User Guide for CHOLMOD: a sparse Cholesky factorization and modification package

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Abstract

CHOLMOD¹ is a set of routines for factorizing sparse symmetric positive definite matrices of the form ${\bf A}$ or ${\bf A}{\bf A}^{\sf T}$, updating/downdating a sparse Cholesky factorization, solving linear systems, updating/downdating the solution to the triangular system ${\bf L}{\bf x}={\bf b}$, and many other sparse matrix functions for both symmetric and unsymmetric matrices. Its supernodal Cholesky factorization relies on LAPACK and the Level-3 BLAS, and obtains a substantial fraction of the peak performance of the BLAS. Both real and complex matrices are supported. It also includes a non-supernodal ${\bf L}{\bf D}{\bf L}^T$ factorization method that can factorize symmetric indefinite matrices if all of their leading submatrices are well-conditioned (${\bf D}$ is diagonal). CHOLMOD is written in ANSI/ISO C, with both C and MATLAB interfaces. This code works on Microsoft Windows and many versions of Unix and Linux.

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¹CHOLMOD is short for CHOLesky MODification, since a key feature of the package is its ability to update/downdate a sparse Cholesky factorization

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1 Overview

CHOLMOD is a set of ANSI C routines for solving systems of linear equations, $\mathbf{Ax} = \mathbf{b}$, when \mathbf{A} is sparse and symmetric positive definite, and \mathbf{x} and \mathbf{b} can be either sparse or dense.² Complex matrices are supported, in two different formats. CHOLMOD includes high-performance left-looking supernodal factorization and solve methods [21], based on LAPACK [3] and the BLAS [12]. After a matrix is factorized, its factors can be updated or downdated using the techniques described by Davis and Hager in [8, 9, 10]. Many additional sparse matrix operations are provided, for both symmetric and unsymmetric matrices (square or rectangular), including sparse matrix multiply, add, transpose, permutation, scaling, norm, concatenation, sub-matrix access, and converting to alternate data structures. Interfaces to many ordering methods are provided, including minimum degree (AMD [1, 2], COLAMD [6, 7]), constrained minimum degree (CSYMAMD, CCOLAMD, CAMD), and graph-partitioning-based nested dissection (METIS [18]). Most of its operations are available within MATLAB via mexFunction interfaces.

CHOLMOD also includes a non-supernodal \mathbf{LDL}^T factorization method that can factorize symmetric indefinite matrices if all of their leading submatrices are well-conditioned (\mathbf{D} is diagonal).

A pair of articles on CHOLMOD has been submitted to the ACM Transactions on Mathematical Software: [4, 11].

CHOLMOD 1.0 replaces chol (the sparse case), symbfact, and etree in MATLAB 7.2 (R2006a), and is used for x=A\b when A is symmetric positive definite [14]. It will replace sparse in a future version of MATLAB.

The C-callable CHOLMOD library consists of 133 user-callable routines and one include file. Each routine comes in two versions, one for int integers and another for long. Many of the routines can support either real or complex matrices, simply by passing a matrix of the appropriate type.

Nick Gould, Yifan Hu, and Jennifer Scott have independently tested CHOLMOD's performance, comparing it with nearly a dozen or so other solvers [17, 16]. Its performance was quite competitive.

²Some support is provided for symmetric indefinite matrices.

2 Primary routines and data structures

Five primary CHOLMOD routines are required to factorize \mathbf{A} or $\mathbf{A}\mathbf{A}^\mathsf{T}$ and solve the related system $\mathbf{A}\mathbf{x} = \mathbf{b}$ or $\mathbf{A}\mathbf{A}^\mathsf{T}\mathbf{x} = \mathbf{b}$, for either the real or complex cases:

- 1. cholmod_start: This must be the first call to CHOLMOD.
- 2. cholmod_analyze: Finds a fill-reducing ordering, and performs the symbolic factorization, either simplicial (non-supernodal) or supernodal.
- 3. cholmod_factorize: Numerical factorization, either simplicial or supernodal, LL^T or LDL^T using either the symbolic factorization from cholmod_analyze or the numerical factorization from a prior call to cholmod_factorize.
- 4. cholmod_solve: Solves $\mathbf{A}\mathbf{x} = \mathbf{b}$, or many other related systems, where \mathbf{x} and \mathbf{b} are dense matrices. The cholmod_spsolve routine handles the sparse case. Any mixture of real and complex \mathbf{A} and \mathbf{b} are allowed.
- 5. cholmod_finish: This must be the last call to CHOLMOD.

Additional routines are also required to create and destroy the matrices \mathbf{A} , \mathbf{x} , \mathbf{b} , and the $\mathbf{L}\mathbf{L}^\mathsf{T}$ or $\mathbf{L}\mathbf{D}\mathbf{L}^\mathsf{T}$ factorization. CHOLMOD has five kinds of data structures, referred to as objects and implemented as pointers to struct's:

- 1. cholmod_common: parameter settings, statistics, and workspace used internally by CHOLMOD. See Section 12 for details.
- 2. cholmod_sparse: a sparse matrix in compressed-column form, either pattern-only, real, complex, or "zomplex." In its basic form, the matrix A contains:
 - A->p, an integer array of size A->ncol+1.
 - A->i, an integer array of size A->nzmax.
 - A->x, a double array of size A->nzmax or twice that for the complex case. This is compatible with the Fortran and ANSI C99 complex data type.
 - A->z, a double array of size A->nzmax if A is zomplex. A zomplex matrix has a z array, thus the name. This is compatible with the MATLAB representation of complex matrices.

For all four types of matrices, the row indices of entries of column j are located in A->i [A->p [j] ... A->p [j+1]-1]. For a real matrix, the corresponding numerical values are in A->x at the same location. For a complex matrix, the entry whose row index is A->i [p] is contained in A->x [2*p] (the real part) and A->x [2*p+1] (the imaginary part). For a zomplex matrix, the real part is in A->x [p] and imaginary part is in A->z [p]. See Section 13 for more details.

3. cholmod_factor: A symbolic or numeric factorization, either real, complex, or zomplex. It can be either an $\mathbf{L}\mathbf{L}^\mathsf{T}$ or $\mathbf{L}\mathbf{D}\mathbf{L}^\mathsf{T}$ factorization, and either simplicial or supernodal. You will normally not need to examine its contents. See Section 14 for more details.

- 4. cholmod_dense: A dense matrix, either real, complex or zomplex, in column-major order. This differs from the row-major convention used in C. A dense matrix X contains
 - X->x, a double array of size X->nzmax or twice that for the complex case.
 - X->z, a double array of size X->nzmax if X is zomplex.

For a real dense matrix x_{ij} is X->x [i+j*d] where d = X->d is the leading dimension of X. For a complex dense matrix, the real part of x_{ij} is X->x [2*(i+j*d)] and the imaginary part is X->x [2*(i+j*d)+1]. For a zomplex dense matrix, the real part of x_{ij} is X->x [i+j*d] and the imaginary part is X->z [i+j*d]. Real and complex dense matrices can be passed to LAPACK and the BLAS. See Section 15 for more details.

- 5. cholmod_triplet: CHOLMOD's sparse matrix (cholmod_sparse) is the primary input for nearly all CHOLMOD routines, but it can be difficult for the user to construct. A simpler method of creating a sparse matrix is to first create a cholmod_triplet matrix, and then convert it to a cholmod_sparse matrix via the cholmod_triplet_to_sparse routine. In its basic form, the triplet matrix T contains
 - T->i and T->j, integer arrays of size T->nzmax.
 - T->x, a double array of size T->nzmax or twice that for the complex case.
 - T->z, a double array of size T->nzmax if T is zomplex.

The kth entry in the data structure has row index T->i [k] and column index T->j [k]. For a real triplet matrix, its numerical value is T->x [k]. For a complex triplet matrix, its real part is T->x [2*k] and its imaginary part is T->x [2*k+1]. For a zomplex matrix, the real part is T->x [k] and imaginary part is T->z [k]. The entries can be in any order, and duplicates are permitted. See Section 16 for more details.

Each of the five objects has a routine in CHOLMOD to create and destroy it. CHOLMOD provides many other operations on these objects as well. A few of the most important ones are illustrated in the sample program in the next section.

3 Simple example program

```
#include "cholmod.h"
int main (void)
   cholmod_sparse *A ;
   cholmod_dense *x, *b, *r ;
    cholmod_factor *L ;
   double one [2] = \{1,0\}, m1 [2] = \{-1,0\};
                                                   /* basic scalars */
   cholmod_common c ;
   cholmod_start (&c) ;
                                                    /* start CHOLMOD */
                                                    /* read in a matrix */
   A = cholmod_read_sparse (stdin, &c);
                                                    /* print the matrix */
   cholmod_print_sparse (A, "A", &c);
   if (A == NULL || A->stype == 0)
                                                    /* A must be symmetric */
        cholmod_free_sparse (&A, &c) ;
       cholmod_finish (&c) ;
       return (0);
   }
   b = cholmod_ones (A->nrow, 1, A->xtype, &c);
                                                   /* b = ones(n,1) */
                                                    /* analyze */
   L = cholmod_analyze (A, &c) ;
   cholmod_factorize (A, L, &c);
                                                    /* factorize */
   x = cholmod_solve (CHOLMOD_A, L, b, &c);
                                                   /* solve Ax=b */
   r = cholmod_copy_dense (b, &c);
                                                    /* r = b */
#ifndef NMATRIXOPS
   cholmod_sdmult (A, 0, m1, one, x, r, &c);
                                                   /* r = r-Ax */
   printf ("norm(b-Ax) %8.1e\n",
            cholmod_norm_dense (r, 0, &c));
                                                    /* print norm(r) */
   printf ("residual norm not computed (requires CHOLMOD/MatrixOps)\n");
#endif
                                                    /* free matrices */
   cholmod_free_factor (&L, &c) ;
   cholmod_free_sparse (&A, &c) ;
   cholmod_free_dense (&r, &c);
    cholmod_free_dense (&x, &c) ;
    cholmod_free_dense (&b, &c) ;
   cholmod_finish (&c);
                                                    /* finish CHOLMOD */
   return (0);
}
```

Purpose: The Demo/cholmod_simple.c program illustrates the basic usage of CHOLMOD. It reads a triplet matrix from a file (in Matrix Market format), converts it into a sparse matrix, creates a linear system, solves it, and prints the norm of the residual.

See the CHOLMOD/Demo/cholmod_demo.c program for a more elaborate example, and CHOLMOD/Demo/cholmod_l_demo.c for its long integer version.

4 Installation of the C-callable library

CHOLMOD requires a suite of external packages, many of which are distributed along with CHOLMOD, but three of which are not. Those included with CHOLMOD are:

- AMD: an approximate minimum degree ordering algorithm, by Tim Davis, Patrick Amestoy, and Iain Duff [1, 2].
- COLAMD: an approximate column minimum degree ordering algorithm, by Tim Davis, Stefan Larimore, John Gilbert, and Esmond Ng [6, 7].
- CCOLAMD: a constrained approximate column minimum degree ordering algorithm, by Tim Davis and Siva Rajamanickam, based directly on COLAMD. This package is not required if CHOLMOD is compiled with the -DNCAMD flag.
- CAMD: a constrained approximate minimum degree ordering algorithm, by Tim Davis and Yanqing Chen, based directly on AMD. This package is not required if CHOLMOD is compiled with the -DNCAMD flag.
- SuiteSparse_config: a single place where all sparse matrix packages authored or co-authored by Davis are configured.

Three other packages are required for optimal performance:

- METIS 5.1.0: a graph partitioning package by George Karypis, Univ. of Minnesota. Not needed if -DNPARTITION is used. See http://www-users.cs.umn.edu/~karypis/metis.
- BLAS: the Basic Linear Algebra Subprograms. Not needed if -DNSUPERNODAL is used. See http://www.netlib.org for the reference BLAS (not meant for production use). For Kazushige Goto's optimized BLAS (highly recommended for CHOLMOD) see http://www.tacc.utexas.edu/~kgoto/ or http://www.cs.utexas.edu/users/flame/goto/. I recommend that you avoid the Intel MKL BLAS; one recent version returns NaN's, where both the Goto BLAS and the standard Fortran reference BLAS return the correct answer. See CHOLMOD/README for more information.
- LAPACK: the Basic Linear Algebra Subprograms. Not needed if -DNSUPERNODAL is used. See http://www.netlib.org.
- CUDA BLAS: CHOLMOD can exploit an NVIDIA GPU by using the CUDA BLAS for large supernodes. This feature is new to CHOLMOD v2.0.0.

You must first obtain and install LAPACK, and the BLAS. METIS 5.1.0 is optional; a copy of it is in SuiteSparse_metis.

CHOLMOD's specific settings are given by the CHOLMOD_CONFIG string:

- -DNCHECK: do not include the Check module.
- -DNCHOLESKY: do not include the Cholesky module.
- -DNPARTITION: do not include the interface to METIS in the Partition module.

- -DCAMD: do not include the interfaces to CAMD, CCOLAMD, and CSYMAMD in the Partition module.
- -DNMATRIXOPS: do not include the MatrixOps module. Note that the Demo requires the MatrixOps module.
- -DNMODIFY: do not include the Modify module.
- -DNSUPERNODAL: do not include the Supernodal module.
- -DNPRINT: do not print anything.

Type make in the CHOLMOD directory. The AMD, COLAMD, CAMD, CCOLAMD, and CHOLMOD libraries will be compiled. No Fortran compiler is required in this case. A short demo program will be compiled and tested on a few matrices. The residuals should all be small. Compare your output with the CHOLMOD/Demo/make.out file.

CHOLMOD is now ready for use in your own applications. You must link your programs with the libcholmod.*, libamd.*, libcolamd.*, LAPACK, and BLAS libraries. Unless you use -DNPARTITION, you must also link with the METIS 5.1.0 library. Unless -DNCAMD is present at compile time, you must link with CAMD/libcamd.*, and CCOLAMD/libccolamd.*.

The make command now copies all of these libraries and include files into a single place: SuiteSparse/lib and SuiteSparse/include. To tell your your compiler where to find them, use -LSuiteSparse/lib and -ISuiteSparse/include.

To install CHOLMOD in /usr/local/lib and /usr/local/include, do make install. If you do this, youdo not need the -L and -I option when compiling your program. Documentation is also installed in /usr/local/doc. The installation location can be changed at the make command line; see the SuiteSparse/README.txt file for details. To remove CHOLMOD, do make uninstall.

5 Using CHOLMOD in MATLAB

CHOLMOD includes a set of m-files and mexFunctions in the CHOLMOD/MATLAB directory. The following functions are provided:

analyze	order and analyze a matrix
bisect	find a node separator
chol2	same as chol
cholmod2	same as $x=A b$ if A is symmetric positive definite
${\tt cholmod_demo}$	a short demo program
${\tt cholmod_make}$	compiles CHOLMOD for use in MATLAB
etree2	same as etree
${\tt graph_demo}$	graph partitioning demo
lchol	L*L' factorization
ldlchol	L*D*L' factorization
$ldl_normest$	estimate norm(A-L*D*L')
ldlsolve	$x = L' \setminus (D \setminus (L \setminus b))$
ldlsplit	split the output of ldlchol into L and D
ldlupdate	update/downdate an L*D*L' factorization
ldlrowmod	$\operatorname{add}/\operatorname{delete}$ a row from an L*D*L' factorization
metis	interface to METIS_NodeND ordering
mread	read a sparse or dense Matrix Market file
mwrite	write a sparse or dense Matrix Market file
nesdis	CHOLMOD's nested dissection ordering
resymbol	recomputes the symbolic factorization
sdmult	S*F where S is sparse and F is dense
spsym	determine symmetry
sparse2	same as sparse
symbfact2	same as symbfact

Each function is described in the next sections.

5.1 analyze: order and analyze

different:

ANALYZE order and analyze a matrix using CHOLMOD's best-effort ordering. [p count] = analyze (A) orders A, using just tril(A) [p count] = analyze (A,'sym') orders A, using just tril(A) [p count] = analyze (A,'row') orders A*A' [p count] = analyze (A,'col') orders A'*A an optional 3rd parameter modifies the ordering strategy: [p count] = analyze (A,'sym',k) orders A, using just tril(A) [p count] = analyze (A,'row',k) orders A*A' [p count] = analyze (A,'col',k) orders A'*A Returns a permutation and the count of the number of nonzeros in each column of L for the permuted matrix A. That is, count is returned as: count = symbfact2 (A (p,p)) if ordering A count = symbfact2 (A (p,:),'row') if ordering A*A' count = symbfact2 (A (:,p),'col') if ordering A'*A CHOLMOD uses the following ordering strategy: k = 0: Try AMD. If that ordering gives a flop count >= 500 * nnz(L) and a fill-in of nnz(L) >= 5*nnz(C), then try METIS_NodeND (where C = A, A*A', or A'*A is the matrix being ordered. Selects the best ordering tried. This is the default. if k > 0, then multiple orderings are attempted. k = 1 or 2: just try AMD k = 3: also try METIS_NodeND k = 4: also try NESDIS, CHOLMOD's nested dissection (NESDIS), with default parameters. Uses METIS's node bisector and CCOLAMD. k = 5: also try the natural ordering (p = 1:n) k = 6: also try NESDIS with large leaves of the separator tree k = 7: also try NESDIS with tiny leaves and no CCOLAMD ordering k = 8: also try NESDIS with no dense-node removal k = 9: also try COLAMD if ordering A'*A or A*A', (AMD if ordering A). k > 9 is treated as k = 9k = -1: just use AMD k = -2: just use METIS k = -3: just use NESDIS The method returning the smallest nnz(L) is used for p and count. k = 4 takes much longer than (say) k = 0, but it can reduce nnz(L) by a typical 5% to 10%. k = 5 to 9 is getting extreme, but if you have lots of time and want to find the best ordering possible, set k = 9.

If METIS is not installed for use in CHOLMOD, then the strategy is

```
k = 1 to 4: just try AMD
k = 5 to 8: also try the natural ordering (p = 1:n)
k = 9: also try COLAMD if ordering A'*A or A*A', (AMD if ordering A).
k > 9 is treated as k = 9

See also METIS, NESDIS, BISECT, SYMBFACT, AMD
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```

5.2 bisect: find a node separator

BISECT computes a node separator based on METIS_ComputeVertexSeparator.

```
Example:

s = bisect(A) bisects A. Uses tril(A) and assumes A is symmetric.

s = bisect(A,'sym') the same as p=bisect(A).

s = bisect(A,'col') bisects A'*A.

s = bisect(A,'row') bisects A*A'.

A must be square for p=bisect(A) and bisect(A,'sym').

s is a vector of length equal to the dimension of A, A'*A, or A*A', depending on the matrix bisected. s(i)=0 if node i is in the left subgraph, s(i)=1 if it is in the right subgraph, and s(i)=2 if node i is in the node separator.

Requires METIS, authored by George Karypis, Univ. of Minnesota. This MATLAB interface, via CHOLMOD, is by Tim Davis.

See also METIS, NESDIS

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```

5.3 chol2: same as chol

```
CHOL2 sparse Cholesky factorization, A=R'R.

Note that A=L*L' (LCHOL) and A=L*D*L' (LDLCHOL) factorizations are faster than R'*R (CHOL2 and CHOL) and use less memory. The LL' and LDL' factorization methods use tril(A). This method uses triu(A), just like the built-in CHOL.

Example:

R = chol2 (A) same as R = chol (A), just faster

[R,p] = chol2 (A) same as [R,p] = chol(A), just faster

[R,p,q] = chol2 (A) factorizes A(q,q) into R'*R, where q is a fill-reducing ordering
```

A must be sparse.

See also LCHOL, LDLCHOL, CHOL, LDLUPDATE.

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5.4 cholmod2: supernodal backslash

CHOLMOD2 supernodal sparse Cholesky backslash, x = A b

```
x = cholmod2 (A,b)
Computes the LL' factorization of A(p,p), where p is a fill-reducing
ordering, then solves a sparse linear system Ax=b. A must be sparse,
symmetric, and positive definite). Uses only the upper triangular part
of A. A second output, [x,stats]=cholmod2(A,b), returns statistics:
                estimate of the reciprocal of the condition number
    stats(1)
   stats(2)
                ordering used:
                0: natural, 1: given, 2:amd, 3:metis, 4:nesdis,
                5:colamd, 6: natural but postordered.
   stats(3)
               nnz(L)
   stats(4)
                flop count in Cholesky factorization. Excludes solution
                of upper/lower triangular systems, which can be easily
                computed from stats(3) (roughly 4*nnz(L)*size(b,2)).
   stats(5)
                memory usage in MB.
The 3rd argument select the ordering method to use. If not present or -1,
the default ordering strategy is used (AMD, and then try METIS if AMD finds
an ordering with high fill-in, and use the best method tried).
Other options for the ordering parameter:
   0 natural (no etree postordering)
   -1 use CHOLMOD's default ordering strategy (AMD, then try METIS)
   -2 AMD, and then try NESDIS (not METIS) if AMD has high fill-in
   -3 use AMD only
   -4 use METIS only
   -5 use NESDIS only
   -6 natural, but with etree postordering
       user permutation (vector of size n, with a permutation of 1:n)
See also CHOL, MLDIVIDE.
```

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5.5 cholmod_demo: a short demo program

CHOLMOD_DEMO a demo for CHOLMOD

Tests CHOLMOD with various randomly-generated matrices, and the west0479 matrix distributed with MATLAB. Random matrices are not good test cases, but they are easily generated. It also compares CHOLMOD and MATLAB on the sparse matrix problem used in the MATLAB BENCH command.

See CHOLMOD/MATLAB/Test/cholmod_test.m for a lengthy test using matrices from the SuiteSparse Matrix Collection.

Example: cholmod_demo See also BENCH Copyright 2006-2022, Timothy A. Davis, All Rights Reserved. SPDX-License-Identifier: GPL-2.0+ L = full (L); L = full (L);

5.6 cholmod_make: compile CHOLMOD in MATLAB

```
CHOLMOD_MAKE compiles the CHOLMOD mexFunctions
```

Example:
 cholmod_make

CHOLMOD relies on AMD and COLAMD, and optionally CCOLAMD, CAMD, and METIS. You must type the cholmod_make command while in the CHOLMOD/MATLAB directory.

See also analyze, bisect, chol2, cholmod2, etree2, lchol, ldlchol, ldlsolve, ldlupdate, metis, spsym, nesdis, septree, resymbol, sdmult, sparse2, symbfact2, mread, mwrite, ldlrowmod

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MATLAB 8.3.0 now has a -silent option to keep 'mex' from burbling too much Determine if METIS is available

5.7 etree2: same as etree

```
ETREE2 sparse elimination tree.
  Finds the elimination tree of A, A'*A, or A*A', and optionaly postorders
   the tree. parent(j) is the parent of node j in the tree, or 0 if j is a
  root. The symmetric case uses only the upper or lower triangular part of
   A (etree2(A) uses the upper part, and etree2(A,'lo') uses the lower part).
  Example:
  parent = etree2 (A)
                              finds the elimination tree of A, using triu(A)
   parent = etree2 (A,'sym')
                              same as etree2(A)
   parent = etree2 (A,'col')
                              finds the elimination tree of A'*A
  parent = etree2 (A,'row')
                              finds the elimination tree of A*A,
  parent = etree2 (A,'lo')
                              finds the elimination tree of A, using tril(A)
   [parent,post] = etree2 (...) also returns a post-ordering of the tree.
   If you have a fill-reducing permutation p, you can combine it with an
   elimination tree post-ordering using the following code. Post-ordering has
  no effect on fill-in (except for lu), but it does improve the performance
   of the subsequent factorization.
  For the symmetric case, suitable for chol(A(p,p)):
       [parent post] = etree2 (A (p,p));
      p = p (post);
  For the column case, suitable for qr(A(:,p)) or lu(A(:,p)):
       [parent post] = etree2 (A (:,p), 'col') ;
      p = p (post);
  For the row case, suitable for qr(A(p,:)') or chol(A(p,:)*A(p,:)'):
       [parent post] = etree2 (A (p,:), 'row') ;
      p = p (post);
   See also TREELAYOUT, TREEPLOT, ETREEPLOT, ETREE
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```

5.8 graph_demo: graph partitioning demo

```
GRAPH_DEMO graph partitioning demo
graph_demo(n) constructs an set of n-by-n 2D grids, partitions them, and
plots them in one-second intervals. n is optional; it defaults to 60.

Example:
    graph_demo

See also DELSQ, NUMGRID, GPLOT, TREEPLOT
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```

5.9 lchol: LL^{T} factorization

LCHOL sparse A=L*L' factorization.

Note that L*L' (LCHOL) and L*D*L' (LDLCHOL) factorizations are faster than R'*R (CHOL2 and CHOL) and use less memory. The LL' and LDL' factorization methods use tril(A). A must be sparse.

Example:

See also CHOL2, LDLCHOL, CHOL.

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5.10 $1dlchol: LDL^{T}$ factorization

LDLCHOL sparse A=LDL' factorization

Note that L*L' (LCHOL) and L*D*L' (LDLCHOL) factorizations are faster than R'*R (CHOL2 and CHOL) and use less memory. The LL' and LDL' factorization methods use tril(A). A must be sparse.

Example:

The output matrix LD contains both L and D. D is on the diagonal of LD, and L is contained in the strictly lower triangular part of LD. The unit-diagonal of L is not stored. You can obtain the L and D matrices with [L,D] = ldlsplit (LD). LD is in the form needed by ldlupdate.

Explicit zeros may appear in the LD matrix. The pattern of LD matches the pattern of L as computed by symbfact2, even if some entries in LD are explicitly zero. This is to ensure that ldlupdate and ldlsolve work properly. You must NOT modify LD in MATLAB itself and then use ldlupdate or ldlsolve if LD contains explicit zero entries; ldlupdate and ldlsolve will fail catastrophically in this case.

You MAY modify LD in MATLAB if you do not pass it back to ldlupdate or ldlsolve. Just be aware that LD contains explicit zero entries, contrary to the standard practice in MATLAB of removing those entries from all sparse matrices. LD = sparse2 (LD) will remove any zero entries in LD.

See also LDLUPDATE, LDLSOLVE, LDLSPLIT, CHOL2, LCHOL, CHOL, SPARSE2 Copyright 2006-2022, Timothy A. Davis, All Rights Reserved. SPDX-License-Identifier: GPL-2.0+

5.11 Idlsolve: solve using an LDL^{T} factorization

```
LDLSOLVE solve LDL'x=b using a sparse LDL' factorization

Example:
    x = ldlsolve (LD,b)

solves the system L*D*L'*x=b for x. This is equivalent to

[L,D] = ldlsplit (LD);
    x = L' \ (D \ (L \ b));

LD is from ldlchol, or as updated by ldlupdate or ldlrowmod. You must not modify LD as obtained from ldlchol, ldlupdate, or ldlrowmod prior to passing it to this function. See ldlupdate for more details.

See also LDLCHOL, LDLUPDATE, LDLSPLIT, LDLROWMOD
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```

5.12 $ldlsplit: split an LDL^T$ factorization

```
Example:
[L,D] = ldlsplit (LD)

LD contains an LDL' factorization, computed with LD = ldlchol(A), for example. The diagonal of LD contains D, and the entries below the diagonal contain L (which has a unit diagonal). This function splits LD into its two components L and D so that L*D*L' = A.

See also LDLCHOL, LDLSOLVE, LDLUPDATE.
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```

5.13 ldlupdate: update/downdate an LDL^{T} factorization

LDLUPDATE multiple-rank update or downdate of a sparse LDL' factorization.

On input, LD contains the LDL' factorization of A (L*D*L'=A or A(q,q)). The unit-diagonal of L is not stored. In its place is the diagonal matrix D. LD can be computed using the CHOLMOD mexFunctions:

```
LD = ldlchol (A) ;
or
[LD,p,q] = ldlchol (A) ;
```

With this LD, either of the following MATLAB statements,

Example:

```
LD = ldlupdate (LD,C)
LD = ldlupdate (LD,C,'+')
```

return the LDL' factorization of A+C*C' or A(q,q)-C*C' if LD holds the LDL' factorization of A(q,q) on input. For a downdate:

```
LD = ldlupdate (LD,C,'-')
```

returns the LDL' factorization of A-C*C' or A(q,q)-C*C'.

LD and C must be sparse and real. LD must be square, and C must have the same number of rows as LD. You must not modify LD in MATLAB (see the WARNING below).

Note that if C is sparse with few columns, most of the time spent in this routine is taken by copying the input LD to the output LD. If MATLAB allowed mexFunctions to safely modify its inputs, this mexFunction would be much faster, since not all of LD changes.

See also LDLCHOL, LDLSPLIT, LDLSOLVE, CHOLUPDATE

MATLAB drops zero entries from its sparse matrices. LD can contain numerically zero entries that are symbolically present in the sparse matrix data structure. These are essential for ldlupdate and ldlsolve to work properly, since they exploit the graph-theoretic structure of a sparse Cholesky factorization. If you modify LD in MATLAB, those zero entries may get dropped and the required graph property will be destroyed. In this case, ldlupdate and ldlsolve will fail catastrophically (possibly with a segmentation fault, terminating MATLAB). It takes much more time to ensure this property holds than the time it takes to do the update/downdate or the solve, so ldlupdate and ldlsolve simply assume the propertly holds.

