

Timing Rayleigh Quotient minimization in R

true

2023-9-6

Abstract

This vignette is simply to record the methods and results for timing various Rayleigh Quotient minimizations with R using different functions and different ways of running the computations, in particular trying Fortran subroutines and the R byte compiler. It 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.

The computational task

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

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

We could also deal with generalized eigenproblems of the form

$$Ax = eBx$$

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.

Note that the objective is scaled by the parameters, in fact by their sum of squares. Alternatively, we may think of requiring 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. However, to keep the time to prepare this vignette with `Sweave` or `knitr` reasonable, many of the timings will be done with only `system.time`.

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.

Loading required package: microbenchmark

##	n	osize	buildi	builddir	eigentime	eigentimr	bfast	bfastr
## 1	50	20216	1187	844	514	252	497	947
## 2	100	80216	3392	550	1566	93	688	103
## 3	150	180216	7449	686	4400	182	1021	39
## 4	200	320216	13124	886	9030	419	1461	41
## 5	250	500216	20422	554	16154	532	2232	314
## 6	300	720216	29423	614	26572	983	2658	730
## 7	350	980216	41365	6279	40595	784	3370	694
## 8	400	1280216	52262	1207	59179	1107	5164	7140
## 9	450	1620216	67713	6887	82452	1092	7431	9652
## 10	500	2000216	82224	1950	111374	1373	6918	7127

osize - matrix size in bytes

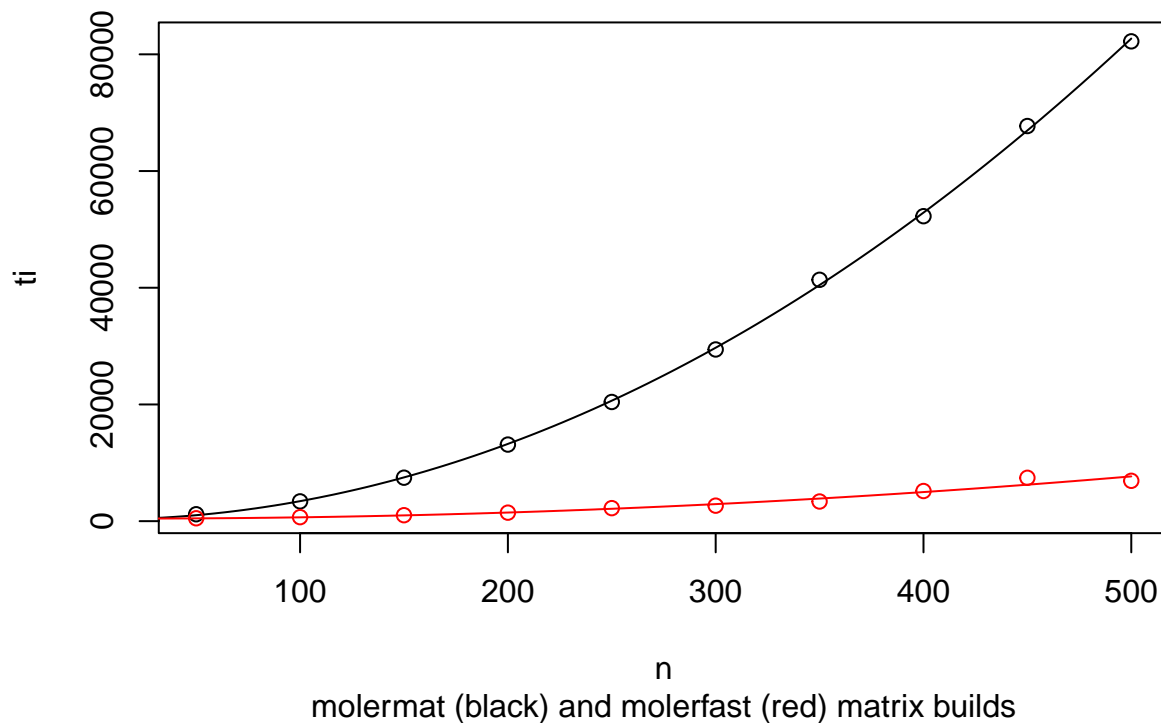
```
## eigentime - all eigensolutions time
## buildi - interpreted build time, range
## bfast - interpreted vectorized build time
## Times converted to milliseconds
```

It does not appear that the compiler has much effect, or else it is being automatically invoked.

