

# Timing Rayleigh Quotient minimization in R

true

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## Abstract

This vignette is simply to record the methods and some results for timing various Rayleigh Quotient minimizations with R using different functions and different ways of running the computations, in particular trying Fortran subroutines. This article has been updated from a 2012 document to reflect changes in R and its packages that make it awkward to reprocess the original document on newer computers and which show that timing profiles of R commands have changed in the interim. In particular, it appears that use of the R byte-compiler is now essentially automatic and manual compilation is not worthwhile.

## The computational task

The maximal and minimal eigensolutions of a symmetric matrix  $A$  are extrema w.r.t.  $x$  of the Rayleigh Quotient

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

Clearly, there may be an infinite set of  $x$  vectors, since multiplication by any signed scale factor will not alter the value of the result. We could also deal with generalized eigenproblems of the form

$$Ax = \lambda Bx$$

where  $B$  is symmetric and positive definite by using the Rayleigh Quotient (RQ)

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

In this document,  $B$  will always be an identity matrix, but some programs we test assume that it is present.

Treating the Rayleigh Quotient as an objective function, we note that it is scaled by the parameters, in fact by their sum of squares. Alternatively, we may think of seeking the **normalized** eigensolution, which is given as

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

## Timings and speedups

In R, execution times can be measured by the function `system.time`, and in particular the third element of the object this function returns. However, various factors influence computing times in a modern computational system, so we generally want to run replications of the times. The R packages `rbenchmark` and `microbenchmark` can be used for this. I have a preference for the latter. To keep the elapsed time to render this document for presentation to a reasonable interval, the number of repetitions may be limited.

There are some ways to speed up R computations.

- The code can be modified to use more efficient language structures. We show some of these below, in particular, to use vector operations.
- We can use the R byte code compiler by Luke Tierney, which has been part of the R distribution since version 2.14.
- We can use compiled code in other languages. Here we show how Fortran subroutines can be used.

## Our example matrix

We will use a matrix called the Moler matrix Nash (1979, Appendix 1). This is a positive definite symmetric matrix with one small eigenvalue. We will show a couple of examples of computing the small eigenvalue solution, but will mainly perform timings using the maximal eigenvalue solution, which we will find by minimizing the RQ of (-1) times the matrix. (The eigenvalue of this matrix is the negative of the maximal eigenvalue of the original, but the eigenvectors are equivalent to within a scaling factor for non-degenerate eigenvalues.)

Here is the code for generating the Moler matrix.

```
molermat<-function(n){
  A<-matrix(NA, nrow=n, ncol=n)
  for (i in 1:n){
    for (j in 1:n) {
      if (i == j) A[i,i]<-i
      else A[i,j]<-min(i,j) - 2
    }
  }
  A
}
```

However, since R is more efficient with vectorized code, the following routine by Ravi Varadhan should do much better.

```
molerfast <- function(n) {
# A fast version of `molermat'
  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
}
```

## Time to build the matrix

Let us see how long it takes to build the Moler matrix of different sizes. In 2012 we used the byte-code compiler, but that now seems to be active by default and NOT to give worthwhile improvements. We also include times for the `eigen()` function that computes the full set of eigensolutions very quickly.

```
##      n   osize buildi buildir eigentime eigentimr bfast bfastr
## 1 100   80216   3820    962    2144      343   869    966
## 2 200  320216  14330   3504    6125     1059  1545   304
## 3 300  720216  31083   4655   11626     1086  2571   649
## 4 400 1280216  53936   4735   20066     1421  4468  5201
## 5 500 2000216  84616   6753   32048     6664  4883   885

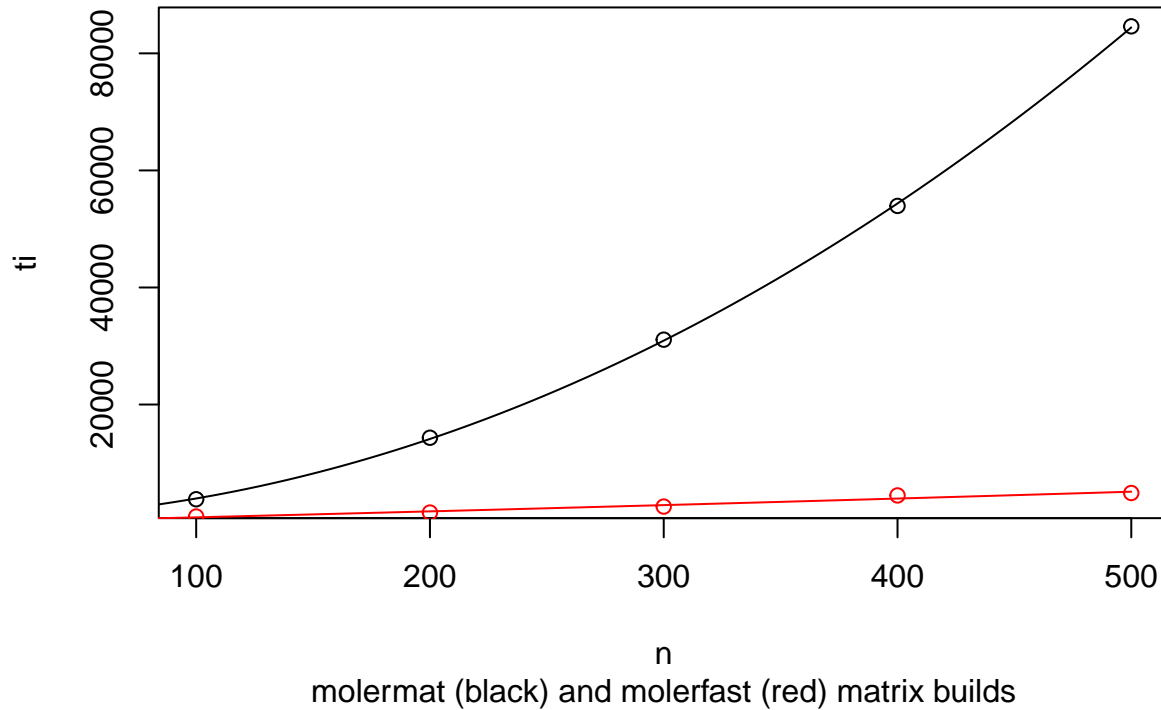
## osize - matrix size in bytes
```

```
## eigentime - all eigensolutions time; eigentimr=std.devn.
## buildi - mean interpreted build time, buildir = std.devn.
## bfast - interpreted vectorized build time; bfastr=std.devn.
## Times converted to microseconds
```