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5.14 ldlrowmod: add/delete a row from an LDL^{T} factorization

LDLROWMOD add/delete a row from a sparse LDL' factorization.

On input, LD contains the LDL' factorization of A (L*D*L'=A or A(q,q)). The unit-diagonal of L is not stored. In its place is the diagonal matrix D. LD can be computed using the CHOLMOD mexFunctions:

```
LD = ldlchol (A) ;
or
[LD,p,q] = ldlchol (A) ;
```

With this LD, either of the following MATLAB statements,

Example:

```
LD = ldlrowmod (LD,k,C) add row k to an LDL' factorization
```

returns the LDL' factorization of S, where S = A except for S(:,k) = C and S(k,:) = C. The kth row of A is assumed to initially be equal to the kth row of identity. To delete a row:

```
LD = ldlrowmod (LD,k) delete row k from an LDL' factorization
```

returns the LDL' factorization of S, where S = A except that S(:,k) and S(k,:) become the kth column/row of speye(n), repespectively.

LD and C must be sparse and real. LD must be square, and C must have the same number of rows as LD. You must not modify LD in MATLAB (see the WARNING below).

Note that if C is sparse with few columns, most of the time spent in this routine is taken by copying the input LD to the output LD. If MATLAB allowed mexFunctions to safely modify its inputs, this mexFunction would be much faster, since not all of LD changes.

See also LDLCHOL, LDLSPLIT, LDLSOLVE, CHOLUPDATE, LDLUPDATE

MATLAB drops zero entries from its sparse matrices. LD can contain numerically zero entries that are symbolically present in the sparse matrix data structure. These are essential for ldlrowmod and ldlsolve to work properly, since they exploit the graph-theoretic structure of a sparse Cholesky factorization. If you modify LD in MATLAB, those zero entries may get dropped and the required graph property will be destroyed. In this case, ldlrowmod and ldlsolve will fail catastrophically (possibly with a segmentation fault, terminating MATLAB). It takes much more time to ensure this property holds than the time it takes to do the row add/delete or the solve, so ldlrowmod and ldlsolve simply assume the propertly holds.

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5.15 mread: read a sparse or dense matrix from a Matrix Market file

MREAD read a sparse matrix from a file in Matrix Market format.

Example:

A = mread (filename)
[A Z] = mread (filename, prefer_binary)

Unlike MMREAD, only the matrix is returned; the file format is not returned. Explicit zero entries can be present in the file; these are not included in A. They appear as the nonzero pattern of the binary matrix Z.

If prefer_binary is not present, or zero, a symmetric pattern-only matrix is returned with A(i,i) = 1 + length(find(A(:,i))) if it is present in the pattern, and A(i,j) = -1 for off-diagonal entries. If you want the original Matrix Market matrix in this case, simply use A = mread (filename,1).

Compare with mmread.m at http://math.nist.gov/MatrixMarket

See also load

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5.16 mwrite: write a sparse or dense matrix to a Matrix Market file

MWRITE write a matrix to a file in Matrix Market form.

Example:

mtype = mwrite (filename, A, Z, comments_filename)

A can be sparse or full.

If present and non-empty, A and Z must have the same dimension. Z contains the explicit zero entries in the matrix (which MATLAB drops). The entries of Z appear as explicit zeros in the output file. Z is optional. If it is an empty matrix it is ignored. Z must be sparse or empty, if present. It is ignored if A is full.

filename is the name of the output file. comments_filename is the file whose contents are include after the Matrix Market header and before the first data line. Ignored if an empty string or not present.

See also mread.

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5.17 metis: order with METIS

METIS nested dissection ordering via METIS_NodeND.

5.18 nesdis: order with CHOLMOD nested dissection

NESDIS nested dissection ordering via CHOLMOD's nested dissection.

sparser than chol(A'*A).

A must be square for p=nesdis(A) or nesdis(A,'sym').

With three output arguments, [p cp cmember] = nesdis(...), the separator tree and node-to-component mapping is returned. cmember(i)=c means that node i is in component c, where c is in the range of 1 to the number of components. length(cp) is the number of components found. cp is the separator tree; cp(c) is the parent of component c, or 0 if c is a root. There can be anywhere from 1 to n components, where n is dimension of A, A*A', or A'*A. cmember is a vector of length n.

An optional 3rd input argument, nesdis (A,mode,opts), modifies the default parameters. opts(1) specifies the smallest subgraph that should not be partitioned (default is 200). opts(2) is 0 by default; if nonzero, connected components (formed after the node separator is removed) are partitioned independently. The default value tends to lead to a more balanced separator tree, cp. opts(3) defines when a separator is kept; it is kept if the separator size is < opts(3) times the number of nodes in the graph being cut (valid range is 0 to 1, default is 1).

opts(4) specifies graph is to be ordered after it is dissected. For the 'sym' case: 0: natural ordering, 1: CAMD, 2: CSYMAMD. For other cases: 0: natural ordering, nonzero: CCOLAMD. The default is 1, to use CAMD for the symmetric case and CCOLAMD for the other cases.

If opts is shorter than length 4, defaults are used for entries that are not present.

NESDIS uses METIS' node separator algorithm to recursively partition the graph. This gives a set of constraints (cmember) that is then passed to CCOLAMD, CSYMAMD, or CAMD, constrained minimum degree ordering algorithms. NESDIS typically takes slightly more time than METIS (METIS_NodeND), but tends to produce better orderings.

Requires METIS, authored by George Karypis, Univ. of Minnesota. This MATLAB interface, via CHOLMOD, is by Tim Davis.

```
See also METIS, BISECT, AMD
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```

5.19 resymbol: re-do symbolic factorization

RESYMBOL recomputes the symbolic Cholesky factorization of the matrix A.

Example: L = resymbol (L, A)

Recompute the symbolic Cholesky factorization of the matrix A. A must be symmetric. Only tril(A) is used. Entries in L that are not in the Cholesky factorization of A are removed from L. L can be from an LL' or LDL' factorization (lchol or ldlchol). resymbol is useful after a series of downdates via ldlupdate or ldlrowmod, since downdates do not remove any entries in L. The numerical values of A are ignored; only its nonzero pattern is used.

See also LCHOL, LDLUPDATE, LDLROWMOD Copyright 2006-2022, Timothy A. Davis, All Rights Reserved. SPDX-License-Identifier: GPL-2.0+

5.20 sdmult: sparse matrix times dense matrix

```
SDMULT sparse matrix times dense matrix

Compute C = S*F or S'*F where S is sparse and F is full (C is also sparse).

S and F must both be real or both be complex. This function is substantially faster than the MATLAB expression C=S*F when F has many columns.

Example:

C = sdmult (S,F);
C = S*F
C = sdmult (S,F,0);
C = S*F
C = sdmult (S,F,1);
C = S'*F

See also MTIMES

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```

5.21 spsym: determine symmetry

```
SPSYM determine if a sparse matrix is symmetric, Hermitian, or skew-symmetric.
  If so, also determine if its diagonal has all positive real entries.
   A must be sparse.
  Example:
  result = spsym (A);
  result = spsym (A,quick);
   If quick = 0, or is not present, then this routine returns:
      1: if A is rectangular
      2: if A is unsymmetric
      3: if A is symmetric, but with one or more A(j,j) \le 0
      4: if A is Hermitian, but with one or more A(j,j) \le 0 or with
           nonzero imaginary part
      5: if A is skew symmetric (and thus the diagonal is all zero as well)
      6: if A is symmetric with real positive diagonal
      7: if A is Hermitian with real positive diagonal
   If quick is nonzero, then the function can return more quickly, as soon as
   it finds a diagonal entry that is \leq 0 or with a nonzero imaginary part.
   In this case, it returns 2 for a square matrix, even if the matrix might
   otherwise be symmetric or Hermitian.
   Regardless of the value of "quick", this function returns 6 or 7 if A is
   a candidate for sparse Cholesky.
   For an MATLAB M-file function that computes the same thing as this
   mexFunction (but much slower), see the get_symmetry function by typing
   "type spsym".
   This spsym function does not compute the transpose of A, nor does it need
   to examine the entire matrix if it is unsymmetric. It uses very little
   memory as well (just size-n workspace, where n = size(A,1)).
   Examples:
      load west0479
      A = west0479;
      spsym (A)
      spsym (A+A')
      spsym (A-A')
      spsym (A+A'+3*speye(size(A,1)))
   See also mldivide.
      function result = get_symmetry (A,quick)
      \mbox{\em GET\_SYMMETRY:} does the same thing as the spsym mexFunction.
      % It's just a lot slower and uses much more memory. This function
      % is meant for testing and documentation only.
       [m n] = size (A) ;
      if (m = n)
           result = 1;
                                   % rectangular
           return
      end
```

```
if (nargin < 2)
         quick = 0;
      end
     d = diag(A);
     posdiag = all (real (d) > 0) & all (imag (d) == 0);
     if (quick & ~posdiag)
         result = 2;
                                 % Not a candidate for sparse Cholesky.
     elseif ("isreal (A) & nnz (A-A') == 0)
         if (posdiag)
                                 % complex Hermitian, with positive diagonal
              result = 7;
          else
             result = 4;
                                 % complex Hermitian, nonpositive diagonal
          end
     elseif (nnz (A-A.') == 0)
         if (posdiag)
             result = 6;
                                 % symmetric with positive diagonal
          else
             result = 3;
                                 % symmetric, nonpositive diagonal
         end
     elseif (nnz (A+A.') == 0)
                                 % skew symmetric
         result = 5;
     else
                                 % unsymmetric
         result = 2;
     end
With additional outputs, spsym computes the following for square matrices:
(in this case "quick" is ignored, and set to zero):
[result xmatched pmatched nzoffdiag nnzdiag] = spsym(A)
  xmatched is the number of nonzero entries for which A(i,j) = conj(A(j,i)).
  pmatched is the number of entries (i,j) for which A(i,j) and A(j,i) are
  both in the pattern of A (the value doesn't matter). nzoffdiag is the
  total number of off-diagonal entries in the pattern. nzdiag is the number
  of diagonal entries in the pattern. If the matrix is rectangular,
 xmatched, pmatched, nzoffdiag, and nzdiag are not computed (all of them are
 returned as zero). Note that a matched pair, A(i,j) and A(j,i) for i != j,
  is counted twice (once per entry).
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```

5.22 sparse2: same as sparse

```
SPARSE2 replacement for SPARSE
  Example:
  S = sparse2 (i,j,s,m,n,nzmax)
  Identical to the MATLAB sparse function (just faster).
  An additional feature is added that is not part of the MATLAB sparse
  function, the Z matrix. With an extra output,
   [S Z] = sparse2 (i,j,s,m,n,nzmax)
  the matrix Z is a binary real matrix whose nonzero pattern contains the
   explicit zero entries that were dropped from S. Z only contains entries
  for the sparse2(i,j,s,...) usage. [S Z]=sparse2(X) where X is full always
   returns Z with nnz(Z) = 0, as does [S Z]=sparse2(m,n). More precisely,
  {\bf Z} is the following matrix (where \dots means the optional m, n, and nzmax
  parameters).
      S = sparse (i,j,s, ...)
      Z = spones (sparse (i,j,1, ...)) - spones (S)
  See also sparse.
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```

```
SYMBFACT2 symbolic factorization
   Analyzes the Cholesky factorization of A, A'*A, or A*A'.
   Example:
   count = symbfact2 (A)
                                       returns row counts of R=chol(A)
   count = symbfact2 (A,'col')
                                       returns row counts of R=chol(A'*A)
   count = symbfact2 (A,'sym')
                                       same as symbfact2(A)
   count = symbfact2 (A,'lo')
                                       same as symbfact2(A'), uses tril(A)
   count = symbfact2 (A,'row')
                                       returns row counts of R=chol(A*A')
  The flop count for a subsequent LL' factorization is sum(count.^2)
   [count, h, parent, post, R] = symbfact2 (...) returns:
      h: height of the elimination tree
      parent: the elimination tree itself
      post: postordering of the elimination tree
      R: a 0-1 matrix whose structure is that of chol(A) for the symmetric
           case, chol(A'*A) for the 'col' case, or chol(A*A') for the
           'row' case.
   symbfact2(A) and symbfact2(A,'sym') uses the upper triangular part of A
   (\operatorname{triu}(A)) and assumes the lower triangular part is the transpose of
   the upper triangular part. symbfact2(A,'lo') uses tril(A) instead.
   With one to four output arguments, symbfact2 takes time almost proportional
   to nnz(A)+n where n is the dimension of R, and memory proportional to
   nnz(A). Computing the 5th argument takes more time and memory, both
   O(nnz(L)). Internally, the pattern of L is computed and R=L' is returned.
   The following forms return L = R' instead of R. They are faster and take
   less memory than the forms above. They return the same count, h, parent,
   and post outputs.
   [count, h, parent, post, L] = symbfact2 (A,'col','L')
   [count, h, parent, post, L] = symbfact2 (A,'sym','L')
   [count, h, parent, post, L] = symbfact2 (A,'lo', 'L')
   [count, h, parent, post, L] = symbfact2 (A,'row','L')
   See also CHOL, ETREE, TREELAYOUT, SYMBFACT
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```

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6 Installation for use in MATLAB

6.1 cholmod_make: compiling CHOLMOD in MATLAB

This is the preferred method, since it allows METIS to be reconfigured to use the MATLAB memory-management functions instead of malloc and free; this avoids the issue of METIS terminating MATLAB if it runs out of memory.

Start MATLAB, cd to the CHOLMOD/MATLAB directory, and type cholmod_make in the MATLAB command window. This will compile the MATLAB interfaces for AMD, COLAMD, CAMD, CCOLAMD, METIS, and CHOLMOD.

7 Using CHOLMOD with OpenMP acceleration

CHOLMOD includes OpenMP acceleration for some operations. In CHOLMOD versions prior to v6.0.0, the number of threads to use was controlled by a compile time parameter. This is now replaced with run-time controls.

Common->nthreads_max defaults to omp_get_max_threads(), or 1 if OpenMP is not in use. This value controls the maximum number of threads that CHOLMOD will use. If zero or less, the default is used. The Common->chunk parameter controls how many threads are used when the work to do is low. If w is a count of operations to perform, c = Common->chunk, and $m = \text{Common->nthreads_max}$, then a parallel region will use $\max(1, \min(\lfloor w/c \rfloor, m))$ threads. These parameters can be revised by the user application at run time.

8 Using CHOLMOD with GPU acceleration

Starting with CHOLMOD v2.0.0, it is possible to accelerate the numerical factorization phase of CHOLMOD using NVIDIA GPUs. Due to the large computational capability of the GPUs, enabling this capability can result in significant performance improvements. Similar to CPU processing, the GPU is better able to accelerate the dense math associated with larger supernodes. Hence the GPU will provide more significant performance improvements for larger matrices that have more, larger supernodes.

In CHOLMOD v2.3.0 this GPU capability has been improved to provide a significant increase in performance and the interface has been expanded to make the use of GPUs more flexible. CHOLMOD can take advantage of a single NVIDIA GPU that supports CUDA and has at least 64MB of memory. (But substantially more memory, typically about 3 GB, is recommended for best performance.)

Only the long integer version of CHOLMOD can leverage GPU acceleration.

8.1 Compiling CHOLMOD with GPU support

In order to support GPU processing, CHOLMOD must be compiled with the preprocessor macro ENABLE_CUDA defined. It is enabled by default but can be disabled by setting this to false when using cmake.

8.2 Enabling GPU acceleration in CHOLMOD

Even if compiled with GPU support, in CHOLMOD v.2.3.0, GPU processing is not enabled by default and must be specifically requested. There are two ways to do this, either in the code calling CHOLMOD or using environment variables.

The code author can specify the use of GPU processing with the Common->useGPU variable. If this is set to 1, CHOLMOD will attempt to use the GPU. If this is set to 0 the use of the GPU will be prohibited. If this is set to -1, which is the default case, then the environment variables (following paragraph) will be queried to determine if the GPU is to be used. Note that the default value of -1 is set when cholmod_start(Common) is called, so the code author must set Common->useGPU after calling cholmod_start.

Alternatively, or if it is not possible to modify the code calling CHOLMOD, GPU processing can invoked using the CHOLMOD_USE_GPU environment variable. This makes it possible for any CHOLMOD user to invoke GPU processing even if the author of the calling program did not consider this. The interpretation of the environment variable CHOLMOD_USE_GPU is that if the string evaluates to an integer other than zero, GPU processing will be enabled. Note that the setting of Common->useGPU takes precedence and the environment variable CHOLMOD_USE_GPU will only be queried if Common->useGPU = -1.

Note that in either case, if GPU processing is requested, but there is no GPU present, CHOLMOD will continue using the CPU only. Consequently it is always safe to request GPU processing.

8.3 Adjustable parameters

There are a number of parameters that have been added to CHOLMOD to control GPU processing. All of these have appropriate defaults such that GPU processing can be used without any modification. However, for any particular combination of CPU/GPU, better performance might be obtained by adjusting these parameters.

From t_cholmod_gpu.c

CHOLMOD_ND_ROW_LIMIT : Minimum number of rows required in a descendant supernode to be eligible for GPU processing during supernode assembly

CHOLMOD_ND_COL_LIMIT : Minimum number of columns in a descendant supernode to be eligible for GPU processing during supernode assembly

CHOLMOD_POTRF_LIMIT : Minimum number of columns in a supernode to be eligible for POTRF and TRSM processing on the GPU

CHOLMOD_GPU_SKIP: Number of small descendant supernodes to assembled on the CPU before querying if the GPU is needs more descendant supernodes queued

From cholmod_core.h

CHOLMOD_HOST_SUPERNODE_BUFFERS: Number of buffers in which to queue descendant supernodes for GPU processing

Programatically

Common->maxGpuMemBytes: Specifies the maximum amount of memory, in bytes, that CHOLMOD can allocate on the GPU. If this parameter is not set, CHOLMOD will allocate as much GPU memory as possible. Hence, the purpose of this parameter is to restrict CHOLMOD's GPU memory use so that CHOLMOD can be used simultaneously with other codes that also use GPU acceleration and require some amount of GPU memory. If the specified amount of GPU memory is not allocatable, CHOLMOD will allocate the available memory and continue.

Common->maxGpuMemFraction: Entirely similar to Common->maxGpuMemBytes but with the memory specified as a fraction of total GPU memory. Note that if both maxGpuMemBytes and maxGpuMemFraction are specified, whichever results in the minimum amount of memory will be used.

Environment variables

CHOLMOD_GPU_MEM_BYTES: Environment variable with a meaning equivalent to Common->maxGpuMemBytes. This will only be queried if Common->useGPU = -1.

CHOLMOD_GPU_MEM_FRACTION: Environment variable with a meaning equivalent to Common->maxGpuMemFracti This will only be queried if Common->useGPU = -1.

9 Integer and floating-point types, and notation used

CHOLMOD supports both int and long integers. CHOLMOD routines with the prefix cholmod_use int integers, cholmod_l_routines use long. All floating-point values are double.

The long integer is redefinable, via SuiteSparse_config.h. That file defines a C preprocessor token SuiteSparse_long which is long on all systems except for Windows-64, in which case it is defined as __int64. The intent is that with suitable compile-time switches, int is a 32-bit integer and SuiteSparse_long is a 64-bit integer. The term long is used to describe the latter integer throughout this document (except in the prototypes).

Two kinds of complex matrices are supported: complex and zomplex. A complex matrix is held in a manner that is compatible with the Fortran and ANSI C99 complex data type. A complex array of size n is a double array x of size 2*n, with the real and imaginary parts interleaved (the real part comes first, as a double, followed the imaginary part, also as a double. Thus, the real part of the kth entry is x[2*k] and the imaginary part is x[2*k+1].

A zomplex matrix of size n stores its real part in one double array of size n called x and its imaginary part in another double array of size n called z (thus the name "zomplex"). This also how MATLAB stores its complex matrices. The real part of the kth entry is x[k] and the imaginary part is z[k].

Unlike UMFPACK, the same routine name in CHOLMOD is used for pattern-only, real, complex, and zomplex matrices. For example, the statement

```
C = cholmod_copy_sparse (A, &Common) ;
```

creates a copy of a pattern, real, complex, or zomplex sparse matrix A. The xtype (pattern, real, complex, or zomplex) of the resulting sparse matrix C is the same as A (a pattern-only sparse matrix contains no floating-point values). In the above case, C and A use int integers. For long integers, the statement would become:

```
C = cholmod_l_copy_sparse (A, &Common) ;
```

The last parameter of all CHOLMOD routines is always &Common, a pointer to the cholmod_common object, which contains parameters, statistics, and workspace used throughout CHOLMOD.

The xtype of a CHOLMOD object (sparse matrix, triplet matrix, dense matrix, or factorization) determines whether it is pattern-only, real, complex, or zomplex.

The names of the int versions are primarily used in this document. To obtain the name of the long version of the same routine, simply replace cholmod_with cholmod_l_.

MATLAB matrix notation is used throughout this document and in the comments in the CHOLMOD code itself. If you are not familiar with MATLAB, here is a short introduction to the notation, and a few minor variations used in CHOLMOD:

- C=A+B and C=A*B, respectively are a matrix add and multiply if both A and B are matrices of appropriate size. If A is a scalar, then it is added to or multiplied with every entry in B.
- a:b where a and b are integers refers to the sequence a, a+1, ... b.
- [A B] and [A,B] are the horizontal concatenation of A and B.
- [A;B] is the vertical concatenation of A and B.

• A(i, j) can refer either to a scalar or a submatrix. For example:

```
A(1,1) a scalar.

A(:,j) column j of A.

A(i,:) row i of A.

