

What kinds of tests can be carried out for such data ?

1. Tests about p_1, \dots, p_k

- ▶ $H_0 : p_1 = \dots = p_k = 1/k$
- ▶ $H_0 : p_1, \dots, p_k$ determined by some parametric distribution (Normal, Poisson etc.)

2. Tests about the factors A and B

- ▶ are A and B dependent ?
- ▶ i.e. does classification by A influence classification by B .

Chi-Squared Test

For one-way tables: suppose that a null hypothesis **completely specifies** p_1, \dots, p_k , that is, we have H_0 of the form

$$H_0 : p_1 = p_1^{(0)}, \dots, p_k = p_k^{(0)}$$

where $p_1^{(0)}, \dots, p_k^{(0)}$ are fixed probabilities. For example, for $k = 3$,

$$H_0 : p_1 = p_2 = p_3 = 1/3$$

or

$$H_0 : p_1 = 1/2, p_2 = p_3 = 1/4$$

To test this hypothesis against the general alternative hypothesis

$$H_a : H_0 \text{ not true.}$$

we use the test statistic

$$X^2 = \sum_{i=1}^k \frac{(n_i - np_i^{(0)})^2}{np_i^{(0)}}$$

If H_0 is true,

$$X^2 \approx \text{Chi-squared}(k - 1).$$

that is, X^2 is approximately distributed as Chi-squared($k - 1$).

In this formula

- ▶ n_i is the **observed** count in cell i
- ▶ $np_i^{(0)}$ is the **expected** count in cell i if H_0 is **true**.

Sometimes the formula is written

$$X^2 = \sum_{i=1}^k \frac{(O_i - E_i)^2}{E_i}$$

where O_i is the observed count, and E_i is the expected count.

If

$$X^2 > \text{Chisq}_\alpha(k - 1)$$

then we reject H_0 at the α significance level, where $\text{Chisq}_\alpha(k - 1)$ is the $1 - \alpha$ (right-hand) tail critical value of the Chi-squared distribution with $k - 1$ degrees of freedom.

This method can be extended in the one-way case to test distribution assumptions, that is, for example

H_0 : Data Normally distributed

or

H_0 : Data Poisson distributed

Unfortunately this facility is not available in SPSS; direct calculation is possible but involved.

For the **two-way** table, we can test the hypothesis

H_0 : Factor A and Factor B levels are assigned independently

that is, classification by factor A is independent of classification by factor B. We use the same test statistic that can be rewritten

$$\chi^2 = \sum_{i=1}^r \sum_{j=1}^c \frac{(n_{ij} - \hat{n}_{ij})^2}{\hat{n}_{ij}}$$

where

$$\hat{n}_{ij} = \frac{n_{i.} n_{.j}}{n} \quad n_{i.} = \sum_{j=1}^c n_{ij} \quad n_{.j} = \sum_{i=1}^r n_{ij}.$$

The terms $n_{i.}$ and $n_{.j}$ are the row and column totals for row i and column j respectively.

If H_0 is true

$$X^2 \approx \text{Chi-squared}((r-1)(c-1))$$

i.e. the degrees of freedom quantity is $(r-1)(c-1)$.
Otherwise the test proceeds as before: we check whether

$$X^2 > \text{Chisq}_\alpha((r-1)(c-1))$$

and if so, we reject H_0 .

Example: DNA Sequence Data.

Counts of Nucleotides A,C,G,T in a genomic segment related to the breast cancer gene BRCA2.

Example: Eye and Hair Colour Data.

The assignment of hair and eye colour in a sample of humans

See handout.

Note: For the Chi-squared test to be valid, we need the expected cell counts

$$np_i^{(0)} \quad i = 1, \dots, k$$

or

$$\hat{n}_{ij} \quad i = 1, \dots, r, j = 1, \dots, c$$

to be sufficiently large. The convention is to require the expected value to be greater than **five**.