

A comparison of `nlsr::nlxb` with `nls` and `minpack::nlsLM`

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R has several tools for estimating nonlinear models and minimizing sums of squares functions. Sometimes we talk of **nonlinear regression** and at other times of **minimizing a sum of squares function**. Many workers conflate these two tasks. Here some of the differences are highlighted by comparing the tools from the package `nlsr`, particularly function `nlxb()` with those from base-R `nls()` and the `nlsLM` function of package `minpack.lm`.

TODO

- shorten example output and put in a table
- add structure of output
- show prediction functions, including where they don't work

Principal differences

The following is a summary of the main differences in the tools.

Derivative information

`nlsr::nlxb()` attempts to use symbolic and algorithmic tools to obtain the derivatives of the model expression that are needed for the **Jacobian** matrix that is used in creating a linearized sub-problem at each iteration of an attempted solution of the minimization of the sum of squared residuals. `nls()` and `minpack.lm::nlsLM()` use a very simple forward-difference approximation for the partial derivatives. For the *i*'th partial derivative of the modelling function with respect to parameter *xx*, they use (in **C**)

```
delta = (xx == 0) ? eps : xx*eps;
```

and

```
REAL(gradient)[start + k] = rDir[i] * (REAL(ans_del)[k] - REAL(ans)[k])/delta;
```

where

```
double eps = sqrt(DOUBLE_EPS), *rDir;
```

and where `DOUBLE_EPS` in `Constants.h` in the **R** source code refers to `DBL_EPSILON` in `float.h` in most C compilers, e.g.,

```
#define DBL_EPSILON 2.2204460492503131e-16
```

Thus `delta` is of the order of 1.490116e-08.

Forward difference approximations are less accurate than central differences, and both are subject to numerical error when the modelling function is “flat”, so that there is a large amount of digit cancellation in the subtraction necessary to compute the derivative approximation.

`minpack.lm::nlsLM` uses the same derivatives as far as I can determine. The loss of information compared to the analytic or algorithmic derivatives of `nlsr::nlxb()` is important in that it can lead to Jacobian

matrices that are computationally singular, where `nls()` will stop with “singular gradient”. (It is actually the Jacobian which is singular here, and I will stay with that terminology.) `minpack.lm::nlsLM()` may fail to get started if the initial Jacobian is singular, but is less susceptible in general, as described in the sub-section on Marquardt stabilization.

Marquardt stabilization

All three of the R functions under consideration try to minimize a sum of squares. If the model is provided in the form

`y ~ (some expression)`

then the residuals are computed by evaluating the difference between `(some expression)` and `y`. My own preference, and that of K F Gauss, is to use `(some expression) - y`. This is to avoid having to be concerned with the negative sign – the derivative of the residual defined in this way is the same as the derivative of the modelling function, and we avoid the chance of a sign error. The Jacobian matrix is made up of elements where element `i, j` is the partial derivative of residual `i` w.r.t. parameter `j`.

`nls()` attempts to minimize a sum of squared residuals by a Gauss-Newton method. If we compute a Jacobian matrix `J` and a vector of residuals `r` from a vector of parameters `x`, then we can define a linearized problem

$$J^T J \delta = -J^T r$$

This leads to an iteration where, from a set of starting parameters `x0`, we compute

$$x_{i+1} = x_i + \delta$$

This is commonly modified to use a step factor `step`

$$x_{i+1} = x_i + step * \delta$$

It is in the mechanisms to choose the size of `step` and to decide when to terminate the iteration that Gauss-Newton methods differ. Indeed, though I have tried several times, I find the very convoluted code behind `nls()` very difficult to decipher. Unfortunately, its authors have now (as far as I am aware) all ceased to maintain the code.

Both `nlsr::nlxb()` and `minpack.lm::nlsLM` use a Levenberg-Marquardt stabilization of the iteration. (Marquardt (1963), Levenberg (1944)), solving

$$(J^T J + \lambda D) \delta = -J^T r$$

where `D` is some diagonal matrix and `lambda` is a number of modest size initially. Clearly for $\lambda = 0$ we have a Gauss-Newton method. Typically, the sum of squares of the residuals calculated at the “new” set of parameters is used as a criterion for keeping those parameter values. If so, the size of λ is reduced. If not, we increase the size of λ and compute a new δ . Note that a new `J`, the expensive step in each iteration, is NOT required.

As for Gauss-Newton methods, the details of how to start, adjust and terminate the iteration lead to many variants, increased by different possibilities for specifying `D`. See Nash (1979).

