

```
> library(knitr)
> opts_chunk$set(
+ concordance=TRUE
+ )
```

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```

Optimization problems constrained by parameter sums

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Abstract

This article presents a discussion of optimization problems where the objective function $f(\mathbf{x})$ has parameters that are constrained by some scaling, so that $q(\mathbf{x}) = \text{constant}$, where this function $q()$ involves a sum of the parameters, their squares, or similar simple function.

1 Background

We consider problems where we want to minimize or maximize a function subject to a constraint that the sum of some function of the parameters, e.g., their sum of squares, must equal some constant. We refer to these problems as **sumscale** optimization problems. We have observed questions about problems like this on the R-help mailing list:

```
Jul 19, 2012 at 10:24 AM, Linh Tran <Tranlm@berkeley.edu> wrote:
> Hi fellow R users,
>
> I am desperately hoping there is an easy way to do this in R.
>
> Say I have three functions:
>
> f(x) = x^2
> f(y) = 2y^2
> f(z) = 3z^2
```

```

>
> constrained such that x+y+z=c (let c=1 for simplicity).
>
> I want to find the values of x,y,z that will minimize
f(x) + f(y) + f(z).

```

If the parameters x , y and z are non-negative, this problem can actually be solved as a Quadratic Program. We revisit this problem at the end of this article.

Other examples of this type of objective function are:

The maximum volume of a regular polyhedron where the sum of the lengths of the sides is fixed.

The minimum negative log likelihood for a multinomial model.

The Rayleigh Quotient for the maximal or minimal eigensolutions of a matrix, where the eigenvectors should be normalized so the square norm of the vector is 1.

For the moment, let us consider a basic example, which is

Problem A : Minimize $(-\prod \mathbf{x})$ subject to $\sum \mathbf{x} = 1$

This is a very simplified version of the multinomial maximum likelihood problem.

Because these problems all have an objective that is dependent on a scaled set of parameters where the scale is defined by a sum, sum of squares, or similar sum of the parameters, we will refer to them as **sumscale** optimization problems.

2 Difficulties using general optimization with sum-scale problems

Let us use the basic example above to consider how we might formulate Problem A for a computational solution in R.

One possibility is to select one of the parameters and solve for it in terms of the others. Let this be the last parameter x_n , so that the set of parameters

to be optimized is $\mathbf{y} = (x_1, x_1, \dots, x_{n-1})$ where n is the original size of our problem. We now have the unconstrained problem

$$\text{minimize}(-(\prod \mathbf{y}) * (1 - \sum y))$$

This is easily coded and tried. We will use a very simple start, namely, the sequence $1, 2, \dots, (n - 1)$ scaled by $1/n^2$. We will also specify that the gradient is to be computed by a central approximation (Nash, 2013).

```
> cat("try loading optimrx\n")

try loading optimrx

> require(optimrx, quietly=TRUE)
> pr <- function(y) {
+ - prod(y)*(1-sum(y))
+ }
> cat("test the simple product for n=5\n")

test the simple product for n=5

> meth <- c("Nelder-Mead", "BFGS")
> n<-5
> st<-1:(n-1)/(n*n)
> ans<-opm(st, pr, gr="grcentral", control=list(trace=0))
> ao<-summary(ans,order=value)
> print(ao)
```

	p1	p2	p3	p4	value	fevals
Nelder-Mead	0.2000034	0.1999983	0.2000017	0.2000021	-3.200000e-04	331
BFGS	NA	NA	NA	NA	8.988466e+307	NA

```

      gevals convergence kkt1 kkt2 xtime
Nelder-Mead      NA           0 TRUE TRUE 0.008
BFGS             NA      9999  NA  NA 0.004
```

While these codes work fine for small n , it is fairly easy to see that there are computational problems as the size of the problem increases. Since the sum of the parameters is constrained to be equal to 1, the parameters are of the order of $1/n$, and the function therefore of the order of $1/(n^n)$, which underflows around $n = 144$ in R.

