# Running funconstrain tests in package optimx

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#### Abstract

The funconstrain package (https://github.com/jlmelville/funconstrain) provides R users with a convenient tool to access the test functions of Moré, Garbow, and Hillstrom (1981). This vignette article describes a program to apply these test functions to solvers in the optimx package (Nash and Varadhan (2011)).

## Background

Numerical optimization of functions of several, namely n, parameters is an important computational task. R (R Development Core Team (2008)) is a major platform for scientific and statistical calculations and has provided tools for numerical optimization and nonlinear least squares since its inception. These have been extended via a number of packages. In particular, the author has been heavily involved in this effort, and in collaboration with others has provided the packate optimx which wraps a number of solvers to allow their invocation by a common calling syntax. Note that optimization in R generally means function minimization, possibly with bounds (or box) constraints on the function parameters.

It is extremely helpful to users to have examples and tests of function minimization. In many situations it is extremely easy to insert an error into code, so easy-to-apply tests allow for the discovery of such errors. There are a number of collections of test functions with many overlaps and minor differences. A well-established and well-documented set of such functions are those of Moré, Garbow, and Hillstrom (1981). These have been translated into R by James Melville in the R package funconstrain (https://github.com/jlmelville/funconstrain). While initially these provided the function and its gradient given a set of suitable input parameters, the present author added code to compute the Hessian for each test function. This allows Newton-like solvers to be applied. funconstrain also provides suggested initial parameter vectors for each of the 35 test functions. However, where there are multiple input possibilities, just one is provided, for example when the test function has a variable number of parameters.

What is then missing is the link between funconstrain and the tools in optimx, which this article aims to provide.

# Function fufn()

Most of the test functions in (More1981TU?) are sums of squares of nonlinear functions. While n is the number of parameters, we may have a different number of functions squared in the summation. Call this m. This may be altered to give different variations of a given function, so m must be provided.

Many of the solvers in optimx are capable of handling bounds constraints on the n parameters. That is parameter i must satisfy

lower[i] <= prm[i] <= upper[i]</pre>

where prm is the parameter vector and lower and upper are vectors of numbers providing lower and upper bounds. Methods in optimx that can handle masks are listed in the character vector bdmeth returned by the function optimx::ctrldefault(n). Note that a number of parameters n must nominally be provided to ctrldefault() but generally n can be specified as 2 to get the default settings for 'optimx. At time of writing

```
bdmeth <- c("L-BFGS-B", "nlminb", "lbfgsb3c", "Rcgmin", "Rtnmin", "nvm",
"Rvmmin", "bobyqa", "nmkb", "hjkb", "hjn", "snewtonm", "ncg",
"slsqp", "tnewt", "nlnm", "snewtm", "spg")</pre>
```

If the upper and lower bound for a parameter are equal, we can say the parameter is **fixed** or **masked**. This may seem to be a silly option, since it essentially reduces the dimensionality of the problem. However, there are many situations where we have evidence that a parameter takes a particular (fixed) value, but know that we may wish to allow optimization over that parameter in later investigations. Masks allow us to avoid having to rewrite the function, gradient and Hessian code. However, only a few optimization solvers handle masks. The function optimx::ctrldefault() returns a value maskmeth with a list of solvers that do handle the situation where lower and upper bounds coincide. At the time of writing this is specified as

```
maskmeth <- c("Rcgmin", "nvm", "hjn", "ncg", "snewtonm", "nlminb", "L-BFGS-B")
```

With the above in mind, the function fufn() was written to access the test functions of funconstrain.

# Calling fufn()

While we can write our own driver for fufn(), I wanted to make the task extremely easy. Thus the script fufnrun.R is provided. This is set up to use a simple text file, RFO.txt, to specify which test functions are to be applied to which solvers. Moreover, a "sink" file name can be specified to save the text output of the run. Note that the specification file is always called RFO.txt with the present incarnation of the fufnrun.R program.

### Test specification file RFO.txt

Let us consider an example.

```
testsink240408A.txt
1, 9, 9, 1, 6:8, 35
c("L-BFGS-B", "lbfgs", "lbfgsb3c", "lbfgs")
FALSE
```

The lines of the above file provide the following information:

- the first line is the name of the text file to use to save the output via sink().
- line 2 says that test functions 1, 6, 7, 8, and 35 are to be used. Note that we can use the colon ":" when giving a contiguous range of function numbers. These numbers by referring back to the vector funnam at the top of function fufn() specify functions "rosen", "jenn-samp", "helical", "bard" and "chebyquad". Using the function numbers. Appendix A lists the numbers and corresponding names. The specification 1:35 uses all test functions. The program removes duplicate problem numbers and sorts the list in ascending order.
- line 3 gives an R character vector of the solver methods to be applied. At the time of writing, there is no check for duplicate entries in the vector.