We can graph the times. The code, which is not echoed here, also models the times and the object size created. These turn out to be almost perfect quadratic models in  $n$ .

```
##
## Call:
## lm(formula = ti ~ n + n2)
##
## Residuals:
##      1      2      3      4      5
## -131.7  209.9  160.3 -423.7  185.1
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 417.60000   831.85397   0.502 0.665476
## n              2.16943     6.33928   0.342 0.764802
## n2             0.33171     0.01037  32.001 0.000975 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 387.9 on 2 degrees of freedom
## Multiple R-squared:  0.9999, Adjusted R-squared:  0.9999
## F-statistic: 1.397e+04 on 2 and 2 DF,  p-value: 7.159e-05
##
## Call:
## lm(formula = tf ~ n + n2)
##
## Residuals:
##      1      2      3      4      5
## 142.1 -202.2 -246.3  530.6 -224.3
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2.436e+02  1.021e+03  -0.238   0.834
## n              9.455e+00  7.784e+00   1.215   0.348
## n2             2.493e-03  1.273e-02   0.196   0.863
##
## Residual standard error: 476.2 on 2 degrees of freedom
## Multiple R-squared:  0.9636, Adjusted R-squared:  0.9272
## F-statistic: 26.46 on 2 and 2 DF,  p-value: 0.03642
```

## Execution time vs matrix size (microsec)



```
## Warning in summary.lm(osize): essentially perfect fit: summary may be
## unreliable
```

```
##
```

```
## Call:
```

```
## lm(formula = os ~ n + n2)
```

```
##
```

```
## Residuals:
```

```
##      1      2      3      4      5
## 4.765e-13 5.479e-12 -1.930e-11 2.025e-11 -6.908e-12
```

```
##
```

```
## Coefficients:
```

```
##              Estimate Std. Error    t value Pr(>|t|)
## (Intercept)  2.160e+02  4.448e-11  4.856e+12  <2e-16 ***
## n           -7.363e-13  3.390e-13 -2.172e+00   0.162
## n2           8.000e+00  5.543e-16  1.443e+16  <2e-16 ***
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

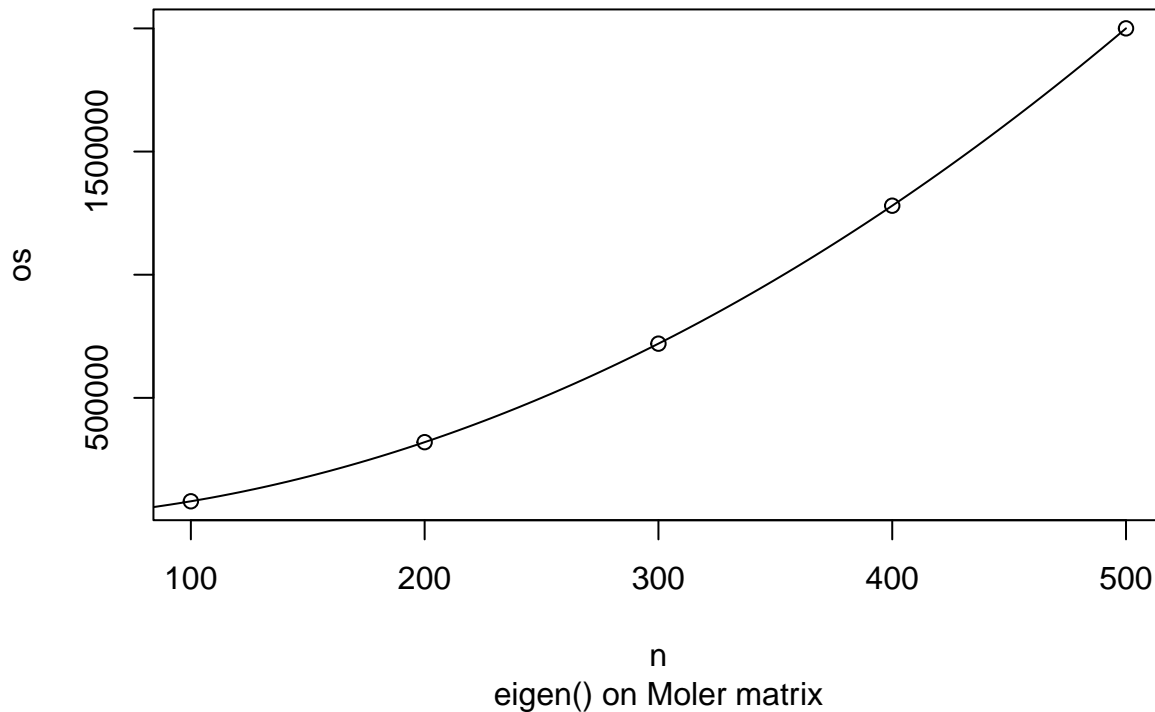
```
##
```

```
## Residual standard error: 2.074e-11 on 2 degrees of freedom
```

```
## Multiple R-squared:  1, Adjusted R-squared:  1
```

```
## F-statistic: 2.782e+33 on 2 and 2 DF, p-value: < 2.2e-16
```

## Execution time vs matrix size (microsec)



## Computing the Rayleigh Quotient

The Rayleigh Quotient requires the quadratic form  $x'Ax$  divided by the inner product  $x'x$ . R lets us form this in several ways. However, if we assume that  $x$  is normalized, then we avoid the need to compute the denominator.

```
rqdir<-function(x, AA){  
  rq<-0.0  
  n<-length(x) # assume x, AA conformable  
  for (i in 1:n) {  
    for (j in 1:n) {  
      rq<-rq+x[i]*AA[[i,j]]*x[j]  
    }  
  }  
  rq  
}
```

Somewhat better (as we shall show below) is

```
ray1<-function(x, AA){  
  rq<- t(x)%*%AA%*%x  
}
```

and (believed) better still is

```
ray2<-function(x, AA){  
  rq<- as.numeric(crossprod(x, crossprod(AA,x)))  
}
```

Note that we could implicitly include the minus sign in these routines to allow for finding the maximal

eigenvalue by minimizing the Rayleigh Quotient of  $-A$ . However, such shortcuts often rebound when the implicit negation is overlooked.

If we already have the inner product  $A * x$  as vector `ax` from some other computation, then we can simply use

```
ray3<-function(x, AA, ax=axftn){
  # ax is a function to form AA%%x
  rq<- - as.numeric(crossprod(x, ax(x, AA)))
}
```

## Matrix-vector products

In generating the RQ, we do not actually need the matrix itself, but simply the inner product with a vector `x`, from which a second inner product with `x` gives us the quadratic form  $x'Ax$ . If `n` is the order of the problem, then for large `n`, we avoid storing and manipulating a very large matrix if we use **implicit inner product** formation. We do this with the following code. For future reference, we include a function for the multiplication by an identity to facilitate programs for the generalized Rayleigh Quotient.

```
ax<-function(x, AA){
  u<- as.numeric(AA%%x)
}

axx<-function(x, AA){
  u<- as.numeric(crossprod(AA, x))
}
```

Note that second argument, supposedly communicating the matrix which is to be used in the matrix-vector product, is ignored in the following implicit product routine. It is present only to provide a common syntax when we wish to try different routines within other computations.