Consequences of different derivative computations

While readers might expect that the precise derivative information of `nlsr::nlxb()` would mean a faster solution, this is quite often not the case. Approximate derivatives may allow faster approach to the solution by “ironing out” wrinkles in the function surface. In my opinion, the main advantage of precise derivative information is in testing that we actually have arrived at a solution.

There are even some cases where the approximation may be helpful, though users may not realize the potential danger. Thanks to Karl Schilling for an example of modelling with the function

```
a * (x ^ b)
```

where x is our data and we wish to estimate a and b . Now the partial derivative of this function w.r.t. b is

```
partialderiv <- D(expression(a * (x ^ b)), "b")
print(partialderiv)
```

```
## a * (x^b * log(x))
```

The danger here is that we may have data values $x = 0$, in which case the **derivative** is not defined, though the model can still be evaluated. Thus `nlsr::nlxb()` will not compute a solution, while `nls()` and `minpack.lm::nlsLM()` will generally proceed. A workaround is to provide a very small value instead of zero for the data, though I find this inelegant. A proper treatment might be to develop the limit of the derivative as the data value goes to zero, but finding general software does not seem worth the effort.

An illustrative problem So we can illustrate some of the issues, let us create some example data for a family of related computational problems.

```
# Here we set up an example problem with data
# Define independent variables
t1 <- 1:20
t0 <- 0:19
t0a <- t0
t0a[1] <- 1e-7
y1 <- 4 * (t1^0.25)
n <- length(t1)
fuzz <- rnorm(n)
range <- max(y1)-min(y1)
y1p <- y1 + 0.1*range*fuzz
y1q <- y1 + 0.2*range*fuzz
y1r <- y1 + 0.5*range*fuzz
edta <- data.frame(t0=t0, t1=t1, t0a=t0a, y1=y1, y1p=y1p, y1q=y1q, y1r=y1r)
```

Let us try our example modelling $y1$ against $t1$. Note that this is a zero-residual problem, so `nls()` should complain or fail, which it appears to do but by exceeding the iteration limit, which is not very communicative of the underlying issue.

```
start1 <- c(a=1, b=1)
try(nlsy1t1 <- nls(formula=y1~a*(t1^b), start=start1, data=edta))
```

```
## Error in nls(formula = y1 ~ a * (t1^b), start = start1, data = edta) :
##   number of iterations exceeded maximum of 50
```

```
try(nlsy1t1 <- nls(formula=y1~a*(t1^b), start=start1, data=edta, control=nls.control(maxiter=10000)))
```

```
## Error in nls(formula = y1 ~ a * (t1^b), start = start1, data = edta, control = nls.control(maxiter =
##   number of iterations exceeded maximum of 10000
```

```

library(nlsr)
nlsry1t1 <- nlxb(formula=y1~a*(t1^b), start=start1, data=edta)
nlsry1t1

## nlsr object: x
## residual sumsquares = 4.1415e-30 on 20 observations
## after 7 Jacobian and 8 function evaluations
## name      coeff      SE      tstat      pval      gradient      JSingval
## a          4      2.396e-16  1.67e+16  3.618e-282  2.262e-15      77.28
## b          0.25    2.446e-17  1.022e+16  2.489e-278  4.099e-14      1.993

library(minpack.lm)
nlsLMy1t1 <- nlsLM(formula=y1~a*(t1^b), start=start1, data=edta)
nlsLMy1t1

## Nonlinear regression model
## model: y1 ~ a * (t1^b)
## data: edta
## a      b
## 4.00 0.25
## residual sum-of-squares: 0
##
## Number of iterations to convergence: 7
## Achieved convergence tolerance: 1.49e-08

try(nlsy1t0 <- nls(formula=y1~a*(t0^b), start=start1, data=edta))
nlsy1t0

## Nonlinear regression model
## model: y1 ~ a * (t0^b)
## data: edta
## a      b
## 4.524 0.209
## residual sum-of-squares: 16.1
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 4.66e-07

library(nlsr)
try(nlsry1t0 <- nlxb(formula=y1~a*(t0^b), start=start1, data=edta))

## Error in model2rjfun(formula, pnun, data = data) : Jacobian contains NaN
##FAIL -- nlsry1t0
library(minpack.lm)
nlsLMy1t0 <- nlsLM(formula=y1~a*(t0^b), start=start1, data=edta)
nlsLMy1t0

## Nonlinear regression model
## model: y1 ~ a * (t0^b)
## data: edta
## a      b
## 4.524 0.209
## residual sum-of-squares: 16.1
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 1.49e-08