We can graph the times. The code, which is not echoed here, also models the times and the object size created as almost perfect quadratic models in n .

```
##
## Call:
## lm(formula = ti ~ n + n2)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -613.95 -281.85  -84.55  112.94  899.14
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 286.966667  655.343394   0.438   0.675
## n           -1.932394    5.473981  -0.353   0.734
## n2             0.333512    0.009699   34.385 4.56e-09 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 557.2 on 7 degrees of freedom
## Multiple R-squared:  0.9997, Adjusted R-squared:  0.9996
## F-statistic: 1.153e+04 on 2 and 7 DF, p-value: 4.863e-13
##
## Call:
## lm(formula = tf ~ n + n2)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -753.49 -203.15   23.92   88.99 1179.56
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 456.23333  678.85119   0.672   0.5231
## n           -1.09221    5.67034  -0.193   0.8527
## n2             0.03105    0.01005   3.090   0.0176 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 577.2 on 7 degrees of freedom
## Multiple R-squared:  0.9599, Adjusted R-squared:  0.9485
## F-statistic: 83.85 on 2 and 7 DF, p-value: 1.288e-05
```

Execution time vs matrix size



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

```
##
```

```
## Call:
```

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

```
##
```

```
## Residuals:
```

```
##      Min       1Q   Median       3Q      Max
## -7.074e-11  2.234e-12  7.557e-12  1.166e-11  3.113e-11
```

```
##
```

```
## Coefficients:
```

```
##              Estimate Std. Error  t value Pr(>|t|)
## (Intercept)  2.160e+02  3.730e-11  5.791e+12  <2e-16 ***
## n            0.000e+00  3.116e-13  0.000e+00      1
## n2           8.000e+00  5.521e-16  1.449e+16  <2e-16 ***
```

```
## ---
```

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

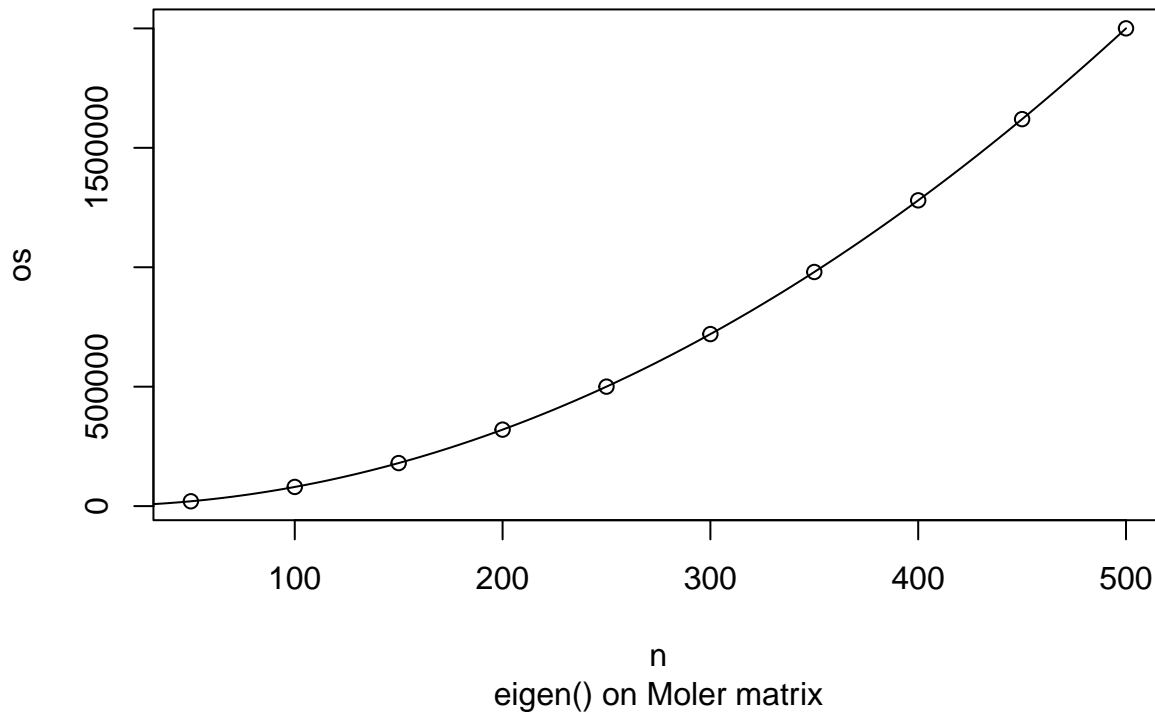
```
##
```

```
## Residual standard error: 3.172e-11 on 7 degrees of freedom
```

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

```
## F-statistic: 2.09e+33 on 2 and 7 DF, p-value: < 2.2e-16
```

Execution time vs matrix size



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.

```
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 $x^T A x$ 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 $x^T A x$. 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 the multiplication by an identity.

