

User's Guide for SuiteSparseQR, a multifrontal multithreaded sparse QR factorization package (with optional GPU acceleration)

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Abstract

SuiteSparseQR is an implementation of the multifrontal sparse QR factorization method. Parallelism is exploited both in the BLAS and across different frontal matrices using Intel's Threading Building Blocks, a shared-memory programming model for modern multicore architectures. It can obtain a substantial fraction of the theoretical peak performance of a multicore computer. The package is written in C++ with user interfaces for MATLAB, C, and C++. Both real and complex sparse matrices are supported.

1 Introduction

The algorithms used in SuiteSparseQR are discussed in a companion paper, [7], and an overview of how to use the software is given in [6]. This document gives detailed information on the installation and use of SuiteSparseQR.

2 Using SuiteSparseQR in MATLAB

The simplest way to use SuiteSparseQR is via MATLAB. Its syntax includes every feature of the MATLAB `qr` in version R2009a and earlier [12], plus additional features not available in MATLAB. It is also a replacement for $\mathbf{x}=\mathbf{A}\backslash\mathbf{b}$ for least-squares problems and underdetermined systems. In addition to substantial gains in performance (10x to 100x is not uncommon, up to 10,000x has been observed), SuiteSparseQR adds new capabilities that are not present in MATLAB. For example, it provides an efficient method for finding the minimum 2-norm solution to an underdetermined system.

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2.1 Installing SuiteSparseQR for use in MATLAB

All packages in SuiteSparse, including SuiteSparseQR and the codes it relies on (AMD, COLAMD, CHOLMOD, METIS, CCAMD, and CCOLAMD) are compiled with a single command typed into the MATLAB Command Window. SuiteSparseQR uses the LAPACK and BLAS libraries provided with MATLAB; you do not need to do anything to use these. Below are step-by-step instructions for compiling all of SuiteSparse (including SuiteSparseQR), and optional instructions on using METIS and/or Intel's Threading Building Blocks (TBB).

2.1.1 Required instructions for Windows

For Windows, you cannot use the `lcc` compiler that ships with MATLAB; it is not a C++ compiler. To compile SuiteSparseQR, you must obtain a C++ compiler; Microsoft Visual Studio C++ Express Edition will work fine. Install this compiler from <http://www.microsoft.com/express/vc/> and then type `mex -setup` in the MATLAB Command Window.

2.1.2 Optional instructions on using METIS for any operating system

SuiteSparse now relies on METIS 5.1.0, which is distributed along with SuiteSparse itself. Its use is optional, however. If you compile with `-DNPARTITION`, or if you delete or move the `SuiteSparse/metis-5.1.0` folder, then SuiteSparse will be compiled without it.

METIS tends to give orderings that are good for the parallelism exploited by TBB, so its use with TBB is recommended. Note however that METIS is not bug-free; it can occasionally cause segmentation faults, particularly if used when finding basic solutions to underdetermined systems with many more columns than rows (SuiteSparseQR does not use METIS, by default, for those systems). This (rare) faulty behavior has been confirmed with valid inputs to the METIS test programs themselves; it is not a bug in the SuiteSparse interface to METIS. Use METIS at your own risk.

2.1.3 Optional instructions for using TBB on Linux/Unix/Mac

If you are using a Debian-based Linux system (such as Ubuntu), you're in luck! You can install TBB via the Synaptic Package Manager. Just search for TBB, select it, and click Apply. This will place the right files in `/usr/lib` and it will create the `/usr/include/tbb` directory. It's by far the simplest way to install TBB. If you do this, skip the rest of this section.

Alternatively, obtain a copy of TBB from <http://www.threadingbuildingblocks.org> as a `tbb*.tgz` file appropriate for your version of Linux/Unix. If you install via the `tbb*.tgz` file, make sure the `libtbb*.so*` and `libtbbmalloc*.so*` files are placed in the `/usr/lib` directory. Make sure the `tbb` directory containing all of the include files is placed in `/usr/include` (for example, the `/usr/include/tbb/task_scheduler_init.h` file must exist).

If you do not have permission to install TBB properly into the `/usr/lib` and `/usr/include` directories, you can place it in your own directory and modify the `tbb_path` variable in

`spqr_make.m` to specify the location of your own copy of TBB (refer to that file for instructions). You must also set your `LD_LIBRARY_PATH` environment variable to include the directory containing the `libtbb*so` files. You need to first determine which subdirectory of the TBB distribution contains the libraries appropriate for your system; in the examples below this is just called `/path`. It must be an absolute path, starting with the `/` character. There must be no spaces in the path name.