```

```
try(nlsy1pt1 <- nls(formula=y1p~a*(t1^b), start=start1, data=edta))
nlsy1pt1
```

```
## Nonlinear regression model
## model: y1p ~ a * (t1^b)
## data: edta
## a b
## 3.994 0.261
## residual sum-of-squares: 3.43
##
## Number of iterations to convergence: 5
## Achieved convergence tolerance: 5.39e-06
```

```
library(nlsr)
nlsry1pt1 <- nlxb(formula=y1p~a*(t1^b), start=start1, data=edta)
nlsry1pt1
```

```
## nlsr object: x
## residual sumsquares = 3.4258 on 20 observations
## after 7 Jacobian and 8 function evaluations
## name coeff SE tstat pval gradient JSingval
## a 3.99385 0.215 18.57 3.446e-13 -1.798e-12 79.49
## b 0.261419 0.02193 11.92 5.625e-10 1.728e-11 2.019
```

```
library(minpack.lm)
nlsLMy1pt1 <- nlsLM(formula=y1p~a*(t1^b), start=start1, data=edta)
nlsLMy1pt1
```

```
## Nonlinear regression model
## model: y1p ~ a * (t1^b)
## data: edta
## a b
## 3.994 0.261
## residual sum-of-squares: 3.43
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 1.49e-08
```

```
try(nlsy1qt1 <- nls(formula=y1q~a*(t1^b), start=start1, data=edta))
nlsy1qt1
```

```
## Nonlinear regression model
## model: y1q ~ a * (t1^b)
## data: edta
## a b
## 3.991 0.272
## residual sum-of-squares: 13.7
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 2.58e-07
```

```
library(nlsr)
nlsry1qt1 <- nlxb(formula=y1q~a*(t1^b), start=start1, data=edta)
nlsry1qt1
```

```
## nlsr object: x
## residual sumsquares = 13.706 on 20 observations
```

```
##      after 8      Jacobian and 9 function evaluations
##  name      coeff      SE      tstat      pval      gradient      JSingval
## a          3.99135      0.4248      9.396  2.307e-08  1.065e-11      81.71
## b          0.272165      0.04324      6.294  6.202e-06  3.889e-07      2.044
```

```
library(minpack.lm)
nlsLMy1qt1 <- nlsLM(formula=y1q~a*(t1^b), start=start1, data=edta)
nlsLMy1qt1
```

```
## Nonlinear regression model
##  model: y1q ~ a * (t1^b)
##  data: edta
##    a      b
## 3.991 0.272
## residual sum-of-squares: 13.7
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 1.49e-08
```

```
try(nlsy1rt1 <- nls(formula=y1r~a*(t1^b), start=start1, data=edta))
nlsy1rt1
```

```
## Nonlinear regression model
##  model: y1r ~ a * (t1^b)
##  data: edta
##    a      b
## 4.003 0.301
## residual sum-of-squares: 85.7
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 2.25e-06
```

```
library(nlsr)
nlsry1rt1 <- nlxb(formula=y1r~a*(t1^b), start=start1, data=edta)
nlsry1rt1
```

```
## nlsr object: x
## residual sumsquares = 85.695 on 20 observations
##      after 8      Jacobian and 9 function evaluations
##  name      coeff      SE      tstat      pval      gradient      JSingval
## a          4.00253      1.026      3.899  0.001051 -9.577e-12      88.34
## b          0.300873      0.1035      2.907  0.009412  4.398e-11      2.116
```

```
library(minpack.lm)
nlsLMy1rt1 <- nlsLM(formula=y1r~a*(t1^b), start=start1, data=edta)
nlsLMy1rt1
```

```
## Nonlinear regression model
##  model: y1r ~ a * (t1^b)
##  data: edta
##    a      b
## 4.003 0.301
## residual sum-of-squares: 85.7
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 1.49e-08
```

Output of the modelling functions

`nls()` and `nlsLM()` return the same structure. Let us examine this for one of our example results (we will choose one that does NOT have small residuals, so that all the functions “work”).

```
str(nlsy1pt1)
```

```
## List of 6
## $ m          :List of 16
## ..$ resid    :function ()
## ..$ fitted    :function ()
## ..$ formula   :function ()
## ..$ deviance  :function ()
## ..$ lhs       :function ()
## ..$ gradient  :function ()
## ..$ conv      :function ()
## ..$ incr      :function ()
## ..$ setVarying:function (vary = rep_len(TRUE, length(useParams)))
## ..$ setPars   :function (newPars)
## ..$ getPars   :function ()
## ..$ getAllPars:function ()
## ..$ getEnv    :function ()
## ..$ trace     :function ()
## ..$ Rmat      :function ()
## ..$ predict   :function (newdata = list(), qr = FALSE)
## ..- attr(*, "class")= chr "nlsModel"
## $ convInfo    :List of 5
## ..$ isConv    : logi TRUE
## ..$ finIter   : int 5
## ..$ finTol    : num 5.39e-06
## ..$ stopCode  : int 0
## ..$ stopMessage: chr "converged"
## $ data        : symbol edta
## $ call        : language nls(formula = y1p ~ a * (t1^b), data = edta, start = start1, algorithm = "d
## $ dataClasses: Named chr "numeric"
## ..- attr(*, "names")= chr "t1"
## $ control     :List of 5
## ..$ maxiter   : num 50
## ..$ tol       : num 1e-05
## ..$ minFactor: num 0.000977
## ..$ printEval: logi FALSE
## ..$ warnOnly  : logi FALSE
## - attr(*, "class")= chr "nls"
```

