Physics 403 Numerical Methods,

Numerical Methods, Maximum Likelihood, and Least Squares

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Last Time

The quadratic approximation of the PDF in more than one dimension:

$$p(\boldsymbol{x}|D, I) \propto \exp\left[(\boldsymbol{x} - \hat{\boldsymbol{x}})^{\top} \boldsymbol{H}(\hat{\boldsymbol{x}})(\boldsymbol{x} - \hat{\boldsymbol{x}})
ight]$$

• The Hessian matrix $H(\hat{x})$ is an $N \times N$ symmetric matrix with components

$$H_{ij} = \frac{\partial^2 L}{\partial x_i \partial x_j} \bigg|_{\hat{x}_i, \hat{x}_j}$$

where

 $L = \ln p$

The covariance matrix Σ is related to the negative of the inverse Hessian matrix:

$$[\boldsymbol{\Sigma}]_{ij} = [-\boldsymbol{H}^{-1}]_{ij}$$

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Geometric Intuition



- The uncertainty contours (in 2D) define an ellipse whose principal axes are the eigenvectors of *H*
- When the covariance cov (x_i, x_j) = 0, the ellipse is aligned with x_i and x_j
- When the covariance is nonzero the ellipse is tilted. In this case, a rotation can remove the covariances; there exists an orthogonal matrix of the eigenvectors of *H* which diagonalizes the Hessian:

PHY 403

Geometric Intuition



▶ When working from a joint distribution p(x, y, ... |D, I), the uncertainty on the estimator x (for example) requires you to calculate

$$p(x|D, I) = \int p(x, y, z, \dots | D, I) \, dy \, dz \, \dots$$

- Remember that this is different from calculating the width of the joint distribution at the maximum
- ► The width of the contour at the maximum will underestimate the width of p(x|D, I)

Estimating μ if μ and σ are Unknown ${}^{\rm Student-{\it t}}$ Distribution

If we have Gaussian data with unknown μ and σ, the resulting marginal distribution for μ is

$$p(\mu|D,I) \propto \left[\sum_{i=1}^{N} (x_i - \mu)^2\right]^{-(N-1)/2}$$

if we use a uniform prior for σ . If we use a Jeffreys prior,

$$p(\mu|D,I) \propto \left[\sum_{i=1}^{N} (x_i - \mu)^2\right]^{-N/2}$$

The width estimator is the usual sample variance

$$s^2 = rac{1}{N-1}\sum_{i=1}^N (x_i - \hat{\mu})^2 = rac{1}{N-1}\sum_{i=1}^N (x_i - \bar{x})^2$$

for the uniform prior, and narrower $(\propto 1/N)$ if using Jeffreys prior

Estimating σ if μ and σ are Unknown $_{\chi^2 \mbox{ Distribution}}$

• If we have Gaussian data with unknown μ and σ , the resulting marginal distribution for σ is

$$p(\sigma|D, I) \propto \sigma^{-(N-1)} \exp\left(-\frac{V}{2\sigma^2}\right), \qquad V = \sum_{i=1}^N (x_i - \bar{x})^2$$

if we use a uniform prior for $\sigma.$ If we use a Jeffreys prior,

$$p(\sigma|D, I) \propto \sigma^{-N} \exp\left(-rac{V}{2\sigma^2}
ight)$$

• $\hat{\sigma}^2 = s^2$, and the reliability of the width estimator is

$$\sigma = \hat{\sigma} \pm \frac{\hat{\sigma}}{\sqrt{2(N-1)}}.$$

The marginal PDF is equivalent to the $\chi^2_{2(N-1)}$ distribution.

