Why are Nonlinear Fits to Data so Challenging?

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Fitting model parameters to experimental data is a common yet often challenging task, especially if the model contains many parameters. Typically, algorithms get lost in regions of parameter space in which the model is unresponsive to changes in parameters, and one is left to make adjustments by hand. We explain this difficulty by interpreting the fitting process as a generalized interpolation procedure. By considering the manifold of all model predictions in data space, we find that cross sections have a hierarchy of widths and are typically very narrow. Algorithms become stuck as they move near the boundaries. We observe that the model manifold, in addition to being tightly bounded, has low extrinsic curvature, leading to the use of geodesics in the fitting process. We improve the convergence of the Levenberg-Marquardt algorithm by adding geodesic acceleration to the usual step.

The estimation of model parameters from experimental data is astonishingly challenging. A nonlinear model with tens of parameters, fit (say) by least squares to experimental data, often demands weeks of human guidance to find a good starting point; even then, the parameters cannot usually be extracted from the data. Both general minimization algorithms and algorithms like the Levenberg-Marquardt algorithm that are designed for least-squares fits routinely get lost in parameter space. This becomes a serious obstacle to progress when one is unsure of the validity of the model, e.g., in systems biology where one wants to automatically generate and explore a variety of alternative models.

Here we use differential geometry to explain why fits are so hard. We first explore the structure of the model manifold $M$, the manifold of predictions embedded in the space of data $D$, and find that it is typically bounded, with cross sections having a hierarchy of widths, so that the overall structure is similar to that of a long, thin ribbon. We explain this hierarchy by viewing the fitting process as a generalized interpolation procedure with few effective model degrees of freedom. We interpret the difficulty in fitting to be due to algorithms getting stuck near the boundary of $M$, where the model is unresponsive to variations in the parameters. We then discuss how geometry motivates algorithms to alleviate this difficulty.

A typical nonlinear least-squares problem fits a model $Y_m(\theta)$ with $N$ parameters $\theta$ to $M$ experimental data points $Y_m$. We define the model manifold $M$ as the parametrized $N$-dimensional surface $\bar{Y}(\theta)$ embedded in Euclidean data space, $D = \mathbb{R}^M$. The best fit to the experiment is given by the point on $M$ with Euclidean distance closest to the data, minimizing the cost $C = \frac{1}{2} (\bar{Y}(\theta) - \bar{y})^2$. The Euclidean metric of data space [with distance between models given by the change in residuals, $\bar{r} = \bar{Y}(\theta) - \bar{y}$] induces a metric on the manifold, $g_{\mu \nu} = \partial_{\mu} \bar{Y} \cdot \partial_{\nu} \bar{Y} = (J^T J)_{\mu \nu}$, where $J_{\mu \nu} = \frac{\partial}{\partial \theta^\nu} Y_m$; $g_{\mu \nu}$ is known as the Fisher information matrix. As an example, the model $Y(t, \theta) = f_0(t) = e^{-\theta_1 t} + e^{-\theta_2 t}$ sampled at three time points is given in Fig. 1. The model manifold has been extensively studied by the information geometry statistics community [1], but they focus on the intrinsic curvature; as the cost is the distance in data space, the embedding and its extrinsic curvature are crucial to finding best fits [2,3].

As seen in Figs. 2 and 3, this model manifold can take the form of a hyper-ribbon, with thinnest direction 4 orders of magnitude smaller than the long axes. To understand this observed hierarchy, consider the special case of analytic models, $f(t, \theta)$, of a single independent variable (time) where the data points are $Y_m = f(t_m)$. Let $R$ be the typical time scale over which the model behavior changes, so that the $n$th term of the Taylor series $f^{(n)}(t)/n! \leq R^{-n}$ (roughly the radius of convergence). If the function is sampled at $n$ time point $(t_1, t_2, \ldots, t_n)$ within this time scale, the Taylor series may be approximated by the unique polynomial of degree $n - 1$, $P_{n-1}(t)$ passing through these points. At a new point, $t_0$, the discrepancy between the interpolation and the function is given by

$$f(t_0) - P_{n-1}(t_0) = \omega_n(t_0) f^{(n)}(\xi)/n!,$$

where $\omega_n(t_0)$ is the $(n+1)$th derivative of the $n$th degree polynomial. (The ribbon-like structure of Fig. 2 emerges only in higher dimensions.)
where $\xi$ lies somewhere in the interval containing $t_0, t_1, \ldots, t_n$ [4]. The polynomial $\omega_n(t)$ has roots at each of the interpolating points $\omega_n(t) = (t - t_1) (t - t_2) \cdots (t - t_n)$. The possible error of the interpolation function bounds the allowed range of behavior, $\Delta f_n$, of the model at $t_0$ after constraining the nearby $n$ data points (i.e., cross sections). Consider the ratio of successive cross sections, $\frac{\Delta f_{n+1}}{\Delta f_n} = (t - t_{n+1}) (n + 1) \frac{\omega_{n+1}(t)}{\omega_n(t)}$. If $n$ is sufficiently large, then $(n + 1) \frac{\omega_{n+1}(t)}{\omega_n(t)} = \frac{1}{R}$; therefore, we find that $\frac{\Delta f_{n+1}}{\Delta f_n} = \frac{t - t_{n+1}}{R} < 1$ by the ratio test. Each cross section is thinner than the last by a roughly constant factor, yielding the observed hierarchy.

