User Guide for SuiteSparse:GraphBLAS

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Abstract

SuiteSparse:GraphBLAS is a full implementation of the Graph-BLAS standard, which defines a set of sparse matrix operations on an extended algebra of semirings using an almost unlimited variety of operators and types. When applied to sparse adjacency matrices, these algebraic operations are equivalent to computations on graphs. GraphBLAS provides a powerful and expressive framework for creating graph algorithms based on the elegant mathematics of sparse matrix operations on a semiring.

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1 Introduction

The GraphBLAS standard defines sparse matrix and vector operations on an extended algebra of semirings. The operations are useful for creating a wide range of graph algorithms.

For example, consider the matrix-matrix multiplication, $\mathbf{C} = \mathbf{AB}$. Suppose \mathbf{A} and \mathbf{B} are sparse n-by-n Boolean adjacency matrices of two undirected graphs. If the matrix multiplication is redefined to use logical AND instead of scalar multiply, and if it uses the logical OR instead of add, then the matrix \mathbf{C} is the sparse Boolean adjacency matrix of a graph that has an edge (i,j) if node i in \mathbf{A} and node j in \mathbf{B} share any neighbor in common. The OR-AND pair forms an algebraic semiring, and many graph operations like this one can be succinctly represented by matrix operations with different semirings and different numerical types. GraphBLAS provides a wide range of built-in types and operators, and allows the user application to create new types and operators without needing to recompile the GraphBLAS library.

For more details on SuiteSparse:GraphBLAS, and its use in LAGraph, see [Dav19, Dav18, DAK19, ACD+20, MDK+19].

A full and precise definition of the GraphBLAS specification is provided in *The GraphBLAS C API Specification* by Aydın Buluç, Timothy Mattson, Scott McMillan, José Moreira, and Carl Yang [BMM+17a, BMM+17b], based on *GraphBLAS Mathematics* by Jeremy Kepner [Kep17]. The GraphBLAS C API Specification is available at http://graphblas.org. This version of Suite-Sparse:GraphBLAS fully conforms to Version 1.2.0 (May 18, 2018) of that specification. In this User Guide, aspects of the GraphBLAS specification that would be true for any GraphBLAS implementation are simply called "GraphBLAS." Details unique to this particular implementation are referred to as SuiteSparse:GraphBLAS.

SPEC: See the tag **SPEC:** for SuiteSparse extensions to the spec. They are also placed in text boxes like this one. All functions, objects, and macros with a name of the form <code>GxB_*</code> are extensions to the spec.

1.1 Release Notes:

• Version 3.3.0 (TODO, 2020). Added complex types (GxB_FC32 and GxB_FC64), many unary operators, binary operators, monoids, and semirings. Added bitwise operators, and their monoids and semirings. Added the predefined monoids and semirings from the v1.3 spec. MATLAB interface: added complex matrices and operators, and changed behavior of integer operations to more closely match the behavior on MATLAB integer matrices. The rules for typecasting large floating point values to integers has changed. The specific object-based GrB_Matrix_wait and related functions have been added

The no-argument GrB_wait() has been deprecated.

Added GrB_getVersion, GrB_Matrix_resize, GrB_Vector_resize, GrB_kronecker, GrB_*_wait, ...

- Version 3.2.0 (Feb 20, 2020). Faster GrB_mxm, GrB_mxv, and GrB_vxm, and faster operations on dense matrices/vectors. Removed compiletime user objects (GxB_*_define), since these were not compatible with the faster matrix operations. Added the ANY and PAIR operators. Added the predefined descriptor, GrB_DESC_*. Added the structural mask option. Changed default chunk size to 65,536. Note that v3.2.0 is not compatible with the MS Visual Studio compiler; use v3.1.2 instead. MATLAB interface modified: GrB.init is now optional.
- Version 3.1.2 (Dec, 2019). Changes to allow SuiteSparse:GraphBLAS to be compiled with the Microsoft Visual Studio compiler. This compiler does not support the _Generic keyword, so the polymorphic functions are not available. Use the equivalent non-polymorphic functions instead, when compiling GraphBLAS with MS Visual Studio. In addition, variable-length arrays are not supported, so user-defined types are limited to 128 bytes in size. These changes have no effect if you have an ANSI C11 compliant compiler.

MATLAB interface modified: GrB.init is now required.

 Version 3.1.0 (Oct 1, 2019). MATLAB interface added. See the GraphBLAS/GraphBLAS folder for details and documentation, and Section 3.1. • Version 3.0 (July 26, 2019), with OpenMP parallelism.

The version number is increased to 3.0, since this version is not backward compatible with V2.x. The GxB_select operation changes; the Thunk parameter was formerly a const void * pointer, and is now a GxB_Scalar. A new parameter is added to GxB_SelectOp_new, to define the expected type of Thunk. A new parameter is added to GxB_init, to specify whether or not the user-provided memory management functions are thread safe.

The remaining changes add new features, and are upward compatible with V2.x. The major change is the addition of OpenMP parallelism. This addition has no effect on the API, except that round-off errors can differ with the number of threads used, for floating-point types. GxB_set can optionally define the number of threads to use (the default is $omp_get_max_threads$). The number of threads can also defined globally, and/or in the $GrB_Descriptor$. The RDIV and RMINUS operators are added, which are defined as f(x,y) = y/x and f(x,y) = y-x, respectively. Additional options are added to GxB_get .

