Exceedingly Simple Gram-Schmidt Code

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1 Problem

The QR decomposition of a rectangular $n \times m$ matrix X of rank m is of the form X = QR, with Q $n \times m$ orthonormal and R non-singular and square upper-triangular of order m. If X has rank r < m we can still make the decomposition, but we allow some columns of Q and some rows of R to be zero.

There are various ways to compute the QR decomposition. In this note we implement the Gram-Schmidt or GS method in both R and C. GS operates on each of the columns of X in turn, and replaces them by the columns of Q.

```
[,1] [,2] [,3]
##
## [1,]
             1
                   1
                        0
## [2,]
             2
                   1
                        1
## [3,]
             3
                  0
                        3
## [4,]
                        3
             4
## [5,]
             5
                        4
                  1
```

2 Example

```
x<-matrix (rnorm(12), 3, 4)
print (b <- solve (x[,1:3], x[,4]))
## [1] -3.68212898 -0.06338116 52.63346992
print(h <- gsrc(x))</pre>
## $q
                         [,2]
                                   [,3]
##
              [,1]
                                                 [,4]
## [1,] 0.18827415 -0.9683207 0.16403643 -1.554312e-15
## [2,] -0.98201034 -0.1880653 0.01694537 2.414735e-15
## [3,] -0.01444101 0.1642758 0.98630873 0.000000e+00
##
## $r
##
           [,1]
                     [,2]
                                [,3]
                                           [,4]
## [1,] 3.100749 0.7108688 0.223148917 0.28268835
## [2,] 0.000000 1.3640832 0.002042439 0.02104348
## [3,] 0.000000 0.0000000 0.040102686 2.11074351
##
## $rank
## [1] 3
h$q[,1:3]
                        [,2]
              [,1]
                                   [,3]
## [1,] 0.18827415 -0.9683207 0.16403643
## [2,] -0.98201034 -0.1880653 0.01694537
## [3,] -0.01444101 0.1642758 0.98630873
```

```
x[,4]
## [1] 0.3790849 -0.2457931 2.0812194

colSums(x[,4]*h$q[,1:3])/b
## [1] -0.07677307 -0.33201473 0.04010269
```

3 Timing

##

##

C 87.9257 112.7299

Q 263.1983 300.5889

```
set.seed (12345)
x<-matrix (rnorm (1000000L), 10000L, 100L)
library (microbenchmark)
mb < -microbenchmark(R = gs(x), C = gsrc(x), Q = qr(x), times = 100L)
mb
## Unit: milliseconds
##
                           lq
                                   mean
                                           median
                                                          uq
                                                                   max neval
##
       R 901.55286 1070.5833 1129.0989 1123.8111 1186.3891 1562.5058
                                                                          100
                    113.7328
##
          90.77601
                              133.6758
                                         129.8622
                                                    143.9768
                                                              280.4501
                                                                          100
##
          88.92902
                    104.5294 115.5634
                                         113.3968
                                                    123.7876
                                                              183.0908
                                                                          100
```

Thus for this example the C code is about 8-10 times as fast as the R code. The QR decomposition that comes with R, based on Householder transformations, is again twice as fast.

In a personal communication Bill Venables pointed out (01/18/16) that the above timing comparisons are somewhat unfavorable to our routines, because the standard qr routines in R still have to dig Q and R out of the qr structure. So an alternative, and perhaps more suitable comparison, is

```
mb<-microbenchmark(R = gs(x), C = gsrc(x), Q = {qrx <- qr(x); list(q = qr.Q(qrx), r = qr
mb

## Unit: milliseconds
## expr min lq mean median uq max neval
## R 880.3235 987.0325 1100.4779 1086.1389 1201.6151 1573.2712 100</pre>
```

334.6637

144.5255

384.1192

244.5871

584.5522

100

100

Now gsrc is faster than qr, which now includes the cost of the copies and assignments. So a completely fair comparison will be somewhere in between the two benchmark results.

131.5274 124.2346

348.3349

4 Appendix: Code

```
dyn.load("gs.so")
gs \leftarrow function (x, eps = 1e-10) {
 n \leftarrow nrow(x)
 m \leftarrow ncol(x)
 q <- matrix (0, n, m)
 r <- matrix (0, m, m)
 h \leftarrow .C("gsc", x = as.double(x), q = as.double(q), r = as.double(r), n = as.integer(n)
  return (list (q = matrix(hq, n, m), r = matrix (hr, m ,m), rank = hrank))
}
#include <math.h>
void
gsc (double *x, double *q, double *r, int *n, int *m, int *rank, double *eps)
                     i, j, l, jn, ln, jm, imax = *n, jmax = *m;
    int
                s = 0.0;
    double
    *rank = 0;
    for (i = 0; i < imax; i++)
        s += *(x + i) * *(x + i);
    if (s > *eps) {
        *rank = 1;
        s = sqrt(s);
        *r = s;
        for (i = 0; i < imax; i++)
            *(q + i) = *(x + i) / s;
    for (j = 1; j < jmax; j++) {
        jn = j * imax;
        jm = j * jmax;
        for (1 = 0; 1 < j; 1++) {
            ln = l * imax;
            s = 0.0;
            for (i = 0; i < imax; i++)
                s += *(q + ln + i) * *(x + jn + i);
            *(r + jm + 1) = s;
            for (i = 0; i < imax; i++)
                *(q + jn + i) += s * *(q + ln + i);
        for (i = 0; i < imax; i++)
```

```
*(q + jn + i) = *(x + jn + i) - *(q + jn + i);
s = 0.0;
for (i = 0; i < imax; i++)
    s += *(q + jn + i) * *(q + jn + i);
if (s > *eps) {
    s = sqrt(s);
    *rank = *rank + 1;
    *(r + jm + j) = s;
    for (i = 0; i < imax; i++)
        *(q + jn + i) /= s;
}
</pre>
```

5 NEWS

6 References