

Chapter 9. Testing hypotheses and assessing goodness of fit

1 Hypotheses testing

Find a rule based on data for choosing between two mutually exclusive hypotheses

null hypothesis H_0 : the effect of interest is zero,

alternative H_1 : the effect of interest is not zero.

H_0 represents an established theory that must be discredited in order to demonstrate some effect H_1 .

Two types of error

type I error = false positive: reject H_0 when it's true,

type II error = false negative: accept H_0 when it's false.

Test result	Negative: do not reject H_0	Positive: reject H_0
If H_0 is true	True negative. Specificity = $1 - \alpha$	False positive. Significance level α
If H_1 is true	False negative $\beta = P(\text{accept } H_0 H_1)$	True positive. Sensitivity = $1 - \beta$

Significance test

Test statistic = a function of the data with distinct typical values under H_0 and H_1 .

Rejection region (RR) of a test = a set of values for the test statistic when H_0 is rejected.

If test statistic and sample size are fixed, then either α or β gets larger when RR is changed.

Significance test approach to choose a rejection region:

fix an appropriate significance level α ,

find a RR from $\alpha = P(\text{test statistic} \in \text{RR}|H_0)$ using the null distribution of the test statistic.

Common significance levels: 5%, 1%, 0.1%

2 Large-sample test for the proportion

Data is modeled by a sample count $Y \sim \text{Bin}(n, p)$. An unbiased point estimate for the population proportion p is the sample proportion $\hat{p} = \frac{Y}{n}$.

For $H_0: p = p_0$ use the test statistic $Z = \frac{Y - np_0}{\sqrt{np_0q_0}} = \frac{\hat{p} - p_0}{\sqrt{p_0q_0/n}}$.

Approximate null distribution: $Z \stackrel{a}{\sim} N(0,1)$. Let $\Phi(z_\alpha) = 1 - \alpha$. Three different rejection regions for three composite alternative hypotheses

one-sided $H_1: p > p_0$, RR = $\{Z \geq z_\alpha\}$,

one-sided $H_1: p < p_0$, RR = $\{Z \leq -z_\alpha\}$,

two-sided $H_1: p \neq p_0$, RR = $\{Z \geq z_{\alpha/2} \text{ or } Z \leq -z_{\alpha/2}\}$.

Power function

The power of the test (sensitivity): $P_w = P(\text{reject } H_0 | H_1 \text{ is true})$.

Let $H_0: p = p_0$, $H_1: p = p_1$, and $p_1 > p_0$. The power function of the one-sided test

$$P_w(p_1) = P\left(\frac{Y - np_0}{\sqrt{np_0q_0}} \geq z_\alpha \mid p = p_1\right) \approx 1 - \Phi\left(\frac{z_\alpha \sqrt{p_0q_0} + \sqrt{n}(p_0 - p_1)}{\sqrt{p_1q_1}}\right), \quad p_1 > p_0.$$

Planning of sample size: given α and β , choose sample size n such that $\sqrt{n} = \frac{z_\alpha \sqrt{p_0q_0} + z_\beta \sqrt{p_1q_1}}{|p_1 - p_0|}$.

Example: extrasensory perception.

ESP test: guess the suits of $n = 100$ cards chosen at random with replacement from a deck of cards with four suits. Number of cards guessed correctly $Y \sim \text{Bin}(100, p)$

$H_0: p = 0.25$ (pure guessing), $H_1: p > 0.25$ (ESP ability).

Rejection region at 5% significance level = $\left\{\frac{\hat{p} - 0.25}{0.0433} \geq 1.645\right\} = \{\hat{p} \geq 0.32\} = \{Y \geq 32\}$.

With a simple alternative $H_1: p = 0.30$ the power of the test is $1 - \Phi\left(\frac{1.645 - 0.0433 - 0.5}{0.0458}\right) = 32\%$.

The sample size required for the 90% power is $n = \left(\frac{1.645 - 0.0433 + 1.28 \cdot 0.0458}{0.05}\right)^2 = 675$.

P-value of the test

P-value is the probability of obtaining a test statistic value as extreme or more extreme than the observed one, given that H_0 is true.

For the significance level α , reject H_0 , if $P \leq \alpha$, and do not reject H_0 , if $P > \alpha$.

Two-sided P-value = $2 \times$ one-sided P-value
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Example: extrasensory perception.