For Linux use this command at the system command line before starting MATLAB, where `/path` should be replaced with the actual full path of the TBB `lib` directory:

```
setenv LD_LIBRARY_PATH /path:$LD_LIBRARY_PATH
```

If you use the `cs` shell, place the command in your `~/.cshrc` file so you don't have to type it each time you start MATLAB.

For the Mac, edit the `/etc/profile` file and add this line to the end of the file:

```
export DYLD_LIBRARY_PATH=/path:$DYLD_LIBRARY_PATH
```

For example, using the 64-bit Linux version placed in my home directory, this command would be placed in my `~/.cshrc` file, prior to starting MATLAB.

```
setenv LD_LIBRARY_PATH /home/davis/tbb21_009oss/em64t/cc4.1.0_libc2.4_kernel2.6.16.21/lib:$LD_LIBRARY_PATH
```

Then, before starting MATLAB, make sure this variable is set by typing this command at the system command line (you only have to do this for this session; whenever you start a new command shell this will be done automatically):

```
source ~/.cshrc
```

For this example, my `spqr_make.m` file would contain this line:

```
tbb_path = '/home/davis/tbb21_009oss' ;
```

2.1.4 Optional instructions for using TBB on Windows

Obtain a copy of TBB from <http://www.threadingbuildingblocks.org> ; for TBB Version 2.1 this file is `tbb21_009oss_win.zip`.

Create a folder and place the `tbb21_009oss_win.zip` file there, and extract the file. In the example below, I extracted to `C:\TBB\tbb21_009oss` (if you do the same then you do not have to edit `spqr_make.m`). There should be no spaces in the path (for example, placing TBB under the `Program Files` directory will not work).

In Windows XP, right-click `My Computer` and select `Properties`. Click the `Advanced` tab. Click `Environment Variables`. Under `System` variables, edit the `Path` to append the name of the folder containing the TBB `bin` folder appropriate for your system, preceded by a semicolon. For example, in a 32-bit Windows system if TBB is installed in `C:\TBB` you would append the string `;C:\TBB\tbb21_009oss\ia32\vc9\bin\` to the end of your system `Path` variable.

For Vista, the instructions are the same, except that you choose **Computer** instead of **My Computer**, and you click the **Advanced System Settings** tab instead of the **Advanced** tab.

Next, edit the `SPQR\MATLAB\spqr_make.m` file. Change the `tbb_path` variable to point to your copy of TBB. For example, if you installed TBB into `C:\TBB` your `spqr_make.m` file would contain this line:

```
tbb_path = 'C:\TBB\tbb21_009oss' ;
```

That line is already in `spqr_make.m`, so if you choose to install TBB in that location, you do not have to edit the file.

2.1.5 Now you're ready to compile (on any operating system)

Type these commands in the MATLAB window:

```
cd SuiteSparse
SuiteSparse_install
```

You will be asked if you want to run some demos. I recommend that you do this to ensure your functions have been installed correctly. Next type the command

```
pathtool
```

and examine your MATLAB path. The various SuiteSparse directories have been placed in your path. Click “save” to save this path for future MATLAB sessions. If this fails, you do not have permission to modify the `pathdef.m` file (it is shared by all users). An alternative is to type the command:

```
path
```