3 Other formulations

Traditionally, statisticians solve maximum likelihood problems by **minimizing** the negative log-likelihood. That is, the objective function is formed as (-1) times the logarithm of the likelihood. This converts our product to a sum. Choosing the first parameter to be the one determined by the summation constraint, we can write the function and gradient quite easily. As programs that try to find the minimum may change the parameters so that logarithms of non-positive numbers are attempted, we have put some safeguards in the function `nll`. At this point we have assumed the gradient calculation is only attempted if the function can be computed satisfactorily, so we have not put safeguards in the gradient.

```
> nll <- function(y) {
+   if ((any(y <= 10*.Machine$double.xmin)) || (sum(y)>1-.Machine$double.eps))
+       .Machine$double.xmax
+   else   - sum(log(y)) - log(1-sum(y))
+ }
> nll.g <- function(y) { - 1/y + 1/(1-sum(y)) } # so far not safeguarded
```

We can easily try several optimization methods using the `optimx` package. Here are the calls, which overall did not perform as well as we would like. Note that we do not ask for `method="ALL"` as we found that some of the methods, in particular those using Powell's quadratic approximation methods, seem to get "stuck".

```
> require(optimrx, quietly=TRUE)
> n<-5
> mset<-c("L-BFGS-B", "BFGS", "CG", "spg", "ucminf", "nlm", "nlminb", "Rvmmin", "Rcgmin")
> a5<-opm(2:n/n^2, nll, gr="grfwd", method=mset, control=list(dowarn=FALSE))
> a5g<-opm(2:n/n^2, nll, nll.g, method=mset, control=list(dowarn=FALSE))
> a5gb<-opm(2:n/n^2, nll, nll.g, lower=0, upper=1, method=mset, control=list(dowarn=FALSE))
> #- a5x <- opm(2:n/n^2, nll, nll.g, method="ALL", control=list(dowarn=FALSE))
> summary(a5,order=value)
```

	p1	p2	p3	p4	value	fevals	gevals
nlminb	0.1910859	0.1858287	0.2032001	0.2296229	8.062194e+00	40	4
Rcgmin	0.1912221	0.1859094	0.2032530	0.2296592	8.062239e+00	2237	136
nlm	0.1925000	0.1866667	0.2037500	0.2300000	8.062829e+00	NA	3
Rvmmin	0.1945455	0.1878788	0.2045455	0.2305455	8.064408e+00	37	4
spg	0.2050000	0.1940741	0.2086111	0.2333333	8.085799e+00	1213	11

ucminf	0.2379245	0.2135849	0.2214151	0.2421132	8.371136e+00	6	6
L-BFGS-B	NA	NA	NA	NA	8.988466e+307	NA	NA
BFGS	NA	NA	NA	NA	8.988466e+307	NA	NA
CG	NA	NA	NA	NA	8.988466e+307	NA	NA

	convergence	kkt1	kkt2	xtime
nlminb	1	FALSE	TRUE	0.000
Rcgmin	1	FALSE	TRUE	0.124
nlm	0	FALSE	TRUE	0.004
Rvmmin	3	FALSE	TRUE	0.004
spg	0	FALSE	TRUE	0.148
ucminf	0	FALSE	TRUE	0.004
L-BFGS-B	9999	NA	NA	0.000
BFGS	9999	NA	NA	0.000
CG	9999	NA	NA	0.000

> summary(a5g,order=value)

	p1	p2	p3	p4	value	fevals	gevals
ucminf	0.2000000	0.2000000	0.2000000	0.2000000	8.047190e+00	14	14
Rvmmin	0.2000000	0.2000000	0.2000000	0.2000000	8.047190e+00	43	12
spg	0.2000000	0.2000000	0.2000000	0.2000000	8.047190e+00	17	13
Rcgmin	0.2000000	0.2000000	0.2000000	0.2000000	8.047190e+00	28	12
CG	0.2000000	0.2000000	0.2000000	0.2000000	8.047190e+00	59	21
nlm	0.2000006	0.1999995	0.2000000	0.2000000	8.047190e+00	NA	11
BFGS	0.2000007	0.1999989	0.2000012	0.1999981	8.047190e+00	33	9
nlminb	0.2000004	0.1999990	0.1999989	0.1999992	8.047190e+00	23	12
L-BFGS-B	NA	NA	NA	NA	8.988466e+307	NA	NA

	convergence	kkt1	kkt2	xtime
ucminf	0	TRUE	TRUE	0.000
Rvmmin	0	TRUE	TRUE	0.008
spg	0	TRUE	TRUE	0.064
Rcgmin	0	TRUE	TRUE	0.000
CG	0	TRUE	TRUE	0.000
nlm	0	TRUE	TRUE	0.004
BFGS	0	TRUE	TRUE	0.000
nlminb	0	TRUE	TRUE	0.000
L-BFGS-B	9999	NA	NA	0.000