Note that this structure has a lot of special functions in the sub-list `m`. By contrast, the `nlsr()` output is much less flamboyant. There are, in fact, no functions as part of the structure.

```
str(nlsry1pt1)
```

```
## List of 11
## $ resid      : num [1:20] 0.4062 -0.702 0.0809 0.0728 0.7917 ...
## ..- attr(*, "gradient")= num [1:20, 1:2] 1 1.2 1.33 1.44 1.52 ...
## .. ..- attr(*, "dimnames")=List of 2
## .. .. ..$ : NULL
## .. .. ..$ : chr [1:2] "a" "b"
## $ jacobian    : num [1:20, 1:2] 1 1.2 1.33 1.44 1.52 ...
```

```
##   ..- attr(*, "dimnames")=List of 2
##   .. ..$ : NULL
##   .. ..$ : chr [1:2] "a" "b"
##   $ feval      : num 8
##   $ jeval      : num 7
##   $ coefficients: Named num [1:2] 3.994 0.261
##   ..- attr(*, "names")= chr [1:2] "a" "b"
##   $ ssquares   : num 3.43
##   $ lower      : num [1:2] -Inf -Inf
##   $ upper      : num [1:2] Inf Inf
##   $ maskidx    : int(0)
##   $ weights    : NULL
##   $ formula    :Class 'formula' language y1p ~ a * (t1^b)
##   .. ..- attr(*, ".Environment")=<environment: R_GlobalEnv>
##   - attr(*, "class")= chr "nlshr"
```

Which of these approaches is “better” can be debated. My preference is for the results of optimization computations to be essentially data, including messages, though some tools within some of my packages will return functions for specific reasons, e.g., to return a function from an expression. However, I prefer to use specified functions such as `predict.nlsr()` below to obtain predictions. I welcome comment and discussion, as this is not, in my view, a closed topic.

Prediction

Let us predict our models at the mean of the data. Because `nlxb()` returns a different structure from that found by `nls()` and `nlsLM()` the code for `predict()` is different. (`minpack.lm` uses `predict.nls` since the output structure of the modelling step is equivalent to that from `nls()`.)

```
nudta <- colMeans(edta)
predict(nlsy1pt1, newdata=nudta)
```

```
## [1] 7.385
```

```
predict(nlsLMy1pt1, newdata=nudta)
```

```
## [1] 7.385
```

```
predict(nlsry1pt1, newdata=nudta)
```

```
## [1] 7.385
## attr(,"class")
## [1] "predict.nlsr"
## attr(,"pkgname")
## [1] "nlshr"
```

Problems that are NOT regressions

Some nonlinear least squares problems are NOT nonlinear regressions. That is, we do not have a formula $y \sim (\text{some function})$ to define the problem. This is another reason to use the residual in the form $(\text{some function}) - y$. In many cases of interest we have no y .