```
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 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<-10
ptable<-matrix(NA, nrow=nmax, ncol=11) # to hold results
# loop over sizes
for (ni in 1:nmax){
```

```

n<-50*ni
x<-runif(n) # generate a vector
ptable[[ni, 1]]<-n
AA<-molermat(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], axr=ptable[,3], axx=ptable[,4],
                  axxr=ptable[,5], axftn=ptable[,6], axftnr=ptable[,7],
                  aximp=ptable[,8], aximpr=ptable[,9],
                  axmfast=ptable[,10], axmfastr=ptable[,11])

print(axtym)

```

```

##      n  ax axr axx axxr axftn axftnr aximp aximpr axmfast axmfastr
## 1   50  51 239  55  249   103   462  1038   1158    500    952
## 2  100  11  21   9    3    29    3  3194    508    647    40
## 3  150  23  54  20   4    61    6  7029    707   1077    40
## 4  200  30  42  35   4   105    3 12498    861   1536    50
## 5  250  33   6  58  24   163    6 19591    418   2285   1129
## 6  300  48  18  80  14   232    5 29601    716   2695    927
## 7  350  66  32 110  21   314    4 38569    903   3287    944
## 8  400  80  16 140  13   410    4 50213   1141   4009   1060
## 9  450 101  17 178  24   520   11 64425   7088   4799   1344
## 10 500 130  51 219  18   637    4 79824   7589   5404   1110

```

```

## 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 milliseconds from microbenchmark

```

From the above output, we see that the `crossprod` variant of the 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 millisecs) adjusted for matrix build

```

```

##      n axbld axxbld axftn aximp
## 1   50  1238   1242   103  1038
## 2  100  3403   3401    29  3194

```



```
## 3 150 7472 7469 61 7029
## 4 200 13154 13159 105 12498
## 5 250 20455 20480 163 19591
## 6 300 29471 29503 232 29601
## 7 350 41431 41475 314 38569
## 8 400 52342 52402 410 50213
## 9 450 67814 67891 520 64425
## 10 500 82354 82443 637 79824
```

Out of all this, we see that the Fortran implicit matrix-vector product is the overall winner at all values of `n`. 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 interpreted matrix-product based on `crossprod` and an actual matrix is good enough, especially if a fast matrix build is used and we have plenty of memory, or
- the interpreted or byte-code compiled implicit matrix-vector multiply `axmolerfast`.

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`.

It seems overkill to show the RQ computation time for all versions and matrices, so we will do the timing simply for a matrix of order 500.

```
## Direct algorithm: 17567.68
## ray1: mat-mult algorithm: 239.4928
## ray2: crossprod algorithm: 234.7876
## ray3: ax Fortran + crossprod: 659.3261
## ray3: ax fast R implicit + crossprod: 5567.37
```

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.

Solution by spg

To actually solve the eigensolution problem we will first use the projected gradient method `spg` from BB. We repeat the RQ function so that it is clear which routine we are using.

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

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)

## Loading required package: 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 =", mean(teig)*0.001,"\\n")

## eigen time = 1558.793

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

## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296

```

```

## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
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## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
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## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537

```

```

## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
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## iter: 30 f-value: 0.4664893 pgrad: 0.627411
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## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806

```

```

## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
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## iter: 30 f-value: 0.4664893 pgrad: 0.627411
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## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
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## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
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## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
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## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384

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## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
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## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694

```

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## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
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## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296

```

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## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
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## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
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## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806

```

```

## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
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## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384

```

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## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
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## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
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## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
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## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
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## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694

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## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384
## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
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## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
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## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
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## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296

```

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## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
## iter: 0 f-value: 3185.5 pgrad: 0.2379617
## iter: 10 f-value: 25.69723 pgrad: 0.5634518
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## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
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## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537

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## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247
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## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
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## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
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## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
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```