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Methods for Automatic Minimization

- Getting the best estimate of a PDF means calculating its maximum.
 Sometimes this cannot be done analytically
- Brute force approach: just plot the PDF on a grid of points and visually pick out the maximum
- Unfortunately, this becomes impractical as the dimensionality of the problem grows
- Issue 1: visualizing a maximum in more than 2D is hard
- Issue 2: computational expense. For a problem with N dimensions, evaluating 10 points on each axis requires 10^N calculations
- ► Issue 3: a regular grid could miss narrow features in the PDF
- So we need other methods to find the maximum of a function. Most popular methods linearize the problem

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Method of Steepest Descent



- How do we automatically minimize a multivariable function f(x), or maximize -f(x)?
- Steepest Descent: given a point a, f(x) decreases fastest in the direction

 $-\nabla f(\boldsymbol{a})$

• Start with a guess x_0 and update:

$$\mathbf{x}_{n+1} = \mathbf{x}_n - \gamma_n \nabla f(\mathbf{x}_n), \ n \ge 0$$

- Control the step size with γ_n
- Keep iterating until (hopefully) x_n converges to a local minimum

Method of Steepest Descent

Known Issues

- ► There are several known issues with the steepest descent algorithm
- For example, if the sequence steps into a "valley" along the minimum it can start zig-zagging along the walls



> This can make the algorithm quite slow as it approaches the minimum

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Method of Steepest Descent Behavior in the Valley

The figure below shows why the steepest descent algorithm oscillates back and forth when you enter a valley [1]



- A step starts off in the local gradient direction perpendicular to the contour lines
- The step traverses a straight line until a local minimum is reached, where the traverse is parallel to the local contour lines
- Next update is perpendicular to the last direction. Result: S-L-O-W

Quadratic Approximation

Suppose we Taylor-expand our function f(x) about some arbitrary point x', so that

$$f(\mathbf{x}) = f(\mathbf{x}') + (\mathbf{x} - \mathbf{x}')^{\top} \nabla f(\mathbf{x}') + \frac{1}{2} (\mathbf{x} - \mathbf{x}')^{\top} \nabla \nabla f(\mathbf{x}') (\mathbf{x} - \mathbf{x}') + \dots$$
$$\approx f(\mathbf{x}') + (\mathbf{x} - \mathbf{x}')^{\top} \nabla f(\mathbf{x}') + \frac{1}{2} (\mathbf{x} - \mathbf{x}')^{\top} \mathbf{H}(\mathbf{x}') (\mathbf{x} - \mathbf{x}')$$

where $H(\mathbf{x}') = \nabla \nabla f(\mathbf{x}')$ is the Hessian matrix of f

• Differentiating f with respect to the $\{x_i\}$ gives

$$abla f(\mathbf{x}) pprox
abla f(\mathbf{x}') + \mathbf{H}(\mathbf{x}')(\mathbf{x} - \mathbf{x}')$$

• If we demand $\nabla f(\hat{x}) = 0$, since we're at an extremum, we obtain

$$\hat{\boldsymbol{x}} \approx \boldsymbol{x}' - [\boldsymbol{H}(\boldsymbol{x}')]^{-1} \nabla f(\boldsymbol{x}')$$

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Newton's Method



This expression suggests an iterative scheme for approaching a minimum:

$$\mathbf{x}_{n+1} = \mathbf{x}_n - [\mathbf{H}(\mathbf{x}_n)]^{-1} \nabla f(\mathbf{x}_n), \quad n \ge 0$$

- Intuition: each iteration approximates
 f(x) by a quadratic function and takes
 a step toward the minimum of the function
- If f(x) is quadratic, the extremum will be found in exactly one step
- When the quadratic approximation is reasonable, this method will converge to the minimum much faster than the steepest descent algorithm

Newton's Method Computational Tricks

► The stability of the iterations can be improved by reducing the step size by some positive factor γ < 1:</p>

$$\mathbf{x}_{n+1} = \mathbf{x}_n - \gamma [\mathbf{H}(\mathbf{x}_n)]^{-1} \nabla f(\mathbf{x}_n), \quad n \ge 0$$

- ▶ Note: in *N* dimensions, inverting *H* takes $\mathcal{O}(N^3)$ operations
- ► Instead of inverting, calculate the vector $\boldsymbol{p}_n = [\boldsymbol{H}(\boldsymbol{x}_n)]^{-1} \nabla f(\boldsymbol{x}_n)$ as the solution to the system of linear equations

$$\boldsymbol{H}(\boldsymbol{x}_n) \cdot \boldsymbol{p}_n = \nabla f(\boldsymbol{x}_n)$$