A([1 2], [1 2]) a 2-by-2 matrix containing the 2-by-2 leading minor of A.
```

If p is a permutation of 1:n, and A is n-by-n, then A(p,p) corresponds to the permuted matrix PAP^{T} .

- tril(A) is the lower triangular part of A, including the diagonal.
- tril(A,k) is the lower triangular part of A, including entries on and below the kth diagonal.
- triu(A) is the upper triangular part of A, including the diagonal.
- triu(A,k) is the upper triangular part of A, including entries on and above the kth diagonal.
- size(A) returns the dimensions of A.
- find(x) if x is a vector returns a list of indices i for which x(i) is nonzero.
- A' is the transpose of A if A is real, or the complex conjugate transpose if A is complex.
- A.' is the array transpose of A.
- diag(A) is the diagonal of A if A is a matrix.
- C=diag(s) is a diagonal matrix if s is a vector, with the values of s on the diagonal of C.
- S=spones(A) returns a binary matrix S with the same nonzero pattern of A.
- nnz(A) is the number of nonzero entries in A.

Variations to MATLAB notation used in this document:

- CHOLMOD uses 0-based notation (the first entry in the matrix is A(0,0)). MATLAB is 1-based. The context is usually clear.
- I is the identity matrix.
- A(:,f), where f is a set of columns, is interpreted differently in CHOLMOD, but just for the set named f. See cholmod_transpose_unsym for details.

10 The CHOLMOD Modules, objects, and functions

CHOLMOD contains a total of 133 int-based routines (and the same number of long routines), divided into a set of inter-related Modules. Each Module contains a set of related functions. The functions are divided into two types: Primary and Secondary, to reflect how a user will typically use CHOLMOD. Most users will find the Primary routines to be sufficient to use CHOLMOD in their programs. Each Module exists as a sub-directory (a folder for Windows users) within the CHOLMOD directory (or folder).

There are seven Modules that provide user-callable routines for CHOLMOD.

- 1. Core: basic data structures and definitions
- 2. Check: prints/checks each of CHOLMOD's objects
- 3. Cholesky: sparse Cholesky factorization
- 4. Modify: sparse Cholesky update/downdate and row-add/row-delete
- 5. MatrixOps: sparse matrix operators (add, multiply, norm, scale)
- 6. Supernodal: supernodal sparse Cholesky factorization
- 7. Partition: graph-partitioning-based orderings, which uses a slightly modified copy of METIS 5.1.0 in the SuiteSparse_metis folder.

Additional directories provide support functions and documentation:

- 1. Include: include files for CHOLMOD and programs that use CHOLMOD
- 2. Demo: simple programs that illustrate the use of CHOLMOD
- 3. Doc: documentation (including this document)
- 4. MATLAB: CHOLMOD's interface to MATLAB
- 5. Tcov: an exhaustive test coverage (requires Linux or Solaris)
- 6. Valgrind: runs the Tcov test under valgrind (requires Linux)
- 7. cmake_modules: how other packages can find CHOLMOD when using cmake.
- 8. Config: a folder containing the input files to create the cholmod.h include file, via cmake.

10.1 Core Module: basic data structures and definitions

CHOLMOD includes five basic objects, defined in the Core Module. The Core Module provides basic operations for these objects and is required by all six other CHOLMOD library Modules:

10.1.1 cholmod_common: parameters, statistics, and workspace

You must call cholmod_start before calling any other CHOLMOD routine, and you must call cholmod_finish as your last call to CHOLMOD (with the exception of cholmod_print_common and cholmod_check_common in the Check Module). Once the cholmod_common object is initialized, the user may modify CHOLMOD's parameters held in this object, and obtain statistics on CHOLMOD's activity.

Primary routines for the cholmod_common object:

- cholmod_start: the first call to CHOLMOD.
- cholmod_finish: the last call to CHOLMOD (frees workspace in the cholmod_common object).

Secondary routines for the cholmod_common object:

- cholmod_defaults: restores default parameters
- cholmod_maxrank: determine maximum rank for update/downdate.
- cholmod_allocate_work: allocate workspace.
- cholmod_free_work: free workspace.
- cholmod_clear_flag: clear Flag array.
- cholmod_error: called when CHOLMOD encounters and error.
- \bullet cholmod_dbound: bounds the diagonal of L or D.
- cholmod_hypot: compute sqrt(x*x+y*y) accurately.
- cholmod_divcomplex: complex divide.

10.1.2 cholmod_sparse: a sparse matrix in compressed column form

A sparse matrix A is held in compressed column form. In the basic type ("packed," which corresponds to how MATLAB stores its sparse matrices), and nrow-by-ncol matrix with nzmax entries is held in three arrays: p of size ncol+1, i of size nzmax, and x of size nzmax. Row indices of nonzero entries in column j are held in i [p[j] ... p[j+1]-1], and their corresponding numerical values are held in x [p[j] ... p[j+1]-1]. The first column starts at location zero (p[0]=0). There may be no duplicate entries. Row indices in each column may be sorted or unsorted (the A->sorted flag must be false if the columns are unsorted). The A->stype determines the storage mode: 0 if the matrix is unsymmetric, 1 if the matrix is symmetric with just the upper triangular part stored, and -1 if the matrix is symmetric with just the lower triangular part stored.

In "unpacked" form, an additional array nz of size ncol is used. The end of column j in i and x is given by p[j]+nz[j]. Columns not need be in any particular order (p[0] need not be zero), and there may be gaps between the columns.

Primary routines for the cholmod_sparse object:

- cholmod_allocate_sparse: allocate a sparse matrix
- cholmod_free_sparse: free a sparse matrix

Secondary routines for the cholmod_sparse object:

- cholmod_reallocate_sparse: change the size (number of entries) of a sparse matrix.
- cholmod_nnz: number of nonzeros in a sparse matrix.
- cholmod_speye: sparse identity matrix.
- cholmod_spzeros: sparse zero matrix.
- cholmod_transpose: transpose a sparse matrix.
- cholmod_ptranspose: transpose/permute a sparse matrix.
- cholmod_transpose_unsym: transpose/permute an unsymmetric sparse matrix.
- cholmod_transpose_sym: transpose/permute a symmetric sparse matrix.
- cholmod_sort: sort row indices in each column of a sparse matrix.
- cholmod_band: extract a band of a sparse matrix.
- cholmod_band_inplace: remove entries not with a band.
- cholmod_aat: C = A*A'.
- cholmod_copy_sparse: C = A, create an exact copy of a sparse matrix.
- cholmod_copy: C = A, with possible change of stype.
- cholmod_add: C = alpha*A + beta*B.
- cholmod_sparse_xtype: change the xtype of a sparse matrix.

10.1.3 cholmod_factor: a symbolic or numeric factorization

A factor can be in **LL**^T or **LDL**^T form, and either supernodal or simplicial form. In simplicial form, this is very much like a packed or unpacked **cholmod_sparse** matrix. In supernodal form, adjacent columns with similar nonzero pattern are stored as a single block (a supernode).

Primary routine for the cholmod_factor object:

• cholmod_free_factor: free a factor

Secondary routines for the cholmod_factor object:

- cholmod_allocate_factor: allocate a factor. You will normally use cholmod_analyze to create a factor.
- cholmod_reallocate_factor: change the number of entries in a factor.
- cholmod_change_factor: change the type of a factor (LDL^T to LL^T, supernodal to simplicial, etc.).
- cholmod_pack_factor: pack the columns of a factor.
- cholmod_reallocate_column: resize a single column of a factor.
- cholmod_factor_to_sparse: create a sparse matrix copy of a factor.
- cholmod_copy_factor: create a copy of a factor.
- cholmod_factor_xtype: change the xtype of a factor.

10.1.4 cholmod_dense: a dense matrix

This consists of a dense array of numerical values and its dimensions.

Primary routines for the cholmod_dense object:

- cholmod_allocate_dense: allocate a dense matrix.
- cholmod_free_dense: free a dense matrix.

Secondary routines for the cholmod_dense object:

- cholmod_zeros: allocate a dense matrix of all zeros.
- cholmod_ones: allocate a dense matrix of all ones.
- cholmod_eye: allocate a dense identity matrix.
- cholmod_sparse_to_dense: create a dense matrix copy of a sparse matrix.
- cholmod_dense_to_sparse: create a sparse matrix copy of a dense matrix.
- cholmod_copy_dense: create a copy of a dense matrix.
- cholmod_copy_dense2: copy a dense matrix (pre-allocated).
- cholmod_dense_xtype: change the xtype of a dense matrix.

10.1.5 cholmod_triplet: a sparse matrix in "triplet" form

The cholmod_sparse matrix is the basic sparse matrix used in CHOLMOD, but it can be difficult for the user to construct. It also does not easily support the inclusion of new entries in the matrix. The cholmod_triplet matrix is provided to address these issues. A sparse matrix in triplet form consists of three arrays of size nzmax: i, j, and x, and a z array for the zomplex case.

Primary routines for the cholmod_triplet object:

- cholmod_allocate_triplet: allocate a triplet matrix.
- cholmod_free_triplet: free a triplet matrix.
- cholmod_triplet_to_sparse: create a sparse matrix copy of a triplet matrix.

Secondary routines for the cholmod_triplet object:

- cholmod_reallocate_triplet: change the number of entries in a triplet matrix.
- cholmod_sparse_to_triplet: create a triplet matrix copy of a sparse matrix.
- cholmod_copy_triplet: create a copy of a triplet matrix.
- cholmod_triplet_xtype: change the xtype of a triplet matrix.

10.1.6 Memory management routines

By default, CHOLMOD uses the ANSI C malloc, free, calloc, and realloc routines. You may use different routines by modifying function pointers in the cholmod_common object.

Primary routines:

- cholmod_malloc: malloc wrapper.
- cholmod_free: free wrapper.

Secondary routines:

- cholmod_calloc: calloc wrapper.
- cholmod_realloc: realloc wrapper.
- cholmod_realloc_multiple: realloc wrapper for multiple objects.

10.1.7 cholmod_version: Version control

The cholmod_version function returns the current version of CHOLMOD.

10.2 Check Module: print/check the CHOLMOD objects

The Check Module contains routines that check and print the five basic objects in CHOLMOD, and three kinds of integer vectors (a set, a permutation, and a tree). It also provides a routine to read a sparse matrix from a file in Matrix Market format (http://www.nist.gov/MatrixMarket). Requires the Core Module.

Primary routines:

- cholmod_print_common: print the cholmod_common object, including statistics on CHOLMOD's behavior (fill-in, flop count, ordering methods used, and so on).
- cholmod_write_sparse: write a sparse matrix to a file in Matrix Market format.
- cholmod_write_dense: write a sparse matrix to a file in Matrix Market format.
- cholmod_read_matrix: read a sparse or dense matrix from a file in Matrix Market format.

Secondary routines:

- cholmod_check_common: check the cholmod_common object
- cholmod_check_sparse: check a sparse matrix
- cholmod_print_sparse: print a sparse matrix
- cholmod_check_dense: check a dense matrix
- cholmod_print_dense: print a dense matrix
- cholmod_check_factor: check a Cholesky factorization
- cholmod_print_factor: print a Cholesky factorization
- cholmod_check_triplet: check a triplet matrix
- cholmod_print_triplet: print a triplet matrix
- cholmod_check_subset: check a subset (integer vector in given range)
- cholmod_print_subset: print a subset (integer vector in given range)
- cholmod_check_perm: check a permutation (an integer vector)
- cholmod_print_perm: print a permutation (an integer vector)
- cholmod_check_parent: check an elimination tree (an integer vector)
- cholmod_print_parent: print an elimination tree (an integer vector)
- cholmod_read_triplet: read a triplet matrix from a file
- cholmod_read_sparse: read a sparse matrix from a file
- cholmod_read_dense: read a dense matrix from a file

10.3 Cholesky Module: sparse Cholesky factorization

The primary routines are all that a user requires to order, analyze, and factorize a sparse symmetric positive definite matrix \mathbf{A} (or $\mathbf{A}\mathbf{A}^\mathsf{T}$), and to solve $\mathbf{A}\mathbf{x} = \mathbf{b}$ (or $\mathbf{A}\mathbf{A}^\mathsf{T}\mathbf{x} = \mathbf{b}$). The primary routines rely on the secondary routines, the Core Module, and the AMD and COLAMD packages. They make optional use of the Supernodal and Partition Modules, the METIS package, the CAMD package, and the CCOLAMD package. The Cholesky Module is required by the Partition Module.

Primary routines:

- cholmod_analyze: order and analyze (simplicial or supernodal).
- cholmod_factorize: simplicial or supernodal Cholesky factorization.
- cholmod_solve: solve a linear system (simplicial or supernodal, dense x and b).
- \bullet cholmod_spsolve: solve a linear system (simplicial or supernodal, sparse \mathbf{x} and \mathbf{b}).

Secondary routines:

- cholmod_analyze_p: analyze, with user-provided permutation or f set.
- cholmod_factorize_p: factorize, with user-provided permutation or f.
- cholmod_analyze_ordering: analyze a permutation
- cholmod_solve2: solve a linear system, reusing workspace.
- cholmod_etree: find the elimination tree.
- cholmod_rowcolcounts: compute the row/column counts of L.
- cholmod_amd: order using AMD.
- cholmod_colamd: order using COLAMD.
- cholmod_rowfac: incremental simplicial factorization.
- cholmod_row_subtree: find the nonzero pattern of a row of L.
- cholmod_row_lsubtree: find the nonzero pattern of a row of L.
- cholmod_row_lsubtree: find the nonzero pattern of $\mathbf{L}^{-1}b$.
- cholmod_resymbol: recompute the symbolic pattern of L.
- cholmod_resymbol_noperm: recompute the symbolic pattern of L, no permutation.
- cholmod_postorder: postorder a tree.
- cholmod_rcond: compute the reciprocal condition number estimate.
- cholmod_rowfac_mask: for use in LPDASA only.

10.4 Modify Module: update/downdate a sparse Cholesky factorization

The Modify Module contains sparse Cholesky modification routines: update, downdate, row-add, and row-delete. It can also modify a corresponding solution to $\mathbf{L}\mathbf{x} = \mathbf{b}$ when L is modified. This module is most useful when applied on a Cholesky factorization computed by the Cholesky module, but it does not actually require the Cholesky module. The Core module can create an identity Cholesky factorization ($\mathbf{L}\mathbf{D}\mathbf{L}^\mathsf{T}$ where $\mathbf{L} = \mathbf{D} = \mathbf{I}$) that can then be modified by these routines. Requires the Core module. Not required by any other CHOLMOD Module.

Primary routine:

• cholmod_updown: multiple rank update/downdate

Secondary routines:

- cholmod_updown_solve: update/downdate, and modify solution to $\mathbf{L}\mathbf{x} = \mathbf{b}$
- $cholmod_updown_mark$: update/downdate, and modify solution to partial $\mathbf{L}\mathbf{x} = \mathbf{b}$
- cholmod_updown_mask: for use in LPDASA only.
- \bullet cholmod_rowadd: add a row to an \mathbf{LDL}^T factorization
- ullet cholmod_rowadd_solve: add a row, and update solution to $\mathbf{L}\mathbf{x} = \mathbf{b}$
- cholmod_rowadd_mark: add a row, and update solution to partial Lx = b
- \bullet cholmod_rowdel: delete a row from an \mathbf{LDL}^T factorization
- \bullet cholmod_rowdel_solve: delete a row, and downdate $\mathbf{L}\mathbf{x} = \mathbf{b}$
- cholmod_rowdel_mark: delete a row, and downdate solution to partial $\mathbf{L}\mathbf{x} = \mathbf{b}$

10.5 MatrixOps Module: basic sparse matrix operations

The MatrixOps Module provides basic operations on sparse and dense matrices. Requires the Core module. Not required by any other CHOLMOD module. In the descriptions below, A, B, and C: are sparse matrices (cholmod_sparse), X and Y are dense matrices (cholmod_dense), s is a scalar or vector, and alpha beta are scalars.

- cholmod_drop: drop entries from A with absolute value \geq a given tolerance.
- cholmod_norm_dense: s = norm (X), 1-norm, infinity-norm, or 2-norm
- cholmod_norm_sparse: s = norm (A), 1-norm or infinity-norm
- cholmod_horzcat: C = [A,B]
- cholmod_scale: A = diag(s)*A, A*diag(s), s*A or diag(s)*A*diag(s).
- cholmod_sdmult: Y = alpha*(A*X) + beta*Y or alpha*(A'*X) + beta*Y.
- cholmod_ssmult: C = A*B

- cholmod_submatrix: C = A (i,j), where i and j are arbitrary integer vectors.
- cholmod_vertcat: C = [A ; B].
- cholmod_symmetry: determine symmetry of a matrix.

10.6 Supernodal Module: supernodal sparse Cholesky factorization

The Supernodal Module performs supernodal analysis, factorization, and solve. The simplest way to use these routines is via the Cholesky Module. This Module does not provide any fill-reducing orderings. It normally operates on matrices ordered by the Cholesky Module. It does not require the Cholesky Module itself, however. Requires the Core Module, and two external packages: LAPACK and the BLAS. Optionally used by the Cholesky Module. All are secondary routines since these functions are more easily used via the Cholesky Module.

Secondary routines:

- cholmod_super_symbolic: supernodal symbolic analysis
- cholmod_super_numeric: supernodal numeric factorization
- \bullet cholmod_super_lsolve: supernodal Lx = b solve
- \bullet cholmod_super_ltsolve: supernodal $\mathbf{L}^\mathsf{T} \mathbf{x} = \mathbf{b}$ solve

10.7 Partition Module: graph-partitioning-based orderings

The Partition Module provides graph partitioning and graph-partition-based orderings. It includes an interface to CAMD, CCOLAMD, and CSYMAMD, constrained minimum degree ordering methods which order a matrix following constraints determined via nested dissection. Requires the Core and Cholesky Modules, and two packages: METIS 5.1.0, CAMD, and CCOLAMD. Optionally used by the Cholesky Module. All are secondary routines since these are more easily used by the Cholesky Module.

Note that METIS does not have a version that uses long integers. If you try to use these routines (except the CAMD, CCOLAMD, and CSYMAMD interfaces) on a matrix that is too large, an error code will be returned.

Secondary routines:

- cholmod_nested_dissection: CHOLMOD nested dissection ordering
- cholmod_metis: METIS nested dissection ordering (METIS_NodeND)
- cholmod_camd: interface to CAMD ordering
- cholmod_ccolamd: interface to CCOLAMD ordering
- cholmod_csymamd: interface to CSYMAMD ordering
- cholmod_bisect: graph partitioner (currently based on METIS)
- cholmod_metis_bisector: direct interface to METIS_NodeComputeSeparator.
- cholmod_collapse_septree: pruned a separator tree from cholmod_nested_dissection.

11 CHOLMOD naming convention, parameters, and return values

All routine names, data types, and CHOLMOD library files use the cholmod_prefix. All macros and other #define statements visible to the user program use the CHOLMOD prefix. The cholmod.h file must be included in user programs that use CHOLMOD:

#include "cholmod.h"

All CHOLMOD routines (in all modules) use the following protocol for return values:

- int: TRUE (1) if successful, or FALSE (0) otherwise. (exception: cholmod_divcomplex).
- long: a value > 0 if successful, or -1 otherwise.
- double: a value ≥ 0 if successful, or -1 otherwise.
- size_t: a value > 0 if successful, or 0 otherwise.
- void *: a non-NULL pointer to newly allocated memory if successful, or NULL otherwise.
- cholmod_sparse *: a non-NULL pointer to a newly allocated sparse matrix if successful, or NULL otherwise.
- cholmod_factor *: a non-NULL pointer to a newly allocated factor if successful, or NULL otherwise.
- cholmod_triplet *: a non-NULL pointer to a newly allocated triplet matrix if successful, or NULL otherwise.
- cholmod_dense *: a non-NULL pointer to a newly allocated dense matrix if successful, or NULL otherwise.

TRUE and FALSE are not defined in cholmod.h, since they may conflict with the user program. A routine that described here returning TRUE or FALSE returns 1 or 0, respectively. Any TRUE/FALSE parameter is true if nonzero, false if zero.

Input, output, and input/output parameters:

- Input parameters appear first in the parameter lists of all CHOLMOD routines. They are not modified by CHOLMOD.
- Input/output parameters (except for Common) appear next. They must be defined on input, and are modified on output.
- Output parameters are listed next. If they are pointers, they must point to allocated space on input, but their contents are not defined on input.
- Workspace parameters appear next. They are used in only two routines in the Supernodal module.

• The cholmod_common *Common parameter always appears as the last parameter (with two exceptions: cholmod_hypot and cholmod_divcomplex). It is always an input/output parameter.

A floating-point scalar is passed to CHOLMOD as a pointer to a double array of size two. The first entry in this array is the real part of the scalar, and the second entry is the imaginary part. The imaginary part is only accessed if the other inputs are complex or zomplex. In some cases the imaginary part is always ignored (cholmod_factor_p, for example).

12 Core Module: cholmod_common object

12.1 Constant definitions

```
/* itype defines the types of integer used: */
#define CHOLMOD_INT 0
                            /* all integer arrays are int32_t */
#define CHOLMOD_INTLONG 1
                              /* most are int32_t, some are int64_t */
#define CHOLMOD_LONG 2
                              /* all integer arrays are int64_t */
/* The itype of all parameters for all CHOLMOD routines must match.
* FUTURE WORK: CHOLMOD_INTLONG is not yet supported.
/* dtype defines what the numerical type is (double or float): */
#define CHOLMOD_SINGLE 1
                              /* all numerical values are float */
/* The dtype of all parameters for all CHOLMOD routines must match.
* Scalar floating-point values are always passed as double arrays of size 2
* (for the real and imaginary parts). They are typecast to float as needed.
* FUTURE WORK: the float case is not supported yet.
*/
/* xtype defines the kind of numerical values used: */
#define CHOLMOD_PATTERN 0
                              /* pattern only, no numerical values */
#define CHOLMOD_REAL 1
                              /* a real matrix */
                             /* a complex matrix (ANSI C99 compatible) */
#define CHOLMOD_COMPLEX 2
#define CHOLMOD_ZOMPLEX 3
                              /* a complex matrix (MATLAB compatible) */
/* The xtype of all parameters for all CHOLMOD routines must match.
* CHOLMOD_PATTERN: x and z are ignored.
* CHOLMOD_DOUBLE: x is non-null of size nzmax, z is ignored.
* CHOLMOD_COMPLEX: x is non-null of size 2*nzmax doubles, z is ignored.
* CHOLMOD_ZOMPLEX: x and z are non-null of size nzmax
* In the real case, z is ignored. The kth entry in the matrix is x [k].
* There are two methods for the complex case. In the ANSI C99-compatible
* CHOLMOD_COMPLEX case, the real and imaginary parts of the kth entry
* are in x [2*k] and x [2*k+1], respectively. z is ignored. In the
* MATLAB-compatible CHOLMOD_ZOMPLEX case, the real and imaginary
* parts of the kth entry are in x [k] and z [k].
* Scalar floating-point values are always passed as double arrays of size 2
* (real and imaginary parts). The imaginary part of a scalar is ignored if
* the routine operates on a real matrix.
* These Modules support complex and zomplex matrices, with a few exceptions:
       Check
                   all routines
       Cholesky
                   all routines
                   all except cholmod_aat, add, band, copy
       Core
       Demo
                   all routines
       Partition all routines
```

```
Supernodal all routines support any real, complex, or zomplex input.
                        There will never be a supernodal zomplex L; a complex
                        supernodal L is created if A is zomplex.
       Tcov
                    all routines
       Valgrind
                    all routines
  These Modules provide partial support for complex and zomplex matrices:
       MATLAB
                    all routines support real and zomplex only, not complex,
                        with the exception of ldlupdate, which supports
                        real matrices only. This is a minor constraint since
                       MATLAB's matrices are all real or zomplex.
                    only norm_dense, norm_sparse, and sdmult support complex
       MatrixOps
                        and zomplex
* These Modules do not support complex and zomplex matrices at all:
       Modify
                    all routines support real matrices only
/* Definitions for cholmod_common: */
#define CHOLMOD_MAXMETHODS 9
                                /* maximum number of different methods that */
                                /* cholmod_analyze can try. Must be >= 9. */
/* Common->status values. zero means success, negative means a fatal error,
* positive is a warning. */
#define CHOLMOD_OK O
                                       /* success */
#define CHOLMOD_NOT_INSTALLED (-1)
                                       /* failure: method not installed */
#define CHOLMOD_OUT_OF_MEMORY (-2)
                                       /* failure: out of memory */
#define CHOLMOD_TOO_LARGE (-3)
                                       /* failure: integer overflow occured */
#define CHOLMOD_INVALID (-4)
                                       /* failure: invalid input */
#define CHOLMOD_GPU_PROBLEM (-5)
                                       /* failure: GPU fatal error */
                                       /* warning: matrix not pos. def. */
#define CHOLMOD_NOT_POSDEF (1)
                                       /* warning: D for LDL' or diag(L) or */
#define CHOLMOD_DSMALL (2)
                                       /* LL' has tiny absolute value */
/* ordering method (also used for L->ordering) */
#define CHOLMOD_NATURAL 0
                               /* use natural ordering */
#define CHOLMOD_GIVEN 1
                               /* use given permutation */
#define CHOLMOD_AMD 2
                               /* use minimum degree (AMD) */
#define CHOLMOD_METIS 3
                               /* use METIS' nested dissection */
#define CHOLMOD_NESDIS 4
                                /* use CHOLMOD's version of nested dissection:*/
                                /* node bisector applied recursively, followed
                                * by constrained minimum degree (CSYMAMD or
                                 * CCOLAMD) */
                                /* use AMD for A, COLAMD for A*A' */
#define CHOLMOD_COLAMD 5
/* POSTORDERED is not a method, but a result of natural ordering followed by a
* weighted postorder. It is used for L->ordering, not method [ ].ordering. */
#define CHOLMOD_POSTORDERED 6 /* natural ordering, postordered. */
/* supernodal strategy (for Common->supernodal) */
#define CHOLMOD_SIMPLICIAL 0
                               /* always do simplicial */
#define CHOLMOD_AUTO 1
                               /* select simpl/super depending on matrix */
#define CHOLMOD_SUPERNODAL 2
                               /* always do supernodal */
```

Purpose: These definitions are used within the cholmod_common object, called Common both here and throughout the code.

12.2 cholmod_common: parameters, statistics, and workspace

```
typedef struct cholmod_common_struct
             /* parameters for symbolic/numeric factorization and update/downdate */
   /* ----- */
   double dbound ;
                      /* Smallest absolute value of diagonal entries of D
                        * for LDL' factorization and update/downdate/rowadd/
       \boldsymbol{\ast} rowdel, or the diagonal of L for an LL' factorization.
       * Entries in the range 0 to dbound are replaced with dbound.
       * Entries in the range -dbound to 0 are replaced with -dbound. No
       * changes are made to the diagonal if dbound <= 0. Default: zero */
   double grow0 ;
                       /* For a simplicial factorization, L->i and L->x can
                        * grow if necessary. grow0 is the factor by which
       * it grows. For the initial space, L is of size MAX (1,grow0) times
       * the required space. If L runs out of space, the new size of L is
       * MAX(1.2,grow0) times the new required space. If you do not plan on
       * modifying the LDL' factorization in the Modify module, set grow0 to
       * zero (or set grow2 to 0, see below). Default: 1.2 */
   double grow1 ;
                       /* For a simplicial factorization, each column j of L
   size_t grow2 ;
                       * is initialized with space equal to
       * grow1*L->ColCount[j] + grow2. If grow0 < 1, grow1 < 1, or grow2 == 0,
       * then the space allocated is exactly equal to L->ColCount[j]. If the
       * column j runs out of space, it increases to grow1*need + grow2 in
       * size, where need is the total # of nonzeros in that column. If you do
       * not plan on modifying the factorization in the Modify module, set
       * grow2 to zero. Default: grow1 = 1.2, grow2 = 5. */
   size_t maxrank ;
                       /* rank of maximum update/downdate. Valid values:
                        * 2, 4, or 8. A value < 2 is set to 2, and a
       * value > 8 is set to 8. It is then rounded up to the next highest
       * power of 2, if not already a power of 2. Workspace (Xwork, below) of
       \boldsymbol{\ast} size nrow-by-maxrank double's is allocated for the update/downdate.
       * If an update/downdate of rank-k is requested, with k > maxrank,
       * it is done in steps of maxrank. Default: 8, which is fastest.
       * Memory usage can be reduced by setting maxrank to 2 or 4.
       */
   double supernodal_switch; /* supernodal vs simplicial factorization */
                              /* If Common->supernodal <= CHOLMOD_SIMPLICIAL</pre>
   int supernodal ;
                               * (0) then cholmod_analyze performs a
       * simplicial analysis. If >= CHOLMOD_SUPERNODAL (2), then a supernodal
       * analysis is performed. If == CHOLMOD_AUTO (1) and
       * flop/nnz(L) < Common->supernodal_switch, then a simplicial analysis
```

```
* is done. A supernodal analysis done otherwise.
    * Default: CHOLMOD_AUTO. Default supernodal_switch = 40 */
                    /* If TRUE, then ignore the other final_* parameters
int final_asis ;
                     * (except for final_pack).
                     * The factor is left as-is when done. Default: TRUE.*/
                    /* If TRUE, leave a factor in supernodal form when
int final_super ;
                     * supernodal factorization is finished. If FALSE,
                     * then convert to a simplicial factor when done.
                     * Default: TRUE */
                    /* If TRUE, leave factor in LL' form when done.
int final_ll ;
                    * Otherwise, leave in LDL' form. Default: FALSE */
                    /* If TRUE, pack the columns when done. If TRUE, and
int final_pack ;
                     * cholmod_factorize is called with a symbolic L, L is
    * allocated with exactly the space required, using L->ColCount. If you
    * plan on modifying the factorization, set Common->final_pack to FALSE,
    * and each column will be given a little extra slack space for future
    * growth in fill-in due to updates. Default: TRUE */
                      /* If TRUE, ensure columns are monotonic when done.
int final_monotonic ;
                     * Default: TRUE */
int final_resymbol ;/* if cholmod_factorize performed a supernodal
                     * factorization, final_resymbol is true, and
    * final_super is FALSE (convert a simplicial numeric factorization),
    * then numerically zero entries that resulted from relaxed supernodal
    * amalgamation are removed. This does not remove entries that are zero
    * due to exact numeric cancellation, since doing so would break the
    * update/downdate rowadd/rowdel routines. Default: FALSE. */
/* supernodal relaxed amalgamation parameters: */
double zrelax [3];
size_t nrelax [3] ;
    /* Let ns be the total number of columns in two adjacent supernodes.
     * Let z be the fraction of zero entries in the two supernodes if they
     * are merged (z includes zero entries from prior amalgamations). The
     * two supernodes are merged if:
          (ns <= nrelax [0]) || (no new zero entries added) ||
          (ns <= nrelax [1] && z < zrelax [0]) ||
          (ns <= nrelax [2] && z < zrelax [1]) || (z < zrelax [2])
     * Default parameters result in the following rule:
          (ns <= 4) || (no new zero entries added) ||
          (ns <= 16 && z < 0.8) || (ns <= 48 && z < 0.1) || (z < 0.05)
int prefer_zomplex ;
                        /* X = cholmod_solve (sys, L, B, Common) computes
                         * x=A\b or solves a related system. If L and B are
     * both real, then X is real. Otherwise, X is returned as
     * CHOLMOD_COMPLEX if Common->prefer_zomplex is FALSE, or
     * CHOLMOD_ZOMPLEX if Common->prefer_zomplex is TRUE. This parameter
```

```
* is needed because there is no supernodal zomplex L. Suppose the
    * caller wants all complex matrices to be stored in zomplex form
    * (MATLAB, for example). A supernodal L is returned in complex form
    * if A is zomplex. B can be real, and thus X = cholmod_solve (L,B)
    * should return X as zomplex. This cannot be inferred from the input
    * arguments L and B. Default: FALSE, since all data types are
    * supported in CHOLMOD_COMPLEX form and since this is the native type
    * of LAPACK and the BLAS. Note that the MATLAB/cholmod.c mexFunction
    * sets this parameter to TRUE, since MATLAB matrices are in
    * CHOLMOD_ZOMPLEX form.
int prefer_upper ;
                      /* cholmod_analyze and cholmod_factorize work
                       * fastest when a symmetric matrix is stored in
    * upper triangular form when a fill-reducing ordering is used. In
    * MATLAB, this corresponds to how x=A\b works. When the matrix is
    * ordered as-is, they work fastest when a symmetric matrix is in lower
    * triangular form. In MATLAB, R=chol(A) does the opposite. This
    * parameter affects only how cholmod_read returns a symmetric matrix.
    * If TRUE (the default case), a symmetric matrix is always returned in
    * upper-triangular form (A->stype = 1). */
int quick_return_if_not_posdef ;
                                  /* if TRUE, the supernodal numeric
                                   * factorization will return quickly if
   * the matrix is not positive definite. Default: FALSE. */
int prefer_binary ;
                      /* cholmod_read_triplet converts a symmetric
                       * pattern-only matrix into a real matrix. If
   * prefer_binary is FALSE, the diagonal entries are set to 1 + the degree
   * of the row/column, and off-diagonal entries are set to -1 (resulting
   * in a positive definite matrix if the diagonal is zero-free). Most
   * symmetric patterns are the pattern a positive definite matrix. If
   * this parameter is TRUE, then the matrix is returned with a 1 in each
   * entry, instead. Default: FALSE. Added in v1.3. */
/* ------ */
/* printing and error handling options */
/* ----- */
                  /* print level. Default: 3 */
int print;
                  /* if TRUE, print 16 digits. Otherwise print 5 */
int precise;
/* CHOLMOD print_function replaced with SuiteSparse_config.print_func */
int try_catch ;
                  /* if TRUE, then ignore errors; CHOLMOD is in the middle
                   * of a try/catch block. No error message is printed
    * and the Common->error_handler function is not called. */
void (*error_handler) (int status, const char *file,
   int line, const char *message) ;
   /* Common->error_handler is the user's error handling routine. If not
    * NULL, this routine is called if an error occurs in CHOLMOD. status
    * can be CHOLMOD_OK (0), negative for a fatal error, and positive for
    * a warning. file is a string containing the name of the source code
```

```
* file where the error occured, and line is the line number in that
    * file. message is a string describing the error in more detail. */
/* ----- */
/* ordering options */
/* ----- */
/* The cholmod_analyze routine can try many different orderings and select
* the best one. It can also try one ordering method multiple times, with
 * different parameter settings. The default is to use three orderings,
 * the user's permutation (if provided), AMD which is the fastest ordering
 * and generally gives good fill-in, and METIS. CHOLMOD's nested dissection
 * (METIS with a constrained AMD) usually gives a better ordering than METIS
 * alone (by about 5% to 10%) but it takes more time.
* If you know the method that is best for your matrix, set Common->nmethods
 * to 1 and set Common->method [0] to the set of parameters for that method.
 * If you set it to 1 and do not provide a permutation, then only AMD will
 * be called.
 * If METIS is not available, the default # of methods tried is 2 (the user
 * permutation, if any, and AMD).
 * To try other methods, set Common->nmethods to the number of methods you
 * want to try. The suite of default methods and their parameters is
 * described in the cholmod_defaults routine, and summarized here:
       Common->method [i]:
       i = 0: user-provided ordering (cholmod_analyze_p only)
       i = 1: AMD (for both A and A*A')
       i = 2: METIS
       i = 3: CHOLMOD's nested dissection (NESDIS), default parameters
       i = 4: natural
       i = 5: NESDIS with nd_small = 20000
       i = 6: NESDIS with nd_small = 4, no constrained minimum degree
       i = 7: NESDIS with no dense node removal
       i = 8: AMD for A, COLAMD for A*A'
 * You can modify the suite of methods you wish to try by modifying
 * Common.method [...] after calling cholmod_start or cholmod_defaults.
 * For example, to use AMD, followed by a weighted postordering:
       Common->nmethods = 1 ;
       Common->method [0].ordering = CHOLMOD_AMD ;
       Common->postorder = TRUE ;
 * To use the natural ordering (with no postordering):
       Common->nmethods = 1 ;
       Common->method [0].ordering = CHOLMOD_NATURAL ;
       Common->postorder = FALSE ;
 * If you are going to factorize hundreds or more matrices with the same
 * nonzero pattern, you may wish to spend a great deal of time finding a
```