The Brown and Dennis test problem (Moré, Garbow, and Hillstom (1981), problem 16) is of this form. Suppose we have m observations, then we create a scaled index t which is the “data” for the function. To run the nonlinear least squares functions that use a formula, we do, however, need a “ y ” variable. Clearly adding zero to the residual will not change the problem, so we set the data for “ y ” as all zeros. Note that `nls()` and `nlsLM()` need some extra iterations to find the solution to this somewhat nasty problem.


```

m <- 20
t <- seq(1, m) / 5
y <- rep(0, m)
library(nlsr)
library(minpack.lm)

bddata <- data.frame(t=t, y=y)
bdform <- y ~ ((x1 + t * x2 - exp(t))^2 + (x3 + x4 * sin(t) - cos(t))^2)
prm0 <- c(x1=25, x2=5, x3=-5, x4=-1)
fbd <- model2ssgrfun(bdform, prm0, bddata)
cat("initial sumsquares=", as.numeric(crossprod(fbd(prm0))), "\n")

## initial sumsquares= 6.2832e+13

## prmstar <-
nlsrbd <- nlxb(bdform, start=prm0, data=bddata, trace=FALSE)
## summary(nlsrbd)
nlsrbd

## nlsr object: x
## residual sumsquares = 85822 on 20 observations
## after 3056 Jacobian and 4279 function evaluations
## name coeff SE tstat pval gradient JSingval
## x1 -11.5944 4.017 -2.886 0.01075 0.02637 176
## x2 13.2036 1.231 10.73 1.025e-08 -0.07207 28.1
## x3 -0.403442 28.08 -0.01437 0.9887 -0.002117 3.917
## x4 0.236777 39.79 0.005951 0.9953 -0.001648 1.624

nlsbd <- try(nls(bdform, start=prm0, data=bddata, trace=FALSE))

## Error in nls(bdform, start = prm0, data = bddata, trace = FALSE) :
## number of iterations exceeded maximum of 50

nlsbd

## [1] "Error in nls(bdform, start = prm0, data = bddata, trace = FALSE) : \n number of iterations exceeded maximum of 50"
## attr(,"class")
## [1] "try-error"
## attr("condition")
## <simpleError in nls(bdform, start = prm0, data = bddata, trace = FALSE): number of iterations exceeded maximum of 50
nlsbd10k <- nls(bdform, start=prm0, data=bddata, trace=FALSE, control=nls.control(maxiter=10000))
nlsbd10k

## Nonlinear regression model
## model: y ~ ((x1 + t * x2 - exp(t))^2 + (x3 + x4 * sin(t) - cos(t))^2)
## data: bddata
## x1 x2 x3 x4
## -11.594 13.204 -0.403 0.237
## residual sum-of-squares: 85822
##
## Number of iterations to convergence: 855
## Achieved convergence tolerance: 9.6e-06

nlsLMbd <- nlsLM(bdform, start=prm0, data=bddata, trace=FALSE)

## Warning in nls.lm(par = start, fn = FCT, jac = jac, control = control, lower = lower, : lmdif: info:

```

```
nlsLMbd
```

```
## Nonlinear regression model
## model: y ~ ((x1 + t * x2 - exp(t))^2 + (x3 + x4 * sin(t) - cos(t))^2)
## data: bddata
##      x1      x2      x3      x4
## 1.630  8.639 -0.154  0.959
## residual sum-of-squares: 252493
##
## Number of iterations till stop: 50
## Achieved convergence tolerance: 1.49e-08
## Reason stopped: Number of iterations has reached `maxiter' == 50.
```

```
nlsLMbd10k <- nlsLM(bdform, start=prm0, data=bddata, trace=FALSE, control=nls.lm.control(maxiter=10000,
```

```
## Warning in nls.lm(par = start, fn = FCT, jac = jac, control = control, lower =
## lower, : resetting `maxiter' to 1024!
```

```
nlsLMbd10k
```

```
## Nonlinear regression model
## model: y ~ ((x1 + t * x2 - exp(t))^2 + (x3 + x4 * sin(t) - cos(t))^2)
## data: bddata
##      x1      x2      x3      x4
## -11.592 13.203 -0.404  0.237
## residual sum-of-squares: 85822
##
## Number of iterations to convergence: 242
## Achieved convergence tolerance: 1.49e-08
```

Let us try predicting the “residual” for some new data.

```
ndata <- data.frame(t=c(5,6), y=c(0,0))
predict(nlsLMbd, newdata=ndata)
```

```
## [1] 10732 122476
```

```
# now nls
```

```
predict(nlsbd10k, newdata=ndata)
```

```
## [1] 8834.9 112764.7
```

```
# now nlsr
```

```
predict(nlsrbd, newdata=ndata)
```

```
## [1] 8834.9 112764.7
```

```
## attr(,"class")
```

```
## [1] "predict.nlsr"
```

```
## attr(,"pkgname")
```

```
## [1] "nlsr"
```

