Goodness of Fit Tests Mathematics 47: Lecture 32

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Goodness of Fit Tests

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• Given some  $\Pi = (\pi_1, \pi_2, \dots, \pi_k)$ , where  $0 \le \pi_i \le 1$  for  $i = 1, 2, \dots, k$ and  $\sum_{i=1}^k \pi_i = 1$ , suppose we wish to test

 $H_0: \mathbf{p} = \mathbf{\Pi}$  $H_A: \mathbf{p} \neq \mathbf{\Pi}.$ 

• If, for 
$$i = 1, 2, \ldots, k$$
, we let

 $Y_i$  = number of outcomes of type i,

then the likelihood function is

$$L(p_1, p_2, \ldots, p_k) = p_1^{Y_1} p_2^{Y_2} \cdots p_k^{Y_k}.$$

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We have seen previously that the maximum likelihood estimator for p<sub>i</sub> is

$$\hat{p}_i = \frac{\mathbf{Y}_i}{n},$$

so the generalized likelihood ratio is

$$\Lambda = \frac{L(\pi_1, \pi_2, \ldots, \pi_k)}{L(\hat{p}_1, \hat{p}_2, \ldots, \hat{p}_k)} = \prod_{i=1}^k \left(\frac{\pi_i}{\hat{p}_i}\right)^{Y_i}.$$

► Hence

$$-2\log(\Lambda) = -2\sum_{i=1}^{k} Y_i \log\left(\frac{\pi_i}{\hat{p}_i}\right)$$
$$= 2\sum_{i=1}^{k} Y_i \log\left(\frac{\hat{p}_i}{\pi_i}\right)$$
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► Note: when H<sub>0</sub> is true, nπ<sub>i</sub> = E[Y<sub>i</sub>], the expected number of observations of outcome i.

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- Note: when H₀ is true, nπ<sub>i</sub> = E[Y<sub>i</sub>], the expected number of observations of outcome i.
- Hence we may remember the final expression for  $-2\log(\Lambda)$  as

$$2\sum$$
(Observed) log  $\left(\frac{Observed}{Expected}\right)$ ,

where the sum extends over all possible outcomes.

Now for large n, −2 log(Λ) is approximately χ<sup>2</sup>(k − 1) (k − 1, not k, because there are only k − 1 parameters since ∑<sup>k</sup><sub>i=1</sub> p<sub>i</sub> = 1).

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- Since the generalized likelihood ratio test rejects H<sub>0</sub> for small values of Λ, we reject H<sub>0</sub> for large values of −2 log(Λ), with, for an observed value λ of Λ, *p*-value equal to P(U ≥ −2 log(λ)), where U is χ<sup>2</sup>(k − 1).

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- Since the generalized likelihood ratio test rejects H<sub>0</sub> for small values of Λ, we reject H<sub>0</sub> for large values of −2 log(Λ), with, for an observed value λ of Λ, *p*-value equal to P(U ≥ −2 log(λ)), where U is χ<sup>2</sup>(k − 1).
- ► Note: A conservative rule of thumb is to assume the approximation to the chi-square distribution is reasonable as long as nπ<sub>i</sub> ≥ 5 for i = 1, 2, ..., k.

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Let

- $p_1 =$  probability of smooth-yellow,
- $p_2 =$  probability of smooth-green,
- $p_3 =$  probability of wrinkled-yellow,
- $p_4 =$  probability of wrinkled-green.

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- $p_4 =$  probability of wrinkled-green.

• Mendel's theory predicted that  $p_1 = \frac{9}{16}$ ,  $p_2 = \frac{3}{16}$ ,  $p_3 = \frac{3}{16}$ , and  $p_4 = \frac{1}{16}$ .

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► Hence we want to test

$$H_0: \mathbf{p} = \left(\frac{9}{16}, \frac{3}{16}, \frac{3}{16}, \frac{1}{16}\right)$$
$$H_A: \mathbf{p} \neq \left(\frac{9}{16}, \frac{3}{16}, \frac{3}{16}, \frac{1}{16}\right)$$

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ight). \end{aligned}$$

The following table contains Mendel's data, along with the expected frequency of each outcome under  $H_0$ :

Туре	<b>Observed Frequency</b>	Expected Frequency
Smooth-yellow	315	312.75
Smooth-green	108	104.25
Wrinkled-yellow	101	104.25
Wrinkled-green	32	34.75
Total	556	556.00

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► We may now compute

$$-2\log(\lambda) = 2\sum_{i=1}^{4} y_i \log\left(\frac{y_i}{n\pi_i}\right) = 0.4754$$

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We may now compute

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 If U is χ<sup>2</sup>(3), then the p-value for this test is P(U ≥ 0.4754) = 0.9242617, giving us no evidence for rejecting the null hypothesis.

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- If U is χ<sup>2</sup>(3), then the p-value for this test is P(U ≥ 0.4754) = 0.9242617, giving us no evidence for rejecting the null hypothesis.
- Indeed, these data provide such a good fit to Mendel's data that it has been suspected that his assistants manipulated the data to fit the hypothesis.



$$f(x) = x \log\left(\frac{x}{a}\right).$$

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• So f(a) = 0, f'(a) = 1, and  $f''(a) = \frac{1}{a}$ .

▶ Hence, for *x* close to *a*,

$$f(x) \approx (x-a) + \frac{1}{2a}(x-a)^2.$$

• Thus, when  $H_0$  is true and *n* is large,

$$y_i \log\left(\frac{y_i}{n\pi_i}\right) \approx (y_i - n\pi_i) + \frac{1}{2n\pi_i}(y_i - n\pi_i)^2.$$

• Thus, when  $H_0$  is true and *n* is large,

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It follows that

$$2\log(\Lambda) = 2\sum_{i=1}^{k} Y_i \log\left(\frac{Y_i}{n\pi_i}\right)$$
$$\approx 2\sum_{i=1}^{k} (Y_i - n\pi_i) + \sum_{i=1}^{k} \frac{(Y_i - n\pi_i)^2}{n\pi_i}$$
$$= 2(n - n) + \sum_{i=1}^{k} \frac{(Y_i - n\pi_i)^2}{n\pi_i}$$
$$= \sum_{i=1}^{k} \frac{(Y_i - n\pi_i)^2}{n\pi_i}.$$

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$$Q = \sum_{i=1}^{k} \frac{(Y_i - n\pi_i)^2}{n\pi_i}$$

Pearson's chi-square statistic.

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We may remember this formula as

$$\sum \frac{(\mathsf{Observed} - \mathsf{Expected})^2}{\mathsf{Expected}}$$

where the sum extends over all possible outcomes.

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We may remember this formula as

$$\sum \frac{(\mathsf{Observed} - \mathsf{Expected})^2}{\mathsf{Expected}},$$

where the sum extends over all possible outcomes.

We may use either −2 log(Λ) or Q to perform the chi-square goodness of fit test.

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► For Mendel's data, we have

$$q = \sum_{i=1}^{4} \frac{(y_i - n\pi_i)^2}{n\pi_i} = 0.4700,$$

differing only slightly from the value of  $-2\log(\lambda)$  computed above.

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Note: If the observed frequencies are in the vector x and the hypothesized probabilities are in the vector pi, then the R command

> chisq.test(x,p=pi)

performs the goodness of fit test using Pearson's chi-square statistic.