> summary(a5gb,order=value)

	p1	p2	p3	p4	value	fevals	gevals
Rvmmin	0.2000000	0.2000000	0.2000000	0.2000000	8.047190e+00	38	14

Rcgmin	0.2000000	0.200000	0.2000000	0.2000000	8.047190e+00	18	10
spg	0.2000000	0.200000	0.2000000	0.2000000	8.047190e+00	18	13
nlminb	0.2000004	0.199999	0.1999989	0.1999992	8.047190e+00	23	12
L-BFGS-B	NA	NA	NA	NA	8.988466e+307	NA	NA
BFGS	NA	NA	NA	NA	8.988466e+307	NA	NA
CG	NA	NA	NA	NA	8.988466e+307	NA	NA
ucminf	NA	NA	NA	NA	8.988466e+307	NA	NA
nlm	NA	NA	NA	NA	8.988466e+307	NA	NA
	convergence	kkt1	kkt2	xtime			
Rvmmin	0	TRUE	TRUE	0.012			
Rcgmin	0	TRUE	TRUE	0.008			
spg	0	TRUE	TRUE	0.072			
nlminb	0	TRUE	TRUE	0.000			
L-BFGS-B	9999	NA	NA	0.000			
BFGS	9999	NA	NA	0.004			
CG	9999	NA	NA	0.000			
ucminf	9999	NA	NA	0.000			
nlm	9999	NA	NA	0.000			

```
> #- summary(a5x,order=value)
```

Most, but not all, of the methods find the solution for the $n = 5$ case. The exception (L-BFGS-B) is due to the optimization method trying to compute the gradient where $\text{sum}(x)$ is greater than 1. We have not tried to determine the source of this particular issue. However, it is almost certainly a consequence of too large a step. The particular form of $\log(1 - \text{sum}(x))$ is undefined once the argument of the logarithm is negative. Indeed, this is the basis of logarithmic barrier functions for constraints. There is a similar issue with the $n - 1$ parameters near zero. Negative values will cause difficulties.

Numerical gradient approximations will similarly fail, particularly as step sizes are often of the order of $1\text{E-}7$ in size. There is generally no special check within numerical gradient routines to apply bounds. Note also that a lower bound of 0 on parameters is not adequate, since $\log(0)$ is undefined. Choosing a bound large enough to avoid the logarithm of a zero or negative argument while still being small enough to allow for parameter optimization is non-trivial.

4 Transformed problems or parameters

When problems give difficulties, it is common to re-formulate them by transformations of the function or the parameters.

4.1 Using a projection

Objective functions defined by $(-1)*\prod \mathbf{x}$ or $(-1)*\sum \log(\mathbf{x})$ will change with the scale of the parameters. Moreover, the constraint $\sum \mathbf{x} = 1$ effectively imposes the scaling $\mathbf{x}_{\text{scaled}} = \mathbf{x} / \sum \mathbf{x}$. The optimizer `spg` from package `BB` allows us to project our search direction to satisfy constraints. Thus, we could use the following approach. Thanks to Ravi Varadhan for the suggestion.

```
> require(BB, quietly=TRUE)
> nllrv <- function(x) {- sum(log(x))}
> nllrv.g <- function(x) {- 1/x }
> proj <- function(x) {x/sum(x)}
> n <- 5
> tspg<-system.time(aspgr <- spg(par=(1:n)/n^2, fn=nllrv, gr=nllrv.g, project=proj))[[1]]

iter: 0 f-value: 11.30689 pgrad: 0.3607565

> tspgn<-system.time(aspgn <- spg(par=(1:n)/n^2, fn=nllrv, project=proj))[[3]]

iter: 0 f-value: 11.30689 pgrad: 0.1333334

> cat("Times: with gradient =",tspg," using numerical approx.=", tspgn,"\n")

Times: with gradient = 0.085 using numerical approx.= 0.065

> cat("F_optimal: with gradient=",aspgr$value," num. approx.=",aspgn$value,"\n")

F_optimal: with gradient= 8.04719 num. approx.= 8.04719

> pbest<-rep(1/n, n)
> cat("fbest = ",nllrv(pbest)," when all parameters = ", pbest[1],"\n")

fbest = 8.04719 when all parameters = 0.2

> cat("deviations: with gradient=",max(abs(aspgr$par-pbest)), " num. approx.=",max(a

deviations: with gradient= 3.81244e-06 num. approx.= 3.81244e-06
```