- Version 2.3.3 (May 2019): Collected Algorithm of the ACM. No changes from V2.3.2 other than the documentation.
- Version 2.3 (Feb 2019) improves the performance of many GraphBLAS operations, including an early-exit for monoids. These changes have a significant impact on breadth-first-search (a performance bug was also fixed in the two BFS Demo codes). The matrix and vector import/export functions were added (Section 5.10), in support of the new LAGraph project (https://github.com/GraphBLAS/LAGraph, see also Section 11.1). LAGraph includes a push-pull BFS in GraphBLAS that is faster than two versions in the Demo folder. GxB_init was added to allow the memory manager functions (malloc, etc) to be specified.
- Version 2.2 (Nov 2018) adds user-defined objects at compile-time, via user *.m4 files placed in GraphBLAS/User, which use the GxB_*_define macros described in Section 10 (NOTE: feature removed in v3.2). The default matrix format is now GxB_BY_ROW. Also added are the GxB_*print methods for printing the contents of each GraphBLAS object (Section 9). PageRank demos have been added to the Demos folder. Prior versions required GraphBLAS to be compiled with OpenMP, for it to be

thread-safe. It can now be compiled with POSIX pthreads. The cmake script automatically detects if OpenMP and/or POSIX pthreads are available. Demos have been added to show how GraphBLAS can be called from a multi-threaded user application.

- Version 2.1 (Oct 2018) was a major update with support for new matrix formats (by row or column, and hypersparse matrices), and MATLAB-like colon notation (I=begin:end or I=begin:inc:end). Some graph algorithms are more naturally expressed with matrices stored by row, and this version includes the new GxB_BY_ROW format. The default format in Version 2.1 and prior versions is by column. New extensions to GraphBLAS in this version include GxB_get, GxB_set, and GxB_AxB_METHOD, GxB_RANGE, GxB_STRIDE, and GxB_BACKWARDS, and their related definitions, described in Sections 5.11, 6, and 7.
- Version 2.0 (March 2018) addressed changes in the GraphBLAS C API Specification and added GxB_kron and GxB_resize.
- Version 1.1 (Dec 2017) primarily improved the performance.
- Version 1.0 was released on Nov 25, 2017.

2 Basic Concepts

Since the *GraphBLAS C API Specification* provides a precise definition of GraphBLAS, not every detail of every function is provided here. For example, some error codes returned by GraphBLAS are self-explanatory, but since a specification must precisely define all possible error codes a function can return, these are listed in detail in the *GraphBLAS C API Specification*. However, including them here is not essential and the additional information on the page might detract from a clearer view of the essential features of the GraphBLAS functions.

This User Guide also assumes the reader is familiar with the MATLAB language, created by Cleve Moler. MATLAB supports only the conventional plus-times semiring on sparse double and complex matrices, but a MATLAB-like notation easily extends to the arbitrary semirings used in GraphBLAS. The matrix multiplication in the example in the Introduction can be written in MATLAB notation as C=A*B, if the Boolean OR-AND semiring is understood. Relying on a MATLAB-like notation allows the description in this User Guide to be expressive, easy to understand, and terse at the same time. The GraphBLAS C API Specification also makes use of some MATLAB-like language, such as the colon notation.

MATLAB notation will always appear here in fixed-width font, such as C=A*B(:,j). In standard mathematical notation it would be written as the matrix-vector multiplication $C=Ab_j$ where b_j is the jth column of the matrix B. The GraphBLAS standard is a C API and SuiteSparse:GraphBLAS is written in C, and so a great deal of C syntax appears here as well, also in fixed-width font. This User Guide alternates between all three styles as needed.

2.1 Graphs and sparse matrices

Graphs can be huge, with many nodes and edges. A dense adjacency matrix \mathbf{A} for a graph of n nodes takes $O(n^2)$ memory, which is impossible if n is, say, a million. Most graphs arising in practice are sparse, however, with only $|\mathbf{A}| = O(n)$ edges, where $|\mathbf{A}|$ denotes the number of edges in the graph, or the number of explicit entries present in the data structure for the matrix \mathbf{A} . Sparse graphs with millions of nodes and edges can easily be created by representing them as sparse matrices, where only explicit values need to be stored. Some graphs are *hypersparse*, with $|\mathbf{A}| << n$. SuiteSparse:GraphBLAS sup-

ports two kinds of sparse matrix formats: a regular sparse format, taking $O(n + |\mathbf{A}|)$ space, and a hypersparse format taking only $O(|\mathbf{A}|)$ space. As a result, creating a sparse matrix of size *n*-by-*n* where $n = 2^{60}$ (about 10^{18}) can be done on quite easily on a commodity laptop, limited only by $|\mathbf{A}|$.

A sparse matrix data structure only stores a subset of the possible n^2 entries, and it assumes the values of entries not stored have some implicit value. In conventional linear algebra, this implicit value is zero, but it differs with different semirings. Explicit values are called *entries* and they appear in the data structure. The *pattern* of a matrix defines where its explicit entries appear. It will be referenced in one of two equivalent ways. It can be viewed as a set of indices (i, j), where (i, j) is in the pattern of a matrix \mathbf{A} if $\mathbf{A}(i, j)$ is an explicit value. It can also be viewed as a Boolean matrix \mathbf{S} where $\mathbf{S}(i, j)$ is true if (i, j) is an explicit entry and false otherwise. In MATLAB notation, $\mathbf{S}=\mathbf{spones}(\mathbf{A})$ or $\mathbf{S}=(\mathbf{A}^{\sim}=\mathbf{0})$, if the implicit value is zero. The (\mathbf{i},\mathbf{j}) pairs, and their values, can also be extracted from the matrix via the MATLAB expression $[\mathbf{I},\mathbf{J},\mathbf{X}]=\mathbf{find}(\mathbf{A})$, where the kth tuple $(\mathbf{I}(\mathbf{k}),\mathbf{J}(\mathbf{k}),\mathbf{X}(\mathbf{k}))$ represents the explicit entry $\mathbf{A}(\mathbf{I}(\mathbf{k}),\mathbf{J}(\mathbf{k}))$, with numerical value $\mathbf{X}(\mathbf{k})$ equal to a_{ij} , with row index $i=\mathbf{I}(\mathbf{k})$ and column index $j=\mathbf{J}(\mathbf{k})$.