We could, of course, try setting up a different formula. However, we discover that the parsing of the model formula for `nls()` and `nlsLM()` fails for this formulation, but `nlxb()` proceeds as usual.

```
bdf2 <- (x1 + t * x2 - exp(t))^2 ~ - (x3 + x4 * sin(t) - cos(t))^2
```

```
nlsbd2 <- try(nls(bdf2, start=prm0, data=bddata, trace=FALSE))
```

```
## Error in eval(formula[[2L]], data, env) : object 'x1' not found
```

```
nlsbd2

## [1] "Error in eval(formula[[2L]], data, env) : object 'x1' not found\n"
## attr(,"class")
## [1] "try-error"
## attr(,"condition")
## <simpleError in eval(formula[[2L]], data, env): object 'x1' not found>
nlsLMbd2 <- try(nlsLM(bdf2, start=prm0, data=bddata, trace=FALSE,
                     control=nls.lm.control(maxiter=10000, maxfev=10000)))
```

```
## Error in eval(formula[[2L]], data, env) : object 'x1' not found
```

```
nlsLMbd2

## [1] "Error in eval(formula[[2L]], data, env) : object 'x1' not found\n"
## attr(,"class")
## [1] "try-error"
## attr(,"condition")
## <simpleError in eval(formula[[2L]], data, env): object 'x1' not found>
```

```
nlsrbd2 <- nlxb(bdf2, start=prm0, data=bddata, trace=FALSE)
##summary(nlsrbd2)
nlsrbd2
```

```
## nlsr object: x
## residual sumsquares = 85822 on 20 observations
## after 3056 Jacobian and 4279 function evaluations
## name      coeff      SE      tstat      pval      gradient      JSingval
## x1         -11.5944    4.017    -2.886    0.01075    0.02637      176
## x2          13.2036    1.231     10.73   1.025e-08   -0.07207     28.1
## x3         -0.403442   28.08    -0.01437   0.9887   -0.002117     3.917
## x4          0.236777   39.79     0.005951  0.9953   -0.001648     1.624
```

We can try “prediction” again for `nlxb()`. The answers are, of course, rather different, as the `predict.nlsr()` function **ONLY** evaluates the right hand side of the formula. We are **mis-applying** the function here, and that is a reason the function has been renamed. It would be good to have more checks on the formula, and I welcome collaboration to do this.

```
nldata <- data.frame(t=c(5,6), y=c(0,0))
predict(nlsrbd2, newdata=nldata)
```

```
## [1] -0.83568 -2.04425
## attr(,"class")
## [1] "predict.nlsr"
## attr(,"pkgname")
## [1] "nlsr"
```

A check on the Brown and Dennis calculation via function minimization

```
## Brown and Dennis Function
##
## Test function 16 from the More', Garbow and Hillstom paper.
##
## The objective function is the sum of \code{m} functions, each of \code{n}
## parameters.
##
## \itemize{
```

```

#' \item Dimensions: Number of parameters \code{n = 4}, number of summand
#' functions \code{m >= n}.
#' \item Minima: \code{f = 85822.2} if \code{m = 20}.
#' }
#'
#' @param m Number of summand functions in the objective function. Should be
#' equal to or greater than 4.
#' @return A list containing:
#' \itemize{
#' \item \code{fn} Objective function which calculates the value given input
#' parameter vector.
#' \item \code{gr} Gradient function which calculates the gradient vector
#' given input parameter vector.
#' \item \code{fg} A function which, given the parameter vector, calculates
#' both the objective value and gradient, returning a list with members
#' \code{fn} and \code{gr}, respectively.
#' \item \code{x0} Standard starting point.
#' }
#'
#' @references
#' More', J. J., Garbow, B. S., & Hillstom, K. E. (1981).
#' Testing unconstrained optimization software.
#' \emph{ACM Transactions on Mathematical Software (TOMS)}, \emph{7}(1), 17-41.
#' \url{https://doi.org/10.1145/355934.355936}
#'
#' Brown, K. M., & Dennis, J. E. (1971).
#' \emph{New computational algorithms for minimizing a sum of squares of
#' nonlinear functions} (Report No. 71-6).
#' New Haven, CT: Department of Computer Science, Yale University.
#'
#' @examples
#' # Use 10 summand functions
#' fun <- brown_den(m = 10)
#' # Optimize using the standard starting point
#' x0 <- fun$x0
#' res_x0 <- stats::optim(par = x0, fn = fun$fn, gr = fun$gr, method =
#' "L-BFGS-B")
#' # Use your own starting point
#' res <- stats::optim(c(0.1, 0.2, 0.3, 0.4), fun$fn, fun$gr, method =
#' "L-BFGS-B")
#'
#' # Use 20 summand functions
#' fun20 <- brown_den(m = 20)
#' res <- stats::optim(fun20$x0, fun20$fn, fun20$gr, method = "L-BFGS-B")
#' @export
#`
brown_den <- function(m = 20) {
  list(
    fn = function(par) {
      x1 <- par[1]
      x2 <- par[2]
      x3 <- par[3]
      x4 <- par[4]