```
aximp<-function(x, AA=1){ # implicit moler A*x
  n<-length(x)
  y<-rep(0,n)
  for (i in 1:n){
    tt<-0.
    for (j in 1:n) {
      if (i == j) tt<-tt+i*x[i]
      else tt<-tt+(min(i,j) - 2)*x[j]
    }
    y[i]<-tt
  }
  y
}

ident<-function(x, B=1) x # identity
```

However, Ravi Varadhan has suggested the following vectorized code for the implicit matrix-vector product.

```
axmolerfast <- function(x, AA=1) {
  # A fast and memory-saving version of A%%x
  # For Moler matrix. Note we need a matrix argument to match other functions
  n <- length(x)
  j <- 1:n
  ax <- rep(0, n)
  for (i in 1:n) {
    term <- x * (pmin(i, j) - 2)
    ax[i] <- sum(term[-i])
  }
}
```

```
ax <- ax + j*x
ax
}
```

We can also use external language routines, for example in Fortran. However, this needs a Fortran **subroutine** which outputs the result as one of the returned components. The subroutine is in file `moler.f`.

```
subroutine moler(n, x, ax)
integer n, i, j
double precision x(n), ax(n), sum
c  return ax = A * x for A = moler matrix
c  A[i,j]=min(i,j)-2 for i<>j, or i for i==j
do 20 i=1,n
  sum=0.0
  do 10 j=1,n
    if (i.eq.j) then
      sum = sum+i*x(i)
    else
      sum = sum+(min(i,j)-2)*x(j)
    endif
  10  continue
  ax(i)=sum
20  continue
return
end
```

This is then compiled in a form suitable for R use by the command (this is a command-line tool, and was run in Ubuntu Linux in a directory containing the file `moler.f` but outside this vignette):

```
R CMD SHLIB moler.f
```

This creates files `moler.o` and `moler.so`, the latter being the dynamically loadable library we need to bring into our R session.

```
dyn.load("moler.so")
cat("Is the mat multiply loaded? ",is.loaded("moler"),"\n")
```

```
## Is the mat multiply loaded? TRUE
```

```
axftn<-function(x, AA=1) { # ignore second argument
  n<-length(x) # could speed up by having this passed
  vout<-rep(0,n) # purely for storage
  res<-(.Fortran("moler", n=as.integer(n), x=as.double(x), vout=as.double(vout)))$vout
}
```

We are aware that `.Fortran` is now considered inefficient compared to `.Call`. There is also the `dotCall164` package. However, for the present examples, we will remain with `.Fortran` though welcome efforts to employ other methods, especially if clean and easy-to-follow examples can be created.

We can also byte compile each of the routines above

Now it is possible to time the different approaches to the matrix-vector product.

```
dyn.load("moler.so")
cat("Is the mat multiply loaded? ",is.loaded("moler"),"\n")
```

```
## Is the mat multiply loaded? TRUE
```

```

require(microbenchmark)
nmax<-5
ptable<-matrix(NA, nrow=nmax, ncol=11) # to hold results
# loop over sizes
for (ni in 1:nmax){
  n<-100*ni
  x<-runif(n) # generate a vector
  ptable[[ni, 1]]<-n
  AA<-molermt(n)
  tax<- microbenchmark(oax<-ax(x, AA), times=mbt)$time
  taxx<-microbenchmark(oaxx<-axx(x, AA), times=mbt)$time
  if (! identical(oax, oaxx)) stop("oaxx NOT correct")
  taxftn<-microbenchmark(oaxftn<-axftn(x, AA=1), times=mbt)$time
  if (! identical(oax, oaxftn)) stop("oaxftn NOT correct")
  taximp<-microbenchmark(oaximp<-aximp(x, AA=1), times=mbt)$time
  if (! identical(oax, oaximp)) stop("oaximp NOT correct")
  taxmfi<-microbenchmark(oaxmfi<-axmolerfast(x, AA=1), times=mbt)$time
  if (! identical(oax, oaxmfi)) stop("oaxmfi NOT correct")
  ptable[[ni, 2]]<-msect(tax); ptable[[ni,3]]<-msecr(tax)
  ptable[[ni, 4]]<-msect(taxx); ptable[[ni, 5]]<-msecr(taxx)
  ptable[[ni, 6]]<-msect(taxftn); ptable[[ni, 7]]<-msecr(taxftn)
  ptable[[ni, 8]]<-msect(taximp); ptable[[ni,9]]<-msecr(taximp)
  ptable[[ni, 10]]<-msect(taxmfi); ptable[[ni,11]]<-msecr(taxmfi)
}

axtym<-data.frame(n=ptable[,1], ax=ptable[,2], sd_ax=ptable[,3], axx=ptable[,4],
  sd_axx=ptable[,5], axftn=ptable[,6], sd_axftn=ptable[,7],
  aximp=ptable[,8], sd_aximp=ptable[,9],
  axmfast=ptable[,10], sd_axmfast=ptable[,11])

print(axtym)

```

```

##      n  ax sd_ax axx sd_axx axftn sd_axftn aximp sd_aximp axmfast sd_axmfast
## 1 100  89   369 233    794   206      831  3898   1270    951      986
## 2 200  35    32  30     22   134       4 13648   1010   1719      228
## 3 300  67    34 295   1175   291       5 30949   1438   2776      924
## 4 400 452  1220  92     19   509       4 55195   1115   3759      969
## 5 500 403  1103 211    252   789       5 86189   1934   5265     1124

```

```

## ax = R matrix * vector  A %*% x
## axx = R crossprod A, x
## axftn = Fortran version of implicit Moler A * x
## aximp = implicit moler A*x in R
## axmfast = A fast and memory-saving version of A %*% x
## Times in microseconds from microbenchmark

```

From the above output, we see that the straightforward matrix-vector product appears to be the fastest. However, we have omitted the time to build the matrix. If we must build the matrix, then we need somehow to include that time. Apportioning “fixed costs” to timings is never a trivial decision. Similarly if, where and how to store large matrices if we do build them, and whether it is worth building them more than once if storage is an issue, are all questions that may need to be addressed if performance becomes important.