and cut-and-paste the paths displayed there into your own `startup.m` file, prepending the command `addpath` to each line. For example, if I installed SuiteSparse into my home directory (`/home/davis`) then my `startup.m` file should look like this:

```
addpath /home/davis/SuiteSparse/SPQR/MATLAB
addpath /home/davis/SuiteSparse/RBio
addpath /home/davis/SuiteSparse/MATLAB_Tools/spok
addpath /home/davis/SuiteSparse/MATLAB_Tools/waitmex
addpath /home/davis/SuiteSparse/MATLAB_Tools/shellgui
addpath /home/davis/SuiteSparse/MATLAB_Tools/GEE
addpath /home/davis/SuiteSparse/LINFACTOR
addpath /home/davis/SuiteSparse/MESHND
addpath /home/davis/SuiteSparse/UFcollection
addpath /home/davis/SuiteSparse/SSMULT
addpath /home/davis/SuiteSparse/KLU/MATLAB
addpath /home/davis/SuiteSparse/BTF/MATLAB
addpath /home/davis/SuiteSparse/LDL/MATLAB
addpath /home/davis/SuiteSparse/CXsparse/MATLAB/UFget
addpath /home/davis/SuiteSparse/CXsparse/MATLAB/Demo
```

```
addpath /home/davis/SuiteSparse/CXSparse/MATLAB/CSparse
addpath /home/davis/SuiteSparse/CAMD/MATLAB
addpath /home/davis/SuiteSparse/CCOLAMD/MATLAB
addpath /home/davis/SuiteSparse/COLAMD/MATLAB
addpath /home/davis/SuiteSparse/AMD/MATLAB
addpath /home/davis/SuiteSparse/CHOLMOD/MATLAB
addpath /home/davis/SuiteSparse/UMFPACK/MATLAB
addpath /home/davis/SuiteSparse
addpath /home/davis/SuiteSparse/MATLAB_Tools
```

On a Windows system, I might see paths like this instead in my `startup.m` file:

```
addpath C:\Documents and Settings\davis\My Documents\SuiteSparse\SPQR\MATLAB
...
```

Your `startup.m` file should appear in the directory in which MATLAB starts. Failing that, every time you start MATLAB, find your `startup.m` file and run it. For more help, type `doc startup` in MATLAB.

The `SuiteSparse_install` script works on any version of MATLAB (Linux/Unix, Mac, or Windows) if you have a C++ compiler. The install script will detect if you have placed the METIS directory in the right place, and will compile it for use with SuiteSparseQR if it finds it there. Otherwise METIS will be skipped (the install script will tell you if it finds METIS or not).

2.1.6 Optional instructions for using TBB on any system

If you have followed the steps above (as I recommend that you do), you have just compiled SuiteSparseQR, but it will not yet be using TBB. To use TBB with SuiteSparseQR, first install TBB as described above. Once TBB is installed, type the following commands in the MATLAB Command Window (assuming you have METIS):

```
cd SuiteSparse/SPQR/MATLAB
spqr_make metis tbb
```

Or if you do not have METIS, do this instead:

```
cd SuiteSparse/SPQR/MATLAB
spqr_make nometis tbb
```

For more options, type `help spqr_make`.

2.2 Functions provided to the MATLAB user

Three primary functions are available:

1. `spqr`, a replacement for the MATLAB `qr`

2. `spqr_solve`, a replacement for $x=A\backslash b$ when A is sparse and rectangular. It works for the square case, too, but $x=A\backslash b$ will be faster (using LU or Cholesky factorization). `spqr_solve` is a good method for ill-conditioned or rank-deficient square matrices, however.
3. `spqr_qmult`, which multiplies Q (stored in Householder vector form) times a matrix x .

Their syntax is described below in the table below. The permutation P is chosen to reduce fill-in and to return R in upper trapezoidal form if A is estimated to have less than full rank. The `opts` parameter provides non-default options (refer to the next section). The output Q can be optionally returned in Householder form, which is far sparser than returning Q as a sparse matrix.

<code>R = spqr (A)</code>	Q-less QR factorization
<code>R = spqr (A,0)</code>	economy variant (<code>size(R,1) = min(m,n)</code>)
<code>R = spqr (A,opts)</code>	as above, with non-default options
<code>[Q,R] = spqr (A)</code>	$A=Q*R$ factorization
<code>[Q,R] = spqr (A,0)</code>	economy variant (<code>size(Q,2) = size(R,1) = min(m,n)</code>)
<code>[Q,R] = spqr (A,opts)</code>	$A=Q*R$, with non-default options
<code>[Q,R,P] = spqr (A)</code>	$A*P=Q*R$ where P reduces fill-in
<code>[Q,R,P] = spqr (A,0)</code>	economy variant (<code>size(Q,2) = size(R,1) = min(m,n)</code>)
<code>[Q,R,P] = spqr (A,opts)</code>	as above, with non-default options
<code>[C,R] = spqr (A,B)</code>	as $R=spqr(A)$, also returns $C=Q'*B$
<code>[C,R] = spqr (A,B,0)</code>	economy variant (<code>size(C,1) = size(R,1) = min(m,n)</code>)
<code>[C,R] = spqr (A,B,opts)</code>	as above, with non-default options
<code>[C,R,P] = spqr (A,B)</code>	as $R=spqr(A*P)$, also returns $C=Q'*B$
<code>[C,R,P] = spqr (A,B,0)</code>	economy variant (<code>size(C,1) = size(R,1) = min(m,n)</code>)
<code>[C,R,P] = spqr (A,B,opts)</code>	as above, with non-default options
<code>x = spqr_solve (A,B)</code>	$x=A\backslash B$
<code>[x,info] = spqr_solve (A,B,opts)</code>	as above, with statistics and non-default parameters
<code>Y = spqr_qmult (Q,X,k)</code>	computes $Q'*X$, $Q*X$, $X*Q'$, or $X*Q$ (selected with k)