#### A driver program for fufn()

The following driver program will run the tests specified by RFO.txt:

```
# fufnrun.R -- J C Nash 2024-4-8
## ?? fixing kkt
# RFO.txt is input file
source("./fufn.R") # ensure fufn() loaded
library(funconstrain) # get the functions
library(optimx)
mycon<-file("RFO.txt", open="r", blocking = TRUE)</pre>
```

<sup>&#</sup>x27; line 4 is TRUE if the experimental bounds constraints are to be applied.

```
sfname <- readLines (mycon, n=1)
if (length(sfname) == 0) {
   cat("no sink file\n")
} else {
  cat("opening sink file ",sfname,"\n")
  sink(sfname, split=TRUE)
} # open sink file
## opening sink file testsink240408A.txt
cat("sink file name=",sfname,"\n")
## sink file name= testsink240408A.txt
lin2 <- readLines(mycon, n=1)</pre>
cat("probs =",lin2,"\n")
## probs = 1, 9, 9, 1, 6:8, 35
if (length(lin2) == 0) stop("Unexpected null probs")
txt<-paste("probc<-c(",lin2,")","")</pre>
tryparse<-eval(parse(text=txt))</pre>
# ?? should we check it worked?
cat("Problem numbers:\n"); print(probc)
## Problem numbers:
## [1] 1 9 9 1 6 7 8 35
print(unique(probc))
## [1] 1 9 6 7 8 35
if (length(unique(probc)) < length(probc)) {</pre>
   cat("Duplicated problem numbers, simplifying\n")
  probc <- unique(probc)</pre>
}
## Duplicated problem numbers, simplifying
probc<-sort(probc)</pre>
cat("Final problem numbers:\n"); print(probc)
## Final problem numbers:
## [1] 1 6 7 8 9 35
# check loop
for (iprob in probc){ # loop over problems
  if ((iprob < 1) | (iprob > 35)) {
    stop('Problem number out of range. Stopping.')
  }
} # end check loop
meths <- readLines(mycon, n=1)</pre>
if (length(meths) == 0) stop("Unexpected null meths")
cat("Methods:\n")
## Methods:
cat(meths,"\n")
```

```
## c("L-BFGS-B", "lbfgs", "lbfgsb3c", "lbfgs")
methvec<-paste("methc<-c(",meths,")","")</pre>
tryparse<-eval(parse(text=methvec))</pre>
if (length(unique(methc)) < length(methc)) {</pre>
  cat("Duplicated methods, simplifying\n")
 methc <- unique(methc)</pre>
}
## Duplicated methods, simplifying
cat("methods in list form:"); print(methc)
## methods in list form:
## [1] "L-BFGS-B" "lbfgs"
                              "lbfgsb3c"
tbounds <- readLines (mycon, n=1)
have.bounds<-FALSE
if (tbounds == "TRUE") have.bounds<-TRUE</pre>
cat("have.bounds:",have.bounds,"\n")
## have.bounds: FALSE
close(mycon)
for (iprob in probc){ # loop over problems
   tfun <- fufn(fnum=iprob)
   # print(tfun)
   cat("Problem:", tfun$fname,"\n")
   x0 <- tfun$x0
   if (have.bounds){
    lo <- tfun$lo</pre>
    up <- tfun$up
   else {
    lo <- -Inf
     up <- Inf
   tfn <- tfun$fffn
   attr(tfn, "fname") <- tfun$fname</pre>
   tgr <- tfun$ffgr
   the <- tfun$ffhe
  nx0 < -length(x0)
# cat("about to call opm\n")
   if (have.bounds) {
     t21 <-opm(x0, tfn, tgr, hess=the, lower=lo, upper=up, method=methc,
               contro=list(trace=0))
   } else {
     t21 <-opm(x0, tfn, tgr, hess=the, method=methc, contro=list(trace=0))
   print(summary(t21, order=value, par.select=1:min(nx0,5)))
   cat("END :", tfun$fname,"\n\n")
}
## Problem: rosen
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
```

```
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
##
                                p2 s2
                                              value fevals gevals hevals conv kkt1
## L-BFGS-B 0.9999997
                         0.999995
                                       2.267577e-13
                                                        47
                                                                47
                                                                        0
                                                                             O TRUE
            1.0000006
                         1.0000012
                                       3.545445e-13
                                                                        0
                                                                             O TRUE
## lbfgs
                                                        45
                                                                45
## lbfgsb3c 1.0000275
                         1.0000560
                                       8.551377e-10
                                                        74
                                                                74
                                                                        0
                                                                             O TRUE
##
            kkt2 xtime
## L-BFGS-B TRUE 0.004
## lbfgs
            TRUE 0.050
## lbfgsb3c TRUE 0.005
## END : rosen
## Problem: jenn_samp
## Error in optim(par = par, fn = efn, gr = egr, method = method, hessian = FALSE, :
     non-finite value supplied by optim
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
##
                                               value fevals gevals hevals conv
                   p1 s1
                                p2 s2
## lbfgs
            0.2578252
                         0.2578252
                                        1.243622e+02
                                                         67
                                                                 67
                                                                         0 -1001
## lbfgsb3c 0.2578252
                         0.2578252
                                        1.243622e+02
                                                                         0
                                                         45
                                                                 45
                                                                               0
## L-BFGS-B
                   NA
                                       8.988466e+307
                                                         24
                                                                 24
                                                                         0 9999
            kkt1 kkt2 xtime
##
## lbfgs
            TRUE TRUE 0.001
## lbfgsb3c TRUE TRUE 0.002
                   NA 0.001
## L-BFGS-B
              NA
## END : jenn_samp
##
## Problem: helical
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
                   p1 s1