```
* good permutation. In this case, try setting Common->nmethods to 9.
 * The time spent in cholmod_analysis will be very high, but you need to
 * call it only once.
 * cholmod_analyze sets Common->current to a value between 0 and nmethods-1.
 * Each ordering method uses the set of options defined by this parameter.
                    /* The number of ordering methods to try. Default: 0.
int nmethods;
                     * nmethods = 0 is a special case. cholmod_analyze
    * will try the user-provided ordering (if given) and AMD. Let fl and
    * lnz be the flop count and nonzeros in L from AMD's ordering. Let
   * anz be the number of nonzeros in the upper or lower triangular part
   * of the symmetric matrix A. If fl/lnz < 500 or lnz/anz < 5, then this
   * is a good ordering, and METIS is not attempted. Otherwise, METIS is
             The best ordering found is used. If nmethods > 0, the
   * tried.
   * methods used are given in the method[] array, below. The first
    * three methods in the default suite of orderings is (1) use the given
    * permutation (if provided), (2) use AMD, and (3) use METIS. Maximum
    * allowed value is CHOLMOD_MAXMETHODS. */
                    /* The current method being tried. Default: 0. Valid
int current ;
                     * range is 0 to nmethods-1. */
int selected;
                   /* The best method found. */
/* The suite of ordering methods and parameters: */
struct cholmod_method_struct
    /* statistics for this method */
    double lnz ;
                       /* nnz(L) excl. zeros from supernodal amalgamation,
                         * for a "pure" L */
                       /* flop count for a "pure", real simplicial LL'
   double fl ;
                         * factorization, with no extra work due to
        * amalgamation. Subtract n to get the LDL' flop count. Multiply
        * by about 4 if the matrix is complex or zomplex. */
    /* ordering method parameters */
    double prune_dense ;/* dense row/col control for AMD, SYMAMD, CSYMAMD,
                         * and NESDIS (cholmod_nested_dissection). For a
        * symmetric n-by-n matrix, rows/columns with more than
        * MAX (16, prune_dense * sqrt (n)) entries are removed prior to
        * ordering. They appear at the end of the re-ordered matrix.
        * If prune_dense < 0, only completely dense rows/cols are removed.
        * This paramater is also the dense column control for COLAMD and
        * CCOLAMD. For an m-by-n matrix, columns with more than
        * MAX (16, prune_dense * sqrt (MIN (m,n))) entries are removed prior
        * to ordering. They appear at the end of the re-ordered matrix.
        * CHOLMOD factorizes A*A', so it calls COLAMD and CCOLAMD with A',
        * not A. Thus, this parameter affects the dense *row* control for
        * CHOLMOD's matrix, and the dense *column* control for COLAMD and
```

```
* CCOLAMD.
    * Removing dense rows and columns improves the run-time of the
    * ordering methods. It has some impact on ordering quality
    * (usually minimal, sometimes good, sometimes bad).
    * Default: 10. */
double prune_dense2 ;/* dense row control for COLAMD and CCOLAMD.
                   * Rows with more than MAX (16, dense2 * sqrt (n))
    * for an m-by-n matrix are removed prior to ordering. CHOLMOD's
    * matrix is transposed before ordering it with COLAMD or CCOLAMD,
    * so this controls the dense *columns* of CHOLMOD's matrix, and
    * the dense *rows* of COLAMD's or CCOLAMD's matrix.
    * If prune_dense2 < 0, only completely dense rows/cols are removed.
    st Default: -1. Note that this is not the default for COLAMD and
    * CCOLAMD. -1 is best for Cholesky. 10 is best for LU. */
                   /* in NESDIS, when a node separator is computed, it
double nd_oksep ;
                     * discarded if nsep >= nd_oksep*n, where nsep is
    * the number of nodes in the separator, and n is the size of the
    * graph being cut. Valid range is 0 to 1. If 1 or greater, the
    * separator is discarded if it consists of the entire graph.
    * Default: 1 */
double other_1 [4] ; /* future expansion */
                    /* do not partition graphs with fewer nodes than
size_t nd_small ;
                     * nd_small, in NESDIS. Default: 200 (same as
                     * METIS) */
size_t other_2 [4] ; /* future expansion */
                   /* Aggresive absorption in AMD, COLAMD, SYMAMD,
int aggressive;
                    * CCOLAMD, and CSYMAMD. Default: TRUE */
int order_for_lu ; /* CCOLAMD can be optimized to produce an ordering
                    * for LU or Cholesky factorization. CHOLMOD only
    * performs a Cholesky factorization. However, you may wish to use
    * CHOLMOD as an interface for CCOLAMD but use it for your own LU
    * factorization. In this case, order_for_lu should be set to FALSE.
    * When factorizing in CHOLMOD itself, you should *** NEVER *** set
    * this parameter FALSE. Default: TRUE. */
                   /* If TRUE, compress the graph and subgraphs before
int nd_compress ;
                     * partitioning them in NESDIS. Default: TRUE */
                   /* If 1, follow the nested dissection ordering
int nd_camd ;
                     * with a constrained minimum degree ordering that
    * respects the partitioning just found (using CAMD). If 2, use
    * CSYMAMD instead. If you set nd_small very small, you may not need
    \ast this ordering, and can save time by setting it to zero (no
    * constrained minimum degree ordering). Default: 1. */
```

```
int nd_components; /* The nested dissection ordering finds a node
                        * separator that splits the graph into two parts,
        * which may be unconnected. If nd_components is TRUE, each of
        * these connected components is split independently. If FALSE,
        * each part is split as a whole, even if it consists of more than
        * one connected component. Default: FALSE */
   /* fill-reducing ordering to use */
    int ordering;
    size_t other_3 [4]; /* future expansion */
} method [CHOLMOD_MAXMETHODS + 1] ;
                   /* If TRUE, cholmod_analyze follows the ordering with a
int postorder;
                    * weighted postorder of the elimination tree. Improves
    * supernode amalgamation. Does not affect fundamental nnz(L) and
    * flop count. Default: TRUE. */
                       /* Default: FALSE. If FALSE, then the default
int default_nesdis ;
                        * ordering strategy (when Common->nmethods == 0)
    st is to try the given ordering (if present), AMD, and then METIS if AMD
    * reports high fill-in. If Common->default_nesdis is TRUE then NESDIS
    * is used instead in the default strategy. */
/* memory management, complex divide, and hypot function pointers moved */
/* Function pointers moved from here (in CHOLMOD 2.2.0) to
   SuiteSparse_config.[ch]. See CHOLMOD/Include/cholmod_back.h
   for a set of macros that can be #include'd or copied into your
   application to define these function pointers on any version of {\tt CHOLMOD}.
   */
/* METIS workarounds */
/* ----- */
\slash\hspace{-0.05cm} These workarounds were put into place for METIS 4.0.1. They are safe
   to use with METIS 5.1.0, but they might not longer be necessary. 
 \ast/
double metis_memory ;
                       /* This is a parameter for CHOLMOD's interface to
                        * METIS, not a parameter to METIS itself. METIS
    * uses an amount of memory that is difficult to estimate precisely
    * beforehand. If it runs out of memory, it terminates your program.
    * All routines in CHOLMOD except for CHOLMOD's interface to METIS
   * return an error status and safely return to your program if they run
    st out of memory. To mitigate this problem, the CHOLMOD interface
   * can allocate a single block of memory equal in size to an empirical
   * upper bound of METIS's memory usage times the Common->metis_memory
   st parameter, and then immediately free it. It then calls METIS. If
   \boldsymbol{\ast} this pre-allocation fails, it is possible that METIS will fail as
   * well, and so CHOLMOD returns with an out-of-memory condition without
```

```
* calling METIS.
   * METIS_NodeND (used in the CHOLMOD_METIS ordering option) with its
   * default parameter settings typically uses about (4*nz+40n+4096)
   * times sizeof(int) memory, where nz is equal to the number of entries
    * in A for the symmetric case or AA' if an unsymmetric matrix is
    * being ordered (where nz includes both the upper and lower parts
    * of A or AA'). The observed "upper bound" (with 2 exceptions),
    * measured in an instrumented copy of METIS 4.0.1 on thousands of
    * matrices, is (10*nz+50*n+4096) * sizeof(int). Two large matrices
    * exceeded this bound, one by almost a factor of 2 (Gupta/gupta2).
   * If your program is terminated by METIS, try setting metis_memory to
   st 2.0, or even higher if needed. By default, CHOLMOD assumes that METIS
   * does not have this problem (so that CHOLMOD will work correctly when
    * this issue is fixed in METIS). Thus, the default value is zero.
    * This work-around is not guaranteed anyway.
    * If a matrix exceeds this predicted memory usage, AMD is attempted
    * instead. It, too, may run out of memory, but if it does so it will
    * not terminate your program.
double metis_dswitch ;
                           /* METIS_NodeND in METIS 4.0.1 gives a seg */
                           /* fault with one matrix of order n = 3005 and
size_t metis_nswitch ;
                            * nz = 6,036,025. This is a very dense graph.
 * The workaround is to use AMD instead of METIS for matrices of dimension
 * greater than Common->metis_nswitch (default 3000) or more and with
 * density of Common->metis_dswitch (default 0.66) or more.
 * cholmod_nested_dissection has no problems with the same matrix, even
 * though it uses METIS_ComputeVertexSeparator on this matrix. If this
 * seg fault does not affect you, set metis_nswitch to zero or less,
 * and CHOLMOD will not switch to AMD based just on the density of the
 * matrix (it will still switch to AMD if the metis_memory parameter
 * causes the switch).
/* workspace */
/* ----- */
/* CHOLMOD has several routines that take less time than the size of
 * workspace they require. Allocating and initializing the workspace would
 * dominate the run time, unless workspace is allocated and initialized
 * just once. CHOLMOD allocates this space when needed, and holds it here
 * between calls to CHOLMOD. cholmod_start sets these pointers to NULL
 * (which is why it must be the first routine called in CHOLMOD).
 * cholmod_finish frees the workspace (which is why it must be the last
 * call to CHOLMOD).
 */
                /* size of Flag and Head */
size_t nrow ;
int64_t mark ;
                 /* mark value for Flag array */
size_t iworksize ; /* size of Iwork. Upper bound: 6*nrow+ncol */
size_t xworksize ; /* size of Xwork, in bytes.
```

```
* maxrank*nrow*sizeof(double) for update/downdate.
                    * 2*nrow*sizeof(double) otherwise */
/* initialized workspace: contents needed between calls to CHOLMOD */
void *Flag ;
                   /* size nrow, an integer array. Kept cleared between
                    * calls to cholmod rouines (Flag [i] < mark) */
                   /* size nrow+1, an integer array. Kept cleared between
void *Head ;
                    * calls to cholmod routines (Head [i] = EMPTY) */
void *Xwork ;
                   /* a double array. Its size varies. It is nrow for
                    * most routines (cholmod_rowfac, cholmod_add,
   * cholmod_aat, cholmod_norm, cholmod_ssmult) for the real case, twice
   * that when the input matrices are complex or zomplex. It is of size
   * 2*nrow for cholmod_rowadd and cholmod_rowdel. For cholmod_updown,
    * its size is maxrank*nrow where maxrank is 2, 4, or 8. Kept cleared
    * between calls to cholmod (set to zero). */
/* uninitialized workspace, contents not needed between calls to CHOLMOD */
void *Iwork ;
                   /* size iworksize, 2*nrow+ncol for most routines,
                    * up to 6*nrow+ncol for cholmod_analyze. */
                   /* If CHOLMOD_LONG, Flag, Head, and Iwork are
int itype ;
                    * int64_t. Otherwise all three are int. */
                   /* double or float */
int dtype ;
    /* Common->itype and Common->dtype are used to define the types of all
     * sparse matrices, triplet matrices, dense matrices, and factors
     * created using this Common struct. The itypes and dtypes of all
     * parameters to all CHOLMOD routines must match. */
int no_workspace_reallocate; /* this is an internal flag, used as a
    * precaution by cholmod_analyze. It is normally false. If true,
    * cholmod_allocate_work is not allowed to reallocate any workspace;
    * they must use the existing workspace in Common (Iwork, Flag, Head,
    * and Xwork). Added for CHOLMOD v1.1 */
/* statistics */
/* ----- */
/* fl and lnz are set only in cholmod_analyze and cholmod_rowcolcounts,
 * in the Cholesky module. modfl is set only in the Modify module. */
int status ;
                       /* error code */
double fl ;
                       /* LL' flop count from most recent analysis */
                       /* fundamental nz in L */
double lnz ;
                       /* nonzeros in tril(A) if A is symmetric/lower,
double anz ;
                        * triu(A) if symmetric/upper, or tril(A*A') if
                        * unsymmetric, in last call to cholmod_analyze. */
                       /* flop count from most recent update/downdate/
double modfl ;
                        * rowadd/rowdel (excluding flops to modify the
                        * solution to Lx=b, if computed) */
                       /* # of objects malloc'ed minus the # free'd*/
size_t malloc_count ;
```

```
/* peak memory usage in bytes */
size_t memory_usage ;
size_t memory_inuse ; /* current memory usage in bytes */
double nrealloc_col ; /* # of column reallocations */
double nrealloc_factor ;/* # of factor reallocations due to col. reallocs */
double ndbounds_hit; /* # of times diagonal modified by dbound */
/* # of flops to compute A(:,f)*A(:,f)' */
double aatfl;
int called_nd ;
               /* TRUE if the last call to
                * cholmod_analyze called NESDIS or METIS. */
               /* FALSE if SUITESPARSE_BLAS_INT overflow;
int blas_ok ;
                  TRUE otherwise */
/* ----- */
/* SuiteSparseQR control parameters: */
/* ----- */
/* ----- */
/* SuiteSparseQR statistics */
/* ----- */
/* was other1 [0:3] */
                    /* flop count for SPQR */
double SPQR_flopcount ;
double SPQR_factorize_time ; /* factorize time in seconds for SPQR */
double SPQR_solve_time ;
                    /* backsolve time in seconds */
/* was SPQR_xstat [0:3] */
/* tolerance used */
double SPQR_tol_used ;
double SPQR_norm_E_fro ;
                    /* Frobenius norm of dropped entries */
/* was SPQR_istat [0:9] */
int64_t SPQR_istat [10] ;
/* ----- */
/* GPU configuration and statistics */
/* ----- */
/* useGPU: 1 if gpu-acceleration is requested */
/*
        0 if gpu-acceleration is prohibited */
/*
        -1 if gpu-acceleration is undefined in which case the */
/*
          environment CHOLMOD_USE_GPU will be queried and used. */
/*
          useGPU=-1 is only used by CHOLMOD and treated as 0 by SPQR */
int useGPU;
/* for CHOLMOD: */
size_t maxGpuMemBytes;
```

```
double maxGpuMemFraction;
    /* for SPQR: */
   size_t gpuMemorySize;
                                /* Amount of memory in bytes on the GPU */
   double gpuKernelTime;
                               /* Time taken by GPU kernels */
    int64_t gpuFlops;
                               /* Number of flops performed by the GPU */
    int gpuNumKernelLaunches;
                              /* Number of GPU kernel launches */
    /* If not using the GPU, these items are not used, but they should be
      present so that the CHOLMOD Common has the same size whether the GPU
       is used or not. This way, all packages will agree on the size of
       the CHOLMOD Common, regardless of whether or not they are compiled
      with the GPU libraries or not */
#ifdef SUITESPARSE_CUDA
    /* in CUDA, these three types are pointers */
   #define CHOLMOD_CUBLAS_HANDLE cublasHandle_t
   #define CHOLMOD_CUDASTREAM
                                 cudaStream_t
    #define CHOLMOD_CUDAEVENT
                                 cudaEvent_t
#else
    /* ... so make them void * pointers if the GPU is not being used */
   #define CHOLMOD_CUBLAS_HANDLE void *
   #define CHOLMOD_CUDASTREAM
                                 void *
    #define CHOLMOD_CUDAEVENT
                                  void *
#endif
   CHOLMOD_CUBLAS_HANDLE cublasHandle ;
    /* a set of streams for general use */
   CHOLMOD_CUDASTREAM
                         gpuStream[CHOLMOD_HOST_SUPERNODE_BUFFERS];
    CHOLMOD_CUDAEVENT
                         cublasEventPotrf [3] ;
    CHOLMOD_CUDAEVENT
                         updateCKernelsComplete;
   CHOLMOD_CUDAEVENT
                         updateCBuffersFree[CHOLMOD_HOST_SUPERNODE_BUFFERS];
                         /* pointer to single allocation of device memory */
    void *dev_mempool;
   size_t dev_mempool_size;
   void *host_pinned_mempool; /* pointer to single allocation of pinned mem */
   size_t host_pinned_mempool_size;
   size_t devBuffSize;
    int
          ibuffer;
    double syrkStart ;
                               /* time syrk started */
   /* run times of the different parts of CHOLMOD (GPU and CPU) */
   double cholmod_cpu_gemm_time ;
   double cholmod_cpu_syrk_time ;
   double cholmod_cpu_trsm_time ;
   double cholmod_cpu_potrf_time ;
   double cholmod_gpu_gemm_time ;
   double cholmod_gpu_syrk_time ;
   double cholmod_gpu_trsm_time ;
   double cholmod_gpu_potrf_time ;
```

```
double cholmod_assemble_time ;
   double cholmod_assemble_time2 ;
    /* number of times the BLAS are called on the CPU and the GPU */
   size_t cholmod_cpu_gemm_calls ;
   size_t cholmod_cpu_syrk_calls ;
   size_t cholmod_cpu_trsm_calls ;
   size_t cholmod_cpu_potrf_calls ;
   size_t cholmod_gpu_gemm_calls ;
   size_t cholmod_gpu_syrk_calls ;
   size_t cholmod_gpu_trsm_calls ;
   size_t cholmod_gpu_potrf_calls ;
   double chunk ;
                        // chunksize for computing # of threads to use.
                        // Given nwork work to do, # of threads is
                        // max (1, min (floor (work / chunk), nthreads_max))
   int nthreads_max ; // max # of threads to use in CHOLMOD. Defaults to
                        // SUITESPARSE_OPENMP_MAX_THREADS.
} cholmod_common ;
```

Purpose: The cholmod_common Common object contains parameters, statistics, and workspace used within CHOLMOD. The first call to CHOLMOD must be cholmod_start, which initializes this object.

12.3 cholmod_start: start CHOLMOD

```
int cholmod_start
(
    cholmod_common *Common
);
int cholmod_l_start (cholmod_common *);
```

Purpose: Sets the default parameters, clears the statistics, and initializes all workspace pointers to NULL. The int/long type is set in Common->itype.

12.4 cholmod_finish: finish CHOLMOD

```
int cholmod_finish
(
    cholmod_common *Common
);
int cholmod_l_finish (cholmod_common *);
```

Purpose: This must be the last call to CHOLMOD.

12.5 cholmod_defaults: set default parameters

```
int cholmod_defaults
(
    cholmod_common *Common
);
int cholmod_l_defaults (cholmod_common *);
```

Purpose: Sets the default parameters.

12.6 cholmod_maxrank: maximum update/downdate rank

Purpose: Returns the maximum rank for an update/downdate.

12.7 cholmod_allocate_work: allocate workspace

Purpose: Allocates workspace in Common. The workspace consists of the integer Head, Flag, and Iwork arrays, of size nrow+1, nrow, and iworksize, respectively, and a double array Xwork of size xworksize. The Head array is normally equal to -1 when it is cleared. If the Flag array is cleared, all entries are less than Common->mark. The Iwork array is not kept in any particular state. The integer type is int or long, depending on whether the cholmod_or cholmod_l_routines are used.

12.8 cholmod_free_work: free workspace

```
int cholmod_free_work
(
    cholmod_common *Common
);
int cholmod_l_free_work (cholmod_common *);
```

Purpose: Frees the workspace in Common.

12.9 cholmod_clear_flag: clear Flag array

```
int64_t cholmod_clear_flag
(
    cholmod_common *Common
);
int64_t cholmod_l_clear_flag (cholmod_common *);
```

Purpose: Increments Common->mark so that the Flag array is now cleared.

12.10 cholmod_error: report error

Purpose: This routine is called when CHOLMOD encounters an error. It prints a message (if printing is enabled), sets Common->status. It then calls the user error handler routine Common->error_handler, if it is not NULL.

12.11 cholmod_dbound: bound diagonal of L

Purpose: Ensures that entries on the diagonal of L for an LL^T factorization are greater than or equal to Common->dbound. For an LDL^T factorization, it ensures that the magnitude of the entries of D are greater than or equal to Common->dbound.

12.12 cholmod_hypot: sqrt(x*x+y*y)

```
double cholmod_hypot
(
    /* ---- input ---- */
    double x, double y
);
double cholmod_l_hypot (double, double);
```

Purpose: Computes the magnitude of a complex number. This routine is the default value for the Common->hypotenuse function pointer. See also hypot, in the standard math.h header. If you have the ANSI C99 hypot, you can use Common->hypotenuse = hypot. The cholmod_hypot routine is provided in case you are using the ANSI C89 standard, which does not have hypot.

12.13 cholmod_divcomplex: complex divide

Purpose: Divides two complex numbers. It returns 1 if a divide-by-zero occurred, or 0 otherwise. This routine is the default value for the Common->complex_divide function pointer. This return value is the single exception to the CHOLMOD rule that states all int return values are TRUE if successful or FALSE otherwise. The exception is made to match the return value of a different complex divide routine that is not a part of CHOLMOD, but can be used via the function pointer.

13 Core Module: cholmod_sparse object

13.1 cholmod_sparse: compressed-column sparse matrix

```
typedef struct cholmod_sparse_struct
   size_t nrow ;
                       /* the matrix is nrow-by-ncol */
   size_t ncol ;
   size_t nzmax ;
                       /* maximum number of entries in the matrix */
    /* pointers to int32_t or int64_t: */
                       /* p [0..ncol], the column pointers */
   void *i :
                       /* i [0..nzmax-1], the row indices */
    /* for unpacked matrices only: */
   void *nz ;
                        /* nz [0..ncol-1], the # of nonzeros in each col. In
                         * packed form, the nonzero pattern of column j is in
        * A->i [A->p [j] ... A->p [j+1]-1]. In unpacked form, column j is in
        * A->i [A->p [j] ... A->p [j]+A->nz[j]-1] instead. In both cases, the
       * numerical values (if present) are in the corresponding locations in
       * the array x (or z if A->xtype is CHOLMOD_ZOMPLEX). */
    /* pointers to double or float: */
   void *x ;
                       /* size nzmax or 2*nzmax, if present */
   void *z ;
                       /* size nzmax, if present */
    int stype ;
                       /* Describes what parts of the matrix are considered:
        * 0: matrix is "unsymmetric": use both upper and lower triangular parts
              (the matrix may actually be symmetric in pattern and value, but
             both parts are explicitly stored and used). May be square or
             rectangular.
       * >0: matrix is square and symmetric, use upper triangular part.
             Entries in the lower triangular part are ignored.
        * <0: matrix is square and symmetric, use lower triangular part.
             Entries in the upper triangular part are ignored.
       * Note that stype>0 and stype<0 are different for cholmod_sparse and
        * cholmod_triplet. See the cholmod_triplet data structure for more
        * details.
        */
                        /* CHOLMOD_INT:
                                           p, i, and nz are int32_t.
    int itype ;
                         * CHOLMOD_INTLONG: p is int64_t,
                                           i and nz are int32_t.
                        * CHOLMOD LONG:
                                           p, i, and nz are int64_t */
   int xtype ;
                       /* pattern, real, complex, or zomplex */
   int dtype ;
                       /* x and z are double or float */
    int sorted;
                       /* TRUE if columns are sorted, FALSE otherwise */
    int packed;
                       /* TRUE if packed (nz ignored), FALSE if unpacked
                        * (nz is required) */
} cholmod_sparse ;
```

Purpose: Stores a sparse matrix in compressed-column form.

13.2 cholmod_allocate_sparse: allocate sparse matrix

```
cholmod_sparse *cholmod_allocate_sparse
   /* ---- input ---- */
                   /* # of rows of A */
   size_t nrow,
   size_t ncol,
                    /* # of columns of A */
   size_t nzmax,
                    /* max # of nonzeros of A */
   int sorted,
                    /* TRUE if columns of A sorted, FALSE otherwise */
                      /* TRUE if A will be packed, FALSE otherwise */
   int packed,
                      /* stype of A */
   int stype,
                     /* CHOLMOD_PATTERN, _REAL, _COMPLEX, or _ZOMPLEX */
   int xtype,
   /* ----- */
   cholmod_common *Common
cholmod_sparse *cholmod_l_allocate_sparse (size_t, size_t, size_t, int, int,
   int, int, cholmod_common *);
```

Purpose: Allocates a sparse matrix. A->i, A->x, and A->z are not initialized. The matrix returned is all zero, but it contains space enough for nzmax entries.

13.3 cholmod_free_sparse: free sparse matrix

```
int cholmod_free_sparse
(
    /* ---- in/out --- */
    cholmod_sparse **A, /* matrix to deallocate, NULL on output */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_free_sparse (cholmod_sparse **, cholmod_common *);
```

Purpose: Frees a sparse matrix.

13.4 cholmod_reallocate_sparse: reallocate sparse matrix

Purpose: Reallocates a sparse matrix, so that it can contain nznew entries.

13.5 cholmod_nnz: number of entries in sparse matrix

```
int64_t cholmod_nnz
(
    /* ---- input ---- */
    cholmod_sparse *A,
    /* ------ */
    cholmod_common *Common
);
int64_t cholmod_l_nnz (cholmod_sparse *, cholmod_common *);
```

Purpose: Returns the number of entries in a sparse matrix.

13.6 cholmod_speye: sparse identity matrix

Purpose: Returns the sparse identity matrix.

13.7 cholmod_spzeros: sparse zero matrix

Purpose: Returns the sparse zero matrix. This is another name for cholmod_allocate_sparse, but with fewer parameters (the matrix is packed, sorted, and unsymmetric).

13.8 cholmod_transpose: transpose sparse matrix

Purpose: Returns the transpose or complex conjugate transpose of a sparse matrix.

13.9 cholmod_ptranspose: transpose/permute sparse matrix

```
cholmod_sparse *cholmod_ptranspose
(
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to transpose */
   int values,
                     /* 0: pattern, 1: array transpose, 2: conj. transpose */
                     /* if non-NULL, F = A(p,f) or A(p,p) */
   int32_t *Perm,
   int32_t *fset,
                     /* subset of 0:(A->ncol)-1 */
                     /* size of fset */
   size_t fsize,
   /* ----- */
   cholmod_common *Common
);
cholmod_sparse *cholmod_l_ptranspose (cholmod_sparse *, int, int64_t *,
   int64_t *, size_t, cholmod_common *);
```

Purpose: Returns A' or A(p,p)' if A is symmetric. Returns A', A(:,f)', or A(p,f)' if A is unsymmetric. See cholmod_transpose_unsym for a discussion of how f is used; this usage deviates from the MATLAB notation. Can also return the array transpose.

13.10 cholmod_sort: sort columns of a sparse matrix

```
int cholmod_sort
(
    /* ---- in/out --- */
    cholmod_sparse *A, /* matrix to sort */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_sort (cholmod_sparse *, cholmod_common *);
```

Purpose: Sorts the columns of the matrix A. Returns A in packed form, even if it starts as unpacked. Removes entries in the ignored part of a symmetric matrix.

13.11 cholmod_transpose_unsym: transpose/permute unsymmetric sparse matrix

```
int cholmod_transpose_unsym
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to transpose */
                      /* 0: pattern, 1: array transpose, 2: conj. transpose */
   int values,
   int32_t *Perm,
                     /* size nrow, if present (can be NULL) */
                     /* subset of 0:(A->ncol)-1 */
   int32_t *fset,
   size_t fsize,
                      /* size of fset */
   /* ---- output --- */
   cholmod_sparse *F, /* F = A', A(:,f)', or A(p,f)' */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_transpose_unsym (cholmod_sparse *, int, int64_t *,
   int64_t *, size_t, cholmod_sparse *, cholmod_common *);
```

Purpose: Transposes and optionally permutes an unsymmetric sparse matrix. The output matrix must be preallocated before calling this routine.

Computes F=A', F=A(:,f)' or F=A(p,f)', except that the indexing by f does not work the same as the MATLAB notation (see below). A->stype is zero, which denotes that both the upper and lower triangular parts of A are present (and used). The matrix A may in fact be symmetric in pattern and/or value; A->stype just denotes which part of A are stored. A may be rectangular.

The integer vector p is a permutation of 0:m-1, and f is a subset of 0:n-1, where A is m-by-n. There can be no duplicate entries in p or f.

Three kinds of transposes are available, depending on the values parameter:

- 0: do not transpose the numerical values; create a CHOLMOD_PATTERN matrix
- 1: array transpose
- 2: complex conjugate transpose (same as 2 if input is real or pattern)

The set f is held in fset and fsize:

- fset = NULL means ":" in MATLAB. fset is ignored.
- fset != NULL means f = fset [0..fsize-1].
- fset != NULL and fsize = 0 means f is the empty set.

Columns not in the set f are considered to be zero. That is, if A is 5-by-10 then F=A(:,[3 4])' is not 2-by-5, but 10-by-5, and rows 3 and 4 of F are equal to columns 3 and 4 of A (the other rows of F are zero). More precisely, in MATLAB notation:

