

## 8.4. STATISTICAL SIGNIFICANCE TESTS

 Table 8.4.1.1. Values of  $\chi^2/\nu$  for which the c.d.f.  $\Psi(\chi^2, \nu)$  has the values given in the column headings, for various values of  $\nu$ 

$\nu$	0.5	0.9	0.95	0.99	0.995
1	0.4549	2.7055	3.8415	6.6349	7.8795
2	0.6931	2.3026	2.9957	4.6052	5.2983
3	0.7887	2.0838	2.6049	3.7816	4.2794
4	0.8392	1.9449	2.3719	3.3192	3.7151
6	0.8914	1.7741	2.0986	2.8020	3.0913
8	0.9180	1.6702	1.9384	2.5113	2.7444
10	0.9342	1.5987	1.8307	2.3209	2.5188
15	0.9559	1.4871	1.6664	2.0385	2.1868
20	0.9669	1.4206	1.5705	1.8783	1.9999
25	0.9735	1.3753	1.5061	1.7726	1.8771
30	0.9779	1.3419	1.4591	1.6964	1.7891
40	0.9834	1.2951	1.3940	1.5923	1.6692
50	0.9867	1.2633	1.3501	1.5231	1.5898
60	0.9889	1.2400	1.3180	1.4730	1.5325
80	0.9917	1.2072	1.2735	1.4041	1.4540
100	0.9933	1.1850	1.2434	1.3581	1.4017
120	0.9945	1.1686	1.2214	1.3246	1.3638
140	0.9952	1.1559	1.2044	1.2989	1.3346
160	0.9958	1.1457	1.1907	1.2783	1.3114
200	0.9967	1.1301	1.1700	1.2472	1.2763

$$\Gamma(x) = \int_0^{\infty} t^{x-1} \exp(-t) dt. \quad (8.4.1.13)$$

Although this function is continuous for all  $x > 0$ , its value is of interest in the context of this analysis only for  $x$  equal to positive, integral multiples of  $1/2$ . It can be shown that  $\Gamma(1/2) = \sqrt{\pi}$ ,  $\Gamma(1) = 1$ , and  $\Gamma(x+1) = x\Gamma(x)$ . It follows that, for a positive integer,  $n$ ,  $\Gamma(n) = (n-1)!$ , and that  $\Gamma(3/2) = \sqrt{\pi}/2$ ,  $\Gamma(5/2) = 3\sqrt{\pi}/4$ , etc. The *beta function* is defined by

$$B(x, y) = \int_0^1 t^{x-1} (1-t)^{y-1} dt. \quad (8.4.1.14)$$

It can be shown (Prince, 1994) that  $B(x, y) = \Gamma(x)\Gamma(y)/\Gamma(x+y)$ . Making the substitution  $t = z_1^2/z_1^2 + z_2^2$ , (8.4.1.12) becomes

$$\begin{aligned} \Phi(s^2) &= \frac{1}{2\pi} \exp\left(-\frac{s^2}{2}\right) \int_0^1 [t(1-t)]^{-1/2} dt \\ &= \frac{1}{2\pi} \exp\left(-\frac{s^2}{2}\right) B(1/2, 1/2) \\ &= \frac{1}{2} \exp\left(-\frac{s^2}{2}\right), \quad s^2 \geq 0. \end{aligned} \quad (8.4.1.15)$$

By a similar procedure, it can be shown that, if  $\chi^2$  is the sum of  $\nu$  terms,  $z_1^2, z_2^2, \dots, z_\nu^2$ , where all are drawn independently from a population with the p.d.f. given in (8.4.1.10),  $\chi^2$  has the p.d.f.

$$\begin{aligned} \Phi(\chi^2, \nu) &= \frac{(\chi^2)^{\nu/2-1}}{2^{\nu/2} \Gamma(\nu/2)} \exp\left(-\frac{\chi^2}{2}\right), \quad \chi^2 > 0, \\ \Phi(\chi^2, \nu) &= 0, \quad \chi^2 \leq 0. \end{aligned} \quad (8.4.1.16)$$

The parameter  $\nu$  is known as the number of *degrees of freedom*, but this use of that term must not be confused with the conventional use in physics and chemistry. The p.d.f. in (8.4.1.16) is the *chi-squared distribution with  $\nu$  degrees of freedom*. Table 8.4.1.1 gives the values of  $\chi^2/\nu$  for which the *cumulative distribution function (c.d.f.)*  $\Psi(\chi^2, \nu)$  has various values for various choices of  $\nu$ . This table is provided to enable verification of computer codes that may be used to generate more extensive tables. It was generated using a program included in

the statistical library DATAPAC (Filliben, unpublished). Fortran code for this program appears in Prince (1994).