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## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
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## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
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## iter: 10 f-value: 25.69723 pgrad: 0.5634518
## iter: 20 f-value: 1.547079 pgrad: 0.5191694
## iter: 30 f-value: 0.4664893 pgrad: 0.627411
## iter: 40 f-value: 0.235088 pgrad: 0.6666011
## iter: 50 f-value: 0.1551073 pgrad: 0.6005118
## iter: 60 f-value: 0.04815044 pgrad: 0.4510603
## iter: 70 f-value: 0.007978361 pgrad: 0.3776179
## iter: 80 f-value: 0.0005602141 pgrad: 0.05667384

```

```

## iter: 90 f-value: 0.0003087274 pgrad: 0.05920482
## iter: 100 f-value: 0.0001525023 pgrad: 0.03077191
## iter: 110 f-value: 0.0001298035 pgrad: 0.1215445
## iter: 120 f-value: 5.644138e-05 pgrad: 0.01918068
## iter: 130 f-value: 4.497743e-05 pgrad: 0.01715069
## iter: 140 f-value: 4.51772e-05 pgrad: 0.09602806
## iter: 150 f-value: 2.331755e-05 pgrad: 0.01249892
## iter: 160 f-value: 1.919963e-05 pgrad: 0.01137753
## iter: 170 f-value: 1.373694e-05 pgrad: 0.04498692
## iter: 180 f-value: 4.433503e-06 pgrad: 0.005594446
## iter: 190 f-value: 3.026455e-06 pgrad: 0.004643491
## iter: 200 f-value: 2.396084e-06 pgrad: 0.004287537
## iter: 210 f-value: 1.304677e-06 pgrad: 0.01095073
## iter: 220 f-value: 6.335952e-07 pgrad: 0.002152797
## iter: 230 f-value: 5.330441e-07 pgrad: 0.00196514
## iter: 240 f-value: 3.234822e-07 pgrad: 0.001548084
## iter: 250 f-value: 2.382761e-07 pgrad: 0.001321637
## iter: 260 f-value: 2.012441e-07 pgrad: 0.001223296
## iter: 270 f-value: 1.704779e-07 pgrad: 0.001126575
## iter: 280 f-value: 1.438116e-07 pgrad: 0.001036892
## iter: 290 f-value: 8.688898e-08 pgrad: 0.0008079247

tmax<-microbenchmark(amax<-spg(x, fn=rqneg, project=proj,
                                control=list(trace=FALSE)), times=mbt)$time
evalmax<-evs$values[1]
evecmax<-evs$vectors[,1]
evecmax<-sign(evecmax[1])*evecmax/sqrt(as.numeric(crossprod(evecmax))) # normalize
emax<-list(evalmax=evalmax, evecmax=evecmax)
# save(emax, file="temax.Rdata")
evalmin<-evs$values[n]
evecmin<-evs$vectors[,n]
evecmin<-sign(evecmin[1])*evecmin/sqrt(as.numeric(crossprod(evecmin)))
avecmax<-amax$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 ",mean(tmin)*0.001,"\n")

## minimal eigensolution: Value= 4.781863e-08 in time 22929917

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

## Eigenvalue - result from eigen= 4.78183e-08 vector max(abs(diff))= 0.0001219604
#print(amin$par)
cat("maximal eigensolution: Value=",amax$value,"in time ",mean(tmax)*0.001,"\n")

## maximal eigensolution: Value= 3934.277 in time 545902.5

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

## Eigenvalue - result from eigen= -3.760964e-06 vector max(abs(diff))= 4.746153e-06
nmax<-10
stable<-matrix(NA, nrow=nmax, ncol=4) # to hold results

```



```
# ===== works to here, but spg is slower than eigen
# loop over sizes
for (ni in 1:nmax){
  ni<-1
  n<-50*ni
  x<-runif(n) # generate a vector
  AA<-molerfast(n) # make sure defined
  stable[[ni, 1]]<-n
  tbld<-microbenchmark(AA<-molerfast(n), times=mbt)
  tspg<-microbenchmark(asp<-spg(x, fn=rqneg, project=proj,
                                control=list(trace=FALSE)), times=mbt)
  teig<-microbenchmark(aseig<-eigen(AA), times=mbt)
  stable[[ni, 2]]<-mean(tspg$time)*0.001
  stable[[ni, 3]]<-mean(tbld$time)*0.001
  stable[[ni, 4]]<-mean(teig$time)*0.001
}
spgty<-data.frame(n=stable[,1], spgrqt=stable[,2], tbld=stable[,3], teig=stable[,4])
print(round(spgty,0))
```