```
[m n] = size (A)
F = A
notf = ones (1,n)
notf (f) = 0
F (:, find (notf)) = 0
F = F'
```

If you want the MATLAB equivalent F=A(p,f) operation, use cholmod_submatrix instead (which does not compute the transpose). F->nzmax must be large enough to hold the matrix F. If F->nz is present then F->nz [j] is equal to the number of entries in column j of F. A can be sorted or unsorted, with packed or unpacked columns. If f is present and not sorted in ascending order, then F is unsorted (that is, it may contain columns whose row indices do not appear in ascending order). Otherwise, F is sorted (the row indices in each column of F appear in strictly ascending order).

F is returned in packed or unpacked form, depending on F->packed on input. If F->packed is FALSE, then F is returned in unpacked form (F->nz must be present). Each row i of F is large enough to hold all the entries in row i of A, even if f is provided. That is, F->i and F->x [F->p [i] .. F->p [i] + F->nz [i] - 1] contain all entries in A(i,f), but F->p [i+1] - F->p [i] is equal to the number of nonzeros in A (i,:), not just A (i,f). The cholmod_transpose_unsym routine is the only operation in CHOLMOD that can produce an unpacked matrix.

13.12 cholmod_transpose_sym: transpose/permute symmetric sparse matrix

Purpose: Computes F = A' or F = A(p,p)', the transpose or permuted transpose, where A->stype is nonzero. A must be square and symmetric. If A->stype > 0, then A is a symmetric matrix where just the upper part of the matrix is stored. Entries in the lower triangular part may be present, but are ignored. If A->stype < 0, then A is a symmetric matrix where just the lower part of the matrix is stored. Entries in the upper triangular part may be present, but are ignored. If F=A', then F is returned sorted; otherwise F is unsorted for the F=A(p,p)' case. There can be no duplicate entries in p.

Three kinds of transposes are available, depending on the values parameter:

- 0: do not transpose the numerical values; create a CHOLMOD_PATTERN matrix
- 1: array transpose
- 2: complex conjugate transpose (same as 2 if input is real or pattern)

For cholmod_transpose_unsym and cholmod_transpose_sym, the output matrix F must already be pre-allocated by the caller, with the correct dimensions. If F is not valid or has the wrong dimensions, it is not modified. Otherwise, if F is too small, the transpose is not computed; the

contents of F->p contain the column pointers of the resulting matrix, where F->p [F->ncol] > F->nzmax. In this case, the remaining contents of F are not modified. F can still be properly freed with cholmod_free_sparse.

13.13 cholmod_band: extract band of a sparse matrix

Purpose: Returns C = tril (triu (A,k1), k2). C is a matrix consisting of the diagonals of A from k1 to k2. k=0 is the main diagonal of A, k=1 is the superdiagonal, k=-1 is the subdiagonal, and so on. If A is m-by-n, then:

- k1=-m means C = tril (A,k2)
- k2=n means C = triu (A,k1)
- k1=0 and k2=0 means C = diag(A), except C is a matrix, not a vector

Values of k1 and k2 less than -m are treated as -m, and values greater than n are treated as n.

A can be of any symmetry (upper, lower, or unsymmetric); C is returned in the same form, and packed. If A->stype > 0, entries in the lower triangular part of A are ignored, and the opposite is true if A->stype < 0. If A has sorted columns, then so does C. C has the same size as A.

C can be returned as a numerical valued matrix (if A has numerical values and mode > 0), as a pattern-only (mode = 0), or as a pattern-only but with the diagonal entries removed (mode < 0).

The xtype of A can be pattern or real. Complex or zomplex cases are supported only if mode is ≤ 0 (in which case the numerical values are ignored).

13.14 cholmod_band_inplace: extract band, in place

Purpose: Same as cholmod_band, except that it always operates in place. Only packed matrices can be converted in place.

13.15 cholmod_aat: compute AA^T

```
cholmod_sparse *cholmod_aat
   /* ---- input ---- */
   cholmod_sparse *A, /* input matrix; C=A*A' is constructed */
   int32_t *fset,
                       /* subset of 0:(A->ncol)-1 */
                       /* size of fset */
   size_t fsize,
                       /* >0: numerical, 0: pattern, <0: pattern (no diag),</pre>
   int mode,
                        * -2: pattern only, no diagonal, add 50%+n extra
                        * space to C */
   /* ----- */
   cholmod_common *Common
);
cholmod_sparse *cholmod_l_aat (cholmod_sparse *, int64_t *, size_t,
   int, cholmod_common *);
```

Purpose: Computes C = A*A or C = A(:,f)*A(:,f). A can be packed or unpacked, sorted or unsorted, but must be stored with both upper and lower parts (A->stype of zero). C is returned as packed, C->stype of zero (both upper and lower parts present), and unsorted. See cholmod_ssmult in the MatrixOps Module for a more general matrix-matrix multiply. The xtype of A can be pattern or real. Complex or zomplex cases are supported only if mode is ≤ 0 (in which case the numerical values are ignored). You can trivially convert C to a symmetric upper/lower matrix by changing C->stype to 1 or -1, respectively, after calling this routine.

13.16 cholmod_copy_sparse: copy sparse matrix

```
cholmod_sparse *cholmod_copy_sparse
(
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to copy */
    /* ----- */
    cholmod_common *Common
);
cholmod_sparse *cholmod_l_copy_sparse (cholmod_sparse *, cholmod_common *);
```

Purpose: Returns an exact copy of the input sparse matrix A.

13.17 cholmod_copy: copy (and change) sparse matrix

```
cholmod_sparse *cholmod_copy_sparse
(
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to copy */
   /* ----- */
   cholmod_common *Common
);
cholmod_sparse *cholmod_l_copy_sparse (cholmod_sparse *, cholmod_common *);
cholmod_sparse *cholmod_copy
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to copy */
                       /* requested stype of C */
                       /* >0: numerical, 0: pattern, <0: pattern (no diag) */</pre>
   int mode,
   cholmod_common *Common
);
cholmod_sparse *cholmod_l_copy (cholmod_sparse *, int, int, cholmod_common *);
```

Purpose: C = A, which allocates C and copies A into C, with possible change of stype. The diagonal can optionally be removed. The numerical entries can optionally be copied. This routine differs from cholmod_copy_sparse, which makes an exact copy of a sparse matrix.

A can be of any type (packed/unpacked, upper/lower/unsymmetric). C is packed and can be of any stype (upper/lower/unsymmetric), except that if A is rectangular C can only be unsymmetric. If the stype of A and C differ, then the appropriate conversion is made.

There are three cases for A->stype:

- < 0, lower: assume A is symmetric with just tril(A) stored; the rest of A is ignored
- 0, unsymmetric: assume A is unsymmetric; consider all entries in A
- > 0, upper: assume A is symmetric with just triu(A) stored; the rest of A is ignored

There are three cases for the requested symmetry of C (stype parameter):

- < 0, lower: return just tril(C)
- 0, unsymmetric: return all of C
- > 0, upper: return just triu(C)

This gives a total of nine combinations:

Equivalent MATLAB statements	Using cholmod_copy
C = A ;	A unsymmetric, C unsymmetric
C = tril (A) ;	A unsymmetric, C lower
C = triu (A) ;	A unsymmetric, C upper
U = triu (A) ; L = tril (U',-1) ; C = L+U ;	A upper, C unsymmetric
C = triu (A);;	A upper, C lower
C = triu (A) ;	A upper, C upper
L = tril (A) ; U = triu (L',1) ; C = L+U ;	A lower, C unsymmetric
C = tril (A) ;	A lower, C lower
C = tril (A);;	A lower, C upper

The xtype of A can be pattern or real. Complex or zomplex cases are supported only if values is FALSE (in which case the numerical values are ignored).

13.18 cholmod_add: add sparse matrices

```
cholmod_sparse *cholmod_add
   /* ---- input ---- */
   cholmod_sparse *A,
                         /* matrix to add */
                          /* matrix to add */
   cholmod_sparse *B,
   double alpha [2],
                          /* scale factor for A */
   double beta [2],
                          /* scale factor for B */
   int values,
                          /* if TRUE compute the numerical values of C */
   int sorted,
                          /* if TRUE, sort columns of C */
   /* ----- */
   cholmod_common *Common
);
cholmod_sparse *cholmod_l_add (cholmod_sparse *, cholmod_sparse *, double *,
   double *, int, int, cholmod_common *);
```

Purpose: Returns C = alpha*A + beta*B. If the stype of A and B match, then C has the same stype. Otherwise, C->stype is zero (C is unsymmetric).

13.19 cholmod_sparse_xtype: change sparse xtype

Purpose: Changes the **xtype** of a sparse matrix, to pattern, real, complex, or zomplex. Changing from complex or zomplex to real discards the imaginary part.

14 Core Module: cholmod_factor object

14.1 cholmod_factor object: a sparse Cholesky factorization

```
typedef struct cholmod_factor_struct
   /* for both simplicial and supernodal factorizations */
   size_t n ;
                  /* L is n-by-n */
   size_t minor ;
                  /* If the factorization failed, L->minor is the column
                   * at which it failed (in the range 0 to n-1). A value
                   * the matrix has not yet been factorized. */
   /* symbolic ordering and analysis */
   /* ----- */
   void *Perm ;
                 /* size n, permutation used */
   void *ColCount ; /* size n, column counts for simplicial L */
                 /* size n, inverse permutation. Only created by
   void *IPerm ;
                  * cholmod_solve2 if Bset is used. */
   /* ----- */
   /* simplicial factorization */
   /* ----- */
   size_t nzmax ;
                 /* size of i and x */
   void *p ;
                  /* p [0..ncol], the column pointers */
   void *i ;
                  /* i [0..nzmax-1], the row indices */
   void *x ;
                  /* x [0..nzmax-1], the numerical values */
   void *z ;
                  /* nz [0..ncol-1], the # of nonzeros in each column.
   void *nz ;
                   * i [p [j] ... p [j]+nz[j]-1] contains the row indices,
                   * and the numerical values are in the same locatins
                   * in x. The value of i [p [k]] is always k. */
   void *next ;
                  /* size ncol+2. next [j] is the next column in i/x */
                  /* size ncol+2. prev [j] is the prior column in i/x.
   void *prev ;
                   * head of the list is ncol+1, and the tail is ncol. */
   /* ----- */
   /* supernodal factorization */
   /* ----- */
   /* Note that L->x is shared with the simplicial data structure. L->x has
    * size L->nzmax for a simplicial factor, and size L->xsize for a supernodal
   * factor. */
   size_t nsuper ;
                 /* number of supernodes */
```

```
size_t maxcsize ;  /* size of largest update matrix */
size_t maxesize; /* max # of rows in supernodes, excl. triangular part */
               /* size nsuper+1, first col in each supernode */
/* size nsuper+1, pointers to integer patterns */
void *super ;
void *pi ;
                 /* size nsuper+1, pointers to real parts */
void *px ;
                 /* size ssize, integer part of supernodes */
void *s ;
/* factorization type */
int ordering ;  /* ordering method used */
int is_monotonic; /* TRUE if columns of L appear in order 0..n-1.
                   * Only applicable to simplicial numeric types. */
/* There are 8 types of factor objects that cholmod_factor can represent
 * (only 6 are used):
 * Numeric types (xtype is not CHOLMOD_PATTERN)
 * simplicial LDL': (is_ll FALSE, is_super FALSE). Stored in compressed
       column form, using the simplicial components above (nzmax, p, i,
       x, z, nz, next, and prev). The unit diagonal of L is not stored,
       and D is stored in its place. There are no supernodes.
 * simplicial LL': (is_ll TRUE, is_super FALSE). Uses the same storage
       scheme as the simplicial LDL', except that D does not appear.
       The first entry of each column of L is the diagonal entry of
       that column of L.
  supernodal LDL': (is_11 FALSE, is_super TRUE). Not used.
       FUTURE WORK: add support for supernodal LDL'
 * supernodal LL': (is_ll TRUE, is_super TRUE). A supernodal factor,
       using the supernodal components described above (nsuper, ssize,
       xsize, maxcsize, maxesize, super, pi, px, s, x, and z).
 * Symbolic types (xtype is CHOLMOD_PATTERN)
 * simplicial LDL': (is_11 FALSE, is_super FALSE). Nothing is present
       except Perm and ColCount.
 * simplicial LL': (is_ll TRUE, is_super FALSE). Identical to the
       simplicial LDL', except for the is_ll flag.
 * supernodal LDL': (is_ll FALSE, is_super TRUE). Not used.
```

```
FUTURE WORK: add support for supernodal LDL'
     * supernodal LL': (is_11 TRUE, is_super TRUE). A supernodal symbolic
           factorization. The simplicial symbolic information is present
            (Perm and ColCount), as is all of the supernodal factorization
           except for the numerical values (x and z).
   int itype ; /* The integer arrays are Perm, ColCount, p, i, nz,
                * next, prev, super, pi, px, and s. If itype is
                * CHOLMOD_INT, all of these are int arrays.
                * CHOLMOD_INTLONG: p, pi, px are int64_t, others int.
                                  all integer arrays are int64_t. */
                * CHOLMOD_LONG:
   int xtype ; /* pattern, real, complex, or zomplex */
   int dtype ; /* x and z double or float */
   int useGPU; /* Indicates the symbolic factorization supports
                * GPU acceleration */
} cholmod_factor ;
```

Purpose: An $\mathbf{L}\mathbf{L}^\mathsf{T}$ or $\mathbf{L}\mathbf{D}\mathbf{L}^\mathsf{T}$ factorization in simplicial or supernodal form. A simplicial factor is very similar to a **cholmod_sparse** matrix. For an $\mathbf{L}\mathbf{D}\mathbf{L}^\mathsf{T}$ factorization, the diagonal matrix \mathbf{D} is stored as the diagonal of \mathbf{L} ; the unit-diagonal of \mathbf{L} is not stored.

14.2 cholmod_free_factor: free factor

```
int cholmod_free_factor
(
    /* ---- in/out --- */
    cholmod_factor **L, /* factor to free, NULL on output */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_free_factor (cholmod_factor **, cholmod_common *);
```

Purpose: Frees a factor.

14.3 cholmod_allocate_factor: allocate factor

Purpose: Allocates a factor and sets it to identity.

14.4 cholmod_reallocate_factor: reallocate factor

Purpose: Reallocates a simplicial factor so that it can contain nznew entries.

14.5 cholmod_change_factor: change factor

```
int cholmod_change_factor
   /* ---- input ---- */
                     /* to CHOLMOD_PATTERN, _REAL, _COMPLEX, _ZOMPLEX */
   int to_xtype,
                      /* TRUE: convert to LL', FALSE: LDL' */
   int to_ll,
                     /* TRUE: convert to supernodal, FALSE: simplicial */
   int to_super,
                      /* TRUE: pack simplicial columns, FALSE: do not pack */
   int to_packed,
   int to_monotonic, /* TRUE: put simplicial columns in order, FALSE: not */
   /* ---- in/out --- */
   cholmod_factor *L, /* factor to modify */
    /* ----- */
   cholmod_common *Common
);
int cholmod_l_change_factor ( int, int, int, int, int, cholmod_factor *,
   cholmod_common *);
```

Purpose: Change the numeric or symbolic, **LL**^T or **LDL**^T, simplicial or super, packed or unpacked, and monotonic or non-monotonic status of a **cholmod_factor** object.

There are four basic classes of factor types:

- 1. simplicial symbolic: Consists of two size-n arrays: the fill-reducing permutation (L->Perm) and the nonzero count for each column of L (L->ColCount). All other factor types also include this information. L->ColCount may be exact (obtained from the analysis routines), or it may be a guess. During factorization, and certainly after update/downdate, the columns of L can have a different number of nonzeros. L->ColCount is used to allocate space. L->ColCount is exact for the supernodal factorizations. The nonzero pattern of L is not kept.
- 2. simplicial numeric: These represent L in a compressed column form. The variants of this type are:
 - LDL^T: L is unit diagonal. Row indices in column j are located in L->i [L->p [j] ... L->p [j] + L->nz [j]], and corresponding numeric values are in the same locations in L->x. The total number of entries is the sum of L->nz [j]. The unit diagonal is not stored; D is stored on the diagonal of L instead. L->p may or may not be monotonic. The order of storage of the columns in L->i and L->x is given by a doubly-linked list (L->prev and L->next). L->p is of size n+1, but only the first n entries are used. For the complex case, L->x is stored interleaved with real and imaginary parts, and is of size 2*lnz*sizeof(double). For the zomplex case, L->x is of size lnz*sizeof(double) and holds the real part; L->z is the same size and holds the imaginary part.
 - $\mathbf{L}\mathbf{L}^{\mathsf{T}}$: This is identical to the $\mathbf{L}\mathbf{D}\mathbf{L}^{\mathsf{T}}$ form, except that the non-unit diagonal of L is stored as the first entry in each column of L.
- 3. supernodal symbolic: A representation of the nonzero pattern of the supernodes for a supernodal factorization. There are L->nsuper supernodes. Columns L->super [k] to L->super [k+1]-1 are in the kth supernode. The row indices for the kth supernode are in L->s [L->pi

[k] ... L->pi [k+1]-1]. The numerical values are not allocated (L->x), but when they are they will be located in L->x [L->px [k] ... L->px [k+1]-1], and the L->px array is defined in this factor type.

For the complex case, L->x is stored interleaved with real/imaginary parts, and is of size 2*L->xsize*sizeof(double). The zomplex supernodal case is not supported, since it is not compatible with LAPACK and the BLAS.

4. supernodal numeric: Always an **LL**^T factorization. L has a non-unit diagonal. L->x contains the numerical values of the supernodes, as described above for the supernodal symbolic factor. For the complex case, L->x is stored interleaved, and is of size 2*L->xsize*sizeof(double). The zomplex supernodal case is not supported, since it is not compatible with LAPACK and the BLAS.

In all cases, the row indices in each column (L->i for simplicial L and L->s for supernodal L) are kept sorted from low indices to high indices. This means the diagonal of L (or D for a LDL^T factorization) is always kept as the first entry in each column. The elimination tree is not kept. The parent of node j can be found as the second row index in the jth column. If column j has no off-diagonal entries then node j is a root of the elimination tree.

The cholmod_change_factor routine can do almost all possible conversions. It cannot do the following conversions:

- Simplicial numeric types cannot be converted to a supernodal symbolic type. This would simultaneously deallocate the simplicial pattern and numeric values and reallocate uninitialized space for the supernodal pattern. This isn't useful for the user, and not needed by CHOLMOD's own routines either.
- Only a symbolic factor (simplicial to supernodal) can be converted to a supernodal numeric factor.

Some conversions are meant only to be used internally by other CHOLMOD routines, and should not be performed by the end user. They allocate space whose contents are undefined:

- converting from simplicial symbolic to supernodal symbolic.
- converting any factor to supernodal numeric.

Supports all xtypes, except that there is no supernodal zomplex L.

The to_xtype parameter is used only when converting from symbolic to numeric or numeric to symbolic. It cannot be used to convert a numeric xtype (real, complex, or zomplex) to a different numeric xtype. For that conversion, use cholmod_factor_xtype instead.

14.6 cholmod_pack_factor: pack the columns of a factor

```
int cholmod_pack_factor
(
    /* ---- in/out --- */
    cholmod_factor *L, /* factor to modify */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_pack_factor (cholmod_factor *, cholmod_common *);
```

Purpose: Pack the columns of a simplicial **LDL**^T or **LL**^T factorization. This can be followed by a call to cholmod_reallocate_factor to reduce the size of L to the exact size required by the factor, if desired. Alternatively, you can leave the size of L->i and L->x the same, to allow space for future updates/rowadds. Each column is reduced in size so that it has at most Common->grow2 free space at the end of the column. Does nothing and returns silently if given any other type of factor. Does not force the columns of L to be monotonic. It thus differs from

cholmod_change_factor (xtype, L->is_ll, FALSE, TRUE, TRUE, L, Common) which packs the columns and ensures that they appear in monotonic order.

14.7 cholmod_reallocate_column: reallocate one column of a factor

Purpose: Reallocates the space allotted to a single column of L.

14.8 cholmod_factor_to_sparse: sparse matrix copy of a factor

Purpose: Returns a column-oriented sparse matrix containing the pattern and values of a simplicial or supernodal numerical factor, and then converts the factor into a simplicial symbolic factor. If L is already packed, monotonic, and simplicial (which is the case when cholmod_factorize uses the simplicial Cholesky factorization algorithm) then this routine requires only a small amount of time and memory, independent of n. It only operates on numeric factors (real, complex, or zomplex). It does not change L->xtype (the resulting sparse matrix has the same xtype as L). If this routine fails, L is left unmodified.

14.9 cholmod_copy_factor: copy factor

```
cholmod_factor *cholmod_copy_factor
(
    /* ---- input ---- */
    cholmod_factor *L, /* factor to copy */
    /* ------ */
    cholmod_common *Common
);
cholmod_factor *cholmod_l_copy_factor (cholmod_factor *, cholmod_common *);
```

Purpose: Returns an exact copy of a factor.

14.10 cholmod_factor_xtype: change factor xtype

Purpose: Changes the xtype of a factor, to pattern, real, complex, or zomplex. Changing from complex or zomplex to real discards the imaginary part. You cannot change a supernodal factor to the zomplex xtype.

15 Core Module: cholmod_dense object

15.1 cholmod_dense object: a dense matrix

```
typedef struct cholmod_dense_struct
   size_t nrow ;
                       /* the matrix is nrow-by-ncol */
   size_t ncol ;
   size_t nzmax ;
                       /* maximum number of entries in the matrix */
                       /* leading dimension (d >= nrow must hold) */
   size_t d ;
                       /* size nzmax or 2*nzmax, if present */
   void *x ;
   void *z ;
                       /* size nzmax, if present */
                       /* pattern, real, complex, or zomplex */
   int xtype ;
                       /* x and z double or float */
   int dtype ;
} cholmod_dense ;
```

Purpose: Contains a dense matrix.

15.2 cholmod_allocate_dense: allocate dense matrix

Purpose: Allocates a dense matrix.

15.3 cholmod_free_dense: free dense matrix

```
int cholmod_free_dense
(
    /* ---- in/out --- */
    cholmod_dense **X, /* dense matrix to deallocate, NULL on output */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_free_dense (cholmod_dense **, cholmod_common *);
```

Purpose: Frees a dense matrix.

15.4 cholmod_ensure_dense: ensure dense matrix has a given size and type

Purpose: Ensures a dense matrix has a given size and type.

15.5 cholmod_zeros: dense zero matrix

Purpose: Returns an all-zero dense matrix.

15.6 cholmod_ones: dense matrix, all ones

Purpose: Returns a dense matrix with each entry equal to one.

15.7 cholmod_eye: dense identity matrix

Purpose: Returns a dense identity matrix.

15.8 cholmod_sparse_to_dense: dense matrix copy of a sparse matrix

```
cholmod_dense *cholmod_sparse_to_dense
(
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to copy */
    /* ------ */
    cholmod_common *Common
);
cholmod_dense *cholmod_l_sparse_to_dense (cholmod_sparse *,
    cholmod_common *);
```

Purpose: Returns a dense copy of a sparse matrix.

15.9 cholmod_dense_to_sparse: sparse matrix copy of a dense matrix

Purpose: Returns a sparse copy of a dense matrix.

15.10 cholmod_copy_dense: copy dense matrix

```
cholmod_dense *cholmod_copy_dense
(
    /* ---- input ---- */
    cholmod_dense *X,    /* matrix to copy */
    /* ------ */
    cholmod_common *Common
);
cholmod_dense *cholmod_l_copy_dense (cholmod_dense *, cholmod_common *);
```

Purpose: Returns a copy of a dense matrix.

15.11 cholmod_copy_dense2: copy dense matrix (preallocated)

```
int cholmod_copy_dense2
(
    /* ---- input ---- */
    cholmod_dense *X,    /* matrix to copy */
    /* ---- output --- */
    cholmod_dense *Y,    /* copy of matrix X */
    /* ------- */
    cholmod_common *Common
);
int cholmod_l_copy_dense2 (cholmod_dense *, cholmod_dense *, cholmod_common *);
```

Purpose: Returns a copy of a dense matrix, placing the result in a preallocated matrix Y.

15.12 cholmod_dense_xtype: change dense matrix xtype

Purpose: Changes the xtype of a dense matrix, to real, complex, or zomplex. Changing from complex or zomplex to real discards the imaginary part.

16 Core Module: cholmod_triplet object

16.1 cholmod_triplet object: sparse matrix in triplet form

```
typedef struct cholmod_triplet_struct
   size_t nrow ;
                       /* the matrix is nrow-by-ncol */
   size_t ncol ;
   size_t nzmax ;
                       /* maximum number of entries in the matrix */
   size_t nnz ;
                       /* number of nonzeros in the matrix */
   void *i ;
                       /* i [0..nzmax-1], the row indices */
   void *;;
                       /* j [0..nzmax-1], the column indices */
                       /* size nzmax or 2*nzmax, if present */
   void *x ;
   void *z ;
                       /* size nzmax, if present */
    int stype ;
                        /* Describes what parts of the matrix are considered:
        * 0: matrix is "unsymmetric": use both upper and lower triangular parts
              (the matrix may actually be symmetric in pattern and value, but
             both parts are explicitly stored and used). May be square or
             rectangular.
        * >0: matrix is square and symmetric. Entries in the lower triangular
             part are transposed and added to the upper triangular part when
             the matrix is converted to cholmod_sparse form.
        * <0: matrix is square and symmetric. Entries in the upper triangular
             part are transposed and added to the lower triangular part when
             the matrix is converted to cholmod_sparse form.
       * Note that stype>0 and stype<0 are different for cholmod_sparse and
       * cholmod_triplet. The reason is simple. You can permute a symmetric
       * triplet matrix by simply replacing a row and column index with their
       * new row and column indices, via an inverse permutation. Suppose
       * P = L \rightarrow Perm is your permutation, and Pinv is an array of size n.
       * Suppose a symmetric matrix A is represent by a triplet matrix T, with
        * entries only in the upper triangular part. Then the following code:
               Ti = T -> i;
               Ti = T->i;
                for (k = 0 ; k < n ; k++) Pinv [P [k]] = k ;
                for (k = 0; k < nz; k++) Ti [k] = Pinv [Ti [k]];
                for (k = 0 ; k < nz ; k++) Tj [k] = Pinv [Tj [k]] ;
       * creates the triplet form of C=P*A*P'. However, if T initially
        * contains just the upper triangular entries (T->stype = 1), after
       * permutation it has entries in both the upper and lower triangular
        * parts. These entries should be transposed when constructing the
        * cholmod_sparse form of A, which is what cholmod_triplet_to_sparse
        * does. Thus:
                C = cholmod_triplet_to_sparse (T, 0, &Common);
        * will return the matrix C = P*A*P'.
        * Since the triplet matrix T is so simple to generate, it's quite easy
```

```
* to remove entries that you do not want, prior to converting T to the
* cholmod_sparse form. So if you include these entries in T, CHOLMOD
* assumes that there must be a reason (such as the one above). Thus,
* no entry in a triplet matrix is ever ignored.
*/

int itype ; /* CHOLMOD_LONG: i and j are int64_t. Otherwise int */
int xtype ; /* pattern, real, complex, or zomplex */
int dtype ; /* x and z are double or float */
} cholmod_triplet ;
```

Purpose: Contains a sparse matrix in triplet form.

16.2 cholmod_allocate_triplet: allocate triplet matrix

```
cholmod_triplet *cholmod_allocate_triplet
   /* ---- input ---- */
                     /* # of rows of T */
   size_t nrow,
                       /* # of columns of T */
   size_t ncol,
                       /* max # of nonzeros of T */
   size_t nzmax,
   int stype,
                       /* stype of T */
                      /* CHOLMOD_PATTERN, _REAL, _COMPLEX, or _ZOMPLEX */
   int xtype,
   cholmod_common *Common
);
cholmod_triplet *cholmod_l_allocate_triplet (size_t, size_t, size_t, int, int,
    cholmod_common *);
```

Purpose: Allocates a triplet matrix.

16.3 cholmod_free_triplet: free triplet matrix

Purpose: Frees a triplet matrix.

16.4 cholmod_reallocate_triplet: reallocate triplet matrix

Purpose: Reallocates a triplet matrix so that it can hold nznew entries.

16.5 cholmod_sparse_to_triplet: triplet matrix copy of a sparse matrix

```
cholmod_triplet *cholmod_sparse_to_triplet
(
    /* ---- input ---- */
    cholmod_sparse *A,    /* matrix to copy */
    /* ------ */
    cholmod_common *Common
);
cholmod_triplet *cholmod_l_sparse_to_triplet (cholmod_sparse *,
    cholmod_common *);
```

Purpose: Returns a triplet matrix copy of a sparse matrix.

16.6 cholmod_triplet_to_sparse: sparse matrix copy of a triplet matrix

Purpose: Returns a sparse matrix copy of a triplet matrix. If the triplet matrix is symmetric with just the lower part present (T->stype < 0), then entries in the upper part are transposed and placed in the lower part when converting to a sparse matrix. Similarly, if the triplet matrix is symmetric with just the upper part present (T->stype > 0), then entries in the lower part are transposed and placed in the upper part when converting to a sparse matrix. Any duplicate entries are summed.

16.7 cholmod_copy_triplet: copy triplet matrix