The entries in the pattern of A can take on any value, including the implicit value, whatever it happens to be. This differs slightly from MAT-LAB, which always drops all explicit zeros from its sparse matrices. This is a minor difference but it cannot be done in GraphBLAS. For example, in the max-plus tropical algebra, the implicit value is negative infinity, and zero has a different meaning. Here, the MATLAB notation used will assume that no explicit entries are ever dropped because their explicit value happens to match the implicit value.

Graph Algorithms in the Language on Linear Algebra, Kepner and Gilbert, eds., provides a framework for understanding how graph algorithms can be expressed as matrix computations [KG11]. For additional background on sparse matrix algorithms, see also [Dav06] and [DRSL16].

2.2 Overview of GraphBLAS methods and operations

GraphBLAS provides a collection of *methods* to create, query, and free its of objects: sparse matrices, sparse vectors, sparse scalars, types, operators, monoids, semirings, and a descriptor object used for parameter settings. Details are given in Section 5. Once these objects are created they can be used in mathematical *operations* (not to be confused with the how the term *oper-*

ator is used in GraphBLAS). A short summary of these operations and their nearest MATLAB analog is given in the table below.

operation	approximate MATLAB analog
matrix multiplication	C=A*B
element-wise operations	C=A+B and C=A.*B
reduction to a vector or scalar	s=sum(A)
apply unary operator	C=-A
transpose	C=A'
submatrix extraction	C=A(I,J)
submatrix assignment	C(I,J)=A

GraphBLAS can do far more than what MATLAB can do in these rough analogs, but the list provides a first step in describing what GraphBLAS can do. Details of each GraphBLAS operation are given in Section 8. With this brief overview, the full scope of GraphBLAS extensions of these operations can now be described.

GraphBLAS has 11 built-in scalar types: Boolean, single and double precision floating-point, and 8, 16, 32, and 64-bit signed and unsigned integers. In addition, user-defined scalar types can be created from nearly any C typedef, as long as the entire type fits in a fixed-size contiguous block of memory (of arbitrary size). All of these types can be used to create Graph-BLAS sparse matrices, vectors, or scalars.

The scalar addition of conventional matrix multiplication is replaced with a monoid. A monoid is an associative and commutative binary operator z=f(x,y) where all three domains are the same (the types of x, y, and z), and where the operator has an identity value id such that f(x,id)=f(id,x)=x. Performing matrix multiplication with a semiring uses a monoid in place of the "add" operator, scalar addition being just one of many possible monoids. The identity value of addition is zero, since x+0=0+x=x. GraphBLAS includes eight built-in operators suitable for use as a monoid: min (with an identity value of positive infinity), max (whose identity is negative infinity), add (identity is zero) multiply (with an identity of one), and four logical operators: AND, OR, exclusive-OR, and Boolean equality. User-created monoids can be defined with any associative and commutative operator that has an identity value.

Finally, a semiring can use any built-in or user-defined binary operator z=f(x,y) as its "multiply" operator, as long as the type of its output, z matches the type of the semiring's monoid. The user application can create

any semiring based on any types, monoids, and multiply operators, as long these few rules are followed.

Just considering built-in types and operators, GraphBLAS can perform C=A*B in 2,291 unique semirings. With typecasting, any of these 2,291 semirings can be applied to matrices C, A, and B of any of the 11 types, in any combination. This gives $2291 \times 13^3 = 5,033,327$ possible kinds of sparse matrix multiplication supported by GraphBLAS, and this is counting just built-in types and operators. By contrast, MATLAB provides just two semirings for its sparse matrix multiplication C=A*B: plus-times-double and plus-times-complex, not counting the typecasting that MATLAB does when multiplying a real matrix times a complex matrix. All of the 1.4 million forms of matrix multiplication methods in SuiteSparse:GraphBLAS are typically just as fast as computing C=A*B in MATLAB using its own native sparse matrix multiplication methods, and often faster when parallelism can be effectively used.

A monoid can also be used in a reduction operation, like s=sum(A) in MATLAB. MATLAB provides the plus, times, min, and max reductions of a real or complex sparse matrix as s=sum(A), s=prod(A), s=min(A), and s=max(A), respectively. In GraphBLAS, any monoid can be used (min, max, plus, times, AND, OR, exclusive-OR, equality, or any user-defined monoid, on any user-defined type).

Element-wise operations are also expanded from what can be done in MATLAB. Consider matrix addition, C=A+B in MATLAB. The pattern of the result is the set union of the pattern of A and B. In GraphBLAS, any binary operator can be used in this set-union "addition." The operator is applied to entries in the intersection. Entries in A but not B, or visa-versa, are copied directly into C, without any application of the binary operator. The accumulator operation for $\mathbf{Z} = \mathbf{C} \odot \mathbf{T}$ described in Section 2.3 is one example of this set-union application of an arbitrary binary operator.

Consider element-wise multiplication, C=A.*B in MATLAB. The operator (multiply in this case) is applied to entries in the set intersection, and the pattern of C just this set intersection. Entries in A but not B, or visa-versa, do not appear in C. In GraphBLAS, any binary operator can be used in this manner, not just scalar multiplication. The difference between element-wise "add" and "multiply" is not the operators, but whether or not the pattern of the result is the set union or the set intersection. In both cases, the operator is only applied to the set intersection.

Finally, GraphBLAS includes a non-blocking mode where operations can

be left pending, and saved for later. This is very useful for submatrix assignment (C(I,J)=A where I and J are integer vectors), or scalar assignment (C(i,j)=x where i and j are scalar integers). Because of how MATLAB stores its matrices, adding and deleting individual entries is very costly. For example, this is very slow in MATLAB, taking $O(nz^2)$ time:

```
A = sparse (m,n);  % an empty sparse matrix
for k = 1:nz
    compute a value x, row index i, and column index j
    A (i,j) = x;
end
```

The above code is very easy read and simple to write, but exceedingly slow. In MATLAB, the method below is preferred and is far faster, taking at most $O(|\mathbf{A}| \log |\mathbf{A}| + n)$ time. It can easily be a million times faster than the method above. Unfortunately the second method below is a little harder to read and a little less natural to write:

GraphBLAS can do both methods. SuiteSparse:GraphBLAS stores its matrices in a format that allows for pending computations, which are done later in bulk, and as a result it can do both methods above equally as fast as the MATLAB sparse function, allowing the user to write simpler code.