```
##      n spgrqt tbld teig
## 1  50 204526  305  477
## 2  NA      NA   NA   NA
## 3  NA      NA   NA   NA
## 4  NA      NA   NA   NA
## 5  NA      NA   NA   NA
## 6  NA      NA   NA   NA
## 7  NA      NA   NA   NA
## 8  NA      NA   NA   NA
## 9  NA      NA   NA   NA
## 10 NA      NA   NA   NA
```

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 more efficient `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)
```

```
## Loading required package: 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){
  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))
}
# require(optplus)
# mset<-c("L-BFGS-B", "BFGS", "CG", "spg", "ucminf", "nlm", "nlminb", "Rummin", "Rcgmin")
mset<-c("L-BFGS-B", "BFGS", "nbg", "spg", "ucminf", "nlm", "nlminb", "nvm")
nmax<-5
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(starttests=FALSE, dowarn=FALSE))
  # optansout(aall, NULL)
  summary(aall, order=value, )
  cat("Above for n=",n," \n")
}

```

```

## 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

```

```

## 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

```

```

## 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

```

```

## Above for n= 20

```

```

## 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

## 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

## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing

## Above for n= 40

## 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

## 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

## Above for n= 60

## 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

```

```
## 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

## 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

## Above for n= 80

## 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

## 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

## Warning in kktchk(ans$par, fn, wgr, hess = NULL, upper = NULL, lower = NULL, :
## kktchk: pHes not symmetric -- symmetrizing

## Above for n= 100
```

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 `ncg`. Below we use the latter to show the times over different matrix sizes.

```
ctable<-matrix(NA, nrow=10, ncol=2)
nmax<-10
for (ni in 1:nmax){
  n<-50*ni
  x<-runif(n) # generate a vector
  AA<-molerfast(n) # define matrix
  tcgu<-microbenchmark(arcgu<-optimr(x, fn=nobj, gr=ngrobj, method="ncg",
```

```

        AA=-AA), times=mbt)
cetable[[ni,1]]<-n
cetable[[ni,2]]<-mean(tcgu$time)*0.001
}
cgtime<-data.frame(n=cetable[,1], tcgmin=cetable[,2])
print(round(cgtime,0))

```

```

##      n tcgmin
## 1   50    517
## 2  100   1204
## 3  150   1960
## 4  200   3227
## 5  250   4434
## 6  300   7064
## 7  350   7714
## 8  400  13370
## 9  450  16049
## 10 500 27099

```

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<-molermat(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] 191829.6335 95915.2284 47958.6432 23981.4826 11995.1147 6006.3298
## [7] 3020.7228 1545.4837 842.9897 561.9923
##
## $RQ
## [1] 8.582807e-06
##
## $ipr
## [1] 52
##
## $msg
## [1] "Small gradient -- done"
cat("Geradin time=",mean(tg$time),"n")

```

```
## Geradin time= 2792948
```

```
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] -73315037617 2477975547 78187075154 151253321889 219209885501
## [6] 279756302565 330848904031 370761681939 398142206049 412069042406
##
## $RQ
## [1] -31.58981
##
## $ipr
## [1] 38
##
## $msg
## [1] "Small gradient -- done"
```

```
cat("Geradin time=",mean(tgn$time),"\n")
```

```
## Geradin time= 511553.9
```

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
  res<-(-1)*(.Fortran("moler", n=as.integer(n), x=as.double(x), vout=as.double(vout)))$vout
}
```

```
require(microbenchmark)
```

```
nmax<-10
```

```
gtable<-matrix(NA, nrow=nmax, ncol=6) # to hold results
# loop over sizes
```

```

for (ni in 1:nmax){
  n<-50*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]]<-mean(tgax$time)
  tgaximp<-microbenchmark(ogaximp<-geradin(x, naximp, ident, AA=1, BB=1, control=list(trace=FALSE)), times=mbt)
  gtable[[ni, 3]]<-mean(tgaximp$time)
  tgaxftn<-microbenchmark(ogaxftn<-geradin(x, naxftn, ident, AA=1, BB=1, control=list(trace=FALSE)), times=mbt)
  gtable[[ni, 4]]<-mean(tgaxftn$time)
}

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