2.3 The `opts` parameter

The `opts` struct provides control over non-default parameters for SuiteSparseQR. Entries not present in `opts` are set to their defaults.

- `opts.tol`: columns that have 2-norm \leq `opts.tol` are treated as zero. The default is $20*(m+n)*eps*sqrt(\max(\text{diag}(A'*A)))$ where $[m \ n]=\text{size}(A)$.
- `opts.econ`: number of rows of R and columns of Q to return. The default is m . Using n gives the standard economy form (as in the MATLAB `qr(A,0)`). A value less than the estimated rank r is set to r , so `opts.econ=0` gives the “rank-sized” factorization, where `size(R,1)==nnz(diag(R))==r`.
- `opts.ordering`: a string describing which column ordering method to use. Let $[m2 \ n2]=\text{size}(S)$ where S is obtained by removing singletons from A . The singleton permutation places $A*P$ in the form $[A11 \ A12 \ ; \ 0 \ S]$ where $A11$ is upper triangular with diagonal entries all greater than `opts.tol`.

The default is to use COLAMD if $m_2 \leq 2 \cdot n_2$; otherwise try AMD. Let f be the flops for $\text{chol}((S \cdot P)' \cdot (S \cdot P))$ with the ordering P found by AMD. Then if $f/\text{nnz}(R) \geq 500$ and $\text{nnz}(R)/\text{nnz}(S) \geq 5$ then try METIS, and take the best ordering found (AMD or METIS); otherwise use AMD without trying METIS. If METIS is not installed then the default ordering is to use COLAMD if $m_2 \leq 2 \cdot n_2$ and to use AMD otherwise.

The available orderings are:

'default': the default ordering.

'amd': use $\text{amd}(S' \cdot S)$.

'colamd': use $\text{colamd}(S)$.

'metis': use $\text{metis}(S' \cdot S)$, only if METIS is installed.

'best': try all three (AMD, COLAMD, METIS) and take the best.

'bestamd': try AMD and COLAMD and take the best.

'fixed': use $P=I$; this is the only option if P is not present in the output.

'natural': singleton removal only.

- `opts.Q`: a string describing how Q is to be returned. The default is 'discard' if Q is not present in the output, or 'matrix' otherwise. If Q is present and `opts.Q` is 'discard', then $Q=[]$ is returned (thus $R=\text{spqr}(A \cdot P)$ is $[Q,R,P]=\text{spqr}(A)$ where spqr finds P but Q is discarded instead). The usage `opts.Q='matrix'` returns Q as a sparse matrix where $A=Q \cdot R$ or $A \cdot P=Q \cdot R$. Using `opts.Q='Householder'` returns Q as a struct containing the Householder reflections applied to A to obtain R , resulting in a far sparser Q than the 'matrix' option.
- `opts.permutation`: a string describing how P is to be returned. The default is 'matrix', so that $A \cdot P=Q \cdot R$. Using 'vector' gives $A(:,P)=Q \cdot R$ instead.
- `opts.spumoni`: an integer k that acts just like `spparms('spumoni',k)`.
- `opts.min2norm`: used by `spqr_solve`; you can use 'basic' (the default), or 'min2norm'. Determines the kind of solution that `spqr_solve` computes for underdetermined systems. Has no effect for least-squares problems; ignored by `spqr` itself.
- `opts.grain`, `opts.small`, `opts.nthreads`: multitasking control (if compiled with TBB). Let f be the total flop count. The analysis phase tries to ensure that all parallel tasks have at least $\max(\text{total_flops}/\text{opts.grain}, \text{opts.small})$ flops. No TBB parallelism is exploited if `opts.grain` ≤ 1 . The parameter `opts.nthreads` gives the number of threads to use for TBB (which is different than the number of threads used by the BLAS). Setting `opts.nthreads` ≤ 0 means to let TBB determine the number of threads (normally equal to the number of cores); otherwise, exactly `opts.nthreads` threads are used. The defaults are `opts.grain=1`, `opts.small=1e6`, and `opts.nthreads=0`, respectively. That is, TBB is disabled by default since it conflicts with BLAS multithreading. If you enable TBB, be sure to disable BLAS multithreading with the MATLAB command `maxNumCompThreads(1)`, or choose