2.3 The accumulator and the mask

Most GraphBLAS operations can be modified via transposing input matrices, using an accumulator operator, applying a mask or its complement, and by clearing all entries the matrix C after using it in the accumulator operator but before the final results are written back into it. All of these steps are optional, and are controlled by a descriptor object that holds parameter settings (see Section 5.11) that control the following options:

- the input matrices A and/or B can be transposed first.
- an accumulator operator can be used, like the plus in the statement C=C+A*B. The accumulator operator can be any binary operator, and an element-wise "add" (set union) is performed using the operator.
- an optional mask can be used to selectively write the results to the output. The mask is a sparse Boolean matrix Mask whose size is the same size as the result. If Mask(i,j) is true, then the corresponding entry in the output can be modified by the computation. If Mask(i,j) is false, then the corresponding in the output is protected and cannot be modified by the computation. The Mask matrix acts exactly like logical matrix indexing in MATLAB, with one minor difference: in GraphBLAS notation, the mask operation is $C\langle M \rangle = Z$, where the mask M appears only on the left-hand side. In MATLAB, it would appear on both sides as C(Mask) = Z(Mask). If no mask is provided, the Mask matrix is implicitly all true. This is indicated by passing the value GrB_NULL in place of the Mask argument in GraphBLAS operations.

This process can be described in mathematical notation as:

```
\mathbf{A} = \mathbf{A}^{\mathsf{T}}, if requested via descriptor (first input option)

\mathbf{B} = \mathbf{B}^{\mathsf{T}}, if requested via descriptor (second input option)
```

T is computed according to the specific operation

 $\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot\mathbf{T}$, accumulating and writing the results back via the mask

The application of the mask and the accumulator operator is written as $\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot\mathbf{T}$ where $\mathbf{Z} = \mathbf{C}\odot\mathbf{T}$ denotes the application of the accumulator operator, and $\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{Z}$ denotes the mask operator via the Boolean matrix \mathbf{M} . The Accumulator Phase, $\mathbf{Z} = \mathbf{C}\odot\mathbf{T}$, is performed as follows:

```
 \begin{aligned} \textbf{Accumulator Phase} \colon & \text{compute } \mathbf{Z} = \mathbf{C} \odot \mathbf{T} ; \\ & \text{if accum is NULL} \\ & \mathbf{Z} = \mathbf{T} \\ & \text{else} \\ & \mathbf{Z} = \mathbf{C} \odot \mathbf{T} \end{aligned}
```

The accumulator operator is \odot in GraphBLAS notation, or accum in the code. The pattern of $\mathbf{C} \odot \mathbf{T}$ is the set union of the patterns of \mathbf{C} and \mathbf{T} , and the operator is applied only on the set intersection of \mathbf{C} and \mathbf{T} . Entries in neither the pattern of \mathbf{C} nor \mathbf{T} do not appear in the pattern of \mathbf{Z} . That is:

```
for all entries (i, j) in \mathbf{C} \cap \mathbf{T} (that is, entries in both \mathbf{C} and \mathbf{T})
z_{ij} = c_{ij} \odot t_{ij}
for all entries (i, j) in \mathbf{C} \setminus \mathbf{T} (that is, entries in \mathbf{C} but not \mathbf{T})
z_{ij} = c_{ij}
for all entries (i, j) in \mathbf{T} \setminus \mathbf{C} (that is, entries in \mathbf{T} but not \mathbf{C})
z_{ij} = t_{ij}
```

The Accumulator Phase is followed by the Mask/Replace Phase, $\mathbf{C}\langle\mathbf{M}\rangle=\mathbf{Z}$ as controlled by the $\mathtt{GrB_REPLACE}$ and $\mathtt{GrB_COMP}$ descriptor options:

```
\begin{aligned} \mathbf{Mask/Replace\ Phase} &: \ \mathbf{compute\ C}\langle \mathbf{M} \rangle = \mathbf{Z}; \\ & \ \mathrm{if\ (GrB\_REPLACE)\ delete\ all\ entries\ in\ C} \\ & \ \mathrm{if\ Mask\ is\ NULL} \\ & \ \mathrm{if\ (GrB\_COMP)} \\ & \ C\ \mathrm{is\ not\ modified} \\ & \ \mathrm{else} \\ & \ C = \mathbf{Z} \\ & \ \mathrm{else} \\ & \ \mathrm{if\ (GrB\_COMP)} \\ & \ C\langle \neg \mathbf{M} \rangle = \mathbf{Z} \\ & \ \mathrm{else} \\ & \ C\langle \mathbf{M} \rangle = \mathbf{Z} \end{aligned}
```

Both phases of the accum/mask process are illustrated in MATLAB notation in Figure 1. A GraphBLAS operation starts with its primary computation, producing a result T; for matrix multiply, T=A*B, or if A is transposed first, T=A'*B, for example. Applying the accumulator, mask (or its complement) to obtain the final result matrix C can be expressed in the MATLAB accum_mask function shown in the figure. This function is an exact, fully functional, and nearly-complete description of the GraphBLAS accumulator/mask operation. The only aspects it does not consider are typecasting (see Section 2.4), and the value of the implicit identity (for those, see another version in the Test folder).