```

```

##      n      ax      aximp      axftn
## 1   50  635554.9  28376217  884638.4
## 2  100  953088.8 106678242 1406820.2
## 3  150 1620909.0 271475287 2852235.7
## 4  200 2318585.0 485753357 4552841.3
## 5  250 2911666.3 649815328 5831532.3
## 6  300 4507873.3 1089595836 9365230.0
## 7  350 5043376.9 1275443208 10811619.6
## 8  400 6763758.6 1614937973 13448639.5
## 9  450 9301660.3 2110496556 17431933.4
## 10 500 14870457.1 3013031882 24622146.2

```

Let us check that the solution for $n = 100$ by Geradin is consistent with the answer via `eigen()`.

```

n<-100
x<-runif(n)
# emax<-load("temax.Rdata")
evalmax<-emax$evalmax
evecmac<-emax$evecmac
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 diff eigenval from eigen result: ",gval-evalmax,"    max(abs(vector diff))=",
    max(abs(diff)), "\n")

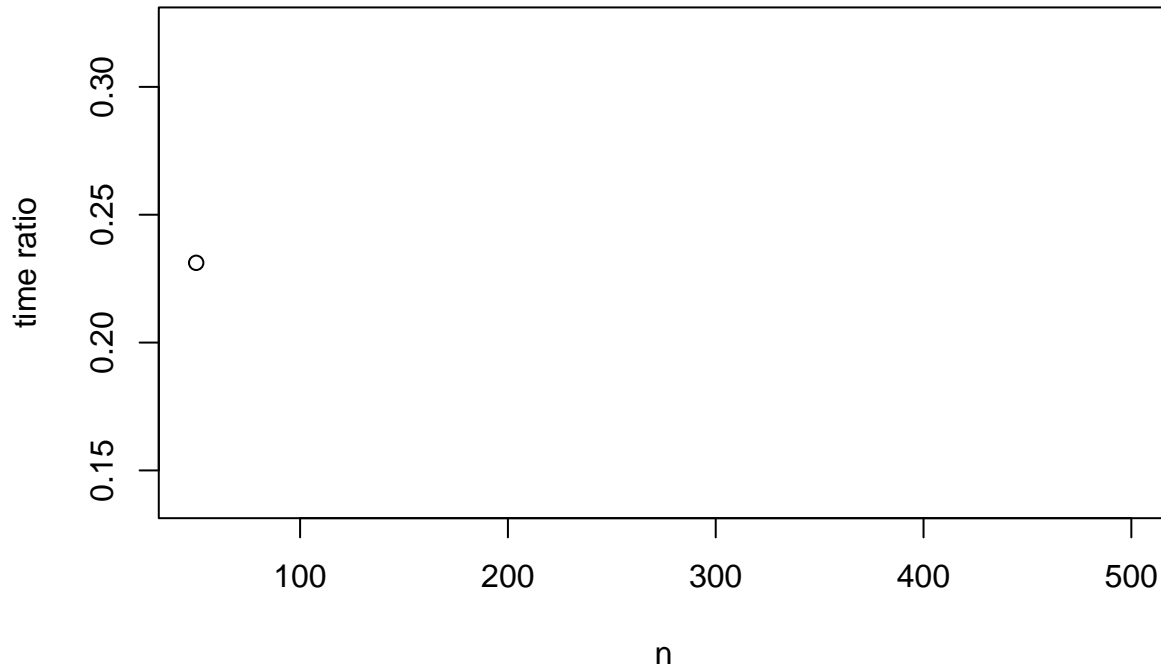
```

```
## Geradin diff eigenval from eigen result: -6.758855e-06    max(abs(vector diff))= 8.150972e-06
```

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.

Ratio of eigensolution times to Geradin routine by matrix size



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

```
## Times in seconds
## Build = 82435948  eigen(): 7295583870  Rcgminu: 286467271  Geradin: 370713834
## Ratios: build= 0.2223708 eigen= 19.67983  Rcgminu= 0.772745
```

Conclusions}

The Rayleigh Quotient minimization approach to eigensolutions has an intuitive appeal and seemingly offers an interesting optimization test problem, especially if we can make it computationally efficient. To improve time efficiency, we can apply the R byte code compiler, use a Fortran (or other compiled language) subroutine, and choose how we set up our objective functions and gradients. To improve memory use, we can consider using a matrix implicitly.

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:

- The R byte code compiler offers a useful gain in speed when our code has statements that access array elements rather than uses them in vectorized form.}
- The `crossprod()` function is very efficient.
- Fortran is not very difficult to use for small subroutines that compute a function such as the implicit matrix-vector product, and it allows efficient computations for such operations.
- 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

```

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
}

```

References

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