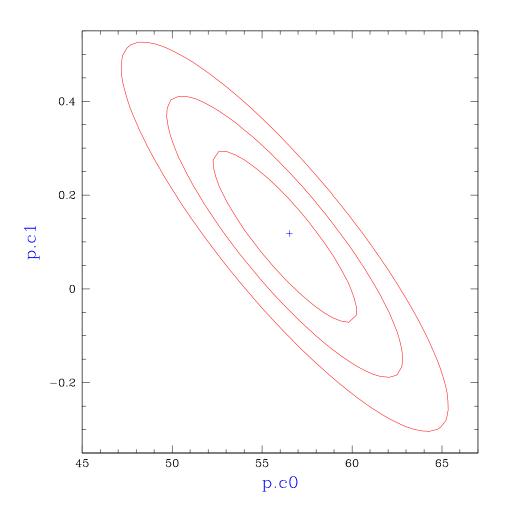
Statistics II: Model Comparison and Parameter Estimation



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Now, Shifting Gears...

A model M has been fit to dataset D and either the maximum of the likelihood function \mathcal{L}_{\max} , the minimum of the χ^2 statistic χ^2_{\min} , or the mode of the posterior distribution $p(\hat{\theta}|D)$ has been determined. What comes next?

- *Model Comparison*. The determination of which of a suite of models (e.g. blackbody, power-law, etc.) best represents the data.
- Parameter Estimation. The characterization of the sampling distribution for each best-fit model parameter (e.g. blackbody temperature and normalization), which allows the errors (i.e. standard deviations) of each parameter to be determined.
- Publication!

Here, we cannot ignore the frequentist/Bayesian divide. Hence we will discuss how frequentists and Bayesians would complete these tasks, separately...

Two models, M_0 and M_1 , have been fit to D. M_0 , the "simpler" of the two models (generally speaking, the model with fewer free parameters) is the *null hypothesis*.

A frequentist would compare these models by:

- constructing a test statistic T from the best-fit statistics of each fit $(e.g. \Delta \chi^2 = \chi_0^2 \chi_1^2)$;
- determining each sampling distributions for T, $p(T|M_0)$ and $p(T|M_1)$;
- determining the *significance*, or Type I error, the probability of selecting M_1 when M_0 is correct:

$$\alpha = \int_{T_{
m obs}}^{\infty} dT p(T|M_0);$$

• and determining the *power*, or Type II error, which is related to the probability β of selecting M_0 when M_1 is correct:

$$1 - \beta = \int_{T_{\text{obs}}}^{\infty} dT p(T|M_1) .$$

 \Rightarrow If α is smaller than a pre-defined threshold (≤ 0.05 , or $\leq 10^{-4}$, etc., with smaller thresholds used for more controversial alternative models), then the frequentist rejects the null hypothesis.

⇒ If there are several model comparison tests to choose from, the frequentist uses the most powerful one!

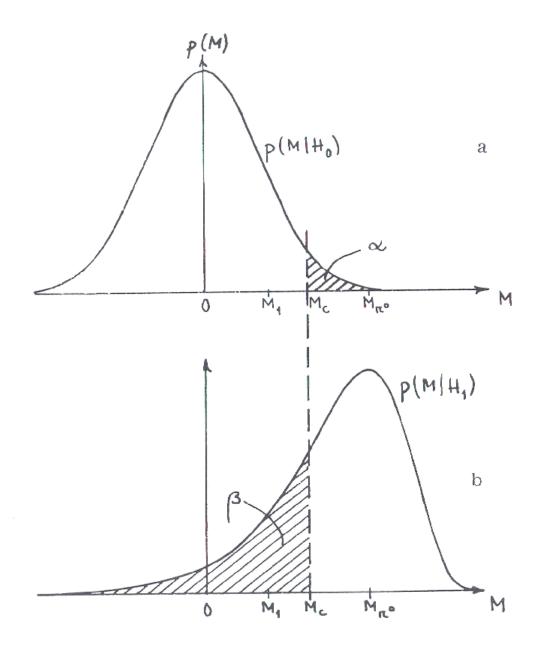


Figure 1: Comparison of distributions $p(T|M_0)$ (from which one determines the significance α) and $p(T|M_1)$ (from which one determines the power of the model comparison test $1-\beta$ (Eadie *et al.* 1971, p. 217).