```

## Times (in microsecs) adjusted for matrix build

##      n axbld axxbld axftn aximp axmfast
## 1 100  3909  4053   206  3898    951

```



```
## 2 200 14365 14360 134 13648 1719
## 3 300 31150 31378 291 30949 2776
## 4 400 54388 54028 509 55195 3759
## 5 500 85019 84827 789 86189 5265
```

Out of all this, we see that the Fortran implicit matrix-vector product is the overall winner at all values of  $n$ . (Why it takes longer for  $n=100$  than  $n=200$  has not been determined.) Moreover, it does NOT require the creation and storage of the matrix. However, using Fortran does involve rather more work for the user, and for most applications it is likely we could live with the use of either

- the implicit matrix-vector multiply `axmolerfast`, or
- the interpreted matrix-product based on standard matrix-vector multiplication with an actual matrix is good enough, especially if a fast matrix build is used and we have plenty of memory>

## RQ computation times

We have set up three versions of a Rayleigh Quotient calculation in addition to the direct form. The third form is set up to use the `axftn` routine that we have already shown is efficient. We could also use this with the implicit matrix-vector product `axmolerfast`.

```
##      n direct matmul crossprod ftncross impcross
## 1 100 2772 1252 1381 3475 3084
## 2 200 2492 109 41 148 2595
## 3 300 5766 126 75 307 3601
## 4 400 10404 178 128 526 4469
## 5 500 16270 235 205 804 5548

## direct - looped a*x
## matmul - A %% x
## crossprod - A * X via crossprod
## ftncross - Fortran + crossprod
## impcross - A * x fast R implicit matrix + crossprod
```

Here we see that the use of either the matrix multiplication in `ray1` or of `crossprod` in `ray2` is very fast, and this is interpreted code. Once again, we note that all timings except those for `ray3` should have some adjustment for the building of the matrix. If storage is an issue, then `ray3`, which uses the implicit matrix-vector product in Fortran, is the approach of choice. My own preference would be to use this option if the Fortran matrix-vector product subroutine is already available for the matrix required. I would not, however, generally choose to write the Fortran subroutine for a “new” problem matrix. The fast implicit matrix-vector tool with `ray3` is also useful and quite fast if we need to minimize memory use. We note once again that the  $n=100$  case seems anomalous. This may reflect some sort of initiation of computations within a loop and that we should run a dummy calculation first before timing this case. Of course, in production computations, we still need to actually start the program, so the overhead is not avoidable.

## Solution by spg

To actually solve the eigensolution problem we will first use the projected gradient method `spg` from BB on the moler matrix for  $n=100$ . We repeat the RQ functions so that it is clear which routine we are using.

```
# spgRQ.R
rqfast<-function(x){
  rq<-as.numeric(t(x) %*% axmolerfast(x))
  rq
}
rqneg<-function(x) { -rqfast(x)}
```

```

proj <- function(x) {sign(x[1]) * x/sqrt(c(crossprod(x))) } # from ravi
# Note that the c() is needed in denominator to avoid error msgs
require(BB)
n<-100
x<-rep(1,n)
x<-x/as.numeric(sqrt(crossprod(x)))
AA<-molerfast(n)
teig<-microbenchmark(evs<-eigen(AA), times=mbt)$time
cat("eigen time =", msect(teig),"sd=",msecr(teig),"\n")

## eigen time = 2140 sd= 111

tmin<-microbenchmark(amin<-spg(x, fn=rqfast, project=proj,
                             control=list(trace=FALSE)), times=mbt)$time
tmax<-microbenchmark(amax<-spg(x, fn=rqneg, project=proj,
                             control=list(trace=FALSE)), times=mbt)$time

evalmax<-evs$values[1]
evecmax<-evs$vectors[,1]
evecmax<-sign(evecmax[1])*evecmax/sqrt(as.numeric(crossprod(evecmax))) # normalize
emax<-list(evalmax=evalmax, evecmax=evecmax)
evalmin<-evs$values[n]
evecmin<-evs$vectors[,n]
evecmin<-sign(evecmin[1])*evecmin/sqrt(as.numeric(crossprod(evecmin)))
emin<-list(evalmin=evalmin, evecmin=evecmin)
avecmax<-amax$par
avecmin<-amin$par
avecmax<-sign(avecmax[1])*avecmax/sqrt(as.numeric(crossprod(avecmax)))
avecmin<-sign(avecmin[1])*avecmin/sqrt(as.numeric(crossprod(avecmin)))
cat("minimal eigensolution: Value=",amin$value,"in time ",
    msect(tmin),"sd=",msecr(tmin),"\n")

## minimal eigensolution: Value= 5.939165e-08 in time 26602150 sd= 89139

cat("(Eigenvalue - result from eigen)=",amin$value-evalmin," vector max(abs(diff))=",
    max(abs(avecmin-evecmin)), "\n")

## (Eigenvalue - result from eigen)= 5.93916e-08 vector max(abs(diff))= 0.000135496
#print(amin$par)
cat("maximal eigensolution: Value=", -amax$value,"in time ",
    msect(tmax),"sd=",msecr(tmax),"\n")

## maximal eigensolution: Value= 3934.277 in time 492479 sd= 7535

cat("(Eigenvalue - result from eigen)=", -amax$value-evalmax," vector max(abs(diff))=",
    max(abs(avecmax-evecmax)), "\n")

## (Eigenvalue - result from eigen)= -3.761099e-06 vector max(abs(diff))= 4.747616e-06

Finding the minimal eigensolution appears to be much more troublesome than the maximal one. Let us
compare timings for the maximal eigensolution over different matrix sizes.

## Times in microsecs using spg() on moler matrix maximal eigensolutions

##      n   spgrqt  tbld  teig
## 1 100   791858   643   2248
## 2 200  3421961  2643   7480
## 3 300  8216368  4147  11888

```

```
## 4 400 16312038 5944 20612
## 5 500 27974462 8263 31099
```

## Solution by other optimizers

We can try other optimizers, but we must note that unlike `spg` they do not take account of the scaling. However, we can build in a transformation, since our function is always the same for all sets of parameters scaled by the square root of the parameter inner product. The function `nobj` forms the quadratic form that is the numerator of the Rayleigh Quotient using the `code{crossprod()}` function

```
rq<- as.numeric(crossprod(y, crossprod(AA,y)))
```

but we first form

```
y<-x/sqrt(as.numeric(crossprod(x)))
```

to scale the parameters.

Since we are running a number of gradient-based optimizers in the wrapper `optimx::opm()`, we have reduced the matrix sizes and numbers.

```
require(optimx)
nobj<-function(x, AA=-AA){
  y<-x/sqrt(as.numeric(crossprod(x)))
  rq<- as.numeric(crossprod(y, crossprod(AA,y)))
}

ngrobj<-function(x, AA=-AA){ # gradient
  y<-x/sqrt(as.numeric(crossprod(x)))
  n<-length(x)
  dd<-sqrt(as.numeric(crossprod(x)))
  T1<-diag(rep(1,n))/dd
  T2<- x%o%x/(dd*dd*dd)
  gt<-T1-T2
  gy<- as.vector(2.*crossprod(AA,y))
  gg<-as.numeric(crossprod(gy, gt))
}

mset<-c("L-BFGS-B", "BFGS", "nbg", "spg", "ucminf", "nlm", "nlminb", "nvm")
for (ni in 1:nmax){
  n<-20*ni
  x<-runif(n) # generate a vector
  AA<-molerfast(n) # make sure defined
  aall<-opm(x, fn=nobj, gr=ngrobj, method=mset, AA=-AA,
    control=list(trace=0,startttests=FALSE, dowarn=FALSE, kkt=FALSE))
  # optansout(aall, NULL)
  summary(aall, order=value, )
}
```

The timings for these matrices of order 20 to 100 are likely too short to be very reliable in detail, but do show that the RQ problem using the scaling transformation and with an analytic gradient can be solved very quickly, especially by the limited memory methods such as L-BFGS-B and `nbg`. Below we use the latter to show the times over different matrix sizes.