- ▶ Methods to solve this equation, like the conjugate gradient (CG) technique [1], require u^T H(x_n)u > 0 for any real nonzero vector u.
- Jargon: the Hessian must be positive definite. This is a useful diagnostic, e.g., it tells you if the iteration converged to a saddle point

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Newton's Method

Known Issues

Because ∇f(x̂) = 0 is just the condition for a stationary point, Newton's method can diverge if x₀ is far from the optimal solution



- In the figure (left) we want to find the maximum of the PDF. It's roughly quadratic so Newton's method converges rapidly
- On the right, if we start out in the tails of the function (outside the dotted lines) the algorithm will not converge to the maximum
- Solution: start with a good first guess. Can use an algorithm that doesn't depend on the gradient, like simplex minimization

Definition of a Simplex

- ► A simplex is basically a "hyper-triangle" in *n* dimensions.
- E.g., the *n*-simplex Δ^n is the subset of \mathbb{R}^{n+1} such that

$$\Delta^n = \{(t_0, \cdots, t_n) \in \mathbb{R}^{n+1} \mid \sum_{i=0}^n t_i = 1$$

and $t_i \ge 0$ for all $i\}$

Simplex/Nelder-Mead Technique [2]: start with N + 1 points p₀ and p_i (i = 1...N) such that

$$\boldsymbol{p}_i = \boldsymbol{p}_0 + \lambda \boldsymbol{e}_i$$

The points define a simplex for your N-dimensional parameter space. Try to move the simplex around and shrink/expand it until it contains the optimal point

Downhill Simplex (Nelder-Mead) Algorithm



- Define the starting point for the simplex
- Pick out the point in the simplex where f(x) is largest
- Reflect this point through the opposite face of the simplex to a lower point
- Shrink or expand the simplex to conserve its volume
- The simplex will crawl, amoeba-like, toward the minimum
- Advantage: no need to calculate the gradient. Use result as a starting point for Newton's method
- Disadvantage: convergence issues if initial simplex is too small

Difficult Problem: Multimodal Parameter Space

 Often you'll find that your parameter space is complex, with multiple minima and maxima



- The algorithms we have discussed so far will run as quickly as possible to the nearest minimum
- There is no way for you to guarantee that you have gotten to the global minimum rather than a local minimum

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Simulated Annealing

Starting from x_n , randomly generate a new point

$$\mathbf{x}_{n+1} = \mathbf{x}_n + \Delta \mathbf{x}$$

Calculate a probability

$$p = \exp\left\{-\frac{f(\mathbf{x}_{n+1}) - f(\mathbf{x}_n)}{kT}\right\}$$

for keeping the point, and generate a random number $u \in [0, 1]$. If u < p, move to x_{n+1} . Otherwise, stay at x_n .

- For large T, the probability of accepting new points (even "bad" moves) is high. For small T, the probability to accept new points is low
- Idea: start with a high T to help you jump out of local minima, then slowly reduce the temperature. Slow cooling helps you find the global minimum energy state, like annealing a piece of metal [3]

Markov Chain Monte Carlo

- The technique of choosing p to sample states of a thermodynamic system is called the Metropolis-Hastings algorithm [4]
- Simulated annealing depends on an annealing schedule for moving T → 0, which you have to tune. Also, there is no guarantee of convergence to the global minimum in a finite time
- Another approach: run a large number of simulations at different temperatures, letting each one randomly walk through the parameter space
- This technique is called Markov Chain Monte Carlo (MCMC), and can be used to simulate exploration of all important parts of a parameter space
- MCMC methods have become central to Bayesian analysis. We'll talk about how and why in a couple of weeks

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Popular Libraries

scipy.optimize

scipy.optimize.minimize

scipy.optimize.minimize(fun, x0, args=(), method=None, jac=None, hess=None, hessp=None, bounds=None, constraints=(), tol=None, callback=None, options=None) [source] Minimization of scalar function of one or more variables New in version 0.11.0. Parameters: fun : callable Objective function. x0 : ndarray Initial guess. args : tuple, optional Extra arguments passed to the objective function and its derivatives (Jacobian, Hessian). method : str or callable, optional Type of solver. Should be one of 'Nelder-Mead' 'Powell' 'CG' • 'BFGS' 'Newton-CG' 'Anneal (deprecated as of scipy version 0.14.0)' 'L-BEGS-B' 'TNC' 'COBYLA' 'SLSOP' 'dogleg' 'trust-ncg' custom - a callable object (added in version 0.14.0) If not given, chosen to be one of BFGS, L-BFGS-B, SLSQP, depending if the problem has

constraints or bounds.

