

CS 4220: Numerical Analysis

Derivative free methods

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Lay of the land

In the landscape of continuous optimization problems, there are three ways for things to be hard:

- Lack of smoothness
- Lack of convexity (or other structure guaranteeing unique global minimizers)
- Lack of information about the function

So far, we have only considered difficulties associated with lack of convexity, and those only superficially – most of our algorithms only give local minimizers, and we haven't talked about finding global optima. We have assumed everything has as many derivatives as we would like, and that we are at least able to compute gradients of our objective.

Today we discuss optimization in one of the hard cases. For this class, we will not deal with the case of problems with very hard global structure, other than to say that this is a land where heuristic methods (simulated annealing, genetic algorithms, and company) may make sense. But there are some useful methods that are available for problems where the global structure is not so hard as to demand heuristics, but the problems are hard in that they are “black box” – that is, we are limited in what we can compute to looking at function evaluations.

Before describing some methods, I make two pleas.

First, consider these only after having thoughtfully weighed the pros and cons of gradient-based methods. If the calculus involved in computing the derivatives is too painful, consider a computer algebra system, or look into a tool for automatic differentiation of computer programs. Alternately, consider whether there are numerical estimates of the gradient (via finite differences) that can be computed more quickly than one might expect by taking advantage of the structure of how the function depends on variables. But if you really have to work with a black box code, or if the pain of computing derivatives (even with a tool) is too great, a gradient-free approach may be for you.

Second, do not fall into the trap of thinking these methods should be simple to implement. Nelder-Mead is perhaps simple to implement, which is one reason why it remains so popular; it also fails to converge on many examples, and rarely converges fast. There are various pattern search methods and model-based methods with more robust convergence or better convergence rates, and there exist good implementations in the world (though Powell himself has passed, his [PDFO suite](#) is still going strong!). And there are much more thorough texts and reviews than this set of lecture notes; a few I like include:

- [Powell, “Direct search algorithms for optimization calculations”](#) – covers a lot of methods, and particularly Powell’s algorithms
- [Powell, “A view of algorithms for optimization without derivatives”](#) – played a big role in this presentation
- [Kolda, Lewis, and Torczon, “Optimization by direct search: new perspectives on some classical and modern methods”](#) – deals with one interesting family of methods (very thoroughly)
- [Conn, Scheinberg, and Vicente, “Introduction to Derivative-Free Optimization”](#) – available for access via the Cornell library subscription to SIAM ebooks
- [Audet and Warren, “Derivative-Free and Blackbox Optimization”](#) – a Springer textbook, but also available courtesy the Cornell library
- [Larson, Menickelly, and Wild, “Derivative-free optimization methods”](#) – a recent Acta Numerica survey, again by people who know what they are doing

Model-based methods

The idea behind Newton’s method is to successively minimize a quadratic *model* of the function behavior based on a second-order Taylor expansion about the most recent guess, i.e. $x^{k+1} = x^k + p$ where

$$\operatorname{argmin}_p \phi(x) + \phi'(x)p + \frac{1}{2}p^T H(x)p.$$

In some Newton-like methods, we use a more approximate model, usually replacing the Hessian with something simpler to compute and factor. In simple gradient-descent methods, we might fall all the way back to a linear model, though in that case we cannot minimize the model globally – we need some other way of controlling step lengths. We can also explicitly incorporate our understanding of the quality of the model by specifying a constraint that keeps us from moving outside a “trust region” where we trust the model to be useful.

In derivative-free methods, we will keep the basic “minimize the model” approach, but we will use models based on interpolation (or regression) in place of the Taylor expansions of the Newton approach. There are several variants.