Here the projection `proj` is the key to success of method `spg`. Other methods do not have the flexibility to impose the projection directly. We would need to carefully build the projection into the function(s) and/or the method codes. This was done by Geradin (1971) for the Rayleigh quotient problem, but requires a number of changes to the program code.

4.2 `log()` transformation of parameters

A common method to ensure parameters are positive is to transform them. In the present case, optimizing over parameters that are the logarithms of the parameters above ensures we have positive arguments to most of the elements of the negative log likelihood. Here is the code. Note that the parameters used in optimization are "lx" and not x.

```
> enll <- function(lx) {
+   x<-exp(lx)
+   fval<- - sum( log( x/sum(x) ) )
+ }
> enll.g <- function(lx){
+   x<-exp(lx)
+   g<-length(x)/sum(x) - 1/x
+   gval<-g*exp(lx)
+ }
```

But where is our constraint? Here we have noted that we could define the objective function only to within the scaling $\mathbf{x}/\sum(\mathbf{x})$. There is a minor nuisance, in that we need to re-scale our parameters after solution to have them in a standard form. This is most noticeable if one uses `optimx` and displays the results of `all.methods`. In the following, we extract the best solution for the 5-parameter problem.

```
> require(optimrx, quietly=TRUE) # just to be sure
> st<-1:5/10 # 5 parameters, crude scaling to start
> a5x<-opm(st, enll, enll.g, method="ALL", control=list(trace=0))
> a5xbyvalue<-summary(a5x, order=value)
> xnor<-a5xbyvalue[1, 1:5] # get the 5 parameters of "best" solution
> xnor<-xnor/sum(xnor)
> cat("normalized parameters:")
```

normalized parameters:

```
> print(xnor)
```

```
      p1 p2 p3 p4 p5
BFGS 0.2 0.2 0.2 0.2 0.2
```

While there are reasons to think that the indeterminacy might upset the optimization codes, in practice, the objective and gradient above are generally well-behaved, though they did reveal that tests of the size of the gradient used, in particular, to decide to terminate iterations in `Rcgmin` were too hasty in stopping progress for problems with larger numbers of parameters. A user-specified tolerance is now allowed; for example `control=list(tol=1e-12)`.

Let us try a larger problem in 100 parameters.

```
> require(Rcgmin, quietly=TRUE)
> st<-1:100/1e3 # large
> stenll<-enll(st)
> cat("Initial function value =",stenll,"\n")
```

```
Initial function value = 460.5587
```

```
> tym<-system.time(acgbig<-Rcgmin(st, enll, enll.g, control=list(trace=0, tol=1e-32))
> cat("Time = ",tym," fval=",acgbig$value,"\n")
```

```
Time = 0.192 fval= 460.517
```

```
> xnor<-acgbig$par/sum(acgbig$par)
> print(xnor)
```

```
[1] 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01
[16] 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01
[31] 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01
[46] 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01
[61] 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01
[76] 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01
[91] 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01
```

One worrying aspect of the solution is that the objective function at the start and end differ by a tiny amount.