```
nmax<-5
ctable<-matrix(NA, nrow=nmax, ncol=3)
for (ni in 1:nmax){
  n<-100*ni
  x<-runif(n) # generate a vector
```

```

AA<-molerfast(n) # define matrix
tcgu<-microbenchmark(arcgu<-optimr(x, fn=nobj, gr=ngrobj, method="nbg",
    AA=-AA), times=mbt)
ctable[[ni,1]]<-n
ctable[[ni,2]]<-msect(tcgu$time)
ctable[[ni,3]]<-msecr(tcgu$time)
}
cat("Times in microseconds for nbg() to find maximal eigensolutions of moler matrix")

```

```

## Times in microseconds for nbg() to find maximal eigensolutions of moler matrix
cgtime<-data.frame(n=ctable[,1], tcgmin=ctable[,2], sdtcgmin=ctable[,3])
print(round(cgtime,0))

```

```

##      n tcgmin sdtcgmin
## 1 100   2361     946
## 2 200   5433     414
## 3 300   9865    3141
## 4 400  10260    2896
## 5 500  16570    4208

```

## A specialized minimizer - Geradin's method

For comparison, let us try the Geradin routine (Appendix 1) as implemented in R by one of us (JN).

```

cat("Test geradin with explicit matrix multiplication\n")

```

```

## Test geradin with explicit matrix multiplication

```

```

n<-10
AA<-molerfast(n)
BB=diag(rep(1,n))
x<-runif(n)
tg<-microbenchmark(ag<-geradin(x, ax, bx, AA=AA, BB=BB,
    control=list(trace=FALSE)), times=mbt)
cat("Minimal eigensolution\n")

```

```

## Minimal eigensolution

```

```

print(ag)

```

```

## $x
## [1] 386618.971 193310.315 96657.231 48332.971 24175.300 12105.330
## [7] 6088.052 3114.812 1698.986 1132.655
##
## $RQ
## [1] 8.582807e-06
##
## $ipr
## [1] 44
##
## $msg
## [1] "Rayleigh Quotient increased in step"

```

```

cat("Geradin time=",msect(tg$time),"sd=",msecr(tg$time),"n")

```

```

## Geradin time= 2670 sd= 9979

```

```
tgn<-microbenchmark(agn<-geradin(x, ax, bx, AA=-AA, BB=BB,
  control=list(trace=FALSE)), times=mbt)
cat("Maximal eigensolution (negative matrix)\n")
```

```
## Maximal eigensolution (negative matrix)
```

```
print(agn)
```

```
## $x
## [1] -228929753806      7726992696   244104578112   472239647405   684417090221
## [6]  873471850096  1033005754381  1157633064483  1243135437891  1286614945412
##
## $RQ
## [1] -31.58981
##
## $ipr
## [1] 35
##
## $msg
## [1] "Small gradient -- done"
```

```
cat("Geradin time=",msec(tgn$time),"sd=",msecr(tgn$time),"\\n")
```

```
## Geradin time= 462 sd= 8
```

Let us time this routine with different matrix vector approaches.

```
naximp<-function(x, A=1){ # implicit moler A*x
  n<-length(x)
  y<-rep(0,n)
  for (i in 1:n){
    tt<-0.
    for (j in 1:n) {
      if (i == j) tt<-tt+i*x[i]
      else tt<-tt+(min(i,j) - 2)*x[j]
    }
    y[i]<- -tt # include negative sign
  }
  y
}
```

```
dyn.load("moler.so")
cat("Is the mat multiply loaded? ",is.loaded("moler"),"\\n")
```

```
## Is the mat multiply loaded? TRUE
```

```
naxftn<-function(x, A) { # ignore second argument
  n<-length(x) # could speed up by having this passed
  vout<-rep(0,n) # purely for storage
  # NEED TO EXPLAIN -1 below
  res<-(-1)*(.Fortran("moler", n=as.integer(n), x=as.double(x), vout=as.double(vout)))$vout
}
```

```
require(microbenchmark)
```

```
## Loading required package: microbenchmark
```

```

nmax<-5
gtable<-matrix(NA, nrow=nmax, ncol=4) # to hold results
# loop over sizes
for (ni in 1:nmax){
  n<-100*ni
  x<-runif(n) # generate a vector
  gtable[[ni, 1]]<-n
  AA<-molermt(n)
  BB<-diag(rep(1,n))
  tgax<-microbenchmark(ogax<-geradin(x, ax, bx, AA=-AA, BB=BB, control=list(trace=FALSE)), times=mbt)
  gtable[[ni, 2]]<-msect(tgax$time)
  tgaximp<-microbenchmark(ogaximp<-geradin(x, naximp, ident, AA=1, BB=1, control=list(trace=FALSE)), times=mbt)
  gtable[[ni, 3]]<-msect(tgaximp$time)
  tgaxftn<-microbenchmark(ogaxftn<-geradin(x, naxftn, ident, AA=1, BB=1, control=list(trace=FALSE)), times=mbt)
  gtable[[ni, 4]]<-msect(tgaxftn$time)
}

gtym<-data.frame(n=gtable[,1], ax=gtable[,2], aximp=gtable[,3], axftn=gtable[,4])
print(gtym)

```

```

##      n      ax      aximp axftn
## 1 100  6057  122044  1655
## 2 200  3626  483309  4280
## 3 300  5275  928223  7287
## 4 400  8908 1768589 13113
## 5 500 14117 2995109 21845

```

Let us check that the eigenvalue approximations by the Geradin are consistent with the answers via `eigen()`.