                                    p2 s2
                                                      p3 s3
                                                                    value fevals
                                           -3.335635e-08
                                                            1.517549e-15
## lbfgs
            1.000000
                         -2.066737e-08
                                                                              34
## L-BFGS-B 0.9999999
                         -8.281311e-07
                                           -1.286731e-06
                                                            2.203966e-12
                                                                              30
                          5.628310e-06
                                            8.770209e-06
## lbfgsb3c 0.9999990
                                                            1.855335e-10
                                                                              53
##
            gevals hevals conv kkt1 kkt2 xtime
                             O TRUE TRUE 0.001
## lbfgs
                        0
                34
## L-BFGS-B
                             O TRUE TRUE 0.001
                30
                        0
## lbfgsb3c
                             O TRUE TRUE 0.001
                        0
                53
```

```
## END : helical
##
## Problem: bard
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
                                                         value fevals gevals hevals
                                p2 s2
                                             p3 s3
                    p1 s1
                                                   0.008214877
## L-BFGS-B 0.08241058
                          1.133036
                                       2.343695
                                                                   24
                                                                           24
                                                                                   0
                                                                   24
                                                                           24
                                                                                   0
## lbfgs
            0.08240992
                          1.133030
                                       2.343698
                                                   0.008214877
## lbfgsb3c 0.08232066
                          1.130012
                                       2.346610
                                                   0.008214943
                                                                   46
                                                                           46
                                                                                   0
##
            conv kkt1 kkt2 xtime
## L-BFGS-B
               O TRUE TRUE 0.001
               O TRUE TRUE 0.001
## lbfgs
## lbfgsb3c
               O TRUE TRUE 0.002
## END : bard
## Problem: gauss
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
##
                                                             value fevals gevals
                   p1 s1
                               p2 s2
                                                p3 s3
                                      1.429070e-20
## lbfgs
            0.3989563
                         1.000018
                                                      1.127976e-08
                                                                         9
                                                                                9
## L-BFGS-B 0.3989646
                         1.000102
                                      7.249508e-21
                                                      1.176721e-08
                                                                         4
                                                                                4
## lbfgsb3c 0.3989646
                         1.000102
                                      7.249508e-21
                                                      1.176721e-08
                                                                                4
##
            hevals conv kkt1 kkt2 xtime
                 Ω
                      O TRUE TRUE 0.001
## lbfgs
## L-BFGS-B
                      O TRUE TRUE 0.001
                 0
## lbfgsb3c
                      O TRUE TRUE 0.001
                 0
## END : gauss
## Problem: chebyquad
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing
##
                                               p3 s3
                                                                          p5 s5
                                 p2 s2
                                                            p4 s4
                    p1 s1
## L-BFGS-B 0.04315278
                          0.1930908
                                        0.2663287
                                                     0.4999999
```

```
## lbfgs
                         0.1930910
                                      0.2663289
                                                   0.4999998
                                                                0.5000002
           0.04315297
                                                                0.5000558
## lbfgsb3c 0.04315505
                         0.1930877
                                      0.2663283
                                                   0.4999442
                 value fevals gevals hevals conv kkt1 kkt2 xtime
## L-BFGS-B 0.003516874
                           28
                                          0
                                               O TRUE TRUE 0.002
                                  28
                           25
                                  25
## lbfgs
         0.003516874
                                          0
                                               O TRUE TRUE 0.001
## lbfgsb3c 0.003516880
                           46
                                  46
                                          0
                                               O TRUE TRUE 0.002
## END : chebyquad
```