Finite difference derivatives

Perhaps the simplest gradient-free approach (though not necessarily the most efficient) takes some existing gradient-based approach and replaces gradients with finite difference approximations. There are two difficulties with this approach:

- If $\phi : \mathbb{R}^n \rightarrow \mathbb{R}$, then computing the $\nabla\phi(x)$ by finite differences involves at least $n + 1$ function evaluations. Thus the typical cost per step ends up being $n + 1$ function evaluations (or more), where methods that are more explicitly designed to live off samples might only use a single function evaluation per step.
- The finite difference approximations depends on a step size h , and their accuracy is a complex function of h . For h too small, the error is dominated by cancellation, revealing roundoff error in the numerical function evaluations. For h large, the error depends on both the step size and the local smoothness of the function.

The first issue (requiring $n + 1$ function evaluations to get the same information content as one function-and-gradient evaluation) is not unique to finite difference computations, and indeed tends to be a limit to a lot of derivative-free methods.

Linear models

A method based on finite difference approximations of gradients might use $n + 1$ function evaluations per step: one to compute a value at some new point, and n more in a local neighborhood to compute values to estimate derivatives. An alternative is to come up with an approximate linear model for the function using $n + 1$ function evaluations that may include some “far away” function evaluations from previous steps.

We insist that the $n + 1$ evaluations form a simplex with nonzero volume; that is, to compute from evaluations at points x_0, \dots, x_n , we want $\{x_j - x_0\}_{j=1}^n$ to be linearly independent vectors. In that case, we can build a model $x \mapsto b^T x + c$ where $b \in \mathbb{R}^n$ and $c \in \mathbb{R}$ are chosen so that the model interpolates the function values. Then, based on this model, we choose a new point.

The following routine computes a simplex estimate $\widehat{\nabla}f(x_0)$ of the gradient $\nabla f(x)$ based on the equations

$$f(x_i) = f(x_0) + (x_i - x_0)^T \widehat{\nabla}f(x_0),$$

Our gradient estimator therefore has the form

$$A \widehat{\nabla}f(x_0) = y, \quad A = \begin{bmatrix} (x_1 - x_0)^T \\ \vdots \\ (x_n - x_0)^T \end{bmatrix}, \quad y = \begin{bmatrix} f(x_1) - f(x_0) \\ \vdots \\ f(x_n) - f(x_0) \end{bmatrix}.$$

Note that if we adapt the simplex by changing one point at a time, we can update the factorization of A in $O(d^2)$ time rather than recomputing every time at a cost of $O(d^3)$. We do not bother to do this here, though.

```

function simplex_gradient(xs, fxs)
    d, npl = size(xs)
    A = (xs[:,2:end] .- xs[:,1])'
    ∇f_s = A \ (fxs[2:end] .- fxs[1])
    ∇f_s, cond(A)
end

```

The true gradient satisfies

$$A\nabla f(x_0) = y + r;$$

where the vector r consist of remainder terms from Taylor's theorem with remainder. If the gradient is Lipschitz with constant M , then the terms satisfy

$$r_i \leq M\|x_i - x_0\|^2/2 \leq Md^2/2$$

where d is the diameter of the simplex. Therefore, we expect

$$\|\hat{\nabla}f(x_0) - \nabla f(x_0)\| \leq \|A^{-1}r\| \leq \|(A/d)^{-1}\|Md/2$$

where $\|(A/d)^{-1}\|$ is scaled so that only the geometry of the simplex matters. Alternately, we have the relative error bound

$$\frac{\|\hat{\nabla}f(x_0) - \nabla f(x_0)\|}{\|\nabla f(x_0)\|} \leq \kappa(A) \frac{Md}{2\|y\|/d}$$

However we write the bounds, a key aspect to these methods is ensuring that the computation of the affine function from the simplex remains well-conditioned.

We give an example of the effects of geometry below by using a simplex gradient approximation of a 2D function where we make the simplex closer and closer to degenerate. As we get closer to degenerate, the error in the gradient estimate increases (Figure 1).