```

for (n in c(100, 200, 300, 400, 500) ) {
  x<-runif(n)
  evalmax<-eigen(x)$evalmax
  # evecmax<-eigen(x)$evecmax
  ogaxftn<-geradin(x, naxftn, ident, AA=1, BB=1, control=list(trace=FALSE))
  gvec<-ogaxftn$x
  gval<-ogaxftn$RQ
  gvec<-sign(gvec[[1]])*gvec/sqrt(as.numeric(crossprod(gvec)))
  diff<-gvec-evecmax
  cat("Geradin eigenvalue - eigen result: ",gval-evalmax,"    max(abs(vector diff))=",
      max(abs(diff)), "\n")
}

## Geradin eigenvalue - eigen result: -3.157309e-05    max(abs(vector diff))= 1.959107e-05
## Geradin eigenvalue - eigen result: 12036.95    max(abs(vector diff))= 0.07695634
## Geradin eigenvalue - eigen result: 32179.59    max(abs(vector diff))= 0.1039409
## Geradin eigenvalue - eigen result: 60427.94    max(abs(vector diff))= 0.1166165
## Geradin eigenvalue - eigen result: 96781.98    max(abs(vector diff))= 0.1236483

```

## Fortran version of Geradin code

Appendix 2 lists a Fortran version of the Geradin Rayleigh Quotient calculations. The algorithm was part of Nash (1979), and in particular A25 in the Fortran version of Nashlib (Nash (1980), see also <https://github.com/pcolsen/Nash-Compact-Numerical-Methods/tree/main/fortran>). We can use this to get an approximation to the All-Fortran timing for the Geradin code, and note that the time to get the minimal

eigensolution is longer than that for the maximal one.

This can be easily compiled on most Linux systems. Here we do so via R.

```
system("gfortran ./a25moler.f")
```

The program takes as input a single line with an integer for the size of the Moler matrix to use (maximum 1600 as per the program declarations) and a number that is either 1.0 for the minimal eigensolution or -1.0 for the maximal one. Though it is slightly clumsy, we have created files for matrices of size 100, 200, 300, 400 and 500.

Let us compare timings for R's `eigen()` with the Fortran Geradin program for these cases.

```
## Geradin fortran version a25moler.f
## eigen(): n=100 build time= 3048   eigen time= 2181
## eigen: Minimal Eigenvalue = 5.04378e-14   RQ= -6.463237e-17
## A25RQM N= 100 matvec ops= 198 Min Est. EV= 2.586285e-18   Gradient= 1.515018e-30   time= 9
## A25RQM N= 100 matvec ops= 13 Max Est. EV= 3934.277   Gradient= 2.441749e-22   time= 2
## eigen: Maximal Eigenvalue = 3934.277   RQ= 3934.277
## eigen(): n=200 build time= 2504   eigen time= 13382
## eigen: Minimal Eigenvalue = -2.491097e-14   RQ= 8.728006e-17
## A25RQM N= 200 matvec ops= 412 Min Est. EV= 2.411243e-19   Gradient= 4.764326e-30   time= 47
## A25RQM N= 200 matvec ops= 9 Max Est. EV= 15971.22   Gradient= 4.973105e-11   time= 2
## eigen: Maximal Eigenvalue = 15971.22   RQ= 15971.22
## eigen(): n=300 build time= 4149   eigen time= 11398
## eigen: Minimal Eigenvalue = -1.716203e-13   RQ= -1.255081e-16
## A25RQM N= 300 matvec ops= 598 Min Est. EV= -5.995543e-20   Gradient= 1.377315e-29   time= 142
## A25RQM N= 300 matvec ops= 11 Max Est. EV= 36113.87   Gradient= 2.340489e-13   time= 4
## eigen: Maximal Eigenvalue = 36113.87   RQ= 36113.87
## eigen(): n=400 build time= 5919   eigen time= 20241
## eigen: Minimal Eigenvalue = -7.201366e-13   RQ= 1.550574e-17
## A25RQM N= 400 matvec ops= 553 Min Est. EV= -1.216296e-19   Gradient= 1.912164e-29   time= 211
## A25RQM N= 400 matvec ops= 12 Max Est. EV= 64362.22   Gradient= 3.036479e-19   time= 6
## eigen: Maximal Eigenvalue = 64362.22   RQ= 64362.22
## eigen(): n=500 build time= 5204   eigen time= 33773
## eigen: Minimal Eigenvalue = 1.949665e-12   RQ= -9.8516e-18
## A25RQM N= 500 matvec ops= 733 Min Est. EV= -3.794744e-18   Gradient= 4.428853e-29   time= 422
## A25RQM N= 500 matvec ops= 12 Max Est. EV= 100716.3   Gradient= 2.030114e-15   time= 8
## eigen: Maximal Eigenvalue = 100716.3   RQ= 100716.3
```

We note that the determination of the minimal eigenvalue is clearly more difficult for all methods. Even the standard `eigen()` method gets a Rayleigh Quotient for the minimal eigenvector that is quite different (in some cases a different sign) from the reported eigenvalue. On the other hand, the Geradin method is highly efficient in finding the maximal eigenvalue.

## Perspective

We can compare the different approaches by looking at the ratio of the best solution time for each method (compiled or interpreted, with best choice of function) to the time for the Geradin approach for the different matrix sizes. In this we will ignore the fact that some approaches do not build the matrix.

```
## Ratios of maximal eigensolution times to Fortran Geradin method
```

```
##   nsize   eigen      spg   rcgmin
## 1   100 1.295468  478.4640 1.4265861
## 2   200 1.431075  799.5236 1.2693925
## 3   300 1.595444 1127.5378 1.3537807
## 4   400 1.530237 1243.9593 0.7824296
## 5   500 1.467063 1280.5888 0.7585260
```

We see here that the `ncg` approach and that of the standard `eigen()` are competitive with Geradin, but `spg()` takes a great deal of time.

To check the value of the Geradin approach in R, let us use a much larger problem, with `n=2000`.

```
## Times in seconds
```

```
## Build = 79513   eigen(): 932744   Rcgminu: 346102   Geradin: 286452
```

```
## Ratios: build= 0.2775788 eigen= 3.256197   Rcgminu= 1.208237
```

## Conclusions

The Rayleigh Quotient minimization approach to eigensolutions has an intuitive appeal and offers an interesting optimization test problem. As an eigensolution tool, given the efficiency of standard methods such as R's `eigen()`, it is of limited utility. On the other hand, in special cases, especially using implicit matrix multiply to avoid storing a large matrix, it may be valuable.

From the tests in this vignette, here is what we may say about these attempts, which we caution are based on a relatively small sample of tests:

- Fortran may be worth using to speed up calculations such as the implicit matrix-vector product, though it does demand much more programming effort. We did not, for example, choose to try `.Call` or `.C64` approaches because of the overhead.
- The `eigen()` routine is a highly effective tool for computing all eigensolutions, even of a large matrix. It is only worth computing a single solution when the matrix is very large, in which case a specialized method such as that of Geradin makes sense and offers significant savings, especially when combined with the Fortran implicit matrix-product routine.

## Acknowledgements

This vignette originated due to a problem suggested by Gabor Grothendieck. Ravi Varadhan has provided inciteful comments and some vectorized functions which greatly altered some of the observations.

## Appendix 1: Geradin routine in R