`opts.nthreads=k` and `maxNumCompThreads(b)` so that the product $k*b$ is equal to the number of cores. Note that these recommendations may change for future versions of TBB. A good value of `opts.grain` is twice that of `opts.nthreads`. If TBB parallelism is enabled, the METIS ordering normally gives the best speedup for large problems.

2.4 Examples on how to use the MATLAB interface

To solve a least-squares problem, or to find the basic solution to an underdetermined system, just use `x = spqr_solve(A,b)` in place of `x=A\b`. To compute the QR factorization, use `[Q,R]=spqr(A)` instead of `[Q,R]=qr(A)`. Better results can be obtained by discarding Q with the usage `R=spqr(A)` (in place of `R=qr(A)`), or by requesting Q in Householder form with `[Q,R]=spqr(A,opts)` where `opts.Q='Householder'`. The latter option is not available in MATLAB. To use a fill-reducing ordering, simply use any of the syntaxes above with P as an output parameter.

The least-squares solution of an overdetermined system $A*x=b$ with $m>n$ (where A has rank n) can be found in one of at least seven ways (in increasing order of efficiency, in time and memory):

<code>x = pinv(full(A)) * b ;</code>	impossible for large A
<code>[Q,R] = spqr (A) ;</code> <code>x = R\ (Q'*b) ;</code>	high fill-in in R, Q costly in matrix form
<code>[Q,R,P] = spqr (A) ;</code> <code>x = P*(R\ (Q'*b)) ;</code>	low fill-in in R, Q costly in matrix form
<code>[Q,R,P] = spqr (A,struct('Q','Householder')) ;</code> <code>x = P*(R\spqr_qmult (Q,b,0)) ;</code>	low fill-in in R, Q in efficient Householder form
<code>[c,R,P] = spqr (A,b) ;</code> <code>x = P*(R\c) ;</code>	Q not kept, P a permutation matrix
<code>[c,R,p] = spqr (A,b,0) ;</code> <code>y = (R\c) ; x(p) = y</code>	Q not kept, p a permutation vector
<code>x = spqr_solve (A,b) ;</code>	less memory and better handling of rank-deficient matrices

The minimum-norm solution of an underdetermined system $A*x=b$ with $m<n$ can be found in one of five ways (in increasing order of efficiency, in time and memory):

<code>x = pinv(full(A)) * b ;</code>	impossible for large A
<code>[Q,R] = spqr (A') ;</code> <code>x = Q*(R'\b) ;</code>	high fill-in in R, Q costly in matrix form
<code>[Q,R,P] = spqr (A') ;</code> <code>x = Q*(R'\(P'*b)) ;</code>	low fill-in in R, Q costly in matrix form
<code>[Q,R,P] = spqr (A',struct('Q','Householder')) ;</code> <code>x = spqr_qmult (Q,R'\(P'*b),1) ;</code>	low fill-in in R, Q in efficient Householder form
<code>opts.solution = 'min2norm' ;</code> <code>x = spqr_solve (A,b,opts) ;</code>	as 4th option above, but faster, less memory, and better handling of rank-deficient matrices

Note that `spqr_solve` uses a fill-reducing ordering, by default. It can be disabled or modified using a non-default `opts` parameter (`opts.ordering`, specifically).

3 Using SuiteSparseQR in C and C++

SuiteSparseQR relies on CHOLMOD for its basic sparse matrix data structure, a compressed sparse column format. CHOLMOD provides interfaces to the AMD, COLAMD, and METIS ordering methods, supernodal symbolic Cholesky factorization (namely, `symbfact` in MATLAB), functions for converting between different data structures, and for basic operations such as transpose, matrix multiply, reading a matrix from a file, writing a matrix to a file, and many other functions.