4.3 Another transformation

A slightly different transformation or projection is inspired by spherical coordinates.

```
> proj2 <- function(theta) {
+   theta2 <- theta^2
+   s2 <- theta2 / (1 + theta2)
+   cumprod(c(1, s2)) * c(1-s2, 1)
+ }
> obj <- function(theta) - sum(log(proj2(theta)))
> n <- 5
> ans <- spg(seq(n-1), obj)

iter: 0 f-value: 11.15175 pgrad: 3
iter: 10 f-value: 8.78015 pgrad: 0.5806909
iter: 20 f-value: 8.04719 pgrad: 3.925749e-06

> proj2(ans$par)

[1] 0.2000000 0.2000007 0.2000002 0.1999996 0.1999995

> n<-100
> ans100 <- spg(seq(n-1), obj, control=list(trace=FALSE), quiet=TRUE)
> proj2( (ans100$par) )

[1] 0.009999999 0.010000001 0.010000000 0.010000000 0.009999999 0.010000003
[7] 0.010000000 0.010000001 0.010000000 0.010000000 0.010000001 0.010000000
[13] 0.010000000 0.010000002 0.010000002 0.010000001 0.009999999 0.010000002
[19] 0.010000000 0.010000002 0.010000000 0.010000000 0.010000002 0.010000001
[25] 0.010000001 0.010000001 0.010000000 0.010000002 0.010000001 0.010000001
[31] 0.010000000 0.010000001 0.010000002 0.010000002 0.010000001 0.010000002
[37] 0.010000001 0.010000000 0.010000001 0.010000001 0.009999999 0.009999996
[43] 0.010000002 0.010000002 0.010000002 0.009999996 0.009999997 0.009999999
[49] 0.010000000 0.009999999 0.009999998 0.010000002 0.010000000 0.009999999
[55] 0.010000000 0.009999996 0.010000002 0.010000002 0.009999998 0.010000000
[61] 0.010000000 0.009999999 0.010000000 0.010000000 0.010000001 0.009999999
[67] 0.010000000 0.010000000 0.010000000 0.010000000 0.009999999 0.010000000
[73] 0.009999999 0.010000000 0.009999999 0.010000000 0.010000000 0.009999999
[79] 0.010000000 0.009999999 0.009999999 0.009999999 0.010000000 0.010000001
[85] 0.009999999 0.009999999 0.010000000 0.010000001 0.010000000 0.009999998
[91] 0.009999998 0.009999999 0.010000000 0.010000000 0.010000000 0.009999997
[97] 0.010000038 0.009999935 0.010000003 0.010000023
```

Since this transformation is embedded into the objective function, we could run all the optimizers in `optimx` as follows. This takes some time, as the derivative-free methods appear to have more difficulty with this formulation. Moreover, `Rcgmin` and `Rvmmmin` are not recommended when an analytic gradient is not provided.

```
> allans<- opm(seq(n-1), obj, gr="grfwd", method="ALL", control=list(dowarn=FALSE))
> summary(allans, order = "list(round(value, 3), fevals)", par.select = FALSE)
```

	value	fevals	gevals	convergence	kkt1	kkt2	xtime
hjkb	4.605170e+02	21008	NA	0	TRUE	TRUE	0.968
bobyqa	4.605170e+02	21493	NA	0	TRUE	TRUE	17.372
hjn	4.605170e+02	23292	NA	0	TRUE	TRUE	1.360
nmkb	7.202616e+02	5045	NA	1	TRUE	FALSE	5.356
Nelder-Mead	7.439745e+02	5002	NA	1	TRUE	FALSE	0.220
Rtnmin	7.556838e+02	8	8	3	TRUE	FALSE	0.128
Rcgmin	7.557004e+02	7	2	0	FALSE	FALSE	0.008
lbfgsb3	7.557324e+02	30	30	0	TRUE	FALSE	0.144
lbfgs	7.557324e+02	NA	NA	-1001	FALSE	FALSE	0.084
Rvmmmin	7.558641e+02	64	4	3	TRUE	FALSE	0.024
nlm	7.558641e+02	NA	2	0	TRUE	FALSE	0.544
nlminb	7.618772e+02	44	44	1	FALSE	FALSE	0.188
ucminf	7.713573e+02	8	8	0	FALSE	FALSE	0.052
spg	7.747860e+02	1033	11	0	TRUE	FALSE	0.248
BFGS	8.988466e+307	NA	NA	9999	NA	NA	0.004
CG	8.988466e+307	NA	NA	9999	NA	NA	0.004
L-BFGS-B	8.988466e+307	NA	NA	9999	NA	NA	0.000
newuoa	8.988466e+307	NA	NA	9999	NA	NA	0.008