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Popular Libraries



ROOT has a C++ version of the "popular" MINUIT non-linear function minimizer. Three minimization algorithms are available:

- 1. Steepest descent (MIGRAD): evaluates gradient and second derivatives (Hessian) numerically. Assumes symmetric Gaussian errors
- 2. MINOS: relaxes error assumption, allows asymmetric error bars
- 3. Simplex: does not require evaluation of derivatives

If you've ever used this before, you know it requires a lot of hand-tuning. The going gets very rough in high-D if the parameter space is bumpy

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Maximum Likelihood Technique

- The method of maximum likelihood is an extremely important technique used in frequentist statistics
- There is no mystery to it. Here is the connection to the Bayesian view: given parameters x and data D, Bayes' Theorem tells us that

$$p(\boldsymbol{x}|\boldsymbol{D},l) \propto p(\boldsymbol{D}|\boldsymbol{x},l) \ p(\boldsymbol{x}|l)$$

where we ignore the marginal evidence p(D|I)

Suppose $p(\mathbf{x}|I) = \text{constant for all } \mathbf{x}$. Then

 $p(\boldsymbol{x}|\boldsymbol{D},l) \propto p(\boldsymbol{D}|\boldsymbol{x},l)$

and the best estimator \hat{x} is simply the value that maximizes the likelihood p(D|x, l)

So the method of maximum likelihood for a frequentist is equivalent to maximizing the posterior p(x|D, l) with uniform priors on the {x_i}.

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Connection to χ^2

Suppose our data *D* are identical independent measurements with Gaussian uncertainties. Then the likelihood is

$$p(D_i|\mathbf{x}, I) = rac{1}{\sqrt{2\pi\sigma_i}} \exp\left[-rac{(F_i - D_i)^2}{2\sigma_i^2}
ight], \quad p(\mathbf{D}|\mathbf{x}, I) = \prod_{i=1}^N p(D_i|\mathbf{x}, I),$$

where we defined the functional relationship between \boldsymbol{x} and the ideal (noiseless) data \boldsymbol{F} as

$$F_i = f(\mathbf{x}, i)$$

• If we define χ^2 as the sum of the squares of the normalized residuals $(F_i - D_i)/\sigma_i$, then

$$\chi^2 = \sum_{i=1}^{N} \frac{(F_i - D_i)^2}{\sigma_i^2} \implies p(\boldsymbol{D}|\boldsymbol{x}, I) \propto \exp\left(-\frac{\chi^2}{2}\right)$$

Maximum Likelihood and Least Squares

▶ With a uniform prior on *x*, the logarithm of the posterior PDF is

$$L = \ln p(\mathbf{x}|\mathbf{D}, I) = \text{constant} - \frac{\chi^2}{2}$$

- The maximum of the posterior (and likelihood) will occur when χ² is a minimum. Hence, the optimal solution x̂ is called the least squares estimate
- Least squares/maximum likelihood is used all the time in data analysis, but...
- Note: there is nothing mysterious or even fundamental about this; least squares is what Bayes' Theorem reduces to if:
 - 1. Your prior on your parameters is uniform
 - 2. The uncertainties on your data are Gaussian
- If one of these conditions isn't met, then use Bayes' Theorem to derive something else

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Maximum Likelihood: Poisson Case

Suppose that our data aren't Gaussian, but a set of Poisson counts n with expectation values v. E.g., we are dealing with binned data in a histogram. Then the likelihood becomes

$$p(\boldsymbol{n}|\boldsymbol{\nu}, l) = \prod_{i=1}^{N} \frac{\nu_i^{n_i} e^{-\nu_i}}{n_i!}$$