We argue that this hyper-ribbon structure will be shared with a wide variety of nonlinear, multiparameter models. Note that the eigenvalues of the metric tensor $g_{\mu \nu}$ in Fig. 3 also form a hierarchy, spanning 8 orders of magnitude—this “sloppiness” has been documented in a number of other models, including 17 in systems biology [5], insect flight and variational quantum wave functions [6], interatomic potentials [7], and a model of the next-generation international linear collider [8]. (Multiparameter models whose parameters are individually measured by the data, and on the other extreme models with sensitive, chaotic dependence on parameters, will likely not fall into this family.) Most parameters in these models have bounded effects—they can be set to limiting values (zero, infinity, etc.) and still have finite model predictions—rates and Michaelis-Menten coefficients in systems biology, Jastrow and determinental factors in variational wave functions, etc. Note that the widths of the model manifold track nicely with these eigenvalues in Fig. 3 (taking the square root to match units): moving parameter combinations along eigendirections of the metric by an order of magnitude (fixed shift in log parameters) exhausts the range of behavior (width). This tracking suggests that the ubiquity of sloppy eigenvalue spectra at best fits implies a ubiquitous hyper-ribbon structure for the model manifold. Our observation that many models are sloppy, presumably sharing this hyper-ribbon model manifold structure, is now explained: multiparameter models are a kind of high-dimensional analytic interpolation scheme, and near degenerate Hessians result whenever multiple data points reside within some generalized radius of convergence. When this is the case, the data points are highly correlated and the model has few effective degrees of freedom. Whenever there are many model parameters for each effective degree of freedom there will be a hierarchy of widths and the model will be sloppy.

Our geometric interpretation explains a number of observations about nonlinear models. First, although parameters cannot be reliably extracted by fitting degenerate models, it is still possible to constrain the outcome of new experiments [5,9]. Because the fitting process is an interpolation scheme, only a few stiff parameter combinations need to be tuned to fit most of the data, since only a few data points constrain the predictions at other times. The remaining unconstrained parameter combinations control the interpolated values, which are already restricted by the analyticity of the model.

Figure 3 also shows that the parameter-effects curvature [2,3,10] and the geodesic extrinsic curvatures vary over twice as many decades as the widths and $\sqrt{\lambda}$; indeed, their formulas include a factor of $1/\lambda$ [11]. Why is the extrinsic curvature so much smaller? The manifold has zero extrinsic curvature if there are equal numbers of parameters as data points, $N = M$ (where the model manifold is a sub-
We use geodesics to construct new polar coordinates \( \gamma^\mu = \gamma^\mu(\theta) \) on \( \mathcal{M} \) that generalize Riemann normal coordinates \([13]\). Since geodesics are nearly straight lines in data space on \( \mathcal{M} \), we find that cost contours in these coordinates are nearly quadratic and isotropic around the best fit, as explicitly computed in Fig. 4. Nonlinear models locally look like linear models with badly chosen parameters, well beyond the harmonic approximation.

Can these nearly straight geodesics inspire algorithms that lead efficiently to the best fit? Integrating the geodesic equation with a single Euler step reproduces the Gauss-Newton step \((\delta \theta^\mu = -g^{\mu\nu}\nabla_\nu C \) in our notation), ineffective due to the large eigenvalues in the inverse metric \(g^{\mu\nu}\); even geodesic motion hits the boundaries of \( \mathcal{M} \). The ribbon is nearly flat, but very thin; the geodesics hit the edges long before finding a good fit.

To improve convergence, we can modify the model manifold to remove the boundaries. One method of doing this is to introduce the model graph \( \mathcal{G} \), which is the \( N \)-dimensional parametric surface drawn by the model embedded in data space crossed with parameter space. Since most boundaries occur at infinite parameter values, the model graph \( \mathcal{G} \) "stretches" these boundaries to infinity in the parameter space portion of the embedding. The metric for the model graph is an interpolation of the data in the parameter space portion of the embedding. The analogy of the Gauss-Newton step on the model graph is the well-known Levenberg-Marquardt step, \( \delta \theta = -(J^T J + \lambda^* I)^{-1} J^T C \) in our notation \([14\text{-}16]\). By dynamically adjusting \( \lambda^* \), the algorithm can shorten its step, removing the danger of the degenerate Hessian, while rotating from the Gauss-Newton direction into the steepest descents direction. Geometrically we understand the superiority of the Levenberg-Marquardt algorithm to be due to the lack of boundaries of the model graph.