```

```

    ti <- (1:m) * 0.2
    l <- x1 + ti * x2 - exp(ti)
    r <- x3 + x4 * sin(ti) - cos(ti)
    f <- l * l + r * r
    sum(f * f)
  },
  gr = function(par) {
    x1 <- par[1]
    x2 <- par[2]
    x3 <- par[3]
    x4 <- par[4]

    ti <- (1:m) * 0.2
    sinti <- sin(ti)
    l <- x1 + ti * x2 - exp(ti)
    r <- x3 + x4 * sinti - cos(ti)
    f <- l * l + r * r
    lf4 <- 4 * l * f
    rf4 <- 4 * r * f
    c(
      sum(lf4),
      sum(lf4 * ti),
      sum(rf4),
      sum(rf4 * sinti)
    )
  },
  fg = function(par) {
    x1 <- par[1]
    x2 <- par[2]
    x3 <- par[3]
    x4 <- par[4]

    ti <- (1:m) * 0.2
    sinti <- sin(ti)
    l <- x1 + ti * x2 - exp(ti)
    r <- x3 + x4 * sinti - cos(ti)
    f <- l * l + r * r
    lf4 <- 4 * l * f
    rf4 <- 4 * r * f

    fsum <- sum(f * f)
    grad <- c(
      sum(lf4),
      sum(lf4 * ti),
      sum(rf4),
      sum(rf4 * sinti)
    )

    list(
      fn = fsum,
      gr = grad
    )
  },

```

```

    x0 = c(25, 5, -5, 1)
  )
}
mbd <- brown_den(m=20)
mbd

## $fn
## function(par) {
##     x1 <- par[1]
##     x2 <- par[2]
##     x3 <- par[3]
##     x4 <- par[4]
##
##     ti <- (1:m) * 0.2
##     l <- x1 + ti * x2 - exp(ti)
##     r <- x3 + x4 * sin(ti) - cos(ti)
##     f <- l * l + r * r
##     sum(f * f)
## }
## <bytecode: 0x5599968f5a00>
## <environment: 0x5599988aaba0>
##
## $gr
## function(par) {
##     x1 <- par[1]
##     x2 <- par[2]
##     x3 <- par[3]
##     x4 <- par[4]
##
##     ti <- (1:m) * 0.2
##     sinti <- sin(ti)
##     l <- x1 + ti * x2 - exp(ti)
##     r <- x3 + x4 * sinti - cos(ti)
##     f <- l * l + r * r
##     lf4 <- 4 * l * f
##     rf4 <- 4 * r * f
##     c(
##         sum(lf4),
##         sum(lf4 * ti),
##         sum(rf4),
##         sum(rf4 * sinti)
##     )
## }
## <bytecode: 0x559999df4ee8>
## <environment: 0x5599988aaba0>
##
## $fg
## function(par) {
##     x1 <- par[1]
##     x2 <- par[2]
##     x3 <- par[3]
##     x4 <- par[4]
##
##     ti <- (1:m) * 0.2

```

```

##      sinti <- sin(ti)
##      l <- x1 + ti * x2 - exp(ti)
##      r <- x3 + x4 * sinti - cos(ti)
##      f <- l * l + r * r
##      lf4 <- 4 * l * f
##      rf4 <- 4 * r * f
##
##      fsum <- sum(f * f)
##      grad <- c(
##          sum(lf4),
##          sum(lf4 * ti),
##          sum(rf4),
##          sum(rf4 * sinti)
##      )
##
##      list(
##          fn = fsum,
##          gr = grad
##      )
##  }
## <bytecode: 0x5599998bad10>
## <environment: 0x5599988aaba0>
##
## $x0
## [1] 25  5 -5  1
mbd$fg(mbd$x0)

## $fn
## [1] 7632895
##
## $gr
## [1] 1127773 1746780 -192836 -120879
bdsolnm <- optim(mbd$x0, mbd$fn, control=list(trace=0))
bdsolnm

## $par
## [1] -11.59488  13.20436 -0.40671  0.23903
##
## $value
## [1] 85822
##
## $counts
## function gradient
##      167      NA
##
## $convergence
## [1] 0
##
## $message
## NULL
bdsolbfgs <- optim(mbd$x0, mbd$fn, method="BFGS", control=list(trace=0))
bdsolbfgs

```

```
## $par
## [1] -11.59444 13.20363 -0.40343 0.23678
##
## $value
## [1] 85822
##
## $counts
## function gradient
##      76      16
##
## $convergence
## [1] 0
##
## $message
## NULL
```

```
library(optimx)
methlist <- c("Nelder-Mead", "BFGS", "Rvmmin", "L-BFGS-B", "Rcgmin", "ucminf")
solo <- opm(mbd$x0, mbd$fn, mbd$gr, method=methlist, control=list(trace=0))
```

```
## Warning in optimr(par, fn, gr, hess = hess, method = meth, lower = lower, : User
## has set control$maximize = FALSE and admissible control$fnscale

## Warning in optimr(par, fn, gr, hess = hess, method = meth, lower = lower, : User
## has set control$maximize = FALSE and admissible control$fnscale

## Warning in optimr(par, fn, gr, hess = hess, method = meth, lower = lower, : User
## has set control$maximize = FALSE and admissible control$fnscale

## Warning in optimr(par, fn, gr, hess = hess, method = meth, lower = lower, : User
## has set control$maximize = FALSE and admissible control$fnscale

## Warning in optimr(par, fn, gr, hess = hess, method = meth, lower = lower, : User
## has set control$maximize = FALSE and admissible control$fnscale

## Warning in optimr(par, fn, gr, hess = hess, method = meth, lower = lower, : User
## has set control$maximize = FALSE and admissible control$fnscale
```

```
summary(solo, order=value)
```