One aspect of GraphBLAS cannot be as easily expressed in a MATLAB sparse matrix: namely, what is the implicit value of entries not in the pattern? To accommodate this difference in the accum_mask MATLAB function, each sparse matrix A is represented with its values A.matrix and its pattern, A.pattern. The latter could be expressed as the sparse matrix A.pattern=spones(A) or A.pattern=(A~=0) in MATLAB, if the implicit

```
function C = accum_mask (C, Mask, accum, T, C_replace, Mask_complement)
[m n] = size (C.matrix) ;
Z.matrix = zeros (m, n) ;
Z.pattern = false (m, n) ;
if (isempty (accum))
   Z = T;
               % no accum operator
  % Z = accum (C,T), like Z=C+T but with an binary operator, accum
  p = C.pattern & T.pattern; Z.matrix (p) = accum (C.matrix (p), T.matrix (p));
  p = C.pattern & ~T.pattern; Z.matrix (p) = C.matrix (p);
  p = ~C.pattern & T.pattern ; Z.matrix (p) = T.matrix (p) ;
  Z.pattern = C.pattern | T.pattern ;
end
% = 1000 apply the mask to the values and pattern
C.matrix = mask (C.matrix, Mask, Z.matrix, C_replace, Mask_complement);
C.pattern = mask (C.pattern, Mask, Z.pattern, C_replace, Mask_complement) ;
end
function C = mask (C, Mask, Z, C_replace, Mask_complement)
% replace C if requested
if (C_replace)
   C(:,:) = 0;
if (isempty (Mask))
                                % if empty, Mask is implicit ones(m,n)
  % implicitly, Mask = ones (size (C))
   if (~Mask_complement)
     C = Z;
                                % this is the default
   else
                                % Z need never have been computed
     C = C;
   end
else
   % apply the mask
   if (~Mask_complement)
     C (Mask) = Z (Mask);
   else
     C (^{\sim}Mask) = Z (^{\sim}Mask);
   end
end
end
```

Figure 1: Applying the mask and accumulator, $\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{C} \odot \mathbf{T}$

value is zero. With different semirings, entries not in the pattern can be 1, +Inf, -Inf, or whatever is the identity value of the monoid. As a result, Figure 1 performs its computations on two MATLAB matrices: the values in A.matrix and the pattern in the logical matrix A.pattern. Implicit values are untouched.

The final computation in Figure 1 with a complemented Mask is easily expressed in MATLAB as C(~Mask)=Z(~Mask) but this is costly if Mask is very sparse (the typical case). It can be computed much faster in MATLAB without complementing the sparse Mask via:

```
R = Z; R (Mask) = C (Mask); C = R;
```

A set of MATLAB functions that precisely compute the $\mathbf{C}\langle\mathbf{M}\rangle=\mathbf{C}\odot\mathbf{T}$ operation according to the full GraphBLAS specification is provided in Suite-Sparse:GraphBLAS as $\mathtt{GB_spec_accum.m}$, which computes $\mathbf{Z}=\mathbf{C}\odot\mathbf{T}$, and $\mathtt{GB_spec_mask.m}$, which computes $\mathbf{C}\langle\mathbf{M}\rangle=\mathbf{Z}$. SuiteSparse:GraphBLAS includes a complete list of $\mathtt{GB_spec_*}$ functions that illustrate every GraphBLAS operation; these are discussed in in Section 8.1.

The methods in Figure 1 rely heavily on MATLAB's logical matrix indexing. For those unfamiliar with logical indexing in MATLAB, here is short summary. Logical matrix indexing in MATLAB is written as A(Mask) where A is any matrix and Mask is a logical matrix the same size as A. The expression x=A(Mask) produces a column vector x consisting of the entries of A where Mask is true. On the left-hand side, logical submatrix assignment A(Mask)=x does the opposite, copying the components of the vector x into the places in A where Mask is true. For example, to negate all values greater than 10 using logical indexing in MATLAB:

```
>> A = magic (4)
A =
    16
                    3
                          13
     5
           11
                   10
                           8
            7
                    6
                          12
      4
           14
                   15
>> A (A>10) =
                  A (A>10)
A =
   -16
             2
                    3
                         -13
                   10
      5
          -11
                           8
     9
            7
                    6
                         -12
          -14
                 -15
                           1
```

In MATLAB, logical indexing with a sparse matrix A and sparse logical matrix Mask is very efficient since MATLAB supports sparse logical matrices. The Mask operator in GraphBLAS works identically as sparse logical indexing in MATLAB, and is equally as fast (or faster) in SuiteSparse:GraphBLAS.

2.4 Typecasting

If an operator z=f(x) or z=f(x,y) is used with inputs that do not match its inputs x or y, or if its result z does not match the type of the matrix it is being stored into, then the values are typecasted. Typecasting in Graph-BLAS extends beyond just operators. Almost all GraphBLAS methods and operations are able to typecast their results, as needed.

If one type can be typecasted into the other, they are said to be *compatible*. All built-in types are compatible with each other. GraphBLAS cannot typecast user-defined types thus any user-defined type is only compatible with itself. When GraphBLAS requires inputs of a specific type, or when one type cannot be typecast to another, the GraphBLAS function returns an error code, Grb_DOMAIN_MISMATCH (refer to Section 4.4 for a complete list of error codes). Typecasting can only be done between built-in types, and it follows the rules of the ANSI C language (not MATLAB) wherever the rules of ANSI C are well-defined. In particular, a large integer outside the range of a smaller one is wrapped, modulo style. This differs from MATLAB.

However, unlike MATLAB, the C language specification states that the results of typecasting a float or double to an integer type is not always defined. In SuiteSparse:GraphBLAS, whenever C leaves the result undefined the rules used in MATLAB are followed. In particular +Inf converts to the largest integer value, -Inf converts to the smallest (zero for unsigned integers), and NaN converts to zero. Positive values outside the range of the integer are converted to the largest positive integer, and negative values less than the most negative integer are converted to that most negative integer. Other than these special cases, SuiteSparse:GraphBLAS trusts the C compiler for the rest of its typecasting.