The quantity  $(n-p)G$  is the sum of  $n$  terms that have mean value  $(n-p)/n$ . Because the process of determining the least-squares fit establishes  $p$  relations among them, however, only  $(n-p)$  of the terms are independent. The number of degrees of freedom is therefore  $\nu = (n-p)$ , and, if the model is correct, and the terms have been properly weighted,  $\chi^2 = (n-p)G^2$  has the chi-squared distribution with  $(n-p)$  degrees of freedom. In crystallography, the number of degrees of freedom tends to be large, and the p.d.f. for  $G$  correspondingly sharp, so that even rather small deviations from  $G^2 = 1$  should cause one or both of the hypotheses of a correct model and appropriate weights to be rejected. It is common practice to assume that the model is correct, and that the weights have correct *relative* values, that is that they have been assigned by  $w_i = k/\sigma_i^2$ , where  $k$  is a number different from, usually greater than, one.  $G$  is then taken to be an estimate of  $k$ , and all elements of  $(A^T W A)^{-1}$  (Section 8.1.2) are multiplied by  $G^2$  to get an estimated variance-covariance matrix. The range of validity of this procedure is limited at best. It is discussed further in Chapter 8.5.

 8.4.2. The *F* distribution

Consider an unconstrained model with  $p$  parameters and a constrained one with  $q$  parameters, where  $q < p$ . We wish to decide whether the constrained model represents an adequate fit to the data, or if the additional parameters in the unconstrained model provide, in some important sense, a better fit to the data. Provided the  $(p-q)$  additional columns of the design matrix,  $A$ , are linearly independent of the previous  $q$  columns, the sum of squared residuals must be reduced by some finite amount by adjusting the additional parameters, but we must decide whether this improved fit would have occurred purely by chance, or whether it represents additional information.

Let  $s_c^2$  and  $s_u^2$  be the weighted sums of squared residuals for the constrained and unconstrained models, respectively. If the constrained and unconstrained models are equally good representations of the data, and the weights have been assigned by  $w_i = 1/\sigma_i^2$ , the expected values of the sums of squares are  $\langle s_c^2 \rangle = (n-q)$  and  $\langle s_u^2 \rangle = (n-p)$ , and, further, they should be distributed as  $\chi^2$  with  $(n-q)$  and  $(n-p)$  degrees of freedom, respectively. Also,  $\langle s_c^2 - s_u^2 \rangle = (p-q)$ , and  $(s_c^2 - s_u^2)$  is distributed as  $\chi^2$  with  $(p-q)$  degrees of freedom.  $s_c^2$  and  $s_u^2$  are not independent, but  $(s_c^2 - s_u^2)$  is the squared magnitude of a vector in a  $(p-q)$ -dimensional subspace that is orthogonal to the  $(n-p)$ -dimensional space of  $s_u^2$ . Therefore,  $s_u^2$  and  $(s_c^2 - s_u^2)$  are independent, random variables, each with a  $\chi^2$  distribution. Let  $\chi_1^2 = (s_c^2 - s_u^2)$ ,  $\chi_2^2 = s_u^2$ ,  $\nu_1 = p-q$ , and  $\nu_2 = n-p$ . The ratio  $F = (\chi_1^2/\nu_1)/(\chi_2^2/\nu_2)$  should have a value close to one, even if the weights have relative rather than absolute values, but we need a measure of how far away from one this ratio can be before we must reject the hypothesis that the two models are equally good representations of the data. The conditional p.d.f. for  $F$ , given a value of  $\chi_2^2$ , is

$$\Phi_C(F|\chi_2^2) = \frac{[(\nu_1/\nu_2)\chi_2^2]^{\nu_1/2} F^{\nu_1/2-1}}{2^{\nu_1/2} \Gamma(\nu_1/2)} \exp[-(\nu_1/\nu_2)\chi_2^2 F/2], \quad (8.4.2.1)$$

and the marginal p.d.f. for  $\chi_2^2$  is

$$\Phi_M(\chi_2^2) = \frac{(\chi_2^2)^{\nu_2/2-1}}{2^{\nu_2/2} \Gamma(\nu_2/2)} \exp(-\chi_2^2/2). \quad (8.4.2.2)$$