• In the limit $N \rightarrow$ large, this becomes

$$p(n_i|\nu_i, I) \propto \exp\left[-\sum_{i=1}^N \frac{(n_i - \nu_i)^2}{2\nu_i}
ight]$$

 \blacktriangleright The corresponding χ^2 statistic is given by

$$\chi^2 = \sum_{i=1}^{N} \frac{(n_i - \nu_i)^2}{\nu_i}$$

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Justifications for Using Least Squares

 \blacktriangleright Nice property: as ${\it N} \to \infty,$ the χ^2 statistic asymptotically approaches the value

 $\chi^2_{\rm M}$ m

where N is the number of data points and m is the number of parameters in x.

- ► I.e., the statistic approximates a \(\chi^2\) distribution with \(N m\) degrees of freedom... if the uncertainties in the data are Gaussian
- ► Our definition of \(\chi^2\) as the quadrature sum (or \(\lambda_2\)-norm) of the residuals makes a lot of calculations easy, but it isn't particularly robust
- ► Note: the *I*₁-norm

$$l_{1}$$
-norm = $\sum_{i=1}^{N} \left| \frac{F_{i} - D_{i}}{\sigma_{i}} \right|$

is much more robust against outliers in the data

Application: Fitting a Straight Line to Data

Example

Suppose we have N measurements y_i with Gaussian uncertainties σ_i measured at positions x_i .



Given the straight line model $y_i = mx_i + b$, what are the best estimators of the parameters *m* and *b*?

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Minimize the χ^2

Letting $F_i = mx_i + b$ and $D_i = y_i$, the χ^2 is

$$\chi^{2} = \sum_{i=1}^{N} \frac{(mx_{i} + b - y_{i})2}{\sigma_{i}^{2}}$$

Minimizing χ^2 as a function of the parameters gives

$$\frac{\partial \chi^2}{\partial m} = \sum_{i=1}^N \frac{2(mx_i + b - y_i)x_i}{\sigma_i^2} \quad \text{and} \quad \frac{\partial \chi^2}{\partial b} = \sum_{i=1}^N \frac{2(mx_i + b - y_i)}{\sigma_i^2}$$

Rewritten as a matrix equation, this becomes

$$\nabla \chi^2 = \begin{pmatrix} A & C \\ C & B \end{pmatrix} \begin{pmatrix} m \\ b \end{pmatrix} - \begin{pmatrix} p \\ q \end{pmatrix} = 0$$
$$A = \sum x_i^2 w_i, \ B = \sum w_i, \ C = \sum x_i w_i, \ p = \sum x_i y_i w_i, \ q = \sum y_i w_i$$

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Best Estimators of a Linear Function

Inverting the matrix, we find that

$$\hat{m} = rac{Bp-Cq}{AB-C^2}$$
 and $\hat{b} = rac{Aq-Cp}{AB-C^2}$

• The covariance matrix is found by evaluating $[2\nabla \nabla \chi^2]^{-1}$:

$$\begin{pmatrix} \sigma_m^2 & \sigma_{mb} \\ \sigma_{mb} & \sigma_b^2 \end{pmatrix} = 2 \begin{pmatrix} A & C \\ C & B \end{pmatrix}^{-1} = \frac{2}{AB - C^2} \begin{pmatrix} B & -C \\ -C & A \end{pmatrix}$$

- ► We note that even though the data {y_i} are independent, the parameters m̂ and b̂ end up anticorrelated due to the off-diagonal terms in the covariance matrix
- This makes a lot of sense, actually; wiggling the slope of the line m clearly changes the y-intercept b

Summary

- Various algorithms available (simplex, Newton, etc.) with trade offs between speed and accuracy
- All algorithms are sensitive, to some degree or another, to the starting position of the minimization
- Maximum likelihood: same as maximizing a posterior PDF when the priors on the parameters are uniform
- Maximizing the likelihood is the same as minimizing χ^2 in the case where the uncertainties on the data are Gaussian
- In case of Gaussian uncertainties, there is asymptotic convergence of the maximum likelihood to the χ² distribution:

$$\chi^2 = -2\ln L \sim \chi^2_{N-m}$$

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