Typecasting to bool is fully defined in the C language specification, even for NaN. The result is false if the value compares equal to zero, and true otherwise. Thus NaN converts to true. This is unlike MATLAB, which does not allow a typecast of a NaN to the MATLAB logical type.

SPEC: the GraphBLAS API states that typecasting follows the rules of ANSI C. Yet C leaves some typecasting undefined. SuiteSparse:GraphBLAS provides a precise definition for all typecasting as an extension to the spec.

2.5 Notation and list of GraphBLAS operations

As a summary of what GraphBLAS can do, the following table lists all GraphBLAS operations (where GxB_* are in SuiteSparse:GraphBLAS only). Upper case letters denote a matrix, lower case letters are vectors, and **AB** denote the multiplication of two matrices over a semiring.

GrB_mxm	matrix-matrix multiply	$\mathbf{C}\langle\mathbf{M} angle=\mathbf{C}\odot\mathbf{AB}$
<pre>GrB_vxm</pre>	vector-matrix multiply	$\mathbf{w}^{T} \langle \mathbf{m}^{T} \rangle = \mathbf{w}^{T} \odot \mathbf{u}^{T} \mathbf{A}$
<pre>GrB_mxv</pre>	matrix-vector multiply	$\mathbf{w}\langle\mathbf{m}\rangle=\mathbf{w}\odot\mathbf{A}\mathbf{u}$
GrB_eWiseMult	element-wise,	$\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{C} \odot (\mathbf{A} \otimes \mathbf{B})$
	set intersection	$\mathbf{w}\langle\mathbf{m}\rangle=\mathbf{w}\odot(\mathbf{u}\otimes\mathbf{v})$
GrB_eWiseAdd	element-wise,	$\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot(\mathbf{A}\oplus\mathbf{B})$
	set union	$\mathbf{w}\langle\mathbf{m}\rangle=\mathbf{w}\odot(\mathbf{u}\oplus\mathbf{v})$
GrB_extract	extract submatrix	$\mathbf{C}\langle\mathbf{M} angle=\mathbf{C}\odot\mathbf{A}(\mathbf{I},\mathbf{J})$
		$\mathbf{w} \langle \mathbf{m} \rangle = \mathbf{w} \odot \mathbf{u}(\mathbf{i})$
GxB_subassign	assign submatrix	$\mathbf{C}(\mathbf{I},\mathbf{J})\langle\mathbf{M} angle = \mathbf{C}(\mathbf{I},\mathbf{J})\odot\mathbf{A}$
	(with submask for $\mathbf{C}(\mathbf{I}, \mathbf{J})$)	$\mathbf{w}(\mathbf{i})\langle\mathbf{m} angle = \mathbf{w}(\mathbf{i})\odot\mathbf{u}$
GrB_assign	assign submatrix	$\mathbf{C}\langle \mathbf{M} angle (\mathbf{I},\mathbf{J}) = \mathbf{C}(\mathbf{I},\mathbf{J})\odot \mathbf{A}$
	(with mask for \mathbf{C})	$\mathbf{w}\langle\mathbf{m} angle(\mathbf{i})=\mathbf{w}(\mathbf{i})\odot\mathbf{u}$
GrB_apply	apply unary operator	$\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot f(\mathbf{A})$
		$\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot f(\mathbf{u})$
GxB_select	apply select operator	$\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot f(\mathbf{A},k)$
		$\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot f(\mathbf{u},k)$
GrB_reduce	reduce to vector	$\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot[\oplus_{j}\mathbf{A}(:,j)]$
	reduce to scalar	$s = s \odot [\oplus_{ij} \mathbf{A}(i,j)]$
GrB_transpose	transpose	$\mathbf{C}\langle\mathbf{M} angle = \mathbf{C}\odot\mathbf{A}^T$
GrB_kronecker	Kronecker product	$\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot \mathrm{kron}(\mathbf{A},\mathbf{B})$

Each operation takes an optional $GrB_Descriptor$ argument that modifies the operation. The input matrices A and B can be optionally transposed, the mask M can be complemented, and C can be cleared of its entries after it is used in $C = C \odot T$ but before the $C \subset M = C \odot T$ assignment. Vectors are never transposed via the descriptor.

Let $\mathbf{A} \oplus \mathbf{B}$ denote the element-wise operator that produces a set union pattern (like A+B in MATLAB). Any binary operator can be used this way in GraphBLAS, not just plus. Let $\mathbf{A} \otimes \mathbf{B}$ denote the element-wise operator

that produces a set intersection pattern (like A.*B in MATLAB); any binary operator can be used this way, not just times.

Reduction of a matrix A to a vector reduces the ith row of A to a scalar w_i . This is like w=sum(A) since by default, MATLAB reduces down the columns, not across the rows.

3 Interfaces to MATLAB, Python, and Julia

The MATLAB interface to SuiteSparse:GraphBLAS is included with this distribution, described in Section 3.1. It is fully polished, and fully tested, but does have some limitations that will be addressed in future releases.

A beta version of a Python interface is now available, as is a Julia interface. These are not part of the SuiteSparse:GraphBLAS distribution. See the links below (see Sections 3.2 and 3.3).

3.1 MATLAB Interface

As of Version 3.1, a MATLAB interface is now available. Refer to the documentation in the GraphBLAS/GraphBLAS folder for details. Start with the README.md file in that directory. An easy-to-read output of the MATLAB demos can be found in GraphBLAS/GraphBLAS/demo/html.

Prior to using an GrB object or method, you must first call GrB.init. The best place to put this is in your MATLAB startup.m file, normally located in the user Documents/MATLAB folder.

The MATLAB interface adds the GrB class, which is an opaque MATLAB object that contains a GraphBLAS matrix, either double or single precision, boolean, or any of the built-in integer types. Complex matrix support will be added in the future. MATLAB sparse and full matrices can be arbitrarily mixed with GraphBLAS matrices. The following overloaded operators and methods all work as you would expect for any matrix. The matrix multiplication A*B uses the conventional PLUS_TIMES semiring.