## 8. REFINEMENT OF STRUCTURAL PARAMETERS

Table 8.4.2.1. Values of the  $F$  ratio for which the c.d.f.  $\Psi(F, \nu_1, \nu_2)$  has the value 0.95, for various choices of  $\nu_1$  and  $\nu_2$

$\nu_1 \backslash \nu_2$	1	2	4	8	15
10	4.9646	4.1028	3.4781	3.0717	2.8450
20	4.3512	3.4928	2.8661	2.4471	2.2033
30	4.1709	3.3158	2.6896	2.2662	2.0148
40	4.0847	3.2317	2.6060	2.1802	1.9245
50	4.0343	3.1826	2.5572	2.1299	1.8714
60	4.0012	3.1504	2.5252	2.0970	1.8364
80	3.9604	3.1108	2.4859	2.0564	1.7932
100	3.9361	3.0873	2.4626	2.0323	1.7675
120	3.9201	3.0718	2.4472	2.0164	1.7505
150	3.9042	3.0564	2.4320	2.0006	1.7335
200	3.8884	3.0411	2.4168	1.9849	1.7167
300	3.8726	3.0259	2.4017	1.9693	1.6998
400	3.8648	3.0183	2.3943	1.9616	1.6914
600	3.8570	3.0107	2.3868	1.9538	1.6831
1000	3.8508	3.0047	2.3808	1.9477	1.6764

Table 8.4.3.1. Values of  $t$  for which the c.d.f.  $\Psi(t, \nu)$  has the values given in the column headings, for various values of  $\nu$

$\nu$	0.75	0.90	0.95	0.99	0.995
1	1.0000	3.0777	6.3138	31.8206	63.6570
2	0.8165	1.8856	2.9200	6.9646	9.9249
3	0.7649	1.6377	2.3534	4.5407	5.8409
4	0.7407	1.5332	2.1319	3.7469	4.6041
6	0.7176	1.4398	1.9432	3.1427	3.7074
8	0.7064	1.3968	1.8596	2.8965	3.3554
10	0.6998	1.3722	1.8125	2.7638	3.1693
12	0.6955	1.3562	1.7823	2.6810	3.0546
14	0.6924	1.3450	1.7613	2.6245	2.9769
16	0.6901	1.3368	1.7459	2.5835	2.9208
20	0.6870	1.3253	1.7247	2.5280	2.8453
25	0.6844	1.3164	1.7081	2.4851	2.7874
30	0.6828	1.3104	1.6973	2.4573	2.7500
35	0.6816	1.3062	1.6896	2.4377	2.7238
40	0.6807	1.3031	1.6839	2.4233	2.7045
50	0.6794	1.2987	1.6759	2.4033	2.6778
60	0.6786	1.2958	1.6707	2.3901	2.6603
80	0.6776	1.2922	1.6641	2.3739	2.6387
100	0.6770	1.2901	1.6602	2.3642	2.6259
120	0.6765	1.2886	1.6577	2.3578	2.6174

The marginal p.d.f. for  $F$  is obtained by integration of the joint p.d.f.,

$$\Phi(F) = \int_0^{\infty} \Phi_C(F|\chi_2^2) \Phi_M(\chi_2^2) d\chi_2^2, \quad (8.4.2.3)$$

yielding the result

$$\Phi(F, \nu_1, \nu_2) = \frac{(\nu_1/\nu_2)F^{\nu_1/2-1}}{B(\nu_1/2, \nu_2/2)[1 + (\nu_1/\nu_2)F]^{(\nu_1+\nu_2)/2}}. \quad (8.4.2.4)$$

This p.d.f. is known as the  $F$  distribution with  $\nu_1$  and  $\nu_2$  degrees of freedom. Table 8.4.2.1 gives the values of  $F$  for which the c.d.f.  $\Psi(F, \nu_1, \nu_2)$  is equal to 0.95 for various choices of  $\nu_1$  and  $\nu_2$ . Fortran code for the program from which the table was generated appears in Prince (1994).

