{n}\right)$  (where  $q_0 = 1 - p_0$ ) so we use z-test with test statistic:

$$z = \frac{\hat{p} - p_0}{\sqrt{p_0 q_0/n}}$$

Example

For small n one can use the binomial distribution to compute probabilities directly (rather than approximating by normal). (Not covered here.)

## Chapter 9.2 Comparing 2 proportions

Example: The Salk polio vaccine trial: compare rate of polio in control and treatment (vaccinated) group. Is this independent samples design or matched pairs?

Sample 1: number of successes  $X \sim Bin(n_1, p_1)$ , observe X = x. Sample 2: number of successes  $Y \sim Bin(n_2, p_2)$ , observe Y = y.

We could compare rates in several ways:

$$p_1 - p_2 \longrightarrow \text{ we'll use this one}$$

$$p_1/p_2 \qquad \text{``relative risk''}$$

$$\left(\frac{p_1}{1-p_1}\right) / \left(\frac{p_2}{1-p_2}\right) \qquad \text{``odds ratio''}$$

For large samples, we'll use the CLT:

$$Z = \frac{\hat{p}_1 - \hat{p}_2 - (p_1 - p_2)}{\sqrt{\frac{\hat{p}_1\hat{q}_1}{n_1} + \frac{\hat{p}_2\hat{q}_2}{n_2}}} \approx N(0, 1)$$

where  $\hat{p_1} = X/n_1$  and  $\hat{p}_2 = Y/n_2$ .

To test

$$H_0: p_1 - p_2 = \delta_0, H_1: p_1 - p_2 = \delta_0,$$

we can just compute zscores, pvalues, and CI. The test statistic is:

$$z = \frac{\hat{p}_1 - \hat{p}_2 - \delta_0}{\sqrt{\frac{\hat{p}_1\hat{q}_1}{n_1} + \frac{\hat{p}_2\hat{q}_2}{n_2}}}.$$

This is a little weird because it really should have terms like " $p_{1,0}q_{1,0}/n_1$ " in the denominator, but we don't have those values under the null hypothesis. So we get an approximation by using  $\hat{p}_1$  and  $\hat{q}_1$  in the denominator.

#### Example

For an independent samples design with small samples, use *Fisher's Exact Test* which uses the Hypergeometric distribution. For a matched pairs design, use *McNemar's Test* which uses the binomial distribution (Both beyond the scope.)

Two Challenges

# 15.075J / ESD.07J Statistical Thinking and Data Analysis Fall 2011

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