A+B A-B A*B A.*B A./B A.\B A.\b A.\b C=A(I,J)
-A +A
$$^{\sim}A$$
 A' A.' A&B A|B b\A C(I,J)=A
A $^{\sim}$ =B A>B A==B A<=B A>=B A

For a list of overloaded operations and static methods, type methods GrB in MATLAB, or help GrB for more details.

Limitations: Some features for MATLAB sparse matrices are not yet available for GraphBLAS matrices. Some of these may be added in future releases.

• Saving a GrB matrix object from MATLAB can be done, but the resulting *.mat file must be read in by the same version of GraphBLAS.

- GrB matrices with dimension larger than 2⁵³ do not display properly in the MATLAB whos command. MATLAB gets this information from size(A), which returns a correct result, but MATLAB rounds it to double before displaying it. The size is displayed correctly with disp or display.
- Non-blocking mode is not exploited; this would require a MATLAB mexFunction to modify its inputs, which is technically possible but not permitted by the MATLAB API.
- Linear indexing, or A(:) for a 2D matrix, and a single output of I=find(A).
- The second output for min and max, and the includenan option.
- Singleton expansion.
- Dynamically growing arrays, where C(i)=x can increase the size of C.
- Saturating element-wise binary and unary operators for integers. For C=A+B with MATLAB uint8 matrices, results saturate if they exceed 255. This is not compatible with a monoid for C=A*B, and thus MATLAB does not support matrix-matrix multiplication with uint8 matrices. In GraphBLAS, uint8 addition acts in a modulo fashion. Saturating binary operators could be added in the future, so that GrB.eadd (A, '+saturate', B) could return the MATLAB result.
- Solvers, so that $x=A\b$ could return a GF(2) solution, for example.
- Sparse matrices with dimension higher than 2. It would be possible to map an N-dimensional matrix to a large 2D hypersparse GraphBLAS matrix.

3.2 Python Interface

See Michel Pelletier's Python interface at https://github.com/michelp/pygraphblas.

3.3 Julia Interface

See Abhinav Mehndiratta's Julia interface at https://github.com/abhinavmehndiratta/SuiteSparseGraphBLAS.jl.

4 GraphBLAS Context and Sequence

A user application that directly relies on GraphBLAS must include the GraphBLAS.h header file:

```
#include "GraphBLAS.h"
```

The GraphBLAS.h file defines functions, types, and macros prefixed with GrB_ and GxB_ that may be used in user applications. The prefix GrB_ denote items that appear in the official *GraphBLAS C API Specification*. The prefix GxB_ refers to SuiteSparse-specific extensions to the GraphBLAS API. Both may be used in user applications but be aware that items with prefixes GxB_ will not appear in other implementations of the GraphBLAS standard.

```
SPEC: The following macros are extensions to the spec.
```

The GraphBLAS.h file includes all the definitions required to use Graph-BLAS, including the following macros that can assist a user application in compiling and using GraphBLAS.

There are two version numbers associated with SuiteSparse:GraphBLAS: the version of the *GraphBLAS C API Specification* it conforms to, and the version of the implementation itself. These can be used in the following manner in a user application:

```
#if GxB_SPEC_VERSION >= GxB_VERSION (2,0,3)
... use features in GraphBLAS specification 2.0.3 ...
#else
... only use features in early specifications
#endif

#if GxB_IMPLEMENTATION > GxB_VERSION (1,4,0)
... use features from version 1.4.0 of a specific GraphBLAS implementation
#endif
```

SuiteSparse:GraphBLAS also defines the following strings with #define. Refer to the GraphBLAS.h file for details.

Macro	purpose
GxB_IMPLEMENTATION_ABOUT	this particular implementation, copyright, and URL
GxB_IMPLEMENTATION_DATE	the date of this implementation
GxB_SPEC_ABOUT	the GraphBLAS specification for this implementation
GxB_SPEC_DATE	the date of the GraphBLAS specification
GxB_IMPLEMENTATION_LICENSE	the license for this particular implementation

Finally, SuiteSparse:GraphBLAS gives itself a unique name of the form GxB_SUITESPARSE_GRAPHBLAS that the user application can use in #ifdef tests. This is helpful in case a particular implementation provides non-standard features that extend the GraphBLAS specification, such as additional predefined built-in operators, or if a GraphBLAS implementation does not yet fully implement all of the GraphBLAS specification. The Suite-Sparse:GraphBLAS name is provided in its GraphBLAS.h file as:

#define GxB_SUITESPARSE_GRAPHBLAS

For example, SuiteSparse:GraphBLAS predefines additional built-in operators not in the specification. If the user application wishes to use these in any GraphBLAS implementation, an #ifdef can control when they are used. Refer to the examples in the GraphBLAS/Demo folder.

As another example, the GraphBLAS API states that an implementation need not define the order in which <code>GrB_Matrix_build</code> assembles duplicate tuples in its <code>[I,J,X]</code> input arrays. As a result, no particular ordering should be relied upon in general. However, SuiteSparse:GraphBLAS does guarantee an ordering, and this guarantee will be kept in future versions of SuiteSparse:GraphBLAS as well. Since not all implementations will ensure a particular ordering, the following can be used to exploit the ordering returned by SuiteSparse:GraphBLAS.

```
#ifdef GxB_SUITESPARSE_GRAPHBLAS
// duplicates in I, J, X assembled in a specific order;
// results are well-defined even if op is not associative.
GrB_Matrix_build (C, I, J, X, nvals, op);
#else
// duplicates in I, J, X assembled in no particular order;
// results are undefined if op is not associative.
GrB_Matrix_build (C, I, J, X, nvals, op);
#endif
```

The remainder of this section describes GraphBLAS functions that create, modify, and destroy the GraphBLAS context, or provide utility methods for dealing with errors:

GraphBLAS function	purpose	Section
GrB_init	start up GraphBLAS	4.1
${\tt GxB_init}$	start up GraphBLAS with different malloc	4.2
<pre>GrB_Info</pre>	status code returned by GraphBLAS functions	4.3
GrB_error	get more details on the last error	4.4
<pre>GrB_finalize</pre>	finish GraphBLAS	4.5

4.1 GrB_init: initialize GraphBLAS

GrB_init must be called before any other GraphBLAS operation. It defines the mode that GraphBLAS will use: blocking or non-blocking. With blocking mode, all operations finish before returning to the user application. With non-blocking mode, operations can be left pending, and are computed only when needed. Non-blocking mode can be much faster than blocking mode, by many orders of magnitude in extreme cases. Blocking mode should be used only when debugging a user application. The mode cannot be changed once it is set by GrB_init.