If the observed sample count is $Y_{\text{obs}} = 30$, then $Z_{\text{obs}} = \frac{0.3 - 0.25}{0.0433} = 1.15$ and a one-sided P-value is $P(Z \geq 1.15) = 12.5\%$. The result is not significant, do not reject H_0 .

3 Small-sample test for the proportion

With $H_0: p = p_0$ the test statistic $Y \sim \text{Bin}(n, p)$ for small n we have to rely on the exact null distribution $Y \sim \text{Bin}(n, p_0)$. Three rejection regions

one-sided $H_1: p > p_0$, $\text{RR} = \{Y \geq y_\alpha\}$,

one-sided $H_1: p < p_0$, $\text{RR} = \{Y \leq y'_\alpha\}$,

two-sided $H_1: p \neq p_0$, $\text{RR} = \{Y \geq y_{\alpha/2} \text{ or } Y \leq y'_{\alpha/2}\}$.

Example: extrasensory perception.

ESP test: guess the suits of $n = 20$ cards. Model: the number of cards guessed correctly is $Y \sim \text{Bin}(20, p)$. For $H_0: p = 0.25$ the null distribution is

Bin(20,0.25) table:	y	8	9	10	11
	$P(Y \geq y)$.101	.041	.014	0.004

One-sided alternative $H_1: p > 0.25$. Rejection region at 5% significance level = $\{Y \geq 9\}$. Notice that the exact significance level = 4.1%. Power function: $P_w(p_1) = P[Y \geq 9 | Y \sim \text{Bin}(20, p_1)]$

p_1	0.27	0.3	0.4	0.5	0.6	0.7
$P_w(p_1)$	0.064	0.113	0.404	0.748	0.934	0.995

Warning for "fishing expeditions": the number of false positives in k tests at level α is $\text{Pois}(k\alpha)$.

4 Tests for the mean

Test $H_0: \mu = \mu_0$ for continuous or discrete data. Large-sample test for mean is used when the population distribution is not necessarily normal but the sample size n is sufficiently large.

$$H_0: \mu = \mu_0, \text{ test statistic } T = \frac{\bar{X} - \mu_0}{s_{\bar{X}}} \text{ with an approximate null distribution } T \stackrel{a}{\sim} N(0,1).$$

The one-sample t-test is used for small n , assuming that the population distribution is normal.

$$H_0: \mu = \mu_0, \text{ test statistic: } T = \frac{\bar{X} - \mu_0}{s_{\bar{X}}} \text{ with an exact null distribution: } T \sim t_{n-1}.$$

CI method of hypotheses testing:

reject $H_0: \mu = \mu_0$ at 5% level if and only if a 95% confidence interval for the mean does not cover μ_0 .

5 Likelihood ratio test

A general method of finding asymptotically optimal tests (having the largest power for a given α).

Two simple hypotheses

For testing $H_0: \theta = \theta_0$ against $H_1: \theta = \theta_1$ use the likelihood ratio $\Lambda = \frac{L(\theta_0)}{L(\theta_1)}$ as a test statistic. Large values of Λ suggest that H_0 explains the data set better than H_1 , while small Λ indicate that H_1 explains the data set better.

$$\text{Likelihood ratio rejection rule: reject } H_0 \text{ for } \Lambda \leq \lambda_\alpha.$$

Neyman-Pearson lemma: the likelihood ratio test is optimal in the case of two simple hypothesis.

Nested hypotheses

With a pair of nested parameter sets $\Omega_0 \subset \Omega$ we get two composite alternatives, $H_0: \theta \in \Omega_0$ and $H_1: \theta \in \Omega \setminus \Omega_0$. Two nested hypotheses $H_0: \theta \in \Omega_0$, $H: \theta \in \Omega$, and two maximum likelihood estimates

$$\hat{\theta}_0 = \text{maximizes likelihood over } \theta \in \Omega_0,$$

$$\hat{\theta} = \text{maximizes likelihood over } \theta \in \Omega.$$

Generalized LRT: reject H_0 for small values of $\frac{L(\hat{\theta}_0)}{L(\hat{\theta})}$ or equivalently

$$\text{GLRT: reject } H_0 \text{ for large values of } \Delta = \log L(\hat{\theta}) - \log L(\hat{\theta}_0).$$

Approximate null distribution: $2\Delta \stackrel{a}{\sim} \chi_{\text{df}}^2$, where $\text{df} = \dim(\Omega) - \dim(\Omega_0)$.