3.1 Installing the C/C++ library on Linux/Unix

Before you compile the SuiteSparseQR library and demo programs, you may wish to edit the `SuiteSparse/SuiteSparse_config/SuiteSparse_config.mk` configuration file. The defaults should be fine on most Linux/Unix systems and on the Mac. It automatically detects what system you have and sets compile parameters accordingly.

Next, type `make` at the Linux/Unix command line, in either the `SuiteSparse` directory (which compiles all of SuiteSparse) or in the `SuiteSparse/SPQR` directory (which just compiles SuiteSparseQR and the libraries it requires). SuiteSparseQR will be compiled, and a set of simple demos will be run (including the one in the next section).

The configuration file defines where the LAPACK and BLAS libraries are to be found. Selecting the right BLAS is critical. There is no standard naming scheme for the name and location of these libraries. The defaults in the `SuiteSparse_config.mk` file use `-llapack` and `-lblas`; the latter may link against the standard Fortran reference BLAS, which will not provide optimal performance. For best results, you should use the OpenBLAS at `openblas.net` (based on the Goto BLAS) [13], or high-performance vendor-supplied BLAS such as the Intel MKL, AMD ACML, or the Sun Performance Library. Selection of LAPACK and the BLAS is done with the `LAPACK=` and `BLAS=` lines in the `SuiteSparse_config.mk` file.

Four compile-time options can be used to modify how SuiteSparseQR is compiled. Select these via the `SPQR_CONFIG=` line in the `SuiteSparse_config.mk` file.

- `-DNPARTITION`: do not compile with METIS, CAMD, or CCOLAMD. These packages are included by default.
- `-DNEXPERT`: do not compile with the “expert” routines in `SuiteSparseQR_expert.cpp`. The expert routines are included by default.
- `-DHAVE_TBB`: enable the Intel Threading Building Blocks, TBB. The use of TBB is disabled by default. It is disabled because not all installations have TBB available. The use of TBB is recommended, however, if you have a multicore computer.

To fully test 100% of the lines of SuiteSparseQR, go to the `Tcov` directory and type `make`. This will work for Linux only.

To install the shared library into `/usr/local/lib` and `/usr/local/include`, do `make install`. To uninstall, do `make uninstall`. For more options, see the `SuiteSparse/README.txt` file.

3.2 C/C++ Example

The C++ interface is written using templates for handling both real and complex matrices. The simplest function computes the MATLAB equivalent of $\mathbf{x}=\mathbf{A}\backslash\mathbf{b}$ and is almost as simple:

```
#include "SuiteSparseQR.hpp"
X = SuiteSparseQR <double> (A, B, cc) ;
```

The C version of this function is almost identical:

```
#include "SuiteSparseQR_C.h"
X = SuiteSparseQR_C_backslash_default (A, B, cc) ;
```

Below is a simple C++ program that illustrates the use of SuiteSparseQR. The program reads in a least-squares problem from `stdin` in MatrixMarket format [4], solves it, and prints the norm of the residual and the estimated rank of \mathbf{A} . The comments reflect the MATLAB equivalent statements. The C version of this program is identical except for the `#include` statement and call to SuiteSparseQR which are replaced with the C version of the statement above, and C-style comments.

```
#include "SuiteSparseQR.hpp"
int main (int argc, char **argv)
{
    cholmod_common Common, *cc ;
    cholmod_sparse *A ;
    cholmod_dense *X, *B, *Residual ;
    double rnorm, one [2] = {1,0}, minusone [2] = {-1,0} ;
    int mtype ;

    // start CHOLMOD
    cc = &Common ;
    cholmod_l_start (cc) ;

    // load A
    A = (cholmod_sparse *) cholmod_l_read_matrix (stdin, 1, &mtype, cc) ;

    // B = ones (size (A,1),1)
    B = cholmod_l_ones (A->nrow, 1, A->xtype, cc) ;

    // X = A\B
    X = SuiteSparseQR <double> (A, B, cc) ;

    // rnorm = norm (B-A*X)
    Residual = cholmod_l_copy_dense (B, cc) ;
    cholmod_l_sdmult (A, 0, minusone, one, X, Residual, cc) ;
    rnorm = cholmod_l_norm_dense (Residual, 2, cc) ;
    printf ("2-norm of residual: %8.1e\n", rnorm) ;
    printf ("rank %ld\n", cc->SPQR_istat [4]) ;

    // free everything and finish CHOLMOD
    cholmod_l_free_dense (&Residual, cc) ;
    cholmod_l_free_sparse (&A, cc) ;
    cholmod_l_free_dense (&X, cc) ;
    cholmod_l_free_dense (&B, cc) ;
```

```

    cholmod_l_finish (cc) ;
    return (0) ;
}