```
ax<-function(x, AA){
  u<-as.numeric(AA%*%x)
}
bx<-function(x, BB){
  v<-as.numeric(BB%*%x)
}
geradin<-function(x, ax, bx, AA, BB, control=list(trace=TRUE, maxit=1000)){
```



```

# Geradin minimize Rayleigh Quotient, Nash CMN Alg 25
# print(control)
trace<-control$trace
n<-length(x)
tol<-n*n*.Machine$double.eps^2
offset<-1e+5 # equality check offset
if (trace) cat("geradin.R, using tol=",tol,"\n")
ipr<-0 # counter for matrix mults
pa<- .Machine$double.xmax
R<-pa
msg<-"no msg"
# step 1 -- main loop
keepgoing<-TRUE
while (keepgoing) {
  avec<-ax(x, AA); bvec<-bx(x, BB); ipr<-ipr+1
  xax<-as.numeric(crossprod(x, avec));
  xbx<-as.numeric(crossprod(x, bvec));
  if (xbx <= tol) {
    keepgoing<-FALSE # not really needed
    msg<-"avoid division by 0 as xbx too small"
    break
  }
  p0<-xax/xbx
  if (p0>pa) {
    keepgoing<-FALSE # not really needed
    msg<-"Rayleigh Quotient increased in step"
    break
  }
  pa<-p0
  g<-2*(avec-p0*bvec)/xbx
  gg<-as.numeric(crossprod(g)) # step 6
  if (trace) cat("Before loop: RQ=",p0," after ",ipr," products, gg=",gg,"\n")
  if (gg<tol) { # step 7
    keepgoing<-FALSE # not really needed
    msg<-"Small gradient -- done"
    break
  }
  t<- -g # step 8
  for (itn in 1:n) { # major loop step 9
    y<-ax(t, AA); z<-bx(t, BB); ipr<-ipr+1 # step 10
    tat<-as.numeric(crossprod(t, y)) # step 11
    xat<-as.numeric(crossprod(x, y))
    xbt<-as.numeric(crossprod(x, z))
    tbt<-as.numeric(crossprod(t, z))
    u<-tat*xbt-xat*tbt
    v<-tat*xbx-xax*tbt
    w<-xat*xbx-xax*xbt
    d<-v*v-4*u*w
    if (d<0) stop("Geradin: imaginary roots not possible") # step 13
    d<-sqrt(d) # step 14
    if (v>0) k<--2*w/(v+d) else k<-0.5*(d-v)/u
    xlast<-x # NOT as in CNM -- can be avoided with loop
    avec<-avec+k*y; bvec<-bvec+k*z # step 15, update
  }
}

```

```

x<-x+k*t
xax<-xax+as.numeric(crossprod(x,avec))
xbx<-xbx+as.numeric(crossprod(x,bvec))
if (xbx<tol) stop("Geradin: xbx has become too small")
chcount<-n - length(which((xlast+offset)==(x+offset)))
if (trace) cat("Number of changed components = ",chcount,"\n")
pn<-xax/xbx # step 17 different order
if (chcount==0) {
  keepgoing<-FALSE # not really needed
  msg<-"Unchanged parameters -- done"
  break
}
if (pn >= p0) {
  if (trace) cat("RQ not reduced, restart\n")
  break # out of itn loop, not while loop (TEST!)
}
p0<-pn # step 19
g<-2*(avec-pn*bvec)/xbx
gg<-as.numeric(crossprod(g))
if (trace) cat("Itn", itn," RQ=",p0," after ",ipr," products, gg=",gg,"\n")
if (gg<tol){ # step 20
  if (trace) cat("Small gradient in iteration, restart\n")
  break # out of itn loop, not while loop (TEST!)
}
xbt<-as.numeric(crossprod(x,z)) # step 21
w<-y-pn*z # step 22
tabt<-as.numeric(crossprod(t,w))
beta<-as.numeric(crossprod(g,(w-xbt*g)))
beta<-beta/tabt # step 23
t<-beta*t-g
} # end loop on itn -- step 24
} # end main loop -- step 25
ans<-list(x=x, RQ=p0, ipr=ipr, msg=msg) # step 26
}

```

## Appendix 2: Geradin routine in Fortran

A modified version of the Fortran particularized to the Moler matrix follows (file 'a25moler.f'):

```

C&&& A25
C  TEST ALG 25 USING GRID (5 POINT)
C  J.C. NASH    JULY 1978, APRIL 1989
      LOGICAL IFR
      INTEGER N,M,NOUT,NIN,KPR,LIMIT,I
      EXTERNAL APR,BPR
C      REAL EPS,P0,X(N),S(N),T(N),U(N),V(N),W(N),Y(N),RNORM
      COMMON /GSZ/ M,IFR,R(1600)
      DOUBLE PRECISION EPS,P0,RNORM,VNORM,RNV, SCALM
      DOUBLE PRECISION S(1600),T(1600),U(1600),V(1600),W(1600),
      X X(1600),Y(1600)
C  I/O CHANNELS
      NIN=5
      NOUT=6
      1  READ(NIN,900)N, SCALM

```

```

900  FORMAT(I6, F10.0)
      LIMIT=1000*N
      WRITE(NOUT,950)N,LIMIT
950  FORMAT(' MOLER MATRIX ORDER',I5,'  LIMIT=',I7)
      IF(N.LE.0)STOP
      IF(SCALM.LT.0.0) WRITE(NOUT,953)
953  FORMAT(' Finding maximal eigenvalue')
      IFR=.FALSE.
C APPROX
      EPS=16.0**(-14)
      KPR=LIMIT
      RNORM=1.0/SQRT(FLOAT(N))
      DO 10 I=1,N
        X(I)=RNORM
10   CONTINUE
      write(NOUT,*) " About to call A25RQM"
      CALL A25RQM(N,X,EPS,KPR,S,T,U,V,W,Y,PO,NOUT,APR,BPR)
CCC   A25RQM(N,X,EPS,KPR,Y,Z,T,G,A,B,PO,IPR,APR,BPR)
      PO=PO*SCALM
      WRITE(NOUT,951)KPR,PO
951  FORMAT(' RETURNED AFTER',I4,' PRODUCTS WITH EV=',1PE16.8)
C    TO GET TIMING STOP HERE
      STOP
      END
      SUBROUTINE BPR(N,X,V)
C J.C. NASH JULY 1978, APRIL 1989
C UNITM MATRIX * X INTO V
      INTEGER N,I
      DOUBLE PRECISION X(N),V(N)
      DO 100 I=1,N
        V(I)=X(I)
100  CONTINUE
      RETURN
      END
      SUBROUTINE APR(N,X,V)
      integer n, i, j
      double precision x(n), V(n), sum
c    return ax = A * x for A = moler matrix
c    A[i,j]=min(i,j)-2 for i<>j, or i for i==j
      do 20 i=1,n
        sum=0.0
        do 10 j=1,n
          if (i.eq.j) then
            sum = sum+i*x(i)
          else
            sum = sum+(min(i,j)-2)*x(j)
          endif
10      continue
        V(i)=sum
C      V(i)=-sum
C use negative to get largest ev
20    continue
      return
      end

```