The cumulative distribution function  $\Psi(F, \nu_1, \nu_2)$  gives the probability that the  $F$  ratio will be less than some value by chance if the models are equally consistent with the data. It is therefore a necessary, but not sufficient, condition for concluding that the unconstrained model gives a significantly better fit to the data that  $\Psi(F, \nu_1, \nu_2)$  be greater than  $1 - \alpha$ , where  $\alpha$  is the desired level of significance. For example, if  $\Psi(F, \nu_1, \nu_2) = 0.95$ , the probability is only 0.05 that a value of  $F$  this large or greater would have been observed if the two models were equally good representations of the data.

Hamilton (1964) observed that the  $F$  ratio could be expressed in terms of the crystallographic weighted  $R$  index, which is defined, for refinement on  $|F|$  (and similarly for refinement on  $|F|^2$ ), by

$$R_w = [\sum w_i(|F_{o_i}| - |F_{c_i}|)^2 / \sum w_i |F_{o_i}|^2]^{1/2}. \quad (8.4.2.5)$$

Denoting by  $R_c$  and  $R_u$  the weighted  $R$  indices for the constrained and unconstrained models, respectively,

$$F = (\nu_2/\nu_1)[(R_c/R_u)^2 - 1], \quad (8.4.2.6)$$

and a c.d.f. for  $R_c/R_u$  can be readily derived from this relation. A significance test based on  $R_c/R_u$  is known as *Hamilton's R-ratio test*; it is entirely equivalent to a test on the  $F$  ratio.

### 8.4.3. Comparison of different models

Tests based on  $F$  or the  $R$  ratio have several limitations. One important one is that they are applicable only when the

parameters of one model form a subset of the parameters of the other. Also, the  $F$  test makes no distinction between improvement in fit as a result of small improvements throughout the entire data set and a large improvement in a small number of critically sensitive data points. A test that can be used for comparing arbitrary pairs of models, and that focuses attention on those data points that are most sensitive to differences in the models, was introduced by Williams & Klot (1953; also Himmelblau, 1970; Prince, 1982).

Consider a set of observations,  $y_{0i}$ , and two models that predict values for these observations,  $y_{1i}$  and  $y_{2i}$ , respectively. We determine the slope of the regression line  $z = \lambda x$ , where  $z_i = [y_{0i} - (1/2)(y_{1i} + y_{2i})]/\sigma_i$ , and  $x_i = (y_{1i} - y_{2i})/\sigma_i$ . Suppose model 1 is a perfect fit to the data, which have been measured with great precision, so that  $y_{0i} = y_{1i}$  for all  $i$ . Under these conditions,  $\lambda = +1/2$ . Similarly, if model 2 is a perfect fit,  $\lambda = -1/2$ . Real experimental data, of course, are subject to random error, and  $|\lambda|$  in general would be expected to be less than  $1/2$ . A least-squares estimate of  $\lambda$  is

$$\hat{\lambda} = \frac{\sum_{i=1}^n z_i x_i}{\sum_{i=1}^n x_i^2}, \quad (8.4.3.1)$$

and it has an estimated variance

$$\hat{\sigma}_{\hat{\lambda}}^2 = \frac{\sum_{i=1}^n z_i^2 - \hat{\lambda}^2 \sum_{i=1}^n x_i^2}{(n-1) \sum_{i=1}^n x_i^2}. \quad (8.4.3.2)$$

The hypothesis that the two models give equally good fits to the data can be tested by considering  $\hat{\lambda}$  to be an unconstrained, one-parameter fit that is to be compared with a constrained, zero-parameter fit for which  $\lambda = 0$ . A p.d.f. for making this comparison can be derived from an  $F$  distribution with  $\nu_1 = 1$  and  $\nu_2 = \nu = (n - 1)$ .

$$\Phi(F, 1, \nu) = \frac{\Gamma[(\nu + 1)/2]}{\sqrt{\pi\nu}\Gamma(\nu/2)(1 + F/\nu)^{(\nu+1)/2}}. \quad (8.4.3.3)$$