```

3.3 C++ Syntax

All features available to the MATLAB user are also available to both the C and C++ interfaces using a syntax that is not much more complicated than the MATLAB syntax. Additional features not available via the MATLAB interface include the ability to compute the symbolic and numeric factorizations separately (for multiple matrices with the same nonzero pattern but different numerical values). The following is a list of user-callable C++ functions and what they can do:

1. `SuiteSparseQR`: an overloaded function that provides functions equivalent to `spqr` and `spqr_solve` in the `SuiteSparseQR` MATLAB interface.
2. `SuiteSparseQR_factorize`: performs both the symbolic and numeric factorizations and returns a QR factorization object such that $A*P=Q*R$. It always exploits singletons.
3. `SuiteSparseQR_symbolic`: performs the symbolic factorization and returns a QR factorization object to be passed to `SuiteSparseQR_numeric`. It does not exploit singletons.
4. `SuiteSparseQR_numeric`: performs the numeric factorization on a QR factorization object, either one constructed by `SuiteSparseQR_symbolic`, or reusing one from a prior call to `SuiteSparseQR_numeric` for a matrix A with the same pattern as the first one, but with different numerical values.
5. `SuiteSparseQR_solve`: solves a linear system using the object returned by `SuiteSparseQR_factorize` or `SuiteSparseQR_numeric`, namely $x=R\backslash b$, $x=P*R\backslash b$, $x=R'\backslash b$, or $x=R'\backslash(P'*b)$.
6. `SuiteSparseQR_qmult`: provides the same function as `spqr_qmult` in the MATLAB interface, computing $Q'*x$, $Q*x$, $x*Q'$, or $x*Q$. It uses either a QR factorization in MATLAB-style sparse matrix format, or the QR factorization object returned by `SuiteSparseQR_factorize` or `SuiteSparseQR_numeric`.
7. `SuiteSparseQR_min2norm`: finds the minimum 2-norm solution to an underdetermined linear system.
8. `SuiteSparseQR_free`: frees the QR factorization object.

3.4 Details of the C/C++ Syntax

For further details of how to use the C/C++ syntax, please refer to the definitions and descriptions in the following files:

1. SuiteSparse/SPQR/Include/SuiteSparseQR.hpp describes each C++ function. Both `double` and `std::complex<double>` matrices are supported.
2. SuiteSparse/SPQR/Include/SuiteSparseQR_definitions.h describes definitions common to both C and C++ functions. For example, each of the ordering methods is given a `#define`'d name. The default is `ordering = SPQR_ORDERING_DEFAULT`, and the default tolerance is given by `tol = SPQR_DEFAULT_TOL`.
3. SuiteSparse/SPQR/Include/SuiteSparseQR_C.h describes the C-callable functions.

Most of the packages in SuiteSparse come in multiple versions with different sized integers. The first is the plain C/C++ `int`. The second the `SuiteSparse_long` integer, defined in the SuiteSparse/SuiteSparse_config/SuiteSparse_config.h file. This integer is `long` except on a Windows-64 platform for which it is the `__int64` type. The intent of `SuiteSparse_long` is that it should be 32-bits on a 32-bit platform, and 64-bits on a 64-bit platform.

By contrast, SuiteSparseQR only provides a `SuiteSparse_long` version. Most users (except Windows-64) can simply use `long` as the basic integer type passed to and returned from SuiteSparseQR.