Standard frequentist model comparison tests include:

• The χ^2 Goodness-of-Fit (GoF) test:

$$egin{array}{lcl} lpha_{\chi^2} &=& \int_{\chi^2_{\min,0}}^{\infty} d\chi^2 p(\chi^2|N-P_0) \ &=& rac{1}{2\Gamma(rac{N-P_0}{2})} \int_{\chi^2_{\min,0}}^{\infty} d\chi^2 \left(rac{\chi^2}{2}
ight)^{rac{N-P_0}{2}-1} e^{-rac{\chi^2}{2}} \, . \end{array}$$

• The Maximum Likelihood Ratio (MLR) test:

$$lpha_{\chi^2
m MLR} \; = \; \int_{\Delta\chi^2}^{\infty} d\chi^2 p(\Delta\chi^2 | \Delta P) \; ,$$

where ΔP is the number of additional freely varying model parameters in model M_1 .

• The F-test:

$$\alpha_F = \int_F^{\infty} dF \ p(F|\Delta P, N - P_1)$$

$$= I_{\frac{N-P_1}{N-P_1 + (\Delta P)F}} \left(\frac{N-P_1}{2}, \frac{\Delta P}{2}\right) ,$$

where P_1 is the total number of thawed parameters in model M_1 , I is the incomplete beta function, and F is the F-statistic

$$F = \frac{\Delta \chi^2}{\Delta P} / \frac{\chi_1^2}{(N - P_1)}.$$

These are standard tests because they allow estimation of the significance without time-consuming simulations!

Notes and caveats regarding these standard tests:

- The GoF test is an "alternative-free" test, as it does not take into account the alternative model M_1 . It is consequently a weak (i.e. not powerful) model comparison test and should not be used!
- Only the version of the F-test which generally has the greatest power is shown above: in principle, one can construct three F statistics out of χ_0^2 , χ_1^2 , and $\Delta \chi^2$.
- The MLR ratio test is generally the most powerful for detecting emission and absorption lines in spectra.

But the most important caveat of all is that...

The F and MLR tests are commonly misused by astronomers! There are two important conditions that must be met so that an estimated derived value α is actually correct, *i.e.* so that it is an accurate approximation of the tail integral of the sampling distribution (Protassov *et al.* 2001):

- M_0 must be nested within M_1 , i.e. one can obtain M_0 by setting the extra ΔP parameters of M_1 to default values, often zero; and
- those default values may not be on a parameter space boundary.

The second condition may not be met, e.g., when one is attempting to detect an emission line, whose default amplitude is zero and whose minimum amplitude is zero. Protassov et al. recommend Bayesian posterior predictive probability values as an alternative, but a discussion of this topic is beyond the scope of this class.

If the conditions for using these tests are not met, then they can still be used, but the significance must be computed via Monte Carlo simulations.

Bayesian Model Comparison

In the previous class, we showed how Bayes' theorem is applied in model fits. It can also be applied to model comparison:

$$p(M|D) = p(M) \frac{p(D|M)}{p(D)}.$$

- p(M) is the prior probability for M;
- ullet p(D) is an ignorable normalization constant; and
- p(D|M) is the average, or global, likelihood:

$$p(D|M) = \int d\theta p(\theta|M) p(D|M,\theta)$$
$$= \int d\theta p(\theta|M) \mathcal{L}(M,\theta).$$

In other words, it is the (normalized) integral of the posterior distribution over all parameter space. Note that this integral may be computed numerically, by brute force, or if the likelihood surface is approximately a multi-dimensional Gaussian (i.e. if $\mathcal{L} \propto \exp[-\chi^2/2]$), by the Laplace approximation:

$$p(D|M) = p(\hat{\theta}|M)(2\pi)^{P/2}\sqrt{{
m det}C}\mathcal{L}_{
m max}$$
,

where C is the covariance matrix (estimated numerically at the mode).

Bayesian Model Comparison

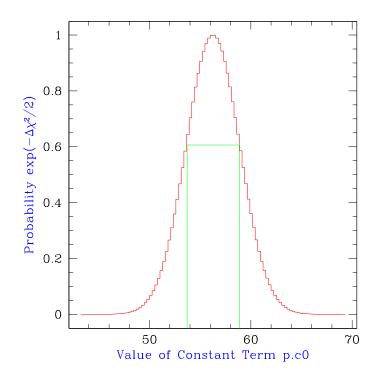
To compare two models, a Bayesian computes the *odds*, or odds ratio:

$$egin{array}{lcl} O_{10} & = & rac{p(M_1|D)}{p(M_0|D)} \ & = & rac{p(M_1)p(D|M_1)}{p(M_0)p(D|M_0)} \ & = & rac{p(M_1)}{p(M_0)}B_{10} \,, \end{array}$$

where B_{10} is the Bayes factor. When there is no a priori preference for either model, $B_{10} = 1$ of one indicates that each model is equally likely to be correct, while $B_{10} \geq 10$ may be considered sufficient to accept the alternative model (although that number should be greater if the alternative model is controversial).