```
cholmod_triplet *cholmod_copy_triplet
(
    /* ---- input ---- */
    cholmod_triplet *T, /* matrix to copy */
    /* ------ */
    cholmod_common *Common
);
cholmod_triplet *cholmod_l_copy_triplet (cholmod_triplet *, cholmod_common *);
```

Purpose: Returns an exact copy of a triplet matrix.

16.8 cholmod_triplet_xtype: change triplet xtype

Purpose: Changes the xtype of a dense matrix, to real, complex, or zomplex. Changing from complex or zomplex to real discards the imaginary part.

17 Core Module: memory management

17.1 cholmod_malloc: allocate memory

Purpose: Allocates a block of memory of size n*size, using the SuiteSparse_config.malloc_func function pointer (default is to use the ANSI C malloc routine). A value of n=0 is treated as n=1. If not successful, NULL is returned and Common->status is set to CHOLMOD_OUT_OF_MEMORY.

17.2 cholmod_calloc: allocate and clear memory

Purpose: Allocates a block of memory of size n*size, using the SuiteSparse_config.calloc_func function pointer (default is to use the ANSI C calloc routine). A value of n=0 is treated as n=1. If not successful, NULL is returned and Common->status is set to CHOLMOD_OUT_OF_MEMORY.

17.3 cholmod_free: free memory

Purpose: Frees a block of memory of size n*size, using the SuiteSparse_config.free_func function pointer (default is to use the ANSI C free routine). The size of the block (n and size) is only required so that CHOLMOD can keep track of its current and peak memory usage. This is a useful statistic, and it can also help in tracking down memory leaks. After the call to cholmod_finish, the count of allocated blocks (Common->malloc_count) should be zero, and the count of bytes in use (Common->memory_inuse) also should be zero. If you allocate a block with one size and free it with another, the Common->memory_inuse count will be wrong, but CHOLMOD will not have a memory leak.

17.4 cholmod_realloc: reallocate memory

Purpose: Reallocates a block of memory whose current size n*size, and whose new size will be nnew*size if successful, using the SuiteSparse_config.calloc_func function pointer (default is to use the ANSI C realloc routine). If the reallocation is not successful, p is returned unchanged and Common->status is set to CHOLMOD_OUT_OF_MEMORY. The value of n is set to nnew if successful, or left unchanged otherwise. A value of nnew=0 is treated as nnew=1.

17.5 cholmod_realloc_multiple: reallocate memory

```
int cholmod_realloc_multiple
   /* ---- input ---- */
                       /* requested # of items in reallocated blocks */
   size_t nnew,
   int nint,
                       /* number of int/int64_t blocks */
   int xtype,
                      /* CHOLMOD_PATTERN, _REAL, _COMPLEX, or _ZOMPLEX */
   /* ---- in/out --- */
                       /* int or int64_t block */
   void **Iblock,
   void **Jblock,
                       /* int or int64_t block */
   void **Xblock,
                       /* complex, double, or float block */
   void **Zblock,
                       /* zomplex case only: double or float block */
   size_t *n,
                       /* current size of the I,J,X,Z blocks on input,
                        * nnew on output if successful */
    /* ----- */
   cholmod_common *Common
);
int cholmod_l_realloc_multiple (size_t, int, int, void **, void **, void **,
   void **, size_t *, cholmod_common *);
```

Purpose: Reallocates multiple blocks of memory, all with the same number of items (but with different item sizes). Either all reallocations succeed, or all are returned to their original size.

18 Core Module: version control

18.1 cholmod_version: return current CHOLMOD version

Purpose: Returns the CHOLMOD version number, so that it can be tested at run time, even if the caller does not have access to the CHOLMOD include files. For example, for a CHOLMOD version 3.2.1, the version array will contain 3, 2, and 1, in that order. This function appears in CHOLMOD 2.1.1 and later. You can check if the function exists with the CHOLMOD_HAS_VERSION_FUNCTION macro, so that the following code fragment works in any version of CHOLMOD:

```
#ifdef CHOLMOD_HAS_VERSION_FUNCTION
v = cholmod_version (NULL);
#else
v = CHOLMOD_VERSION;
#endif
```

Note that cholmod_version and cholmod_l_version have identical prototypes. Both use int's. Unlike all other CHOLMOD functions, this function does not take the Common object as an input parameter, and it does not use any definitions from any include files. Thus, the caller can access this function even if the caller does not include any CHOLMOD include files.

The above code fragment does require the #include "cholmod.h", of course, but cholmod_version can be called without it, if necessary.

19 Check Module routines

No CHOLMOD routines print anything, except for the cholmod_print_* routines in the Check Module, and the cholmod_error routine. The SuiteSparse_config.printf_function is a pointer to printf by default; you can redirect the output of CHOLMOD by redefining this pointer. If the function pointer is NULL, CHOLMOD does not print anything.

The Common->print parameter determines how much detail is printed. Each value of Common->print listed below also prints the items listed for smaller values of Common->print:

- 0: print nothing; check the data structures and return TRUE or FALSE.
- 1: print error messages.
- 2: print warning messages.
- 3: print a one-line summary of the object.
- 4: print a short summary of the object (first and last few entries).
- 5: print the entire contents of the object.

Values less than zero are treated as zero, and values greater than five are treated as five.

19.1 cholmod_check_common: check Common object

```
int cholmod_check_common
(
    cholmod_common *Common
);
int cholmod_l_check_common (cholmod_common *);
```

Purpose: Check if the Common object is valid.

19.2 cholmod_print_common: print Common object

```
int cholmod_print_common
(
    /* ---- input ---- */
    const char *name,    /* printed name of Common object */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_print_common (const char *, cholmod_common *);
```

Purpose: Print the Common object and check if it is valid. This prints the CHOLMOD parameters and statistics.

19.3 cholmod_check_sparse: check sparse matrix

```
int cholmod_check_sparse
(
    /* ---- input ---- */
    cholmod_sparse *A, /* sparse matrix to check */
    /* ------- */
    cholmod_common *Common
);
int cholmod_l_check_sparse (cholmod_sparse *, cholmod_common *);
```

Purpose: Check if a sparse matrix is valid.

19.4 cholmod_print_sparse: print sparse matrix

```
int cholmod_print_sparse
(
    /* ---- input ---- */
    cholmod_sparse *A, /* sparse matrix to print */
    const char *name, /* printed name of sparse matrix */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_print_sparse (cholmod_sparse *, const char *, cholmod_common *);
```

Purpose: Print a sparse matrix and check if it is valid.

19.5 cholmod_check_dense: check dense matrix

```
int cholmod_check_dense
(
    /* ---- input ---- */
    cholmod_dense *X,    /* dense matrix to check */
    /* ------- */
    cholmod_common *Common
);
int cholmod_l_check_dense (cholmod_dense *, cholmod_common *);
```

Purpose: Check if a dense matrix is valid.

19.6 cholmod_print_dense: print dense matrix

```
int cholmod_print_dense
(
    /* ---- input ---- */
    cholmod_dense *X,    /* dense matrix to print */
    const char *name,    /* printed name of dense matrix */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_print_dense (cholmod_dense *, const char *, cholmod_common *);
```

Purpose: Print a dense matrix and check if it is valid.

19.7 cholmod_check_factor: check factor

```
int cholmod_check_factor
(
    /* ---- input ---- */
    cholmod_factor *L, /* factor to check */
    /* ------- */
    cholmod_common *Common
);
int cholmod_l_check_factor (cholmod_factor *, cholmod_common *);
```

Purpose: Check if a factor is valid.

19.8 cholmod_print_factor: print factor

```
int cholmod_print_factor
(
    /* ---- input ---- */
    cholmod_factor *L, /* factor to print */
    const char *name, /* printed name of factor */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_print_factor (cholmod_factor *, const char *, cholmod_common *);
```

Purpose: Print a factor and check if it is valid.

19.9 cholmod_check_triplet: check triplet matrix

```
int cholmod_check_triplet
(
    /* ---- input ---- */
    cholmod_triplet *T, /* triplet matrix to check */
    /* ------- */
    cholmod_common *Common
);
int cholmod_l_check_triplet (cholmod_triplet *, cholmod_common *);
```

Purpose: Check if a triplet matrix is valid.

19.10 cholmod_print_triplet: print triplet matrix

```
int cholmod_print_triplet
   /* ---- input ---- */
   cholmod_triplet *T, /* triplet matrix to print */
   const char *name, /* printed name of triplet matrix */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_print_triplet (cholmod_triplet *, const char *, cholmod_common *);
/* ------ */
/* cholmod_check_subset: check a subset */
/* ----- */
int cholmod_check_subset
   /* ---- input ---- */
               /* Set [0:len-1] is a subset of 0:n-1. Duplicates OK */
   int *Set,
   int64_t len,
                 /* size of Set (an integer array) */
   size_t n,
                 /* 0:n-1 is valid range */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_check_subset (int64_t *, int64_t, size_t,
   cholmod_common *);
```

Purpose: Print a triplet matrix and check if it is valid.

19.11 cholmod_check_subset: check subset

Purpose: Check if a subset is valid.

19.12 cholmod_print_subset: print subset

Purpose: Print a subset and check if it is valid.

19.13 cholmod_check_perm: check permutation

```
int cholmod_check_perm
   /* ---- input ---- */
                   /* Perm [0:len-1] is a permutation of subset of 0:n-1 */
   int32_t *Perm,
                   /* size of Perm (an integer array) */
   size_t len,
                   /* 0:n-1 is valid range */
   size_t n,
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_check_perm (int64_t *, size_t, size_t, cholmod_common *);
/* cholmod_print_perm: print a permutation vector */
/* ------ */
int cholmod_print_perm
   /* ---- input ---- */
   int32_t *Perm, /* Perm [0:len-1] is a permutation of subset of 0:n-1 */
                    /* size of Perm (an integer array) */
   size_t len,
   size_t n,
                    /* 0:n-1 is valid range */
   const char *name, /* printed name of Perm */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_print_perm (int64_t *, size_t, size_t, const char *,
   cholmod_common *);
```

Purpose: Check if a permutation is valid.

19.14 cholmod_print_perm: print permutation

Purpose: Print a permutation and check if it is valid.

19.15 cholmod_check_parent: check elimination tree

Purpose: Check if an elimination tree is valid.

19.16 cholmod_print_parent: print elimination tree

Purpose: Print an elimination tree and check if it is valid.

19.17 cholmod_read_triplet: read triplet matrix from file

Purpose: Read a sparse matrix in triplet form, using the the coord Matrix Market format (http://www.nist.gov/MatrixMarket). Skew-symmetric and complex symmetric matrices are returned with both upper and lower triangular parts present (an stype of zero). Real symmetric and complex Hermitian matrices are returned with just their upper or lower triangular part, depending on their stype. The Matrix Market array data type for dense matrices is not supported (use cholmod_read_dense for that case).

If the first line of the file starts with %%MatrixMarket, then it is interpreted as a file in Matrix Market format. The header line is optional. If present, this line must have the following format:

%%MatrixMarket matrix coord type storage

where type is one of: real, complex, pattern, or integer, and storage is one of: general, hermitian, symmetric, or skew-symmetric. In CHOLMOD, these roughly correspond to the xtype (pattern, real, complex, or zomplex) and stype (unsymmetric, symmetric/upper, and symmetric/lower). The strings are case-insensitive. Only the first character (or the first two for skew-symmetric) is significant. The coord token can be replaced with array in the Matrix Market format, but this format not supported by cholmod_read_triplet. The integer type is converted to real. The type is ignored; the actual type (real, complex, or pattern) is inferred from the number of tokens in each line of the file (2: pattern, 3: real, 4: complex). This is compatible with the Matrix Market format.

A storage of general implies an stype of zero (see below). A storage of symmetric and hermitian imply an stype of -1. Skew-symmetric and complex symmetric matrices are returned with an stype of 0. Blank lines, any other lines starting with "%" are treated as comments, and are ignored.

The first non-comment line contains 3 or 4 integers:

nrow ncol nnz stype

where stype is optional (stype does not appear in the Matrix Market format). The matrix is nrow-by-ncol. The following nnz lines (excluding comments) each contain a single entry. Duplicates are permitted, and are summed in the output matrix.

If stype is present, it denotes the storage format for the matrix.

- stype = 0 denotes an unsymmetric matrix (same as Matrix Market general).
- stype = -1 denotes a symmetric or Hermitian matrix whose lower triangular entries are stored. Entries may be present in the upper triangular part, but these are ignored (same as Matrix Market symmetric for the real case, hermitian for the complex case).

• stype = 1 denotes a symmetric or Hermitian matrix whose upper triangular entries are stored. Entries may be present in the lower triangular part, but these are ignored. This format is not available in the Matrix Market format.

If neither the stype nor the Matrix Market header are present, then the stype is inferred from the rest of the data. If the matrix is rectangular, or has entries in both the upper and lower triangular parts, then it is assumed to be unsymmetric (stype=0). If only entries in the lower triangular part are present, the matrix is assumed to have stype = -1. If only entries in the upper triangular part are present, the matrix is assumed to have stype = 1.

Each nonzero consists of one line with 2, 3, or 4 entries. All lines must have the same number of entries. The first two entries are the row and column indices of the nonzero. If 3 entries are present, the 3rd entry is the numerical value, and the matrix is real. If 4 entries are present, the 3rd and 4th entries in the line are the real and imaginary parts of a complex value.

The matrix can be either 0-based or 1-based. It is first assumed to be one-based (compatible with Matrix Market), with row indices in the range 1 to nool and column indices in the range 1 to nrow. If a row or column index of zero is found, the matrix is assumed to be zero-based (with row indices in the range 0 to nool-1 and column indices in the range 0 to nrow-1). This test correctly determines that all Matrix Market matrices are in 1-based form.

For symmetric pattern-only matrices, the kth diagonal (if present) is set to one plus the degree of the row k or column k (whichever is larger), and the off-diagonals are set to -1. A symmetric pattern-only matrix with a zero-free diagonal is thus converted into a symmetric positive definite matrix. All entries are set to one for an unsymmetric pattern-only matrix. This differs from the MatrixMarket format (A = mmread ('file') returns a binary pattern for A for symmetric pattern-only matrices). To return a binary format for all pattern-only matrices, use A = mread('file', 1).

Example matrices that follow this format can be found in the CHOLMOD/Demo/Matrix and CHOLMOD/Tcov/Matrix directories. You can also try any of the matrices in the Matrix Market collection at http://www.nist.gov/MatrixMarket.

19.18 cholmod_read_sparse: read sparse matrix from file

Purpose: Read a sparse matrix in triplet form from a file (using cholmod_read_triplet) and convert to a CHOLMOD sparse matrix. The Matrix Market format is used. If Common->prefer_upper is TRUE (the default case), a symmetric matrix is returned stored in upper-triangular form (A->stype is 1). Otherwise, it is left in its original form, either upper or lower.

19.19 cholmod_read_dense: read dense matrix from file

Purpose: Read a dense matrix from a file, using the the array Matrix Market format (http://www.nist.gov/MatrixMarket).

19.20 cholmod_read_matrix: read a matrix from file

```
void *cholmod_read_matrix
   /* ---- input ---- */
                       /* file to read from, must already be open */
   int prefer,
                       /* If 0, a sparse matrix is always return as a
                               cholmod_triplet form. It can have any stype
                                (symmetric-lower, unsymmetric, or
                                symmetric-upper).
                        * If 1, a sparse matrix is returned as an unsymmetric
                                cholmod_sparse form (A->stype == 0), with both
                                upper and lower triangular parts present.
                                This is what the MATLAB mread mexFunction does,
                                since MATLAB does not have an stype.
                         * If 2, a sparse matrix is returned with an stype of 0
                                or 1 (unsymmetric, or symmetric with upper part
                         * This argument has no effect for dense matrices.
    /* ---- output---- */
    int *mtype,
                       /* CHOLMOD_TRIPLET, CHOLMOD_SPARSE or CHOLMOD_DENSE */
    cholmod_common *Common
);
void *cholmod_l_read_matrix (FILE *, int, int *, cholmod_common *);
```

Purpose: Read a sparse or dense matrix from a file, in Matrix Market format. Returns a void pointer to either a cholmod_triplet, cholmod_sparse, or cholmod_dense object.

19.21 cholmod_write_sparse: write a sparse matrix to a file

Purpose: Write a sparse matrix to a file in Matrix Market format. Optionally include comments, and print explicit zero entries given by the pattern of the Z matrix. If not NULL, the Z matrix must have the same dimensions and stype as A.

Returns the symmetry in which the matrix was printed (1 to 7) or -1 on failure. See the cholmod_symmetry function for a description of the return codes.

If A and Z are sorted on input, and either unsymmetric (stype = 0) or symmetric-lower (stype ; 0), and if A and Z do not overlap, then the triplets are sorted, first by column and then by row index within each column, with no duplicate entries. If all the above holds except stype ; 0, then the triplets are sorted by row first and then column.

19.22 cholmod_write_dense: write a dense matrix to a file

Purpose: Write a dense matrix to a file in Matrix Market format. Optionally include comments. Returns ¿ 0 if successful, -1 otherwise (1 if rectangular, 2 if square). A dense matrix is written in "general" format; symmetric formats in the Matrix Market standard are not exploited.

20 Cholesky Module routines

20.1 cholmod_analyze: symbolic factorization

```
cholmod_factor *cholmod_analyze
(
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to order and analyze */
    /* ------ */
    cholmod_common *Common
);
cholmod_factor *cholmod_l_analyze (cholmod_sparse *, cholmod_common *);
```

Purpose: Orders and analyzes a matrix (either simplicial or supernodal), in preparation for numerical factorization via cholmod_factorize or via the "expert" routines cholmod_rowfac and cholmod_super_numeric.

In the symmetric case, A or A(p,p) is analyzed, where p is the fill-reducing ordering. In the unsymmetric case, A*A' or A(p,:)*A(p,:)' is analyzed. The cholmod_analyze_p routine can be given a user-provided permutation p (see below).

The default ordering strategy is to first try AMD. The ordering quality is analyzed, and if AMD obtains an ordering where nnz(L) is greater than or equal to 5*nnz(tril(A)) (or 5*nnz(tril(A*A')) if A is unsymmetric) and the floating-point operation count for the subsequent factorization is greater than or equal to 500*nnz(L), then METIS is tried (if installed). For cholmod_analyze_p, the user-provided ordering is also tried. This default behavior is obtained when Common->nmethods is zero. In this case, methods 0, 1, and 2 in Common->method[...] are reset to user-provided, AMD, and METIS, respectively. The ordering with the smallest nnz(L) is kept.

If Common->default_nesdis is true (nonzero), then CHOLMOD's nested dissection (NESDIS) is used for the default strategy described above, in place of METIS.

Other ordering options can be requested. These include:

- 1. natural: A is not permuted to reduce fill-in.
- 2. user-provided: a permutation can be provided to cholmod_analyze_p.
- 3. AMD: approximate minimum degree (AMD for the symmetric case, COLAMD for the A*A' case).
- 4. METIS: nested dissection with METIS_NodeND
- 5. NESDIS: CHOLMOD's nested dissection using METIS_NodeComputeSeparator, followed by a constrained minimum degree (CAMD or CSYMAMD for the symmetric case, CCOLAMD for the A*A' case). This is typically slower than METIS, but typically provides better orderings.

Multiple ordering options can be tried (up to 9 of them), and the best one is selected (the one that gives the smallest number of nonzeros in the simplicial factor L). If one method fails, cholmod_analyze keeps going, and picks the best among the methods that succeeded. This routine fails (and returns NULL) if either the initial memory allocation fails, all ordering methods fail, or the supernodal analysis (if requested) fails. Change Common->nmethods to the number of methods you wish to try. By default, the 9 methods available are:

- 1. user-provided permutation (only for cholmod_analyze_p).
- 2. AMD with default parameters.
- 3. METIS with default parameters.
- 4. NESDIS with default parameters: stopping the partitioning when the graph is of size nd_small = 200 or less, remove nodes with more than max (16, prune_dense * sqrt (n)) nodes where prune_dense = 10, and follow partitioning with constrained minimum degree ordering (CAMD for the symmetric case, CCOLAMD for the unsymmetric case).
- 5. natural ordering (with weighted postorder).
- 6. NESDIS, $nd_small = 20000$, $prune_dense = 10$.
- 7. NESDIS, nd_small = 4, prune_dense = 10, no constrained minimum degree.
- 8. NESDIS, $nd_small = 200$, $prune_dense = 0$.
- 9. COLAMD for A*A' or AMD for A

You can modify these 9 methods and the number of methods tried by changing parameters in the Common argument. If you know the best ordering for your matrix, set Common->nmethods to 1 and set Common->method[0].ordering to the requested ordering method. Parameters for each method can also be modified (refer to the description of cholmod_common for details).

Note that it is possible for METIS to terminate your program if it runs out of memory. This is not the case for any CHOLMOD or minimum degree ordering routine (AMD, COLAMD, CCOLAMD, or CSYMAMD). Since NESDIS relies on METIS, it too can terminate your program.

The selected ordering is followed by a weighted postorder of the elimination tree by default (see cholmod_postorder for details), unless Common->postorder is set to FALSE. The postorder does not change the number of nonzeros in **L** or the floating-point operation count. It does improve performance, particularly for the supernodal factorization. If you truly want the natural ordering with no postordering, you must set Common->postorder to FALSE.

The factor L is returned as simplicial symbolic if Common->supernodal is CHOLMOD_SIMPLICIAL (zero) or as supernodal symbolic if Common->supernodal is CHOLMOD_SUPERNODAL (two). If Common->supernodal is CHOLMOD_AUTO (one), then L is simplicial if the flop count per nonzero in L is less than Common->supernodal_switch (default: 40), and supernodal otherwise. In both cases, L->xtype is CHOLMOD_PATTERN. A subsequent call to cholmod_factorize will perform a simplicial or supernodal factorization, depending on the type of L.

For the simplicial case, L contains the fill-reducing permutation (L->Perm) and the counts of nonzeros in each column of L (L->ColCount). For the supernodal case, L also contains the nonzero pattern of each supernode.

If a simplicial factorization is selected, it will be \mathbf{LDL}^T by default, since this is the kind required by the Modify Module. CHOLMOD does not include a supernodal \mathbf{LDL}^T factorization, so if a supernodal factorization is selected, it will be in the form \mathbf{LL}^T . The \mathbf{LDL}^T method can be used to factorize positive definite matrices and indefinite matrices whose leading minors are well-conditioned (2-by-2 pivoting is not supported). The \mathbf{LL}^T method is restricted to positive definite matrices. To factorize a large indefinite matrix, set Common->supernodal to CHOLMOD_SIMPLICIAL,

and the simplicial \mathbf{LDL}^T method will always be used. This will be significantly slower than a supernodal \mathbf{LL}^T factorization, however.

Refer to cholmod_transpose_unsym for a description of f.

20.2 cholmod_factorize: numeric factorization

```
int cholmod_factorize
(
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to factorize */
    /* ---- in/out --- */
    cholmod_factor *L, /* resulting factorization */
    /* ------ */
    cholmod_common *Common
) ;
int cholmod_l_factorize (cholmod_sparse *, cholmod_factor *, cholmod_common *) ;
```

Purpose: Computes the numerical factorization of a symmetric matrix. The inputs to this routine are a sparse matrix A and the symbolic factor L from cholmod_analyze or a prior numerical factor L. If A is symmetric, this routine factorizes A(p,p). where p is the fill-reducing permutation (L->Perm). If A is unsymmetric, A(p,:)*A(p,:)' is factorized. The nonzero pattern of the matrix A must be the same as the matrix passed to cholmod_analyze for the supernodal case. For the simplicial case, it can be different, but it should be the same for best performance.

A simplicial factorization or supernodal factorization is chosen, based on the type of the factor L. If L->is_super is TRUE, a supernodal LL^T factorization is computed. Otherwise, a simplicial numeric factorization is computed, either LL^T or LDL^T , depending on Common->final_11 (the default for the simplicial case is to compute an LDL^T factorization).

Once the factorization is complete, it can be left as is or optionally converted into any simplicial numeric type, depending on the Common->final_* parameters. If converted from a supernodal to simplicial type, and Common->final_resymbol is TRUE, then numerically zero entries in L due to relaxed supernodal amalgamation are removed from the simplicial factor (they are always left in the supernodal form of L). Entries that are numerically zero but present in the simplicial symbolic pattern of L are left in place (the graph of L remains chordal). This is required for the update/downdate/rowadd/rowdel routines to work properly.

If the matrix is not positive definite the routine returns TRUE, but Common->status is set to CHOLMOD_NOT_POSDEF and L->minor is set to the column at which the failure occurred. Columns L->minor to L->n-1 are set to zero.

Supports any xtype (pattern, real, complex, or zomplex), except that the input matrix A cannot be pattern-only. If L is simplicial, its numeric xtype matches A on output. If L is supernodal, its xtype is real if A is real, or complex if A is complex or zomplex. CHOLMOD does not provide a supernodal zomplex factor, since it is incompatible with how complex numbers are stored in LAPACK and the BLAS.

20.3 cholmod_analyze_p: symbolic factorization, given permutation

```
cholmod_factor *cholmod_analyze_p
    /* ---- input ---- */
   cholmod_sparse *A, /* matrix to order and analyze */
   int32_t *UserPerm, /* user-provided permutation, size A->nrow */
   int32_t *fset,
                    /* subset of 0:(A->ncol)-1 */
   size_t fsize,
                     /* size of fset */
    /* ----- */
    cholmod_common *Common
);
cholmod_factor *cholmod_l_analyze_p (cholmod_sparse *, int64_t *,
    int64_t *, size_t, cholmod_common *);
cholmod_factor *cholmod_analyze_p2
    /* ---- input ---- */
                       /* FOR_SPQR
                                      (0): for SPQR but not GPU-accelerated
   int for_whom,
                          FOR_CHOLESKY (1): for Cholesky (GPU or not)
                          FOR_SPQRGPU (2): for SPQR with GPU acceleration */
   cholmod_sparse *A, /* matrix to order and analyze */
   int32_t *UserPerm, /* user-provided permutation, size A->nrow */
   int32_t *fset,
                      /* subset of 0:(A->ncol)-1 */
                      /* size of fset */
   size_t fsize,
    /* ----- */
    cholmod_common *Common
);
cholmod_factor *cholmod_l_analyze_p2 (int, cholmod_sparse *, int64_t *,
   int64_t *, size_t, cholmod_common *);
```

Purpose: Identical to cholmod_analyze, except that a user-provided permutation p can be provided, and the set f for the unsymmetric case can be provided. The matrices A(:,f)*A(:,f)' or A(p,f)*A(p,f)' can be analyzed in the the unsymmetric case.

20.4 cholmod_factorize_p: numeric factorization, given permutation

```
int cholmod_factorize_p
    /* ---- input ---- */
   cholmod_sparse *A, /* matrix to factorize */
   double beta [2],
                      /* factorize beta*I+A or beta*I+A'*A */
    int32_t *fset,
                      /* subset of 0:(A->ncol)-1 */
   size_t fsize,
                      /* size of fset */
    /* ---- in/out --- */
   cholmod_factor *L, /* resulting factorization */
    /* ----- */
   cholmod_common *Common
);
int cholmod_l_factorize_p (cholmod_sparse *, double *, int64_t *,
   size_t, cholmod_factor *, cholmod_common *);
```

Purpose: Identical to cholmod_factorize, but with additional options. The set f can be provided for the unsymmetric case; A(p,f)*A(p,f)' is factorized. The term beta*I can be added to the matrix before it is factorized, where beta is real. Only the real part, beta[0], is used.

20.5 cholmod_solve: solve a linear system

Purpose: Returns a solution X that solves one of the following systems:

```
system
                   sys parameter
                                            system
                                                           sys parameter
Ax = b
                   0: CHOLMOD_A
LDL^{\mathsf{T}}x = b
                                            \mathbf{L}^\mathsf{T}\mathbf{x} = \mathbf{b}
                  1: CHOLMOD_LDLt
                                                         5: CHOLMOD_Lt
LDx = b
                                            \mathbf{D}\mathbf{x} = \mathbf{b}
                   2: CHOLMOD_LD
                                                           6: CHOLMOD_D
DL^Tx = b
                                            x = Pb
                   3: CHOLMOD_DLt
                                                           7: CHOLMOD_P
                                            \mathbf{x} = \mathbf{P}^\mathsf{T}\mathbf{b} 8: CHOLMOD_Pt
Lx = b
                   4: CHOLMOD_L
```

The factorization can be simplicial \mathbf{LDL}^T , simplicial \mathbf{LL}^T , or supernodal \mathbf{LL}^T . For an \mathbf{LL}^T factorization, \mathbf{D} is the identity matrix. Thus CHOLMOD_LD and CHOLMOD_L solve the same system if an \mathbf{LL}^T factorization was performed, for example. This is one of the few routines in CHOLMOD for which the xtype of the input arguments need not match. If both L and B are real, then X is returned real. If either is complex or zomplex, X is returned as either complex or zomplex, depending on the Common->prefer_zomplex parameter (default is complex).

This routine does not check to see if the diagonal of L or D is zero, because sometimes a partial solve can be done with an indefinite or singular matrix. If you wish to check in your own code, test $L->\min$. If $L->\min$ == L->n, then the matrix has no zero diagonal entries. If $k = L->\min$ < L->n, then L(k,k) is zero for an LL^T factorization, or D(k,k) is zero for an LDL^T factorization.

Iterative refinement is not performed, but this can be easily done with the MatrixOps Module. See Demo/cholmod_demo.c for an example.

20.6 cholmod_spsolve: solve a linear system

Purpose: Identical to cholmod_solve, except that B and X are sparse. This function converts B to full format, solves the system, and then converts X back to sparse. If you want to solve with a sparse B and get just a partial solution back in X (corresponding to the pattern of B), use cholmod_solve2 below.

20.7 cholmod_solve2: solve a linear system, reusing workspace

```
int cholmod_solve2
                       /* returns TRUE on success, FALSE on failure */
   /* ---- input ---- */
   int sys,
                                   /* system to solve */
   cholmod_factor *L,
                                   /* factorization to use */
   cholmod_dense *B,
                                   /* right-hand-side */
   cholmod_sparse *Bset,
   /* ---- output --- */
   cholmod_dense **X_Handle,
                                   /* solution, allocated if need be */
   cholmod_sparse **Xset_Handle,
   /* ---- workspace */
   cholmod_dense **Y_Handle,
                                   /* workspace, or NULL */
   cholmod_dense **E_Handle,
                                   /* workspace, or NULL */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_solve2 (int, cholmod_factor *, cholmod_dense *, cholmod_sparse *,
   cholmod_dense **, cholmod_sparse **, cholmod_dense **, cholmod_dense **,
   cholmod_common *);
```

Purpose: Solve a linear system, optionally reusing workspace from a prior call to cholmod_solve2. The inputs to this function are the same as cholmod_solve, with the addition of three parameters: X, Y, and E. The dense matrix X is the solution on output. On input, &X can point to a NULL matrix, or be the wrong size. If that is the case, it is freed and allocated to be the proper size. If X has the right size and type on input, then the allocation is skipped. In contrast, the cholmod_solve function always allocates its output X. This cholmod_solve2 function allows you to reuse the memory space of a prior X, thereby saving time.

The two workspace matrices Y and E can also be reused between calls. You must free X Y, and E yourself, when your computations are done. Below is an example of usage. Note that X Y, and E must be defined on input (either NULL, or valid dense matrices).