6 Pearson's chi-square test

Data: each observation belongs to one of J classes. A null hypothesis proposing a model for the data

$$H_0: (p_1, \dots, p_J) = (p_1(\lambda), \dots, p_J(\lambda)) \text{ with unknown parameter } \lambda = (\lambda_1, \dots, \lambda_r), \dim(\Omega_0) = r.$$

Test how well a model fits the data using the MLE $\hat{\lambda}$ of λ describing H_0 . Data is summarized as the vector of observed counts (O_1, \dots, O_J) .

$$\text{Chi-square test statistic: } X^2 = \sum_{j=1}^J \frac{(O_j - E_j)^2}{E_j}, \text{ expected cell counts } E_j = n \cdot p_j(\hat{\lambda}).$$

Generalized likelihood ratio test approach: reject H_0 for large values of $2\Delta \approx X^2$ having an approximate null distribution χ_{J-1-r}^2 .

$$\text{df} = (\text{number of cells}) - 1 - (\text{number of independent parameters estimated from the data})$$

Since the chi-square test is approximate, all expected counts are recommended to be at least 5. If not, combine small cells and recalculate df.

Example: bird hops.

H_0 : number of hops that a bird does between flights has a $\text{Geom}(p)$ distribution. Using a MLE $\hat{p} = 0.358$ and $J = 7$ we obtain $X^2 = 1.86$. With $\text{df} = 5$ and $P\text{-value} = 0.87$ we do not reject the geometric distribution model for number of bird hops.

Example: gender ratio.

In a study made in Germany in 1889 the gender ratios for $n = 6115$ families with 12 children were recorded. The data give Y_1, \dots, Y_n numbers of boys in each family. Each Y_i has $J = 13$ possible values. Here we discuss two models for the gender ratio.

Model 1. A symmetric binomial model: $Y \sim \text{Bin}(12, 0.5)$ corresponds to a simple null hypothesis $H_0: p_j = \binom{12}{j} \cdot 2^{-12}, j = 0, 1, \dots, 12$. Expected cell counts $E_j = 6115 \cdot \binom{12}{j} \cdot 2^{-12}$.

cell j	O_j	E_j model 1	$\frac{(O_j - E_j)^2}{E_j}$	E_j model 2	$\frac{(O_j - E_j)^2}{E_j}$
0	7	1.5	20.2	2.3	9.6
1	45	17.9	41.0	26.1	13.7
2	181	98.5	69.1	132.8	17.5
3	478	328.4	68.1	410.0	11.3
4	829	739.0	11.0	854.2	0.7
5	1112	1182.4	4.2	1265.6	18.6
6	1343	1379.5	1.0	1367.3	0.4
7	1033	1182.4	18.9	1085.2	2.5
8	670	739.0	6.4	628.1	2.8
9	286	328.4	5.5	258.5	2.9
10	104	98.5	0.3	71.8	14.4
11	24	17.9	2.1	12.1	11.7
12	3	1.5	1.5	0.9	4.9
Total	6115	6115	249.2	6115	110.5

Model 1 results: $X^2 = 249.2$, $\text{df} = 12$, $\chi_{12}^2(0.005) = 28.3$, reject H_0 at 0.5% level.

Model 2. More flexible model: $Y \sim \text{Bin}(12, p)$ with an unspecified p . It leads to a composite null hypothesis $H_0: p_j = \binom{12}{j} \cdot p^j (1-p)^{12-j}, j = 0, \dots, 12, 0 \leq p \leq 1$. The MLE and expected cell counts

$$\hat{p} = \frac{\text{number of boys}}{\text{number of children}} = \frac{1 \cdot 45 + 2 \cdot 181 + \dots + 12 \cdot 3}{6115 \cdot 12} = 0.4808, \quad E_j = 6115 \cdot \binom{12}{j} \cdot \hat{p}^j \cdot (1 - \hat{p})^{12-j} .$$

Model 2 results: observed test statistic $X^2 = 110.5$, $r = 1$, $\text{df} = 11$, $\chi_{11}^2(0.005) = 26.76$, reject H_0 at 0.5% level.

Conclusion: even more flexible model is needed to address large variation in the observed cell counts. Suggestion: let the probability of a male child p to differ from family to family.