One should speak of *confidence* or *credible intervals* or *regions* rather than "errors."

- A frequentist derives confidence intervals and regions.
- A Bayesians derives credible intervals and regions.
- An interval is a range (or ranges) of values of a parameter θ that has probability p_{int} of containing the parameter's true value θ_o . (A region is simply the multi-dimensional analogue of an interval.)
- A infinite number of intervals can be defined for a given parameter: here, we'll speak of intervals that contain the *most probable* parameter values.



Instead of the integrated probability p_{int} , many speak of "numbers of σ ." One can convert from $n\sigma$ to p_{int} using the following equation:

$$p_{\mathrm{int}} = \frac{1}{\sqrt{2\pi}\sigma} \int_{-n\sigma}^{+n\sigma} dx \exp\left(-\frac{x^2}{2\sigma^2}\right) = \mathrm{erf}\left(\frac{n}{\sqrt{2}}\right)$$

$p_{ m int}$	σ
68.3%	1.0
90.0%	1.6
95.5%	2.0
99.0%	2.6
99.7%	3.0

Note: this conversion between p_{int} and σ , while strictly true only if the sampling distribution is a one-dimensional Gaussian, is used by many astronomers in casual conversation regardless of the actual distribution shape or dimensionality.

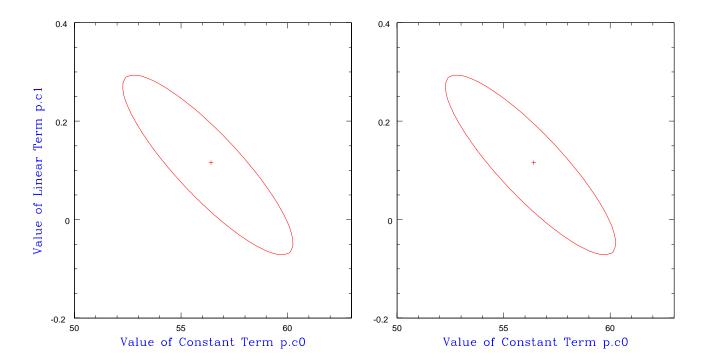
- Tables showing $\Delta \chi^2$ as a function of integrated probability $p_{\rm int}$ and number of degrees of freedom $\nu = N P$ can cause confusion. For instance:
 - "I have two free parameters in my model. Hence I should compute 68.3% confidence intervals for each parameter using $\Delta\chi^2=2.30$, right?"
 - "No."

			ν			
$p_{ m int}$	1	2	3	4	5	6
68.3%	1.00	2.30	3.53	4.72	5.89	7.04
90%	2.71	4.61	6.25	7.78	9.24	10.6
95.4%	4.00	6.17	8.02	9.70	11.3	12.8
99%	6.63	9.21	11.3	13.3	15.1	16.8
99.73%	9.00	11.8	14.2	16.3	18.2	20.1
99.99%	15.1	18.4	21.1	23.5	25.7	27.8

 $\Delta \chi^2$ as a Function of Confidence Level and Degrees of Freedom (Based on Press et al. 1986, p. 536.)

- To find the $n\sigma$ confidence interval for one parameter, use $\Delta \chi^2$ for $\nu = 1$ (or n^2).
- To find the $n\sigma$ joint confidence region for m parameters, use $\Delta \chi^2$ for $\nu = m$.
- To find either an interval or region using the likelihood function \mathcal{L} , use $\Delta \log \mathcal{L} = \Delta \chi^2/2$.

Never project a (properly estimated) region onto a parameter axis to estimate an interval! This always overestimates the size of the interval.



Frequentist Parameter Estimation

To determine confidence intervals and regions, a frequentist generally must simulate and fit new datasets to determine the sampling distributions for each model parameter.

- If the true parameter values are unknown (which is usually the case), then a grid of model parameter values should be constructed, with a large number of datasets sampled at each grid point.
- But the usual choice is to appeal to asymptotic behavior and sample datasets using $M(\hat{\theta})$. This method may only be useful in limited circumstances, as ≥ 100 datasets should be sampled and fit for accurate results.