```

      SUBROUTINE A25RQM(N,X,EPS,KPR,Y,Z,T,G,A,B,PO,IPR,APR,BPR)
C  STEP 0
      INTEGER N,LP,IPR,ITN,I,LIM,COUNT
      DOUBLE PRECISION X(N),T(N),G(N),Y(N),Z(N),PN,A(N),B(N)
      DOUBLE PRECISION EPS,TOL,PO,PA,XAX,XBX,XAT,XBT,TAT,TBT,W,K,
      X  D,V,GG,BETA,TABT,U
C  ALGORITHM 25 RAYLEIGH QUOTIENT MINIMIZATION BY CONJUGATE GRADIENTS
C  J.C. NASH  JULY 1978, FEBRUARY 1980, APRIL 1989
C  N  = ORDER OF PROBLEM
C  X  = INITIAL (APPROXIMATE?) EIGENVECTOR
C  EPS = MACHINE PRECISION
C&&& for Microsoft test replace with actual names
C  APR,BPR ARE NAMES OF SUBROUTINES WHICH FORM THE PRODUCTS
C      V= A*X    VIA  CALL APR(N,X,V)
C      T= B*X    VIA  CALL BPR(N,X,T)
C  KPR = LIMIT ON THE NUMBER OF PRODUCTS (INPUT) (TAKES ROLE OF IPR)
C      = PRODUCTS USED (OUTPUT)
C  Y,Z,T,G,A,B RE WORKING VECTORS IN AT LEAST N ELEMENTS
C  PO  = APPROXIMATE EIGENVALUE (OUTPUT)
C  IPR = PRINT CHANNEL  PRINTING IF IPR.GT.0
C  IBM VALUE  - APPROX. LARGEST NUMBER REPRESENTABLE.
C&&&      PA=R1MACH(2)
      write(6, 960) N
960  FORMAT(' In A25RQM Geradin Rayleigh Quotient Min for N=', I5)
      PA=1E+35
      LIM=KPR
      KPR=0
      TOL=N*N*EPS*EPS
C  STEP 1
10   KPR=KPR+1
      IF(KPR.GT.LIM)RETURN
C  FIND LIMIT IN ORIGINAL PROGRAMS
      CALL APR(N,X,A)
      CALL BPR(N,X,B)
C  STEP 2
      XAX=0.0
      XBX=0.0
      DO 25 I=1,N
          XAX=XAX+X(I)*A(I)
          XBX=XBX+X(I)*B(I)
25   CONTINUE
C  STEP 3
      IF(XBX.LT.TOL)STOP
C  STEP 4
      PO=XAX/XBX
      IF(PO.GE.PA)RETURN
      IF(IPR.GT.0)WRITE(IPR,963)KPR,PO
963  FORMAT( 1H ,I4,' PRODUCTS, EST. EIGENVALUE=',1PE16.8)
C  STEP 5
      PA=PO
C  STEP 6
      GG=0.0
      DO 65 I=1,N
          G(I)=2.0*(A(I)-PO*B(I))/XBX

```

```

        GG=GG+G(I)**2
65  CONTINUE
C  STEP 7
        IF(IPR.GT.0)WRITE(IPR,964)GG
964  FORMAT(' GRADIENT NORM SQUARED=',1PE16.8)
        IF(GG.LT.TOL)RETURN
C  STEP 8
        DO 85 I=1,N
            T(I)=-G(I)
85  CONTINUE
C  STEP 9
        DO 240 ITN=1,N
C  STEP 10
            KPR=KPR+1
            IF(KPR.GT.LIM)RETURN
            CALL APR(N,T,Y)
            CALL BPR(N,T,Z)
C  STEP 11
            TAT=0.0
            TBT=0.0
            XAT=0.0
            XBT=0.0
            DO 115 I=1,N
                TAT=TAT+T(I)*Y(I)
                XAT=XAT+X(I)*Y(I)
                TBT=TBT+T(I)*Z(I)
                XBT=XBT+X(I)*Z(I)
115  CONTINUE
C  STEP 12
            U=TAT*XBT-XAT*TBT
            V=TAT*XBX-XAX*TBT
            W=XAT*XBX-XAX*XBT
            D=V*V-4.0*U*W
C  STEP 13
            IF(D.LT.0)STOP
C  MAY NOT WISH TO STOP
C  STEP 14
            D=SQRT(D)
            IF(V.GT.0.0)GOTO 145
            K=0.5*(D-V)/U
            GOTO 150
145  K=-2.0*W/(D+V)
150  COUNT=0
C  STEP 15
            XAX=0.0
            XBX=0.0
            DO 155 I=1,N
                A(I)=A(I)+K*Y(I)
                B(I)=B(I)+K*Z(I)
                W=X(I)
                X(I)=W+K*T(I)
                IF(W.EQ.X(I))COUNT=COUNT+1
                XAX=XAX+X(I)*A(I)
                XBX=XBX+X(I)*B(I)

```

```

155    CONTINUE
C  STEP 16
      IF(XBX.LT.TOL)STOP
      PN=XAX/XBX
C  STEP 17
      IF(COUNT.LT.N)GOTO 180
      IF(ITN.EQ.1)RETURN
      GOTO 10
C  STEP 18
180    IF(PN.LT.PO)GOTO 190
      IF(ITN.EQ.1)RETURN
      GOTO 10
C  STEP 19
190    PO=PN
      GG=0.0
      DO 195 I=1,N
        G(I)=2.0*(A(I)-PN*B(I))/XBX
        GG=GG+G(I)**2
195    CONTINUE
C  STEP 20
      IF(GG.LT.TOL)GOTO 10
C  STEP 21
      XBT=0.0
      DO 215 I=1,N
        XBT=XBT+X(I)*Z(I)
215    CONTINUE
C  STEP 22
      TABT=0.0
      BETA=0.0
      DO 225 I=1,N
        W=Y(I)-PN*Z(I)
        TABT=TABT+T(I)*W
        BETA=BETA+G(I)*(W-G(I)*XBT)
225    CONTINUE
C  STEP 23
      BETA=BETA/TABT
      DO 235 I=1,N
        T(I)=BETA*T(I)-G(I)
235    CONTINUE
C  STEP 24
240    CONTINUE
C  STEP 25
      GOTO 10
C  NO STEP 26 - HAVE USED RETURN INSTEAD
      END

```

## References

- Nash, John C. 1979. *Compact Numerical Methods for Computers: Linear Algebra and Function Minimisation*. Bristol: Adam Hilger.
- . 1980. "NASHLIB: Algorithms for Compact Numerical Methods, now available in FORTRAN."