```
cholmod_dense *X = NULL, *Y = NULL, *E = NULL;
...
cholmod_l_solve2 (sys, L, B1, NULL, &X, NULL, &Y, &E, Common);
cholmod_l_solve2 (sys, L, B2, NULL, &X, NULL, &Y, &E, Common);
```

```
cholmod_l_solve2 (sys, L, B3, NULL, &X, NULL, &Y, &E, Common) ;
  cholmod_l_free_dense (&X, Common) ;
  cholmod_l_free_dense (&Y, Common) ;
  cholmod_l_free_dense (&E, Common) ;

The equivalent when using cholmod_solve is:

  cholmod_dense *X = NULL, *Y = NULL, *E = NULL ;
    ...
  X = cholmod_l_solve (sys, L, B1, Common) ;
  cholmod_l_free_dense (&X, Common) ;
  X = cholmod_l_solve (sys, L, B2, Common) ;
  cholmod_l_free_dense (&X, Common) ;
  X = cholmod_l_solve (sys, L, B3, Common) ;
  cholmod_l_free_dense (&X, Common) ;
  cholmod_l_free_dense (&X, Common) ;
  cholmod_l_free_dense (&X, Common) ;
```

Both methods work fine, but in the second method with cholmod_solve, the internal workspaces (Y and E) and the solution (X) are allocated and freed on each call.

The cholmod_solve2 function can also solve for a subset of the solution vector X, if the optional Bset parameter is non-NULL. The right-hand-side B must be a single column vector, and its complexity (real, complex, zomplex) must match that of L. The vector B is dense, but it is assumed to be zero except for row indices specified in Bset. The vector Bset must be a sparse column vector, of dimension the same as B. Only the pattern of Bset is used. The solution X (a dense column vector) is modified on output, but is defined only in the rows defined by the sparse vector Xset. The entries in Bset are a subset of Xset (except if sys is CHOLMOD_P or CHOLMOD_Pt).

No memory allocations are done if the outputs and internal workspaces (X, Xset, Y, and E) have been allocated by a prior call (or if allocated by the user). To let cholmod_solve2 allocate these outputs and workspaces for you, simply initialize them to NULL (as in the example above). Since it is possible for this function to reallocate these 4 arrays, you should always re-acquire the pointers to their internal data (X->x for example) after calling cholmod_solve2), since they may change. They normally will not change except in the first call to this function.

On the first call to cholmod_solve2 when Bset is NULL, the factorization is converted from supernodal to simplicial, if needed. The inverse permutation is also computed and stored in the factorization object, L. This can take a modest amount of time. Subsequent calls to cholmod_solve2 with a small Bset are very fast (both asymptotically and in practice).

You can find an example of how to use cholmod_solve2 in the two demo programs, cholmod_demo and cholmod_l_demo.

20.8 cholmod_etree: find elimination tree

```
int cholmod_etree
(
    /* ---- input ---- */
    cholmod_sparse *A,
    /* ---- output --- */
    int32_t *Parent,    /* size ncol. Parent [j] = p if p is the parent of j */
    /* ------ */
```

```
cholmod_common *Common
);
int cholmod_l_etree (cholmod_sparse *, int64_t *, cholmod_common *);
```

Purpose: Computes the elimination tree of A or A'*A. In the symmetric case, the upper triangular part of A is used. Entries not in this part of the matrix are ignored. Computing the etree of a symmetric matrix from just its lower triangular entries is not supported. In the unsymmetric case, all of A is used, and the etree of A'*A is computed. Refer to [20] for a discussion of the elimination tree and its use in sparse Cholesky factorization.

20.9 cholmod_rowcolcounts: nonzeros counts of a factor

```
int cholmod_rowcolcounts
    /* ---- input ---- */
   cholmod_sparse *A, /* matrix to analyze */
    int32_t *fset,
                       /* subset of 0:(A->ncol)-1 */
    size_t fsize,
                       /* size of fset */
                       /* size nrow. Parent [i] = p if p is the parent of i */
    int32_t *Parent,
    int32_t *Post,
                       /* size nrow. Post [k] = i if i is the kth node in
                        * the postordered etree. */
    /* ---- output --- */
    int32_t *RowCount, /* size nrow. RowCount [i] = # entries in the ith row of
                        * L, including the diagonal. */
    int32_t *ColCount, /* size nrow. ColCount [i] = # entries in the ith
                        * column of L, including the diagonal. */
                       /* size nrow. First [i] = k is the least postordering
   int32_t *First,
                        * of any descendant of i. */
                       /* size nrow. Level [i] is the length of the path from
    int32_t *Level,
                        * i to the root, with Level [root] = 0. */
    cholmod_common *Common
);
int cholmod_l_rowcolcounts (cholmod_sparse *, int64_t *, size_t,
    int64_t *, int64_t *, int64_t *,
    int64_t *, int64_t *, int64_t *,
    cholmod_common *);
```

Purpose: Compute the row and column counts of the Cholesky factor L of the matrix A or A*A'. The etree and its postordering must already be computed (see cholmod_etree and cholmod_postorder) and given as inputs to this routine. For the symmetric case $(\mathbf{LL}^\mathsf{T} = \mathbf{A})$, A must be stored in symmetric/lower form (A->stype = -1). In the unsymmetric case, A*A' or A(:,f)*A(:,f)' can be analyzed. The fundamental floating-point operation count is returned in Common->f1 (this excludes extra flops due to relaxed supernodal amalgamation). Refer to cholmod_transpose_unsym for a description of f. The algorithm is described in [13, 15].

20.10 cholmod_analyze_ordering: analyze a permutation

```
int cholmod_analyze_ordering
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to analyze */
                     /* ordering method used */
   int ordering,
   int32_t *Perm,
                     /* size n, fill-reducing permutation to analyze */
   int32_t *fset,
                     /* subset of 0:(A->ncol)-1 */
   size_t fsize,
                      /* size of fset */
   /* ---- output --- */
   int32_t *Parent,
                       /* size n, elimination tree */
   int32_t *Post,
                       /* size n, postordering of elimination tree */
   int32_t *ColCount, /* size n, nnz in each column of L */
   /* ---- workspace */
                      /* size nworkspace for cholmod_postorder */
   int32_t *First,
                       /* size n workspace for cholmod_postorder */
   int32_t *Level,
   cholmod_common *Common
);
int cholmod_l_analyze_ordering (cholmod_sparse *, int, int64_t *,
   int64_t *, size_t, int64_t *, int64_t *, int64_t *, int64_t *, int64_t *,
   cholmod_common *);
```

Purpose: Given a matrix A and its fill-reducing permutation, compute the elimination tree, its (non-weighted) postordering, and the number of nonzeros in each column of L. Also computes the flop count, the total nonzeros in L, and the nonzeros in tril(A) (Common->fl, Common->lnz, and Common->anz). In the unsymmetric case, A(p,f)*A(p,f)' is analyzed, and Common->anz is the number of nonzero entries in the lower triangular part of the product, not in A itself.

Refer to cholmod_transpose_unsym for a description of f.

The column counts of L, flop count, and other statistics from cholmod_rowcolcounts are not computed if ColCount is NULL.

20.11 cholmod_amd: interface to AMD

Purpose: CHOLMOD interface to the AMD ordering package. Orders A if the matrix is symmetric. On output, Perm [k] = i if row/column i of A is the kth row/column of P*A*P'. This

corresponds to A(p,p) in MATLAB notation. If A is unsymmetric, cholmod_amd orders A*A' or A(:,f)*A(:,f)'. On output, Perm [k] = i if row/column i of A*A' is the kth row/column of P*A*A'*P'. This corresponds to A(p,:)*A(p,:)' in MATLAB notation. If f is present, A(p,f)*A(p,f)' is the permuted matrix. Refer to cholmod_transpose_unsym for a description of f.

Computes the flop count for a subsequent LL^{T} factorization, the number of nonzeros in L, and the number of nonzeros in the matrix ordered (A, A*A' or A(:,f)*A(:,f)'). These statistics are returned in Common->fl, Common->lnz, and Common->anz, respectively.

20.12cholmod_colamd: interface to COLAMD

```
int cholmod_colamd
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to order */
   int32_t *fset,
                       /* subset of 0:(A->ncol)-1 */
   size_t fsize,
                       /* size of fset */
                       /* if TRUE, follow with a coletree postorder */
   int postorder,
   /* ---- output --- */
   int32_t *Perm,
                       /* size A->nrow, output permutation */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_colamd (cholmod_sparse *, int64_t *, size_t, int,
   int64_t *, cholmod_common *);
```

CHOLMOD interface to the COLAMD ordering package. Finds a permutation p such that the Cholesky factorization of P*A*A'*P' is sparser than A*A', using COLAMD. If the postorder input parameter is TRUE, the column elimination tree is found and postordered, and the COLAMD ordering is then combined with its postordering (COLAMD itself does not perform this postordering). A must be unsymmetric (A->stype = 0).

cholmod_rowfac: row-oriented Cholesky factorization 20.13

```
int cholmod_rowfac
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to factorize */
   cholmod_sparse *F, /* used for A*A' case only. F=A' or A(:,fset)' */
   double beta [2], /* factorize beta*I+A or beta*I+A'*A */
   size_t kstart,
                     /* first row to factorize */
   size_t kend,
                      /* last row to factorize is kend-1 */
   /* ---- in/out --- */
   cholmod_factor *L,
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_rowfac (cholmod_sparse *, cholmod_sparse *, double *, size_t,
```

```
size_t, cholmod_factor *, cholmod_common *);
int cholmod_rowfac_mask
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to factorize */
    cholmod_sparse *F, /* used for A*A' case only. F=A' or A(:,fset)' */
   double beta [2],
                       /* factorize beta*I+A or beta*I+A'*A */
   size_t kstart,
                       /* first row to factorize */
   size_t kend,
                       /* last row to factorize is kend-1 */
    int32_t *mask,
                       /* if mask[i] >= 0, then set row i to zero */
    int32_t *RLinkUp,
                       /* link list of rows to compute */
    /* ---- in/out --- */
   cholmod_factor *L,
    /* ----- */
    cholmod_common *Common
);
int cholmod_l_rowfac_mask (cholmod_sparse *, cholmod_sparse *, double *, size_t,
   size_t, int64_t *, int64_t *, cholmod_factor *,
   cholmod_common *);
int cholmod_rowfac_mask2
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to factorize */
    cholmod_sparse *F, /* used for A*A' case only. F=A' or A(:,fset)' */
   double beta [2],
                       /* factorize beta*I+A or beta*I+A'*A */
                       /* first row to factorize */
   size_t kstart,
                       /* last row to factorize is kend-1 */
   size_t kend,
   int32_t *mask,
                       /* if mask[i] >= maskmark, then set row i to zero */
   int32_t maskmark,
    int32_t *RLinkUp,
                      /* link list of rows to compute */
    /* ---- in/out --- */
    cholmod_factor *L,
    /* ----- */
    cholmod_common *Common
);
int cholmod_l_rowfac_mask2 (cholmod_sparse *, cholmod_sparse *, double *,
    size_t, size_t, int64_t *, int64_t, int64_t *,
    cholmod_factor *, cholmod_common *);
```

Purpose: Full or incremental numerical **LDL**^T or **LL**^T factorization (simplicial, not supernodal). cholmod_factorize is the "easy" wrapper for this code, but it does not provide access to incremental factorization. The algorithm is the row-oriented, up-looking method described in [5]. See also [19]. No 2-by-2 pivoting (or any other pivoting) is performed.

cholmod_rowfac computes the full or incremental \mathbf{LDL}^T or \mathbf{LL}^T factorization of A+beta*I (where A is symmetric) or A*F+beta*I (where A and F are unsymmetric and only the upper triangular part of A*F+beta*I is used). It computes L (and D, for \mathbf{LDL}^T) one row at a time. The input scalar beta is real; only the real part (beta[0]) is used.

L can be a simplicial symbolic or numeric (L->is_super must be FALSE). A symbolic factor is converted immediately into a numeric factor containing the identity matrix.

For a full factorization, use kstart = 0 and kend = nrow. The existing nonzero entries (nu-

merical values in L->x and L->z for the zomplex case, and indices in L->i) are overwritten.

To compute an incremental factorization, select kstart and kend as the range of rows of L you wish to compute. Rows kstart to kend-1 of L will be computed. A correct factorization will be computed only if all descendants of all nodes kstart to kend-1 in the elimination tree have been factorized by a prior call to this routine, and if rows kstart to kend-1 have not been factorized. This condition is **not** checked on input.

In the symmetric case, A must be stored in upper form (A->stype is greater than zero). The matrix F is not accessed and may be NULL. Only columns kstart to kend-1 of A are accessed.

In the unsymmetric case, the typical case is F=A'. Alternatively, if F=A(:,f)', then this routine factorizes the matrix S = beta*I + A(:,f)*A(:,f)'. The product A*F is assumed to be symmetric; only the upper triangular part of A*F is used. F must be of size A->ncol by A->nrow.

20.14 cholmod_rowfac_mask: row-oriented Cholesky factorization

```
int cholmod_rowfac_mask
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to factorize */
   cholmod_sparse *F, /* used for A*A' case only. F=A' or A(:,fset)' */
   double beta [2],
                       /* factorize beta*I+A or beta*I+A'*A */
                       /* first row to factorize */
   size_t kstart,
   size_t kend,
                       /* last row to factorize is kend-1 */
    int32_t *mask,
                       /* if mask[i] >= 0, then set row i to zero */
    int32_t *RLinkUp, /* link list of rows to compute */
    /* ---- in/out --- */
    cholmod_factor *L,
    /* ----- */
    cholmod_common *Common
);
int cholmod_l_rowfac_mask (cholmod_sparse *, cholmod_sparse *, double *, size_t,
   size_t, int64_t *, int64_t *, cholmod_factor *,
    cholmod_common *);
int cholmod_rowfac_mask2
    /* ---- input ---- */
   cholmod_sparse *A, /* matrix to factorize */
   cholmod_sparse *F, /* used for A*A' case only. F=A' or A(:,fset)' */
   double beta [2], /* factorize beta*I+A or beta*I+A'*A */
   size_t kstart,
                       /* first row to factorize */
                       /* last row to factorize is kend-1 */
   size_t kend,
    int32_t *mask,
                       /* if mask[i] >= maskmark, then set row i to zero */
    int32_t maskmark,
    int32_t *RLinkUp,
                       /* link list of rows to compute */
    /* ---- in/out --- */
    cholmod_factor *L,
    /* ----- */
    cholmod_common *Common
);
int cholmod_l_rowfac_mask2 (cholmod_sparse *, cholmod_sparse *, double *,
    size_t, size_t, int64_t *, int64_t, int64_t *,
```

```
cholmod_factor *, cholmod_common *);
```

Purpose: For use in LPDASA only.

20.15 cholmod_row_subtree: pattern of row of a factor

Purpose: Compute the nonzero pattern of the solution to the lower triangular system

```
L(0:k-1,0:k-1) * x = A (0:k-1,k)
```

if A is symmetric, or

```
L(0:k-1,0:k-1) * x = A (0:k-1,:) * A (:,k)
```

if A is unsymmetric. This gives the nonzero pattern of row k of L (excluding the diagonal). The pattern is returned postordered, according to the subtree of the elimination tree rooted at node k.

The symmetric case requires A to be in symmetric-upper form.

The result is returned in R, a pre-allocated sparse matrix of size nrow-by-1, with R->nzmax >= nrow. R is assumed to be packed (Rnz [0] is not updated); the number of entries in R is given by Rp [0].

20.16 cholmod_row_lsubtree: pattern of row of a factor

Purpose: Identical to cholmod_row_subtree, except the elimination tree is found from L itself, not Parent. Also, F=A' is not provided; the nonzero pattern of the kth column of F is given by Fi and fnz instead.

20.17 cholmod_resymbol: re-do symbolic factorization

Purpose: Recompute the symbolic pattern of L. Entries not in the symbolic pattern of the factorization of A(p,p) or F*F', where F=A(p,f) or F=A(:,f), are dropped, where p = L->Perm is used to permute the input matrix A.

Refer to cholmod_transpose_unsym for a description of f.

If an entry in L is kept, its numerical value does not change.

This routine is used after a supernodal factorization is converted into a simplicial one, to remove zero entries that were added due to relaxed supernode amalgamation. It can also be used after a series of downdates to remove entries that would no longer be present if the matrix were factorized from scratch. A downdate (cholmod_updown) does not remove any entries from L.

20.18 cholmod_resymbol_noperm: re-do symbolic factorization

```
int cholmod_resymbol_noperm
(
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to analyze */
    int32_t *fset, /* subset of 0:(A->ncol)-1 */
    size_t fsize, /* size of fset */
    int pack, /* if TRUE, pack the columns of L */
    /* ---- in/out --- */
    cholmod_factor *L, /* factorization, entries pruned on output */
```

Purpose: Identical to cholmod_resymbol, except that the fill-reducing ordering L->Perm is not used.

20.19 cholmod_postorder: tree postorder

Purpose: Postorder a tree. The tree is either an elimination tree (the output from cholmod_etree) or a component tree (from cholmod_nested_dissection).

An elimination tree is a complete tree of n nodes with Parent [j] > j or Parent [j] = -1 if j is a root. On output Post [0..n-1] is a complete permutation vector; Post [k] = j if node j is the kth node in the postordered elimination tree, where k is in the range 0 to n-1.

A component tree is a subset of 0:n-1. Parent [j] = -2 if node j is not in the component tree. Parent [j] = -1 if j is a root of the component tree, and Parent [j] is in the range 0 to n-1 if j is in the component tree but not a root. On output, Post [k] is defined only for nodes in the component tree. Post [k] = j if node j is the kth node in the postordered component tree, where k is in the range 0 to the number of components minus 1. Node j is ignored and not included in the postorder if Parent [j] < -1. As a result, cholmod_check_parent (Parent, ...) and cholmod_check_perm (Post, ...) fail if used for a component tree and its postordering.

An optional node weight can be given. When starting a postorder at node j, the children of j are ordered in decreasing order of their weight. If no weights are given (Weight is NULL) then children are ordered in decreasing order of their node number. The weight of a node must be in the range 0 to n-1. Weights outside that range are silently converted to that range (weights < 0 are treated as zero, and weights $\ge n$ are treated as n-1).

20.20 cholmod_rcond: reciprocal condition number

Purpose: Returns a rough estimate of the reciprocal of the condition number: the minimum entry on the diagonal of L (or absolute entry of D for an \mathbf{LDL}^T factorization) divided by the maximum entry. L can be real, complex, or zomplex. Returns -1 on error, 0 if the matrix is singular or has a zero or NaN entry on the diagonal of L, 1 if the matrix is 0-by-0, or min(diag(L))/max(diag(L)) otherwise. Never returns NaN; if L has a NaN on the diagonal it returns zero instead.

21 Modify Module routines

21.1 cholmod_updown: update/downdate

Purpose: Updates/downdates the \mathbf{LDL}^T factorization (symbolic, then numeric), by computing a new factorization of

 $\overline{\mathbf{LDL}}^\mathsf{T} = \mathbf{LDL}^\mathsf{T} \pm \mathbf{CC}^\mathsf{T}$

where $\overline{\mathbf{L}}$ denotes the new factor. C must be sorted. It can be either packed or unpacked. As in all CHOLMOD routines, the columns of L are sorted on input, and also on output. If L does not contain a simplicial numeric \mathbf{LDL}^T factorization, it is converted into one. Thus, a supernodal \mathbf{LL}^T factorization can be passed to $\mathsf{cholmod_updown}$. A symbolic L is converted into a numeric identity matrix. If the initial conversion fails, the factor is returned unchanged.

If memory runs out during the update, the factor is returned as a simplicial symbolic factor. That is, everything is freed except for the fill-reducing ordering and its corresponding column counts (typically computed by cholmod_analyze).

Note that the fill-reducing permutation $L-\operatorname{Perm}$ is not used. The row indices of C refer to the rows of L, not A. If your original system is $\mathbf{LDL}^\mathsf{T} = \mathbf{PAP}^\mathsf{T}$ (where $\mathbf{P} = L-\operatorname{Perm}$), and you want to compute the \mathbf{LDL}^T factorization of $\mathbf{A} + \mathbf{CC}^\mathsf{T}$, then you must permute \mathbf{C} first. That is, if

$$\mathbf{PAP}^\mathsf{T} = \mathbf{LDL}^\mathsf{T}$$

is the initial factorization, then

$$\mathbf{P}(\mathbf{A} + \mathbf{C}\mathbf{C}^\mathsf{T})\mathbf{P}^\mathsf{T} = \mathbf{P}\mathbf{A}\mathbf{P}^\mathsf{T} + \mathbf{P}\mathbf{C}\mathbf{C}^\mathsf{T}\mathbf{P}^\mathsf{T} = \mathbf{L}\mathbf{D}\mathbf{L}^\mathsf{T} + (\mathbf{P}\mathbf{C})(\mathbf{P}\mathbf{C})^\mathsf{T} = \mathbf{L}\mathbf{D}\mathbf{L}^\mathsf{T} + \overline{\mathbf{C}\mathbf{C}}^\mathsf{T}$$

where $\overline{\mathbf{C}} = \mathbf{PC}$.

You can use the cholmod_submatrix routine in the MatrixOps Module to permute C, with:

```
Cnew = cholmod_submatrix (C, L->Perm, L->n, NULL, -1, TRUE, TRUE, Common) ;
```

Note that the sorted input parameter to cholmod_submatrix must be TRUE, because cholmod_updown requires C with sorted columns. Only real matrices are supported. The algorithms are described in [8, 9].

21.2 cholmod_updown_solve: update/downdate

Purpose: Identical to cholmod_updown, except the system $\mathbf{L}\mathbf{x} = \mathbf{b}$ is also updated/downdated. The new system is $\overline{\mathbf{L}}\overline{\mathbf{x}} = \mathbf{b} + \Delta \mathbf{b}$. The old solution \mathbf{x} is overwritten with $\overline{\mathbf{x}}$. Note that as in the update/downdate of \mathbf{L} itself, the fill-reducing permutation L->Perm is not used. The vectors \mathbf{x} and \mathbf{b} are in the permuted ordering, not your original ordering. This routine does not handle multiple right-hand-sides.

21.3 cholmod_updown_mark: update/downdate

Purpose: Identical to cholmod_updown_solve, except that only part of L is used in the update of the solution to Lx = b. For more details, see the source code file CHOLMOD/Modify/cholmod_updown.c. This routine is meant for use in the LPDASA linear program solver only, by Hager and Davis.

21.4 cholmod_updown_mask: update/downdate

```
int cholmod_updown_mask
   /* ---- input ---- */
                     /* TRUE for update, FALSE for downdate */
   int update,
   cholmod_sparse *C, /* the incoming sparse update */
   int32_t *colmark, /* array of size n. See cholmod_updown.c */
   int32_t *mask,
                     /* size n */
   /* ---- in/out --- */
   cholmod_factor *L, /* factor to modify */
   cholmod_dense *X, /* solution to Lx=b (size n-by-1) */
   cholmod_dense *DeltaB, /* change in b, zero on output */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_updown_mask (int, cholmod_sparse *, int64_t *,
   int64_t *, cholmod_factor *, cholmod_dense *, cholmod_dense *,
   cholmod_common *);
int cholmod_updown_mask2
   /* ---- input ---- */
                     /* TRUE for update, FALSE for downdate */
   int update,
   cholmod_sparse *C, /* the incoming sparse update */
   int32_t *mask,
                      /* size n */
   int32_t maskmark,
   /* ---- in/out --- */
   cholmod_factor *L, /* factor to modify */
   cholmod_dense *X, /* solution to Lx=b (size n-by-1) */
   cholmod_dense *DeltaB, /* change in b, zero on output */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_updown_mask2 (int, cholmod_sparse *, int64_t *,
   int64_t *, int64_t, cholmod_factor *, cholmod_dense *,
   cholmod_dense *, cholmod_common *);
```

Purpose: For use in LPDASA only.

21.5 cholmod_rowadd: add row to factor

```
cholmod_common *);
```

Purpose: Adds a row and column to an $\mathbf{LDL}^{\mathsf{T}}$ factorization. The kth row and column of L must be equal to the kth row and column of the identity matrix on input. Only real matrices are supported. The algorithm is described in [10].

21.6 cholmod_rowadd_solve: add row to factor

Purpose: Identical to cholmod_rowadd, except the system $\mathbf{L}\mathbf{x} = \mathbf{b}$ is also updated/downdated, just like cholmod_updown_solve.

21.7 cholmod_rowdel: delete row from factor

Purpose: Deletes a row and column from an **LDL**^T factorization. The kth row and column of L is equal to the kth row and column of the identity matrix on output. Only real matrices are supported.

21.8 cholmod_rowdel_solve: delete row from factor

Purpose: Identical to cholmod_rowdel, except the system $\mathbf{L}\mathbf{x} = \mathbf{b}$ is also updated/downdated, just like cholmod_updown_solve. When row/column k of \mathbf{A} is deleted from the system $\mathbf{A}\mathbf{y} = \mathbf{b}$, this can induce a change to \mathbf{x} , in addition to changes arising when \mathbf{L} and \mathbf{b} are modified. If this is the case, the kth entry of \mathbf{y} is required as input (yk). The algorithm is described in [10].

21.9 cholmod_rowadd_mark: add row to factor

```
int cholmod_rowadd_mark
(
   /* ---- input ---- */
   size_t k,
                       /* row/column index to add */
   cholmod_sparse *R, /* row/column of matrix to factorize (n-by-1) */
                       /* kth entry of the right hand side, b */
   double bk [2],
   int32_t *colmark, /* array of size n. See cholmod_updown.c */
   /* ---- in/out --- */
   cholmod_factor *L, /* factor to modify */
   cholmod_dense *X, /* solution to Lx=b (size n-by-1) */
   cholmod_dense *DeltaB, /* change in b, zero on output */
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_rowadd_mark (size_t, cholmod_sparse *, double *,
   int64_t *, cholmod_factor *, cholmod_dense *, cholmod_dense *,
   cholmod_common *);
```

Purpose: Identical to cholmod_rowadd_solve, except that only part of L is used in the update of the solution to Lx = b. For more details, see the source code file CHOLMOD/Modify/cholmod_rowadd.c. This routine is meant for use in the LPDASA linear program solver only.

21.10 cholmod_rowdel_mark: delete row from factor