Frequentist Parameter Estimation

One can estimate confidence intervals without having to use simulations if the χ^2 or $\log \mathcal{L}$ surface in parameter space is "well-behaved," *i.e.* if

- the surface is approximately shaped like a multidimensional paraboloid; and
- the best-fit point is sufficiently far from parameterspace boundaries.

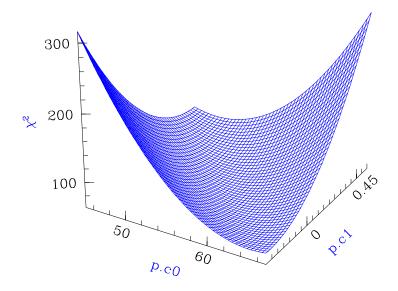
Three common ways of determining $n\sigma$ intervals are:

• varying a parameter's value, while holding the values of all other parameters at their best-fit values, until

$$-\chi^2 = \chi_o^2 + n^2$$
; or $-\log \mathcal{L} = \log \mathcal{L}_o - \frac{n^2}{2}$;

- the same as above, but allowing the values of all other parameters are allowed to float to new best-fit values; and
- computing $n\sqrt{C_{i,i}}$, where the covariance matrix $C_{i,j} = I_{i,j}^{-1}$, and I, the information matrix computed at the best-fit point, is

$$I_{i,j} \equiv \frac{1}{2} \frac{\partial^2 \chi^2}{\partial \theta_i \partial \theta_j} \text{ or } \frac{\partial^2 \log \mathcal{L}}{\partial \theta_i \partial \theta_j}.$$



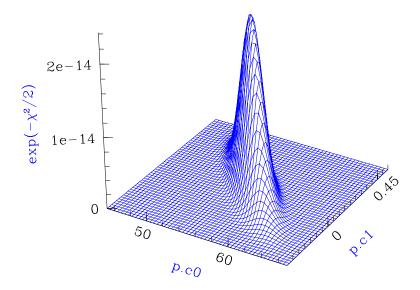


Figure 2: Example of a "well-behaved" statistical surface in parameter space, viewed as a multi-dimensional paraboloid (χ^2, top) , and as a multi-dimensional Gaussian $(\exp(-\chi^2/2) \approx \mathcal{L}, bottom)$.

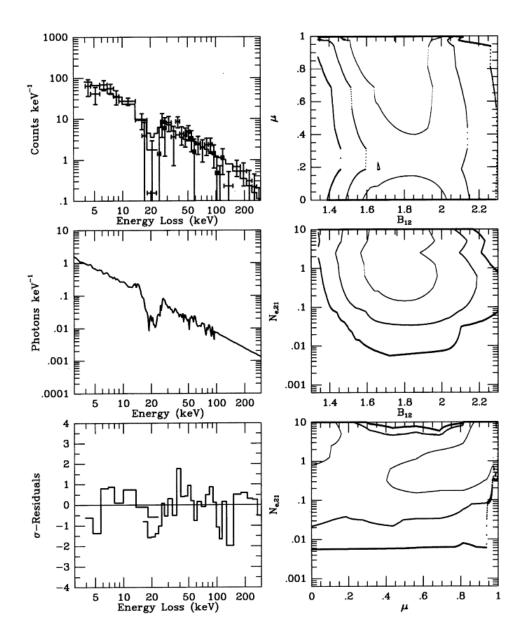


Figure 3: On the right, 1, 2, and 3σ contours determined for a statistical surface that is not "well-behaved" in parameter space. With such a surface, rigorous parameter estimation involves simulations (frequentist approach) or numerical integration of the surface (Bayesian approach). From Freeman *et al.* (1999).

Frequentist Parameter Estimation

Things to keep in mind about these confidence interval estimators (dubbed UNCERTAINTY, PROJECTION, and COVARIANCE in *Sherpa*, respectively):

- The first method will *always* underestimate the interval if the value of the parameter of interest is correlated with other model parameter values.
- The second method (which is relatively slow) is in a rigorous sense no more accurate than the third method (which is fast), but it does provide a means of visualizing the statistical surface.
- A statistical surface is "well-behaved" if the second and third methods give the same interval estimates.
- The condition that the best-fit point be sufficiently far from parameter-space boundaries means that these methods are *not* appropriate for determining upper or lower limits.