Getting a good gradient estimate gives a sense of which way to go, but in order to get good convergence we also need a globalization strategy, which gets more complicated. For example, line search with sufficient decrease can fail because of an inaccurate gradient estimate, in which case we may need to reconstruct a new simplex with a good geometry and a small diameter. There are also trust-region methods that use linear approximations based on interpolation over a simplex. One of the most popular of this family of method is Powell's COBYLA algorithm (Constrained Optimization BY Linear Approximation).

Quadratic models

One can build quadratic models of a function from only function values, but to fit a quadratic model in n -dimensional space, we usually need $(n+2)(n+1)/2$ function evaluations – one for each of the $n(n+1)/2$ distinct second partials, and $n+1$ for the linear part. Hence, purely

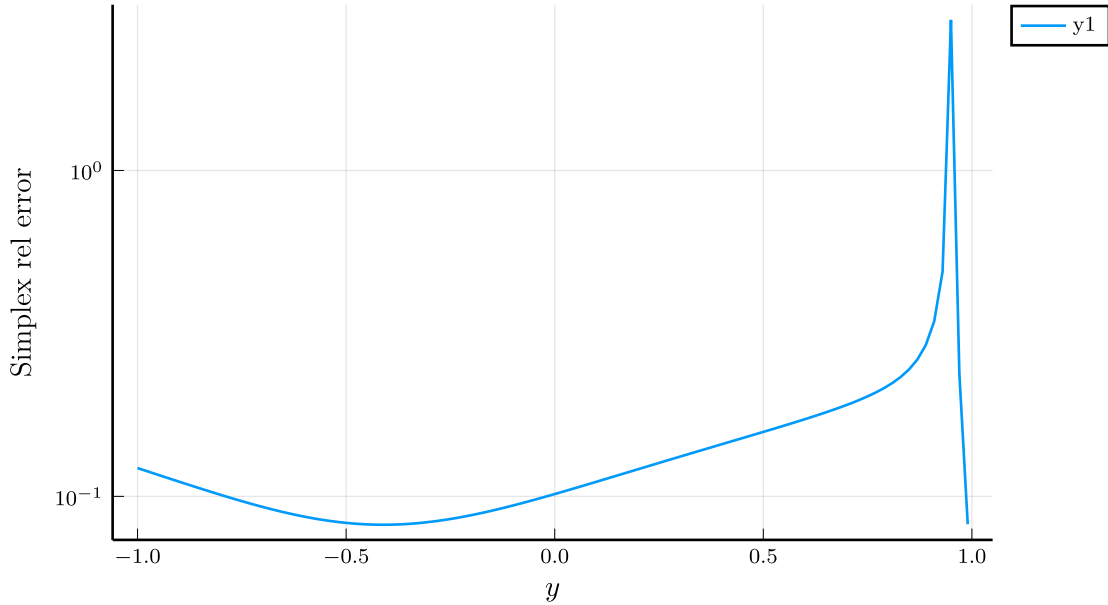


Figure 1: Simplex gradient estimate error. As y approaches 1, the simplex approaches degeneracy (and the linear system we have to solve approaches singularity).

function-based methods that use quadratic models tend to be limited to low-dimensional spaces. However, there are exceptions. The NEWUOA method (again by Powell) uses $2n + 1$ samples to build a quadratic model of the function with a diagonal matrix at second order, and then updates that matrix on successive steps in a Broyden-like way.

Surrogates and response surfaces

Polynomial approximations are useful, but they are far from the only methods for approximating objective functions in high-dimensional spaces. One popular approach is to use *kernel-based approximations*; for example, we might write a model

$$s(x) = \sum_{j=1}^m c_j \phi(\|x - x_j\|)$$

where the coefficients c_j are chosen to satisfy m interpolation conditions at points x_1, \dots, x_m . If we add a polynomial term, you'll recognize this as the form of the interpolants from project 1, but the spline interpretation is only one way of thinking about this class of approximators. Another option is to interpret this as a Gaussian process model; this is used, for example, in most Bayesian Optimization (BO) methods. There are a variety of other surfaces one might consider, as well.