The C/C++ options corresponding to the MATLAB `opts` parameters and the contents of the optional `info` output of `spqr_solve` are described below. Let `cc` be the CHOLMOD Common object, containing parameter settings and statistics. All are of type `double`, except for `SPQR_istat` which is `SuiteSparse_long`, `cc->memory_usage` which is `size_t`, and `cc->SPQR_nthreads` which is `int`. Parameters include:

<code>cc->SPQR_grain</code>	the same as <code>opts.grain</code> in the MATLAB interface
<code>cc->SPQR_small</code>	the same as <code>opts.small</code> in the MATLAB interface
<code>cc->SPQR_nthreads</code>	the same as <code>opts.nthreads</code> in the MATLAB interface

Other parameters, such as `opts.ordering` and `opts.tol`, are input parameters to the various C/C++ functions. Others such as `opts.solution='min2norm'` are separate functions in the C/C++ interface. Refer to the files listed above for details. Output statistics include:

<code>cc->SPQR_flopcount_bound</code>	an upper bound on the flop count
<code>cc->SPQR_tol_used</code>	the tolerance used (<code>opts.tol</code>)
<code>cc->SPQR_istat</code> [0]	upper bound on <code>nnz(R)</code>
<code>cc->SPQR_istat</code> [1]	upper bound on <code>nnz(H)</code>
<code>cc->SPQR_istat</code> [2]	number of frontal matrices
<code>cc->SPQR_istat</code> [3]	number of TBB tasks
<code>cc->SPQR_istat</code> [4]	estimate of the rank of <code>A</code>
<code>cc->SPQR_istat</code> [5]	number of column singletons
<code>cc->SPQR_istat</code> [6]	number of row singletons
<code>cc->SPQR_istat</code> [7]	ordering used
<code>cc->memory_usage</code>	memory used, in bytes

The upper bound on the flop count is found in the analysis phase, which ignores the numerical values of `A` (the same analysis phase operates on both real and complex matrices). Thus, if you are factorizing a complex matrix, multiply this statistic by 4.

4 GPU acceleration

As of version 2.0.0, SuiteSparseQR now includes GPU acceleration. It can exploit a single NVIDIA GPU, via CUDA. To enable GPU acceleration, you must compile SuiteSparseQR with non-default options. See the `SuiteSparse_config_GPU_gcc.mk` file in the `SuiteSparse_config` directory for details. The packages `SuiteSparse_GPURuntime` and `GPUQREngine` are also required (they should appear in the `SuiteSparse` directory, along with `SPQR`).

At run time, you must also enable the GPU by setting `Common->useGPU` to `true`. Before calling any SuiteSparseQR function, you must poll the GPU to set the available memory. Below is a sample code that initializes `CHOLMOD` and then polls the GPU for use in SuiteSparseQR.

```
size_t total_mem, available_mem ;
cholmod_common *cc, Common ;
cc = &Common ;
cholmod_l_start (cc) ;
cc->useGPU = true ;
cholmod_l_gpu_memorysize (&total_mem, &available_mem, cc) ;
cc->gpuMemorySize = available_mem ;
if (cc->gpuMemorySize <= 1)
{
    printf ("no GPU available\n") ;
}

// Subsequent calls to SuiteSparseQR will use the GPU, if available
```

See `Demo/qrdemo_gpu.cpp` for an extended example, which can be compiled via `make gpu` in the `Demo` directory.

GPU acceleration is not yet available via the MATLAB `mexFunction` interface. We expect to include this in a future release.

For a detailed technical report on the GPU-accelerated algorithm, see `qrgpu_paper.pdf` in the `Doc` directory.

5 Requirements and Availability

SuiteSparseQR requires four prior Collected Algorithms of the ACM: `CHOLMOD` [5, 10] (version 1.7 or later), `AMD` [1, 2], and `COLAMD` [8, 9] for its ordering/analysis phase and for its basic sparse matrix data structure, and the `BLAS` [11] for dense matrix computations on its frontal matrices; also required is `LAPACK` [3] for its Householder reflections. An efficient implementation of the `BLAS` is strongly recommended, either vendor-provided (such as the Intel MKL, the AMD ACML, or the Sun Performance Library) or other high-performance `BLAS` such as those of [13].

The use of Intel's Threading Building Blocks is optional [15], but without it, only parallelism within the `BLAS` can be exploited (if available). SuiteSparseQR can optionally use

METIS 4.0.1 [14] and two constrained minimum degree ordering algorithms, CCOLAMD and CAMD [5], for its fill-reducing ordering options. SuiteSparseQR can be compiled without these ordering methods and without TBB.

In addition to appearing as Collected Algorithm 8xx of the ACM, SuiteSparseQR is available at <http://www.suitesparse.com> and at MATLAB Central in the user-contributed File Exchange (<http://www.mathworks.com/matlabcentral>). See SPQR/Doc/License.txt for the license. Alternative licenses are also available; contact the author for details.

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