Inspired by the results in Fig. 4, we further improve the standard Levenberg-Marquardt algorithm. Interpreting the Levenberg-Marquardt step as a velocity, \( v^\mu = -g^{\mu\nu}\nabla_\nu C \), where \( g \) is the metric on the model graph, the geodesic acceleration (giving the parameter-effects curvature) is given by \( a^\mu = -g^{\mu\nu}\partial_\nu \gamma \cdot \partial_\alpha \partial_\beta \gamma^\alpha u^\beta \), giving a step \( \delta \theta^\mu = v^\mu + \frac{1}{2} a^\mu \). The geodesic acceleration is very cheap to calculate, requiring only a directional second derivative, which can be estimated from three (cheap) function evaluations (one or two additional function evaluations) at each step with no extra (expensive) Jacobians. The geodesic acceleration serves two purposes. First, it provides an estimate for the trust region in which the linearization approximation (from which the Levenberg-Marquardt algorithm is traditionally derived) is valid. At each step, we adjust \( \lambda \) until the acceleration is smaller than the velocity, which we find is more effective at avoiding model boundaries than either tuning until a downhill step is found \([14,16]\) or considering the reduction ratio \([15]\). The second benefit

![FIG. 4 (color online). Geodesics can be used to construct polar coordinates on \( \mathcal{M} \) (top). In these new geodesic coordinates, the cost contours are nearly perfect, isotropic circles (bottom).](image_url)
of the acceleration occurs when the algorithm must follow a long narrow canyon to the best fit. In these scenarios convergence may be sped up by approximating the path with a parabola instead of a line. The utility of the geodesic acceleration is seen in Table I, where the performance of several algorithms on a test problem is summarized. More extensive comparisons and further refined algorithms are in preparation [19].

Just as the special sum of squares form of the cost function gives an approximate Hessian using only first derivatives of the residuals, \( H = J^T J \), it has (together with the low extrinsic curvature) allowed us to approximate the \textit{cubic} correction using only a directional second derivative. All other algorithms that seek to improve the Levenberg-Marquardt algorithm use second derivative information only to calculate the correction \( \delta H_{\mu \nu} = \tilde{r} \cdot \partial_{\mu} \partial_{\nu} \tilde{r} \) to the Hessian [20–23]. This correction is negligible if the nonlinearities are primarily parameter-effects curvature; since the unfit data are nearly perpendicular to the surface of the model manifold while the nonlinearities are tangent to the model manifold, the dot product vanishes. Qualitatively, this means that the approximate Hessian is very accurate and that the bending of the local ellipses (due to the third order terms in the cost that we consider) is the most important correction.

By interpreting the fitting process as a generalized interpolation scheme, we have seen that the difficulties in fitting are due to the narrow boundaries on the model manifold \( \mathcal{M} \). These boundaries form a hierarchy of widths dual to the hierarchy of Hessian eigenvalues characteristic of nonlinear model fits. Additionally, we both observe and argue that the model manifold is remarkably flat (low extrinsic curvature), which leads us to the use of geodesics in the fitting process. The modified Gauss-Newton and Levenberg-Marquardt algorithms are understood to be Euler approximations to the geodesic equation on the model manifold and model graph, respectively. The geodesic acceleration improves convergence of the Levenberg-Marquardt algorithm by providing a more accurate trust region while reducing computation time. Data fits are both practically important and theoretically elegant.

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[11] K = \|P \partial_\mu \partial_{\nu} \tilde{v} \|^2 / \|v^\alpha \tilde{r} \| R_{\alpha \beta} \delta_{\beta \mu}, where \( P \) is a projection operator projecting out of the tangent space for extrinsic curvature and into the tangent space for parameter-effects curvature.
[12] If we choose \( N \) independent data points as our parametrization, then the interpolating polynomial \( P_{N-1}(t) \) in Eq. (1) is a linear function of the parameters. As discussed below that equation, the manifold in each additional direction will be constrained to within \( \epsilon = \Delta f_{N+1} \) of \( P_{N-1}(t) \). Presuming that this deviation from flatness smoothly varies along the \( j \)th largest width \( W_j \sim \Delta f_j \) of the manifold (i.e., no complex or sensitive dependence on parameters), the geodesic extrinsic curvature is \( \epsilon / W_j^2 \), explaining the range of curvatures in Fig. 3. The ratio of the curvature to the inverse width should then be \( \epsilon / W_j \sim \Delta f_{N+1} / \Delta f_j \sim (\Delta t / R)^{N+1-j} \), comparable to the widths along the sloppiest direction and with twice the slope. It appears that the scale of the parameter-effects curvature is set by the stiffest width (Fig. 3), leading to the 3 order of magnitude difference between the two curvatures.