| ## | p1 | p2 | p3 | p4 | value | fevals | gevals | convergence |
|----------------|---------|--------|----------|---------|-------|--------|--------|-------------|
| ## Rcgmin | -11.594 | 13.204 | -0.40344 | 0.23678 | 85822 | 2237 | 295 | 1 |
| ## ucminf | -11.594 | 13.204 | -0.40344 | 0.23678 | 85822 | 33 | 33 | 0 |
| ## Rvmmin | -11.594 | 13.204 | -0.40344 | 0.23678 | 85822 | 53 | 20 | 0 |
| ## BFGS | -11.594 | 13.204 | -0.40343 | 0.23678 | 85822 | 76 | 16 | 0 |
| ## L-BFGS-B | -11.594 | 13.204 | -0.40348 | 0.23687 | 85822 | 24 | 24 | 0 |
| ## Nelder-Mead | -11.595 | 13.204 | -0.40671 | 0.23903 | 85822 | 167 | NA | 0 |
| ## | kkt1 | kkt2 | xtime | | | | | |
| ## Rcgmin | TRUE | TRUE | 0.016 | | | | | |
| ## ucminf | TRUE | TRUE | 0.004 | | | | | |
| ## Rvmmin | TRUE | TRUE | 0.003 | | | | | |
| ## BFGS | TRUE | TRUE | 0.001 | | | | | |
| ## L-BFGS-B | TRUE | TRUE | 0.001 | | | | | |
| ## Nelder-Mead | TRUE | TRUE | 0.001 | | | | | |

Small residuals

nls() documentation warns

"Warning

Do not use nls on artificial “zero-residual” data."

It goes on to recommend that users add “error” to the data to avoid such problems. I feel this is a very unsatisfactory kludge. It is NOT due to a genuine mathematical issue, but due to the relative offset convergence criterion used to terminate the method. This is also used in `nlsr`, but I do not see it in the Fortran code that underlies `minpack.lm`, though I must caution that these codes are very convoluted (R calling C calling Fortran, with multiple subsidiary routines).

```
x <- (1:m)
y <- 2.5 * (x ^ 0.33)
fmla <- y ~ a * (x^b)
edta <- data.frame(x=x, y=y)
library(nlsr)
library(minpack.lm)
enls <- try(nls(fmla, start=c(a=1, b=1), data=edta))

## Error in nls(fmla, start = c(a = 1, b = 1), data = edta) :
##   number of iterations exceeded maximum of 50

enlxb <- try(nlxb(fmla, start=c(a=1, b=1), data=edta))
enlxb

## nlsr object: x
## residual sumsquares = 1.3353e-25 on 20 observations
##   after 6   Jacobian and 7 function evaluations
##   name      coeff      SE      tstat      pval      gradient      JSingval
## a           2.5      3.909e-14  6.395e+13  1.149e-238  -4.018e-13      60
## b           0.33      6.273e-15  5.261e+13  3.86e-237   2.548e-12      2.177

enlsLM <- try(nlsLM(fmla, start=c(a=1, b=1), data=edta))
enlsLM

## Nonlinear regression model
##   model: y ~ a * (x^b)
##   data: edta
##   a      b
## 2.50 0.33
## residual sum-of-squares: 0
##
## Number of iterations to convergence: 6
## Achieved convergence tolerance: 1.49e-08
```

References

- Levenberg, Kenneth. 1944. “A Method for the Solution of Certain Non-Linear Problems in Least Squares.” *Quarterly of Applied Mathematics* 2: 164–68.
- Marquardt, Donald W. 1963. “An Algorithm for Least-Squares Estimation of Nonlinear Parameters.” *SIAM Journal on Applied Mathematics* 11 (2): 431–41.
- Moré, Jorge J., Burton S. Garbow, and Kenneth E. Hillstom. 1981. “Testing Unconstrained Optimization Software.” *J-Toms* 7 (1): 17–41.

Nash, J. C. 1979. *Compact Numerical Methods for Computers : Linear Algebra and Function Minimisation*.
Book. Hilger, Bristol :