## Appendix A: function numbers and names

```
1
        rosen
2
        freud_roth
        powell_bs
3
4
        brown_bs
5
        beale
6
        jenn_samp
7
        helical
8
        bard
9
        gauss
10
        meyer
11
        gulf
12
        box 3d
13
        powell_s
14
        wood
15
        kow_osb
16
        brown_den
17
        osborne_1
18
        biggs_exp6
19
        osborne 2
20
        watson
21
        ex_rosen
22
        ex_powell
23
        penalty_1
24
        penalty_2
25
        var_dim
26
        trigon
27
        brown_al
28
        disc_bv
29
        disc ie
30
        broyden_tri
31
        broyden band
32
        linfun_fr
33
        linfun r1
34
        linfun_r1z
35
        chebyquad
```

## Appendix B: the fufn.R code

```
fufn <- function(fnum=NULL){
    # return list with tfn=function, tgr=gradient given fn number and n
    if (is.null(fnum)) stop("ffn needs a function number fnum")
    if ((fnum < 1) || (fnum > 35)) stop("fnum must be in [1, 35]")
```

```
# cat("entering ffn, fnum=",fnum,"\n")
  # select function
  funnam <- c("rosen", "freud_roth", "powell_bs", "brown_bs", "beale",</pre>
               "jenn_samp", "helical", "bard", "gauss", "meyer", "gulf",
               "box_3d", "powell_s", "wood", "kow_osb", "brown_den",
              "osborne_1", "biggs_exp6", "osborne_2", "watson", "ex_rosen",
               "ex_powell", "penalty_1", "penalty_2", "var_dim", "trigon",
               "brown al", "disc bv", "disc ie", "broyden tri", "broyden band",
               "linfun_fr", "linfun_r1", "linfun_r1z", "chebyquad")
# print(str(funnam))
 fname <- funnam[as.integer(fnum)]</pre>
# cat("fname:", fname,"\n")
  while (fnum %in% 1:35) {
    ameth <- optimx::ctrldefault(2)$bdmeth # Choose only bounded methods
    ameth <- ameth[ameth != "lbfgsb3c"] ## ?? Temporarily remove lbfgsb3c</pre>
    ameth <- c(ameth, "L-BFGS-B")</pre>
    # ?? may want to test allmeth to check that inappropriate methods are captured
    # cat("in while, fnum=", fnum); tmp <- readline("cont.")</pre>
    mm <- 0 # in case m value needed
  if (fnum == 1) {
     n <- 2 # fixed
    mm <- 2
    tt <- rosen()
     if (is.function(tt$x0)) {
      xx0<-tt$x0(n)
     else xx0 <- tt$x0
     lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
     break }
  if (fnum == 2) {
     n <- 2 # fixed
     mm <- 2
    tt <- freud_roth()</pre>
     if (is.function(tt$x0)) {
      xx0<-tt$x0(n)
     else xx0 <- tt$x0</pre>
     lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
    break }
  if (fnum == 3) {
    n <- 2 # fixed
    mm <- 2
     tt <- powell_bs()</pre>
     if (is.function(tt$x0)) {
      xx0<-tt$x0(n)
     else xx0 <- tt$x0</pre>
    lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
```

```
break }
  if (fnum == 4) {
     n <- 2 # fixed
     mm <- 3
     tt <- brown_bs()</pre>
     if (is.function(tt$x0)) {
       xx0<-tt$x0(n)
     else xx0 <- tt$x0</pre>
## BAD -- reset 20240323
    lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
     lo <- -1e20
     up <- -lo
     break }
  if (fnum == 5) {
     n <- 2 # fixed
     mm <- 3
     tt <- beale()</pre>
     if (is.function(tt$x0)) {
      xx0<-tt$x0(n)
     else xx0 <- tt$x0
     lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
     break }
  if (fnum == 6) {
     n <- 2 # fixed
     mm <- 10
     tt <- jenn_samp()</pre>
     if (is.function(tt$x0)) {
       xx0<-tt$x0(n)
     else xx0 <- tt$x0</pre>
     lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
     break }
  if (fnum == 7) {
     n <- 3 # fixed
     tt <- helical()
     if (is.function(tt$x0)) {
       xx0<-tt$x0(n)
     else xx0 <- tt$x0</pre>
     lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
     break }
  if (fnum == 8) {
```

```
n <- 3 # fixed
   mm <- 15
   tt <- bard()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 9) {
   n <- 3 # fixed
   mm <- 15
   tt <- gauss()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 10) {
   n <- 3 # fixed
   m <- 16 # ?? how to return
   tt <- meyer()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 11) {
   n <- 3
   mm <- 99
   tt <- gulf()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   }
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 12) {
  n <- 3