In addition to fitting a surface that interpolates known function values, there are also methods that use *regression* to fit some set of known function values in a least squares sense. This is particularly useful when the function values have noise.

Pattern search and simplex

So far, the methods we have described are explicit in building a model that approximates the function. However, there are also methods that use a systematic search procedure in which a model does not explicitly appear. These sometimes go under the heading of “direct search” methods.

Nelder-Mead

The Nelder-Mead algorithm is one of the most popular derivative-free optimizers around. For example, this is the default algorithm used for derivative free optimization with `Optim.jl`. As with methods like COBYLA, the Nelder-Mead approach maintains a simplex of $n + 1$ function evaluation points that it updates at each step. In Nelder-Mead, one updates the simplex based on function values at the simplex corners, the centroid, and one other point; or one contracts the simplex.

Visualizations of Nelder-Mead are often quite striking: the simplex appears to crawl downhill like some sort of mathematical amoeba. But there are examples of functions where Nelder-Mead is not guaranteed to converge to a minimum at all.

Hook-Jeeves and successors

The basic idea of *pattern search* methods is to either

- Keep going if a promising direction turns out good, or
- Poll points in a pattern around the current iterate to find a new direction

For example, in the Hook-Jeeves approach (one of the earliest pattern search methods), one would at each polling move evaluate $\phi(x^{(k)} \pm \Delta e_j)$ for each of the n coordinate directions e_j . If one of the new points is better than $x^{(k)}$, it becomes $x^{(k+1)}$ (and we may increase Δ if we already took a step in this direction to get from $x^{(k-1)}$ to $x^{(k)}$). If $x^{(k)}$ is better than any surrounding point, we decrease Δ and try again.

```

function hooke_jeeves( $\phi$ , x;  $\Delta=1.0$ ,  $\Delta\text{tol}=1e-3$ , maxevals=1000,
                    monitor=(x, $\Delta$ ,poll)->nothing)

 $\phi_0 = \phi(x)$       # Current best value found
poll = true        # Poll step or keep going?
evals = 0          # Number of function evaluations
k = 0              # Current search direction

# Possible search directions (+/- in each coordinate direction)
d = length(x)
P = [Matrix(I,d,d) -Matrix(I,d,d)]

while  $\Delta > \Delta\text{tol}$  && evals < maxevals
    monitor(x,  $\Delta$ , poll)
    if poll

        # Poll in neighborhood. Pick the best descent direction;
        # if we get no descent, cut the radius and try again.
         $\phi_{\text{Pmin}}$ , kmin = findmin( $\phi(x + \Delta*p)$  for p in eachcol(P))
        evals += 2*d
        if  $\phi_0 \geq \phi_{\text{Pmin}}$ 
            k = kmin
            poll = false
        else
             $\Delta = \Delta/2$ 
        end

    else

        # Take a step, accept if decrease, poll otherwise
        xnew = x +  $\Delta * P[:,k]$ 
         $\phi_{\text{new}} = \phi(x_{\text{new}})$ 
        evals += 1
        if  $\phi_{\text{new}} < \phi_0$ 
            x[:] = xnew
             $\phi_0 = \phi_{\text{new}}$ 
        else
            poll = true
        end

    end

end
end

```

```
x, Δ  
end
```

We give an example of the convergence of Hooke-Jeeves on a quadratic model problem with the code below (Figure 2).

