# Test for the probability of a binomial distribution

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For an i.i.d sample from a bernoulli distribution

$$x_1, \cdots, x_n \overset{\text{i.i.d.}}{\sim} Bernoulli(p),$$

The likelihood of the data is

$$f(x_1, \dots, x_n) = \prod_{i=1}^n p^{x_i} (1-p)^{1-x_i} = p^{\sum x_i} (1-p)^{n-\sum x_i}.$$

MLE for p is  $\bar{x} = \frac{1}{n} \sum x_i$  and

$$\sum_{i=1}^{n} x_i \sim Binom(n, p).$$

So here are mainly two situations: One is to test the probability p against some given value  $p_0$ . The other is to compare the probability between two independent random samples  $x_1, \dots, x_n$  and  $y_1, \dots, y_m$ .

Case1: One sample  $x_1, \dots, x_n$  from Bernoulli(p), and test p against a given  $p_0$ .

Case 2: Two samples:  $x_1, \dots, x_2$  from  $Bernoulli(p_1)$  and  $y_1, \dots, y_m$  from  $Bernoulli(p_2)$ . And test whether  $p_1 = p_2$ .

## 1 Normal approximation

### 1.1 Case 1

Note that

$$EX = p$$
,  $VarX = p(1-p)$ .

Then by CLT we have

$$\bar{x} \stackrel{\text{asymp}}{\sim} N\left(p, \frac{p\left(1-p\right)}{n}\right).$$

For  $H_0$ :  $p = p_0$ , we propose a test statistic

$$Z = \frac{\bar{x} - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}.$$

Then Z is asymptotically standard normal under  $H_0$ .

Also we know that under  $H_1$ :

$$Z = \frac{\bar{x} - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}$$

$$= \frac{\bar{x} - p}{\sqrt{\frac{p(1 - p)}{n}}} \cdot \sqrt{\frac{p(1 - p)}{p_0(1 - p_0)}} + \frac{p - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}$$

$$\sim N \left(\frac{p - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}, \quad \frac{p(1 - p)}{p_0(1 - p_0)}\right).$$

So the power of the test can be easily computed.

#### 1.2 Case 2

So we have

$$\bar{x} \stackrel{\text{asymp}}{\sim} N\left(p_1, \frac{p_1(1-p_1)}{n}\right), \text{ and } \bar{y} \stackrel{\text{asymp}}{\sim} N\left(p_2, \frac{p_2(1-p_2)}{m}\right).$$

A test statistic can be

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\hat{p}(1-\hat{p})\left(\frac{1}{n} + \frac{1}{m}\right)}},$$

where  $\hat{p} = \frac{n\bar{x} + m\bar{y}}{n+m}$ . This test statistic can be found at

https://stats.stackexchange.com/questions/361015/

proof-of-the-standard-error-of-the-distribution-between-two-normal-distributions/ 361048#361048

https://stats.stackexchange.com/questions/113602/

test-if-two-binomial-distributions-are-statistically-different-from-each-other

Here this  $\hat{p}(1-\hat{p})$  can be seen as an estimate for the variance p(1-p) when  $H_0$  is true by directly plugging in  $\hat{p}$ . This is **NOT** a pooled variance for these two samples, which should always be no greater than  $\hat{p}(1-\hat{p})$ .

The power of this test statistic is hard to compute under  $H_1$ .

**Note:** One can also use the same idea in the "t-test.pdf" notes and propose the test statistic

$$T = \frac{\bar{x} - p_0}{\sqrt{S_x/n}},$$

where  $S_x = \frac{1}{n-1} \sum_{i=1}^{n} (x_i - \bar{x})$  for Case 1.

And for Case 2

$$T = \frac{\bar{x} - \bar{y} - \Delta}{\sqrt{\left(\frac{1}{n} + \frac{1}{m}\right)S_p}},$$

where  $S_p^2 = \frac{(n-1)S_x^2 + (m-1)S_y^2}{n+m-2}$  and  $\Delta = p_1 - p_2$ . But again it is hard to evaluate the testing power of these statistics.

# 2 Chi-square approximation

See the notes of "chisq\_test.pdf" for details.

### 3 Exact test

### 3.1 Case 1: Clopper-Pearson test

The Clopper-Pearson method is an early method. It's called exact method because it's directly based on p.m.f of binomial distribution. Let  $X = \sum_{i=1}^{n} x_i$ . Then  $X \sim Binom(n, p)$  and the p.m.f is

$$f(x;p) = P(X = x|p) = C_n^x p^x (1-p)^{n-x},$$
(1)

for  $x = 0, 1, \dots, n$ . One thing to point out is that though the |p| notation, (1) is frequenist's point of view, not bayesian's. Now let's recall that p-value is the probability under  $H_0$  that something **as or more extreme than** what we have observed happens. Then after observing  $X = x_0$ , for one-sided test:

•  $H_0: p \leq p_0$  against  $H_1: p > p_0$  for some given  $p_0$ . The p-value at this observed  $x_0$  is

$$p_{val}(x_0) = \sum_{x=x_0}^{n} f(x; p_0).$$
 (2)

•  $H_0: p \ge p_0$  against  $H_1: p < p_0$  for some given  $p_0$ . The p-value at this observed  $x_0$  is

$$p_{val}(x_0) = \sum_{x=0}^{x_0} f(x; p_0).$$
 (3)

For the two-sided test. This is a little complicated. Let index set

$$\mathcal{I} = \{x | P(X = x | p_0) \le P(X = x_0 | p_0), \quad 0 \le x \le n\}.$$

Then  $\mathcal{I}$  contains all possible realizations of X with its probability no greater than the probability of our observation. Then the p-value of  $H_0: p = p_0$  at this observed  $x_0$  is given by

$$p_{val}(x_0) = \sum_{x \in \mathcal{I}} f(x; p_0).$$
(4)

#### 3.1.1 Power analysis

The probability to reject  $H_0$  of Clopper-Pearson test at given underlying p can be computed by

$$P\left(\text{Reject } H_0|p\right) = \sum_{x=0}^{n} P\left(X = x|p\right) \cdot I_{\{p_{val}(x) \le \alpha\}} = \sum_{x=0}^{n} f\left(x;p\right) \cdot I_{\{p_{val}(x) \le \alpha\}}, \tag{5}$$

where  $\alpha$  is the significant level of the test and  $p_{val}(x)$  is computed for different types of  $H_0$  based on (2), (3) and (4).

#### 3.1.2 Confidence interval

First for the one-sided intervals:

•  $(P_L, 1]$ : From (2),  $H_0: p \leq p_0$  is rejected when probability of observing  $x_0$  or more number of success at  $p_0$  is small enough. Therefore the reject area

Rejet Area: 
$$\left\{x_0: \sum_{x=x_0}^n f(x; p_0) \le \alpha\right\}$$
.

Hence the accept area

Accept Area: 
$$\left\{ x_0 : \sum_{x=x_0}^n f(x; p_0) > \alpha \right\}$$

Then we can construct the one-sided CI by increasing  $p_0$  from 0 such that the first  $p_0$  that satisfies this Accept area rule. Then that is the  $P_L$ . Therefore

$$\sum_{x=x_0}^n f(x; P_L) = \alpha. \tag{6}$$

•  $[0, P_U)$ : Similar idea, from (3) we can construct the accept area

Accept Area: 
$$\left\{x_0; \sum_{x=0}^{x_0} f(x; p_0) > \alpha\right\}$$
.

Therefore we decrease  $p_0$  from 1 to find the first  $P_U$  that satisfies this Accept area rule. Therefore

$$\sum_{x=0}^{x_0} f(x; P_U) = \alpha. \tag{7}$$

Now for the two-sided intervals  $(P_L, P_U)$ : we apply the **equal-tail rule** and find  $P_L$  and  $P_U$  such that

$$\sum_{x=x_0}^{n} f(x; P_L) = \alpha/2$$

$$\sum_{x=0}^{x_0} f(x; P_U) = \alpha/2.$$
(8)

This interval can also be expressed as

$$S < \cap S >$$
,

or equivalently

$$(\inf S_{\geq}, \sup S_{\leq}),$$

where

$$S_{\leq} \triangleq \left\{ \theta \middle| P\left(Binomial\left(n, \theta\right) \leq x\right) > \frac{\alpha}{2} \right\}$$
$$S_{\geq} \triangleq \left\{ \theta \middle| P\left(Binomial\left(n, \theta\right) \geq x\right) > \frac{\alpha}{2} \right\}.$$

One can utilize the relationship between the Binomial cumulative distribution function and **regularized incomplete beta function**, i.e. for  $k = 0, \dots, n$ 

$$\begin{split} &P\left(X \leq k\right) = \sum_{i=0}^{k} C_{n}^{i} p^{i} \left(1-p\right)^{n-i} \\ &= \frac{\Gamma\left(n+1\right)}{\Gamma\left(n-k\right) \Gamma\left(k+1\right)} \int_{0}^{1-p} t^{n-k-1} \left(1-t\right)^{k} \mathrm{d}t = pBeta \left(1-p; n-k, k+1\right) \\ &= \frac{\Gamma\left(n+1\right)}{\Gamma\left(n-k\right) \Gamma\left(k+1\right)} \int_{p}^{1} t^{k} \left(1-t\right)^{n-k-1} \mathrm{d}t = 1-pBeta \left(p; k+1, n-k\right), \end{split}$$

where  $pBeta(x; \alpha, \beta)$  represents the cumulative probability of  $Beta(\alpha, \beta)$  distribution, cumulated from 0 to x. And it satisfies

$$pBeta(x; \alpha, \beta) = 1 - pBeta(1 - x; \beta, \alpha), \quad \forall x \in [0, 1].$$