4.4 Use the gradient equations

Another approach is to "solve" the gradient equations. We can do this with a sum of squares minimizer, though the `nls` function in R is specifically NOT useful as it cannot deal with small or zero residuals. However, `nlfb` from package `nlmrt` is capable of dealing with such problems. Unfortunately, it will be slow as it has to generate the Jacobian by numerical approximation unless we can provide a function to prepare the Jacobian analytically. Moreover, the determination of the Jacobian is still subject to the unfortunate scaling issues we have been confronting throughout this article.

5 The Rayleigh Quotient

The maximal and minimal eigensolutions of a symmetric matrix A are extrema of the Rayleigh Quotient

$$R(x) = (x'Ax)/(x'x)$$

We can also deal with generalized eigenproblems of the form

$$Ax = eBx$$

where B is symmetric and positive definite by using the Rayleigh Quotient

$$R_g(x) = (x'Ax)/(x'Bx)$$

Once again, the objective is scaled by the parameters, this time by their sum of squares. Alternatively, we may think of requiring the **normalized** eigensolution, which is given as

$$x_{normalized} = x/\sqrt{x'x}$$

We will first try the projected gradient method `spg` from `BB`. Below is the code, where our test uses a matrix called the Moler matrix (Nash, 1979, Appendix 1). We caution that there are faster ways to compute this matrix in R (Nash, 2012) where different approaches to speed up R computations are discussed. Here we are concerned with getting the solutions correctly rather than the speed of so doing. Note that to get the solution with the most-positive eigenvalue, we minimize the Rayleigh quotient of the matrix multiplied by -1. This is solution `tmax`.

```
> molerbuidl<-function(n){ # Create the moler matrix of order n
+   # A[i,j] = i for i=j, min(i,j)-2 otherwise
+   A <- matrix(0, nrow = n, ncol = n)
+   j <- 1:n
+   for (i in 1:n) {
+     A[i, 1:i] <- pmin(i, 1:i) - 2
+   }
+   A <- A + t(A)
+   diag(A) <- 1:n
+   A
+ }
> raynum<-function(x, A){
+   rayquo<-as.numeric((t(x)%*%A)%*%x)
+ }
> proj<-function(x) { x/sqrt(crossprod(x)) }
> require(BB, quietly=TRUE)
> n<-10
```

```

> x<-rep(1,n)
> A<-molerbuild(n)
> tmin<-system.time(asprqmin<-spg(x, fn=raynum, project=proj, A=A))[[3]]

iter: 0 f-value: 205 pgrad: 3.089431e-09

> tmax<-system.time(asprqmax<-spg(x, fn=raynum, project=proj, A=-A))[[3]]

iter: 0 f-value: -205 pgrad: 0.6324555

> cat("maximal eigensolution: Value=",asprqmax$value,"in time ",tmax,"\n")

maximal eigensolution: Value= -205 in time 0.7

> print(asprqmax$par)

[1] 0.3162278 0.3162278 0.3162278 0.3162278 0.3162278 0.3162278 0.3162278
[8] 0.3162278 0.3162278 0.3162278

> cat("minimal eigensolution: Value=",asprqmin$value,"in time ",tmin,"\n")

minimal eigensolution: Value= 205 in time 0.078

> print(asprqmin$par)

[1] 0.3162278 0.3162278 0.3162278 0.3162278 0.3162278 0.3162278 0.3162278
[8] 0.3162278 0.3162278 0.3162278

```

For the record, these results compare well with eigenvalues from `eigen()`.

If we ignore the constraint, and simply perform the optimization, we can get satisfactory solutions, though comparisons require that we normalize the parameters post-optimization. We can check if the scale of the eigenvectors is becoming large by computing the norm of the final parameter vector. In tests on the Moler matrix up to dimension 100, none grew to a worrying size.

For comparison, we also ran a specialized Geradin routine as implemented in R by one of us (JN). This gave equivalent answers, albeit more efficiently. For those interested, the Geradin routine is available as referenced in (Nash, 2012).