```
int cholmod_rowdel_mark
   /* ---- input ---- */
                     /* row/column index to delete */
   size_t k,
   cholmod_sparse *R, /* NULL, or the nonzero pattern of kth row of L */
   double yk [2], /* kth entry in the solution to A*y=b */
   int32_t *colmark, /* array of size n. See cholmod_updown.c */
    /* ---- in/out --- */
   cholmod_factor *L, /* factor to modify */
   cholmod_dense *X, /* solution to Lx=b (size n-by-1) */
   cholmod_dense *DeltaB, /* change in b, zero on output */
   cholmod_common *Common
);
int cholmod_l_rowdel_mark (size_t, cholmod_sparse *, double *,
   int64_t *, cholmod_factor *, cholmod_dense *, cholmod_dense *,
   cholmod_common *);
```

Purpose: Identical to cholmod_rowadd_solve, except that only part of L is used in the update of the solution to Lx = b. For more details, see the source code file CHOLMOD/Modify/cholmod_rowdel.c. This routine is meant for use in the LPDASA linear program solver only.

22 MatrixOps Module routines

22.1 cholmod_drop: drop small entries

Purpose: Drop small entries from A, and entries in the ignored part of A if A is symmetric. No CHOLMOD routine drops small numerical entries from a matrix, except for this one. NaN's and Inf's are kept.

Supports pattern and real matrices; complex and zomplex matrices are not supported.

22.2 cholmod_norm_dense: dense matrix norm

Purpose: Returns the infinity-norm, 1-norm, or 2-norm of a dense matrix. Can compute the 2-norm only for a dense column vector. All xtypes are supported.

22.3 cholmod_norm_sparse: sparse matrix norm

Purpose: Returns the infinity-norm of 1-norm of a sparse matrix. All xtypes are supported.

22.4 cholmod_scale: scale sparse matrix

```
#define CHOLMOD_SCALAR 0
                               /* A = s*A */
#define CHOLMOD_ROW 1
                               /* A = diag(s)*A */
#define CHOLMOD_COL 2
                               /* A = A*diag(s) */
#define CHOLMOD_SYM 3
                               /* A = diag(s)*A*diag(s) */
int cholmod_scale
    /* ---- input ---- */
   cholmod_dense *S, /* scale factors (scalar or vector) */
    int scale,
                       /* type of scaling to compute */
    /* ---- in/out --- */
   cholmod_sparse *A, /* matrix to scale */
    /* ----- */
   cholmod_common *Common
);
int cholmod_l_scale (cholmod_dense *, int, cholmod_sparse *, cholmod_common *);
```

Purpose: Scales a matrix: A = diag(s)*A, A*diag(s), s*A, or diag(s)*A*diag(s).

A can be of any type (packed/unpacked, upper/lower/unsymmetric). The symmetry of A is ignored; all entries in the matrix are modified.

If A is m-by-n unsymmetric but scaled symmetrically, the result is

```
A = diag (s (1:m)) * A * diag (s (1:n))
```

Row or column scaling of a symmetric matrix still results in a symmetric matrix, since entries are still ignored by other routines. For example, when row-scaling a symmetric matrix where just the upper triangular part is stored (and lower triangular entries ignored) A = diag(s)*triu(A) is performed, where the result A is also symmetric-upper. This has the effect of modifying the implicit lower triangular part. In MATLAB notation:

```
U = diag(s)*triu(A) ;
L = tril (U',-1)
A = L + U ;
```

The scale parameter determines the kind of scaling to perform and the size of S:

scale	operation	size of S
${\tt CHOLMOD_SCALAR}$	s[0]*A	1
${\tt CHOLMOD_ROW}$	diag(s)*A	nrow-by-1 or 1-by-nrow
${\tt CHOLMOD_COL}$	A*diag(s)	ncol-by-1 or 1-by-ncol
$CHOLMOD_SYM$	<pre>diag(s)*A*diag(s)</pre>	<pre>max(nrow,ncol)-by-1, or 1-by-max(nrow,ncol)</pre>
0 1 1 .		

Only real matrices are supported.

22.5 cholmod_sdmult: sparse-times-dense matrix

```
int cholmod_sdmult
    /* ---- input ---- */
   cholmod_sparse *A, /* sparse matrix to multiply */
   int transpose,
                      /* use A if O, or A' otherwise */
   double alpha [2], /* scale factor for A */
                      /* scale factor for Y */
   double beta [2],
   cholmod_dense *X, /* dense matrix to multiply */
    /* ---- in/out --- */
    cholmod_dense *Y, /* resulting dense matrix */
    /* ----- */
   cholmod_common *Common
) ;
int cholmod_l_sdmult (cholmod_sparse *, int, double *, double *,
    cholmod_dense *, cholmod_dense *Y, cholmod_common *);
```

Purpose: Sparse matrix times dense matrix: Y = alpha*(A*X) + beta*Y or Y = alpha*(A'*X) + beta*Y, where A is sparse and X and Y are dense. When using A, X has A->ncol rows and Y has A->nrow rows. When using A', X has A->nrow rows and Y has A->ncol rows. If transpose = 0, then A is used; otherwise, A' is used (the complex conjugate transpose). The transpose parameter is ignored if the matrix is symmetric or Hermitian. (the array transpose A.' is not supported). Supports real, complex, and zomplex matrices, but the xtypes of A, X, and Y must all match.

22.6 cholmod_ssmult: sparse-times-sparse matrix

Purpose: Computes C = A*B; multiplying two sparse matrices. C is returned as packed, and either unsorted or sorted, depending on the sorted input parameter. If C is returned sorted, then either C = (B'*A')' or C = (A*B)' is computed, depending on the number of nonzeros in A, B, and C. The stype of C is determined by the stype parameter. Only pattern and real matrices are supported. Complex and zomplex matrices are supported only when the numerical values are not computed (values is FALSE).

22.7 cholmod_submatrix: sparse submatrix

```
cholmod_sparse *cholmod_submatrix
    /* ---- input ---- */
   cholmod_sparse *A, /* matrix to subreference */
   int32_t *rset,
                      /* set of row indices, duplicates OK */
                       /* size of r; rsize < 0 denotes ":" */</pre>
   int64_t rsize,
                       /* set of column indices, duplicates OK */
   int32_t *cset,
                       /* size of c; csize < 0 denotes ":" */</pre>
   int64_t csize,
    int values,
                       /* if TRUE compute the numerical values of C */
    int sorted,
                       /* if TRUE then return C with sorted columns */
   cholmod_common *Common
);
cholmod_sparse *cholmod_l_submatrix (cholmod_sparse *, int64_t *,
    int64_t, int64_t *, int64_t, int, int,
    cholmod_common *);
```

Purpose: Returns C = A (rset,cset), where C becomes length(rset)-by-length(cset) in dimension. rset and cset can have duplicate entries. A must be unsymmetric. C unsymmetric and is packed. If sorted is TRUE on input, or rset is sorted and A is sorted, then C is sorted; otherwise C is unsorted.

If rset is NULL, it means "[]" in MATLAB notation, the empty set. The number of rows in the result C will be zero if rset is NULL. Likewise if cset means the empty set; the number of columns in the result C will be zero if cset is NULL. If rsize or csize is negative, it denotes ":" in MATLAB notation. Thus, if both rsize and csize are negative C = A(:,:) = A is returned.

For permuting a matrix, this routine is an alternative to cholmod_ptranspose (which permutes and transposes a matrix and can work on symmetric matrices).

The time taken by this routine is O(A->nrow) if the Common workspace needs to be initialized, plus O(C->nrow + C->ncol + nnz (A (:,cset))). Thus, if C is small and the workspace is not initialized, the time can be dominated by the call to cholmod_allocate_work. However, once the workspace is allocated, subsequent calls take less time.

Only pattern and real matrices are supported. Complex and zomplex matrices are supported only when values is FALSE.

22.8 cholmod_horzcat: horizontal concatenation

Purpose: Horizontal concatenation, returns C = [A,B] in MATLAB notation. A and B can have any stype. C is returned unsymmetric and packed. A and B must have the same number of rows. C is sorted if both A and B are sorted. A and B must have the same numeric xtype, unless values is FALSE. A and B cannot be complex or zomplex, unless values is FALSE.

22.9 cholmod_vertcat: vertical concatenation

Purpose: Vertical concatenation, returns C = [A;B] in MATLAB notation. A and B can have any stype. C is returned unsymmetric and packed. A and B must have the same number of columns. C is sorted if both A and B are sorted. A and B must have the same numeric xtype, unless values is FALSE. A and B cannot be complex or zomplex, unless values is FALSE.

22.10 cholmod_symmetry: compute the symmetry of a matrix

```
int cholmod_symmetry
(
    /* ---- input ---- */
    cholmod_sparse *A,
    int option,
    /* ---- output ---- */
    int32_t *xmatched,
    int32_t *pmatched,
    int32_t *nzoffdiag,
    int32_t *nzdiag,
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_symmetry (cholmod_sparse *, int, int64_t *,
    int64_t *, int64_t *, int64_t *,
    cholmod_common *);
```

Purpose:

Determines if a sparse matrix is rectangular, unsymmetric, symmetric, skew-symmetric, or Hermitian. It does so by looking at its numerical values of both upper and lower triangular parts of a CHOLMOD "unsymmetric" matrix, where A-¿stype == 0. The transpose of A is NOT constructed.

If not unsymmetric, it also determines if the matrix has a diagonal whose entries are all real and positive (and thus a candidate for sparse Cholesky if A-¿stype is changed to a nonzero value).

Note that a Matrix Market "general" matrix is either rectangular or unsymmetric.

The row indices in the column of each matrix MUST be sorted for this function to work properly (A-¿sorted must be TRUE). This routine returns EMPTY if A-¿stype is not zero, or if A-¿sorted is FALSE. The exception to this rule is if A is rectangular.

If option == 0, then this routine returns immediately when it finds a non-positive diagonal entry (or one with nonzero imaginary part). If the matrix is not a candidate for sparse Cholesky, it returns the value CHOLMOD_MM_UNSYMMETRIC, even if the matrix might in fact be symmetric or Hermitian

This routine is useful inside the MATLAB backslash, which must look at an arbitrary matrix (A- $\dot{\xi}$ stype == 0) and determine if it is a candidate for sparse Cholesky. In that case, option should be 0.

This routine is also useful when writing a MATLAB matrix to a file in Rutherford/Boeing or Matrix Market format. Those formats require a determination as to the symmetry of the matrix, and thus this routine should not return upon encountering the first non-positive diagonal. In this case, option should be 1.

If option is 2, this function can be used to compute the numerical and pattern symmetry, where 0 is a completely unsymmetric matrix, and 1 is a perfectly symmetric matrix. This option is used when computing the following statistics for the matrices in the SuiteSparse Matrix Collection.

numerical symmetry: number of matched off-diagonal nonzeros over the total number of off-diagonal entries. A real entry a_{ij} , $i \neq j$, is matched if $a_{ji} = a_{ij}$, but this is only counted if both a_{ji}

and a_{ij} are nonzero. This does not depend on Z. (If A is complex, then the above test is modified; a_{ij} is matched if $\operatorname{conj}(a_{ji}) = a_{ij}$.

Then numeric symmetry = xmatched / nzoffdiag, or 1 if nzoffdiag = 0.

pattern symmetry: number of matched offdiagonal entries over the total number of offdiagonal entries. An entry a_{ij} , $i \neq j$, is matched if a_{ji} is also an entry.

Then pattern symmetry = pmatched / nzoffdiag, or 1 if nzoffdiag = 0.

The symmetry of a matrix with no offdiagonal entries is equal to 1.

A workspace of size nool integers is allocated; EMPTY is returned if this allocation fails.

Summary of return values:

EMPTY (-1) out of memory, stype not zero, A not sorted

CHOLMOD_MM_RECTANGULAR 1 A is rectangular CHOLMOD_MM_UNSYMMETRIC 2 A is unsymmetric

CHOLMOD_MM_SYMMETRIC 3 A is symmetric, but with non-pos. diagonal A is Hermitian, but with non-pos. diagonal

CHOLMOD_MM_SKEW_SYMMETRIC 5 A is skew symmetric

CHOLMOD_MM_SYMMETRIC_POSDIAG 6 A is symmetric with positive diagonal CHOLMOD_MM_HERMITIAN_POSDIAG 7 A is Hermitian with positive diagonal See also the spsym mexFunction, which is a MATLAB interface for this code.

If the matrix is a candidate for sparse Cholesky, it will return a result

CHOLMOD_MM_SYMMETRIC_POSDIAG if real, or CHOLMOD_MM_HERMITIAN_POSDIAG if complex. Otherwise, it will return a value less than this. This is true regardless of the value of the option parameter.

23 Supernodal Module routines

23.1 cholmod_super_symbolic: supernodal symbolic factorization

```
int cholmod_super_symbolic
    /* ---- input ---- */
   cholmod_sparse *A, /* matrix to analyze */
   cholmod_sparse *F, /* F = A' or A(:,f)' */
                     /* elimination tree */
   int32_t *Parent,
    /* ---- in/out --- */
   cholmod_factor *L, /* simplicial symbolic on input,
                        * supernodal symbolic on output */
   cholmod_common *Common
) ;
int cholmod_l_super_symbolic (cholmod_sparse *, cholmod_sparse *,
    int64_t *, cholmod_factor *, cholmod_common *);
int cholmod_super_symbolic2
    /* ---- input ---- */
   int for_whom,
                       /* FOR_SPQR
                                       (0): for SPQR but not GPU-accelerated
                          FOR_CHOLESKY (1): for Cholesky (GPU or not)
                          FOR_SPQRGPU (2): for SPQR with GPU acceleration */
   cholmod_sparse *A, /* matrix to analyze */
   cholmod_sparse *F, /* F = A' or A(:,f)' */
                       /* elimination tree */
   int32_t *Parent,
    /* ---- in/out --- */
    cholmod_factor *L, /* simplicial symbolic on input,
                        * supernodal symbolic on output */
    /* ----- */
    cholmod common *Common
);
int cholmod_l_super_symbolic2 (int, cholmod_sparse *, cholmod_sparse *,
    int64_t *, cholmod_factor *, cholmod_common *);
```

Purpose: Supernodal symbolic analysis of the LL^T factorization of A, A*A', or A(:,f)*A(:,f)'. This routine must be preceded by a simplicial symbolic analysis (cholmod_rowcolcounts). See Cholesky/cholmod_analyze.c for an example of how to use this routine. The user need not call this directly; cholmod_analyze is a "simple" wrapper for this routine. A can be symmetric (upper), or unsymmetric. The symmetric/lower form is not supported. In the unsymmetric case F is the normally transpose of A. Alternatively, if F=A(:,f)' then F*F' is analyzed. Requires Parent and L->ColCount to be defined on input; these are the simplicial Parent and ColCount arrays as computed by cholmod_rowcolcounts. Does not use L->Perm; the input matrices A and F must already be properly permuted. Allocates and computes the supernodal pattern of L (L->super, L->pi, L->px, and L->s). Does not allocate the real part (L->x).

23.2 cholmod_super_numeric: supernodal numeric factorization

Purpose: Computes the numerical Cholesky factorization of A+beta*I or A*F+beta*I. Only the lower triangular part of A+beta*I or A*F+beta*I is accessed. The matrices A and F must already be permuted according to the fill-reduction permutation L->Perm. cholmod_factorize is an "easy" wrapper for this code which applies that permutation. The input scalar beta is real; only the real part (beta[0] is used.

Symmetric case: A is a symmetric (lower) matrix. F is not accessed and may be NULL. With a fill-reducing permutation, A(p,p) should be passed for A, where is p is L->Perm.

Unsymmetric case: A is unsymmetric, and F must be present. Normally, F=A'. With a fill-reducing permutation, A(p,f) and A(p,f)' should be passed as the parameters A and F, respectively, where f is a list of the subset of the columns of A.

The input factorization L must be supernodal (L->is_super is TRUE). It can either be symbolic or numeric. In the first case, L has been analyzed by cholmod_analyze or cholmod_super_symbolic, but the matrix has not yet been numerically factorized. The numerical values are allocated here and the factorization is computed. In the second case, a prior matrix has been analyzed and numerically factorized, and a new matrix is being factorized. The numerical values of L are replaced with the new numerical factorization.

 $L->is_11$ is ignored on input, and set to TRUE on output. This routine always computes an LL^T factorization. Supernodal LDL^T factorization is not supported.

If the matrix is not positive definite the routine returns TRUE, but sets Common->status to $CHOLMOD_NOT_POSDEF$ and L->minor is set to the column at which the failure occurred. Columns L->minor to L->n-1 are set to zero.

If L is supernodal symbolic on input, it is converted to a supernodal numeric factor on output, with an xtype of real if A is real, or complex if A is complex or zomplex. If L is supernodal numeric on input, its xtype must match A (except that L can be complex and A zomplex). The xtype of A and A must match.

23.3 cholmod_super_lsolve: supernodal forward solve

```
int cholmod_super_lsolve
(
    /* ---- input ---- */
    cholmod_factor *L, /* factor to use for the forward solve */
    /* ---- output ---- */
    cholmod_dense *X, /* b on input, solution to Lx=b on output */
    /* ---- workspace */
    cholmod_dense *E, /* workspace of size nrhs*(L->maxesize) */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_super_lsolve (cholmod_factor *, cholmod_dense *, cholmod_dense *,
    cholmod_common *);
```

Purpose: Solve Lx = b for a supernodal factorization. This routine does not apply the permutation L->Perm. See cholmod_solve for a more general interface that performs that operation. Only real and complex xtypes are supported. L, X, and E must have the same xtype.

23.4 cholmod_super_ltsolve: supernodal backsolve

Purpose: Solve $\mathbf{L}^\mathsf{T}\mathbf{x} = \mathbf{b}$ for a supernodal factorization. This routine does not apply the permutation L->Perm. See cholmod_solve for a more general interface that performs that operation. Only real and complex xtypes are supported. L, X, and E must have the same xtype.

24 Partition Module routines

24.1 cholmod_nested_dissection: nested dissection ordering

```
int64_t cholmod_nested_dissection
                                        /* returns # of components */
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to order */
    int32_t *fset,
                        /* subset of 0:(A->ncol)-1 */
    size_t fsize,
                        /* size of fset */
    /* ---- output --- */
    int32_t *Perm,
                        /* size A->nrow, output permutation */
    int32_t *CParent,
                        /* size A->nrow. On output, CParent [c] is the parent
                        * of component c, or EMPTY if c is a root, and where
                         * c is in the range 0 to # of components minus 1 */
                        /* size A->nrow. Cmember [j] = c if node j of A is
    int32_t *Cmember,
                         * in component c */
    cholmod_common *Common
);
int64_t cholmod_l_nested_dissection (cholmod_sparse *,
    int64_t *, size_t, int64_t *, int64_t *,
    int64_t *, cholmod_common *);
```

Purpose: CHOLMOD's nested dissection algorithm: using its own compression and connected-components algorithms, an external graph partitioner (METIS), and a constrained minimum degree ordering algorithm (CAMD, CCOLAMD, or CSYMAMD). Typically gives better orderings than METIS_NodeND (about 5% to 10% fewer nonzeros in L).

This method uses a node bisector, applied recursively (but using a non-recursive implementation). Once the graph is partitioned, it calls a constrained minimum degree code (CAMD or CSYMAMD for A+A', and CCOLAMD for A*A') to order all the nodes in the graph - but obeying the constraints determined by the separators. This routine is similar to METIS_NodeND, except for how it treats the leaf nodes. METIS_NodeND orders the leaves of the separator tree with MMD, ignoring the rest of the matrix when ordering a single leaf. This routine orders the whole matrix with CAMD, CSYMAMD, or CCOLAMD, all at once, when the graph partitioning is done.

24.2 cholmod_metis: interface to METIS nested dissection

```
int cholmod_metis
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to order */
   int32_t *fset,
                    /* subset of 0:(A->ncol)-1 */
   size_t fsize,
                     /* size of fset */
   int postorder,
                     /* if TRUE, follow with etree or coletree postorder */
   /* ---- output --- */
                  /* size A->nrow, output permutation */
   int32_t *Perm,
   /* ----- */
   cholmod_common *Common
);
int cholmod_l_metis (cholmod_sparse *, int64_t *, size_t, int,
   int64_t *, cholmod_common *);
```

Purpose: CHOLMOD wrapper for the METIS_NodeND ordering routine. Creates A+A', A*A' or A(:,f)*A(:,f)' and then calls METIS_NodeND on the resulting graph. This routine is comparable to cholmod_nested_dissection, except that it calls METIS_NodeND directly, and it does not return the separator tree.

24.3 cholmod_camd: interface to CAMD

Purpose: CHOLMOD interface to the CAMD ordering routine. Finds a permutation p such that the Cholesky factorization of A(p,p) is sparser than A. If A is unsymmetric, A*A' is ordered. If Cmember[i]=c then node i is in set c. All nodes in set 0 are ordered first, followed by all nodes in set 1, and so on.

24.4 cholmod_ccolamd: interface to CCOLAMD

```
int cholmod_ccolamd
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to order */
   int32_t *fset,
                       /* subset of 0:(A->ncol)-1 */
   size_t fsize,
                       /* size of fset */
   int32_t *Cmember,
                       /* size A->nrow. Cmember [i] = c if row i is in the
                        * constraint set c. c must be >= 0. The # of
                         * constraint sets is max (Cmember) + 1. If Cmember is
                        * NULL, then it is interpretted as Cmember [i] = 0 for
                         * all i */
   /* ---- output --- */
   int32_t *Perm,
                        /* size A->nrow, output permutation */
   cholmod_common *Common
);
int cholmod_l_ccolamd (cholmod_sparse *, int64_t *, size_t,
   int64_t *, int64_t *, cholmod_common *);
```

Purpose: CHOLMOD interface to the CCOLAMD ordering routine. Finds a permutation p such that the Cholesky factorization of A(p,:)*A(p,:)' is sparser than A*A'. The column elimination is found and postordered, and the CCOLAMD ordering is then combined with its postordering. A must be unsymmetric. If Cmember[i]=c then node i is in set c. All nodes in set 0 are ordered first, followed by all nodes in set 1, and so on.

24.5 cholmod_csymamd: interface to CSYMAMD

```
int cholmod_csymamd
(
    /* ---- input ---- */
    cholmod_sparse *A, /* matrix to order */
    /* ---- output --- */
    int32_t *Cmember, /* size nrow. see cholmod_ccolamd above */
    int32_t *Perm, /* size A->nrow, output permutation */
    /* ------ */
    cholmod_common *Common
);
int cholmod_l_csymamd (cholmod_sparse *, int64_t *,
    int64_t *, cholmod_common *);
```

Purpose: CHOLMOD interface to the CSYMAMD ordering routine. Finds a permutation p such that the Cholesky factorization of A(p,p) is sparser than A. The elimination tree is found and postordered, and the CSYMAMD ordering is then combined with its postordering. If A is unsymmetric, A+A' is ordered (A must be square). If Cmember[i]=c then node i is in set c. All nodes in set 0 are ordered first, followed by all nodes in set 1, and so on.

24.6 cholmod_bisect: graph bisector

```
int64_t cholmod_bisect /* returns # of nodes in separator */
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to bisect */
                       /* subset of 0:(A->ncol)-1 */
   int32_t *fset,
   size_t fsize,
                       /* size of fset */
                       /* if TRUE, compress the graph first */
   int compress,
   /* ---- output --- */
   int32_t *Partition, /* size A->nrow. Node i is in the left graph if
                        * Partition [i] = 0, the right graph if 1, and in the
                        * separator if 2. */
   /* ----- */
   cholmod_common *Common
);
int64_t cholmod_l_bisect (cholmod_sparse *, int64_t *,
   size_t, int, int64_t *, cholmod_common *);
```

Purpose: Finds a node bisector of A, A*A', A(:,f)*A(:,f)': a set of nodes that partitions the graph into two parts. Compresses the graph first, ensures the graph is symmetric with no diagonal entries, and then calls METIS.

24.7 cholmod_metis_bisector: interface to METIS node bisector

```
int64_t cholmod_metis_bisector /* returns separator size */
   /* ---- input ---- */
   cholmod_sparse *A, /* matrix to bisect */
                       /* size A->nrow, node weights, can be NULL, */
   int32_t *Anw,
                       /* which means the graph is unweighted. */
   int32_t *Aew,
                       /* size nz, edge weights (silently ignored). */
                       /* This option was available with METIS 4, but not */
                       /* in METIS 5. This argument is now unused, but */
                       /* it remains for backward compatibilty, so as not */
                       /* to change the API for cholmod_metis_bisector. */
   /* ---- output --- */
   int32_t *Partition, /* size A->nrow */
   /* ----- */
   cholmod_common *Common
);
int64_t cholmod_l_metis_bisector (cholmod_sparse *,
   int64_t *, int64_t *, int64_t *,
   cholmod_common *);
```

Purpose: Finds a set of nodes that bisects the graph of A or A*A' (a direct interface to METIS_NodeComputeSeparator).

The input matrix A must be square, symmetric (with both upper and lower parts present) and with no diagonal entries. These conditions are not checked. Use cholmod_bisect to check these conditions.

24.8 cholmod_collapse_septree: prune a separator tree

Purpose: Prunes a separator tree obtained from cholmod_nested_dissection.

References

- [1] P. R. Amestoy, T. A. Davis, and I. S. Duff. An approximate minimum degree ordering algorithm. SIAM J. Matrix Anal. Applic., 17(4):886–905, 1996.
- [2] P. R. Amestoy, T. A. Davis, and I. S. Duff. Algorithm 837: AMD, an approximate minimum degree ordering algorithm. *ACM Trans. Math. Softw.*, 30(3):381–388, 2004.
- [3] E. Anderson, Z. Bai, C. Bischof, S. Blackford, J. Demmel, J. Dongarra, J. Du Croz, A. Green-baum, S. Hammarling, A. McKenny, and D. Sorensen. *LAPACK Users' Guide*, 3rd ed. SIAM, 1999.
- [4] Y. Chen, T. A. Davis, W. W. Hager, and S. Rajamanickam. Algorithm 8xx: CHOLMOD, supernodal sparse Cholesky factorization and update/downdate. *ACM Trans. Math. Softw.*, submitted in 2006.
- [5] T. A. Davis. Algorithm 849: A concise sparse Cholesky algorithm. *ACM Trans. Math. Softw.*, 31(4):587–591, 2005.
- [6] T. A. Davis, J. R. Gilbert, S. I. Larimore, and E. G. Ng. Algorithm 836: COLAMD, a column approximate minimum degree ordering algorithm. ACM Trans. Math. Softw., 30(3):377–380, 2004.
- [7] T. A. Davis, J. R. Gilbert, S. I. Larimore, and E. G. Ng. A column approximate minimum degree ordering algorithm. *ACM Trans. Math. Softw.*, 30(3):353–376, 2004.
- [8] T. A. Davis and W. W. Hager. Modifying a sparse Cholesky factorization. SIAM J. Matrix Anal. Applic., 20(3):606–627, 1999.
- [9] T. A. Davis and W. W. Hager. Multiple-rank modifications of a sparse Cholesky factorization. SIAM J. Matrix Anal. Applic., 22(4):997–1013, 2001.
- [10] T. A. Davis and W. W. Hager. Row modifications of a sparse Cholesky factorization. SIAM J. Matrix Anal. Applic., 26(3):621–639, 2005.
- [11] T. A. Davis and W. W. Hager. Dynamic supernodes in sparse Cholesky update/downdate and triangular solves. *ACM Trans. Math. Softw.*, submitted in 2006.
- [12] J. J. Dongarra, J. Du Croz, I. S. Duff, and S. Hammarling. A set of level-3 basic linear algebra subprograms. *ACM Trans. Math. Softw.*, 16(1):1–17, 1990.
- [13] J. R. Gilbert, X. S. Li, E. G. Ng, and B. W. Peyton. Computing row and column counts for sparse QR and LU factorization. *BIT*, 41(4):693–710, 2001.
- [14] J. R. Gilbert, C. Moler, and R. Schreiber. Sparse matrices in MATLAB: design and implementation. SIAM J. Matrix Anal. Applic., 13(1):333–356, 1992.
- [15] J. R. Gilbert, E. G. Ng, and B. W. Peyton. An efficient algorithm to compute row and column counts for sparse Cholesky factorization. SIAM J. Matrix Anal. Applic., 15(4):1075–1091, 1994.

- [16] N. I. M. Gould, Y. Hu, and J. A. Scott. Complete results from a numerical evaluation of sparse direct solvers for the solution of large sparse, symmetric linear systems of equations. Technical Report Internal report 2005-1 (revision 1), CCLRC, Rutherford Appleton Laboratory, 2005.
- [17] N. I. M. Gould, Y. Hu, and J. A. Scott. A numerical evaluation of sparse direct solvers for the solution of large sparse, symmetric linear systems of equations. *ACM Trans. Math. Softw.*, to appear.
- [18] G. Karypis and V. Kumar. A fast and high quality multilevel scheme for partitioning irregular graphs. SIAM J. Sci. Comput., 20(1):359–392, 1998.
- [19] J. W. H. Liu. A compact row storage scheme for Cholesky factors using elimination trees. *ACM Trans. Math. Softw.*, 12(2):127–148, 1986.
- [20] J. W. H. Liu. The role of elimination trees in sparse factorization. SIAM J. Matrix Anal. Applic., 11(1):134–172, 1990.
- [21] E. Ng and B. Peyton. Block sparse Cholesky algorithms on advanced uniprocessor computers. SIAM J. Sci. Comput., 14:1034–1056, 1993.