Similarly, for the quantile function  $qBeta(p; \alpha, \beta)$ , we can show that

$$1 - qBeta(p; \alpha, \beta) = qBeta(1 - p; \beta, \alpha), \quad \forall p \in [0, 1].$$

So we can see that  $P_L$  and  $P_U$  are actually satisfing

$$1 - \alpha/2 = (\ge)P(X \le x_0 - 1|P_L) = pBeta(1 - P_L; n - x_0 + 1, x_0)$$

$$\implies 1 - P_L = (\le)qBeta(1 - \alpha/2; n - x_0 + 1, x_0)$$

$$\implies P_L = (\ge)qBeta(\alpha/2; x_0, n - x_0 + 1)$$

For  $P_L$ , it's taking inf, therefore

$$P_L = qBeta(\alpha/2; x_0, n - x_0 + 1).$$
 (9)

Similarly, we have for  $P_U$ :

$$\alpha/2 = P(X \le x_0|P_U) = pBeta(1 - P_U; n - x_0, x_0 + 1)$$
  
 $\implies 1 - P_U = qBeta(\alpha/2; n - x_0, x_0 + 1)$   
 $\implies P_U = qBeta(1 - \alpha/2; x_0 + 1, n - x_0)$ 

Therefore

$$P_U = qBeta (1 - \alpha/2; x_0 + 1, n - x_0). \tag{10}$$

Also, note that this cumulative probability is also related to F-distribution via

$$P(X \le x_0) = F\left(x = \frac{1-p}{p} \frac{x_0+1}{n-x_0}; d_1 = 2(n-x_0), d_2 = 2(x_0+1)\right)$$

where  $F(x; d_1, d_2)$  is the cumulative probability function of a F-distribution with degree of freedom  $d_1$  and  $d_2$ , cumulated from 0 to x. The we have

$$P_{L} = \left(1 + \frac{n - x_{0} + 1}{x_{0} \times qF\left(\frac{\alpha}{2}; 2x_{0}, 2\left(n - x_{0} + 1\right)\right)}\right)^{-1}$$

$$P_{U} = \left(1 + \frac{n - x_{0}}{\left(x_{0} + 1\right) \times qF\left(1 - \frac{\alpha}{2}; 2\left(x_{0} + 1\right), 2\left(n - x_{0}\right)\right)}\right)^{-1}$$
(11)

where  $qF(\alpha; d_1, d_2)$  is the quantile function of F-distribution.

### 3.2 Case 2: Fisher's exact test

Fisher's exact test is a method for testing proportion difference. A toy example of a  $2 \times 2$  contingency table is shown in Table 1. When the margin of this table is fixed  $(n_x, n_y, n_1$  and  $n_2)$ , the probability for observing this table follows the hyper-genometric distribution

$$P(\#\{\text{Sample 1, Success}\} = a) \frac{C_{n_x}^a C_{n_y}^{n_1 - a}}{C_n^{n_1}}.$$

	Success	Failure	Total
Sample 1	a	b	$n_x = a + b$
Sample 2	С	d	$n_y = c + d$
Total	$n_1 = a + c$	$n_0 = b + d$	n = a + b + c + d

Table 1: Data sample

We can compute the p-value based on the same idea from Section 3.1. Note that here the p-value is a **conditional** one since it is conditional on the fixed marginal values. To simplify the notation, denote X the number in cell (Sample 1, Success) and

$$f(x; n_x, n_y, n_1) = P(X = x | n_x, n_y, n_1) = \frac{C_{n_x}^x C_{n_y}^{n_1 - x}}{C_{n_x + n_y}^{n_1}}$$

Then for a observation with  $X = x_0$  and fixed  $n_x, n_y, n_1$ :

•  $H_0: p_x \ge p_y$  against  $H_1: p_x < p_y$ . The p-value can be computed as

$$p_{val} = \sum_{i=0}^{x_0} f(i; n_x, n_y, n_1).$$
 (12)

•  $H_0: p_x \leq p_y$  against  $H_1: p_x > p_y$ . The p-value can be computed as

$$p_{val} = \sum_{i=x_0}^{n_1} f(i; n_x, n_y, n_1).$$
 (13)

•  $H_0: p_x = p_y$  against  $H_1: p_x \neq p_y$ . The p-value can be computed as

$$p_{val} = \sum_{i=a_I}^{a_U} f(i; n_x, n_y, n_1) I_{\{f(i; n_x, n_y, n_1) \le f(x_0; n_x, n_y, n_1) \delta\}}, \tag{14}$$

where the summation limits  $a_L = \max(0, n_1 - n_y)$  and  $a_U = \min(n_x, n_1)$ . Note that in an ideal world the red  $\delta$  is just 1 in (14). But in actuality, since the computation involves large factorials, especially when sample size is large, the numerical results might be inaccurate. To ensure a convervative test,  $\delta$  is set to 1.0000001 in R.