   mm <- 20
   tt <- box_3d()
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
```

```
else xx0 <- tt$x0</pre>
   lo <- rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 13) {
   n <- 4
   tt <- powell_s()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 14) {
   n <- 4
   tt <- wood()
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 15) {
   mm <- 11
   n <- 4
   tt <- kow_osb()
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   }
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 16) {
   mm <- 20
   n <- 4
   tt <- brown_den()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 17) {
```

```
mm <- 33
   n <- 5
   tt <- osborne 1()
   ameth<-ameth[-which(ameth=="L-BFGS-B")] # remove L-BFGS-B from this case
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   lo[4] <- 0
   lo[5] <- 0
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 18) {
   mm <- 20
   n <- 6
   tt <- biggs_exp6()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   }
   else xx0 <- tt$x0
   lo \leftarrow rep((min(xx0)-0.1), n)
   up - rep((max(xx0)+0.1), n)
   break }
if (fnum == 19) {
   mm <- 65
   n <- 11
   tt <- osborne_2()
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up - rep((max(xx0)+0.1), n)
   break }
if (fnum == 20) {
   n <-8
   mm <- 31
   tt <- watson()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 21) {
   n <- 10
   tt <- ex_rosen()</pre>
```

```
if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo <- rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 22) {
   n <- 20
   tt <- ex_powell()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 23) {
   n <- 10
   mm \leftarrow n + 1
   tt <- penalty_1()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up - rep((max(xx0)+0.1), n)
   break }
if (fnum == 24) {
   n <- 10
   mm \leftarrow n + 1
   tt <- penalty_2()
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 25) {
   n <- 6
   mm \leftarrow n + 2
   tt <- var_dim()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
```

```
break }
if (fnum == 26) {
   n <- 8
   tt <- trigon()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 27) {
   n <- 8
   mm <- n
   tt <- brown_al()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   }
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 28) {
   n <- 6
   mm <- n
   tt <- disc_bv()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 29) {
   n <- 8
   mm <- n
   tt <- disc_ie()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo <- rep((min(xx0)-0.1), n)
   up - rep((max(xx0)+0.1), n)
   break }
if (fnum == 30) {
   n <- 8
   mm <- n
   tt <- broyden_tri()</pre>
```

```
if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo <- rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 31) {
   n <- 8
   mm <- n
   tt <- broyden_band()</pre>
   if (is.function(tt$x0)) {
     xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo \leftarrow rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 32) {
   mm <- 10
   n <- 8
   tt <- linfun_fr()</pre>
   if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
   else xx0 <- tt$x0</pre>
   lo <- rep((min(xx0)-0.1), n)
   up \leftarrow rep((max(xx0)+0.1), n)
   break }
if (fnum == 33) {
  mm <- 10
  n <- 8
  tt <- linfun_r1()</pre>
  if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
  }
  else xx0 <- tt$x0
  lo <- rep((min(xx0)-0.1), n)
  up \leftarrow rep((max(xx0)+0.1), n)
  break }
if (fnum == 34) {
 mm <- 10
 n <- 8
  tt <- linfun_r1z()</pre>
  if (is.function(tt$x0)) {
    xx0<-tt$x0(n)
  }
  else xx0 <- tt$x0</pre>
  lo \leftarrow rep((min(xx0)-0.1), n)
```

```
up \leftarrow rep((max(xx0)+0.1), n)
    break }
  if (fnum == 35) {
     n <- 8
     m <- n
     tt <- chebyquad()
     if (is.function(tt$x0)) {
       xx0<-tt$x0(n)
     else xx0 <- tt$x0</pre>
     lo \leftarrow rep((min(xx0)-0.1), n)
     up \leftarrow rep((max(xx0)+0.1), n)
     break }
  }
# NOTE: bounds are experimental only
  mask <- rep(1L, n) # masks set to "free" (not masked)</pre>
  val <- list(npar = n, fffn=tt$fn, ffgr=tt$gr, x0=xx0, lo=lo, up=up,</pre>
               mask=mask, fname=fname, ameth=ameth)
# cat("val:"); print(val); tmp<-readline('exit ffn')</pre>
  val
} # end fufn
```

#### References

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