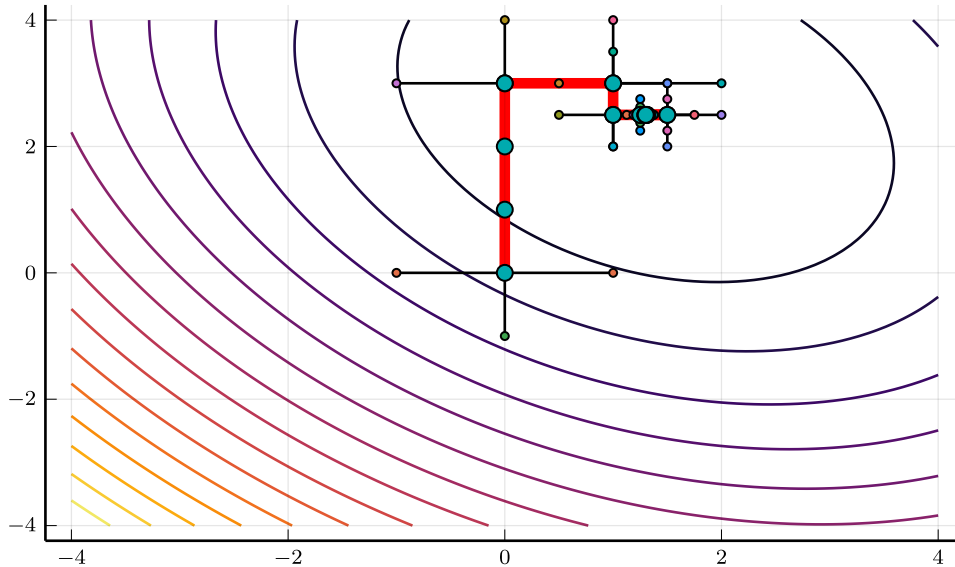


Figure 2: Convergence of the Hooke-Jeeves algorithm for a quadratic model problem.

More generally, we would evaluate $\phi(x^{(k)} + d)$ for $d \in \mathcal{G}(\Delta)$, a *generating set* of directions with some scale factor Δ . There are many variants on the “search-and-poll” strategy of the general pattern search; for example,

- We can do a random selection from the pattern directions and choose the first promising one (rather than polling all directions).
- We can use a model to suggest some extra promising directions in addition to the pattern, or to modify from a fixed pattern.

Simple methods like Hooke-Jeeves converge for smooth objectives, but may fail in the nonsmooth case. However, the mesh adaptive direct search (MADS) class of methods can converge even in this case.

Summarizing thoughts

Direct search methods have been with us for more than half a century: the original Hook-Jeeves paper was from 1961, and the Nelder-Mead paper goes back to 1965. These methods are attractive in that they require only the ability to compute objective function values, and can be used with “black box” codes – or even with evaluations based on running a physical experiment! Computing derivatives requires some effort, even when automatic differentiation and related tools are available, and so gradient-free approaches may also be attractive because of ease-of-use.

Gradient-free methods often work well in practice for solving optimization problems with modest accuracy requirements. This is true even of methods like Nelder-Mead, for which there are examples of very nice functions (smooth and convex) for which the method is guaranteed to mis-converge. But though the theoretical foundations for these methods have gradually improved with time, the theory for gradient-free methods is much less clear-cut than the theory for gradient-based methods. Gradient-based methods also have a clear advantage at higher accuracy requirements.

Gradient-free methods do *not* free a user from the burden of finding a good initial guess. Methods like Nelder-Mead and pattern search will, at best, converge to local minima. Many heuristic methods for finding global minimizers are gradient-free; I include among these methods like simulated annealing, genetic algorithms, and Bayesian optimization techniques. On the other hand, branch-and-bound methods that yield provable global minimizers are often heavily dependent on derivatives (or bounds proved with the help of derivatives).

Just because a method does not explicitly use gradients does not mean it doesn't rely on them implicitly. Gradient-free methods may have just as much difficulty with functions that are discontinuous, or that have large Lipschitz constants – particularly those methods that implicitly build a local linear or quadratic model.

In many areas in numerics, an ounce of analysis pays for a pound of computation. If the computation is to be done repeatedly, or must be done to high accuracy, then it is worthwhile to craft an approach that takes advantage of specific problem structure. On the other hand, sometimes one just wants to do a cheap exploratory computation to get started, and the effort of using a specialized approach may not be warranted. An overview of the options that are available is useful for approaching these tradeoffs intelligently.