6 The R-help example

As a final example, let us use our present techniques to solve the problem posed by Lanh Tran on R-help. We will use only a method that scales the parameters directly inside the objective function and not bother with gradients for this small problem.

```
> ssums<-function(x){
+   n<-length(x)
+   tt<-sum(x)
+   ss<-1:n
+   xx<-(x/tt)*(x/tt)
+   sum(ss*xx)
+ }
> cat("Try penalized sum\n")
```

Try penalized sum

```
> require(optimx)
> st<-runif(3)
> aos<-opm(st, ssums, gr="grcentral", method="ALL")
> # rescale the parameters
> nsol<-dim(aos)[1]
> for (i in 1:nsol){
+   tpar<-aos[i,1:3]
+   ntpar<-sum(tpar)
+   tpar<-tpar/ntpar
+   # cat("Method ",aos[i, "meth"]," gives fval =", ssums(tpar))
+   aos[i, 1:3]<-tpar
+ }
> summary(aos,order=value)[1:5,]
```

	p1	p2	p3	value	fevals	gevals	convergence
hjn	0.5454545	0.2727273	0.1818182	0.5454545	238	NA	0
newuoa	0.5454545	0.2727273	0.1818182	0.5454545	56	NA	0
bobyqa	0.5454546	0.2727273	0.1818182	0.5454545	50	NA	0
hjkb	0.5454545	0.2727273	0.1818181	0.5454545	326	NA	0
Nelder-Mead	0.5455187	0.2726870	0.1817943	0.5454546	88	NA	0
	kkt1	kkt2	xtime				
hjn	TRUE	FALSE	0.008				

```

newuoa      TRUE FALSE 0.004
bobyqa      TRUE FALSE 0.000
hjk        TRUE FALSE 0.008
Nelder-Mead TRUE FALSE 0.000

> ssum<-function(x){
+   n<-length(x)
+   ss<-1:n
+   xx<-x*x
+   sum(ss*xx)
+ }
> proj.simplex <- function(y) {
+ # project an n-dim vector y to the simplex Dn
+ # Dn = { x : x n-dim, 1 >= x >= 0, sum(x) = 1}
+ # Ravi Varadhan, Johns Hopkins University
+ # August 8, 2012
+
+ n <- length(y)
+ sy <- sort(y, decreasing=TRUE)
+ csy <- cumsum(sy)
+ rho <- max(which(sy > (csy - 1)/(1:n)))
+ theta <- (csy[rho] - 1) / rho
+ return(pmax(0, y - theta))
+ }
> as<-spg(st, ssum, project=proj.simplex)

iter:  0 f-value:  1.691001 pgrad:  0.8434151

> cat("Using project.simplex with spg: fmin=",as$value," at \n")

Using project.simplex with spg: fmin= 0.5454545  at

> print(as$par)

[1] 0.5454559 0.2727254 0.1818187

```

Apart from the parameter rescaling, this is an entirely "doable" problem. Note that we can also solve the problem as a Quadratic Program using the quadprog package.

```

> library(quadprog)
> Dmat<-diag(c(1,2,3))

```



```

> Amat<-matrix(c(1, 1, 1), ncol=1)
> bvec<-c(1)
> meq=1
> dvec<-c(0, 0, 0)
> ans<-solve.QP(Dmat, dvec, Amat, bvec, meq=0, factorized=FALSE)
> ans

$solution
[1] 0.5454545 0.2727273 0.1818182

$value
[1] 0.2727273

$unconstrained.solution
[1] 0 0 0

$iterations
[1] 2 0

$Lagrangian
[1] 0.5454545

$iact
[1] 1

```

7 Conclusion

Sumscale problems can present difficulties for optimization (or function minimization) codes. These difficulties are by no means insurmountable, but they do require some attention.

While specialized approaches are "best" for speed and correctness, a general user is more likely to benefit from a simpler approach of embedding the scaling in the objective function and rescaling the parameters before reporting them. Another choice is to use the projected gradient via `spg` from package `BB`.

References

M. Geradin. The computational efficiency of a new minimization algorithm for eigenvalue analysis. *J. Sound Vib.*, 19:319–331, 1971.

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