Example with a Well-Behaved Parameter Space

sherpa> fit powll: v1.2

powll: initial function value = 8.22297E+01
powll: converged to minimum = 6.27050E+01 at iteration = 7

powll: final function value = 6.27050E+01

p.c0 56.2579 p.c1 0.11117 p.c2 -0.00119999

sherpa> uncertainty

Computed for uncertainty.sigma = 1

Parameter Name	Best-Fit	Lower Bound	Upper Bound
p.c0	56.2579	-0.865564	+0.864461
p.c1	0.11117	-0.0148228	+0.0148038
p.c2	-0.00119999	-0.000189496	+0.000189222

sherpa> projection

Computed for projection.sigma = 1

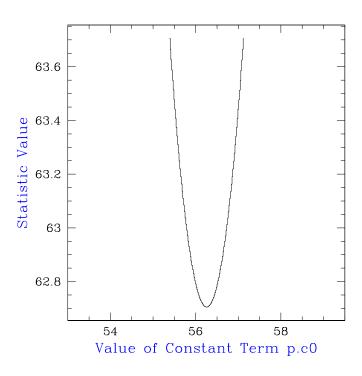
Parameter Name	Best-Fit	Lower Bound	Upper Bound
p.c0	000_000	-2.64465	+2.64497
p.c1	0.11117	-0.120684	+0.120703
p.c2	-0.00119999	-0.00115029	+0.00114976

sherpa> covariance

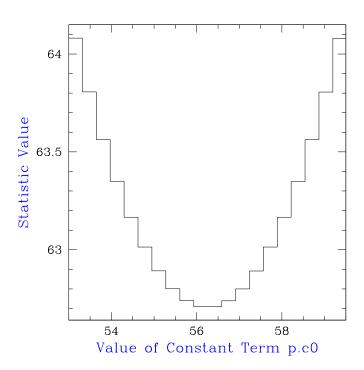
Computed for covariance.sigma = 1

Parameter Name	Best-Fit	Lower Bound	Upper Bound
p.c0	56.2579	-2.64786	+2.64786
p.c1	0.11117	-0.121023	+0.121023
p.c2	-0.00119999	-0.00115675	+0.00115675

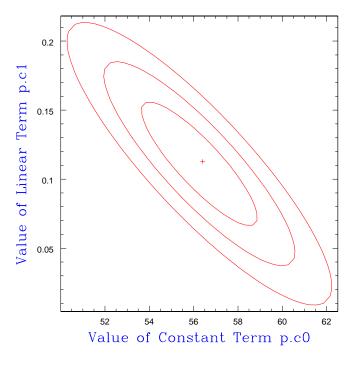
Interval - Uncertainty



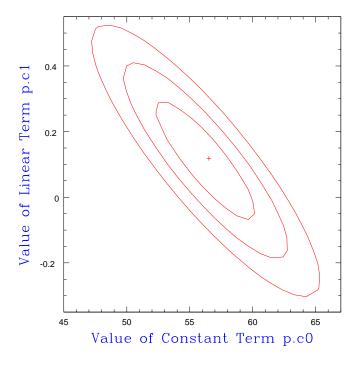
Interval - Projection



Confidence Region - Uncertainty



Confidence Region - Projection



Bayesian Parameter Estimation

A Bayesian estimates credible intervals and regions by marginalizing (integrating) the parameter posterior distribution over the space of nuisance (uninteresting) parameters. For instance:

$$p(\theta_1|D) = \int_{\theta_2} d\theta_2 \cdots \int_{\theta_P} d\theta_P \ p(\theta|D).$$

The central 68% of the distribution $p(\theta_1|D)$ is the 1σ credible interval.

Marginalization may be done by brute-force integration or, for higher dimensional problems ($N \ge 10$), by adaptive integration. However, if the statistical surface is "well-behaved," one can also estimate credible intervals using the Laplace Approximation:

$$p(\theta_1|D) = p(\hat{\theta}_2, \dots, \hat{\theta}_P)(2\pi)^{(P-1)/2} \times \sqrt{\det C(\theta_1, \hat{\theta}_2, \dots, \hat{\theta}_P)} \mathcal{L}(\theta_1, \hat{\theta}_2, \dots, \hat{\theta}_P).$$

If the value of parameter θ_1 is correlated with other parameter values, then when computing $p(\theta_1|D)$, the values of parameters $(\theta_2, \dots, \theta_P)$ should be allowed to *float* to new best-fit values.

Selected References

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