GraphBLAS objects are opaque to the user application. This allows GraphBLAS to postpone operations and then do them later in a more efficient manner by rearranging them and grouping them together. In non-blocking mode, the computations required to construct an opaque GraphBLAS object might not be finished when the GraphBLAS method or operation returns to the user. However, user-provided arrays are not opaque, and GraphBLAS methods and operations that read them (such as GrB_Matrix_build) or write to them (such as GrB_Matrix_extractTuples) always finish reading them, or creating them, when the method or operation returns to the user application.

In addition, all methods and operations that extract values from a Graph-BLAS object and return them into non-opaque user arrays always ensure that the computations for that object are completed when the method returns, namely: GrB_*_nvals, GrB_*_extractElement, GrB_*_extractTuples, and GrB_*_reduce (to scalar). These methods only ensure that the computations for a single object are completed.

SuiteSparse:GraphBLAS is multithreaded internally, via OpenMP, and it is also safe to use in a multithreaded user application. See Section 12 for details.

User threads must not operate on the same matrices at the same time, with one exception. Multiple threads can use the same matrices or vectors as read-only inputs to GraphBLAS operations or methods, but only if they have no pending operations (use <code>GrB_Matrix_nvals</code> first). User threads cannot simultaneously modify a matrix or vector via any GraphBLAS operation or method.

With multiple user threads, exactly one user thread must call <code>GrB_init</code> before any user thread may call any <code>GrB_*</code> or <code>GxB_*</code> function. When the user application is finished, exactly one user thread must call <code>GrB_finalize</code>, after which no user thread may call any <code>GrB_*</code> or <code>GxB_*</code> function.

You can query the mode of a GraphBLAS session with the following (see Section 6), which returns the mode passed to GrB_init:

```
GrB_mode mode ;
GxB_get (GxB_MODE, &mode) ;
```

4.2 GxB init: initialize with alternate malloc

GxB_init is identical to GrB_init, except that it also redefines the memory management functions that SuiteSparse:GraphBLAS will use. Giving the user application control over this is particularly important when using the GxB_*import and GxB_*export functions described in Section 5.10, since they require the user application and GraphBLAS to use the same memory manager.

These functions can only be set once, when GraphBLAS starts. Either GrB_init or GxB_init must be called before any other GraphBLAS operation, but not both. The last argument to GxB_init informs GraphBLAS as to whether or not the functions are thread-safe. The ANSI C and Intel TBB functions are thread-safe, but the MATLAB mxMalloc and related functions are not thread-safe. If not thread-safe, GraphBLAS calls the functions from inside an OpenMP critical section.

The following usage is identical to GrB_init(mode):

```
GxB_init (mode, malloc, calloc, realloc, free, true);
```

SuiteSparse:GraphBLAS can be compiled as normal (outside of MAT-LAB) and then linked into a MATLAB mexFunction. However, a mexFunction should use the MATLAB memory managers. To do this, use the following instead of GrB_init(mode) in a MATLAB mexFunction.

```
#include "mex.h"
#include "GraphBLAS.h"
...
GxB_init (mode, mxMalloc, mxCalloc, mxRealloc, mxFree, false);
```

Passing in the last parameter as false requires that GraphBLAS be compiled with OpenMP. Internally, SuiteSparse:GraphBLAS never calls any memory management function inside a parallel region. Results are undefined if all three of the following conditions hold: (1) the user application calls GraphBLAS in parallel from multiple user-level threads, (2) the memory functions are not thread-safe, and (3) GraphBLAS is not compiled with OpenMP. Safety is guaranteed if at least one of those conditions is true.

As another example, the scalable Intel TBB memory manager can be used:

SPEC: GxB_init is an extension to the spec.

4.3 GrB_Info: status code returned by GraphBLAS

Each GraphBLAS method and operation returns its status to the caller as its return value, an enumerated type (an enum) called GrB_Info. The first two values in the following table denote a successful status, the rest are error codes.

GrB_SUCCESS	0	the method or operation was successful
GrB_NO_VALUE	1	A(i,j) requested but not there. Its value
		is implicit.
GrB_UNINITIALIZED_OBJECT	2	object has not been initialized
GrB_INVALID_OBJECT	3	object is corrupted
GrB_NULL_POINTER	4	input pointer is NULL
GrB_INVALID_VALUE	5	generic error code; some value is bad
<pre>GrB_INVALID_INDEX</pre>	6	a row or column index is out of bounds; for
		indices passed as scalars, not in a list.
GrB_DOMAIN_MISMATCH	7	object domains are not compatible
GrB_DIMENSION_MISMATCH	8	matrix dimensions do not match
GrB_OUTPUT_NOT_EMPTY	9	output matrix already has values in it
GrB_OUT_OF_MEMORY	10	out of memory
GrB_INSUFFICIENT_SPACE	11	output array not large enough
<pre>GrB_INDEX_OUT_OF_BOUNDS</pre>	12	a row or column index is out of bounds; for
		indices in a list of indices.
GrB_PANIC	13	unrecoverable error.

Not all GraphBLAS methods or operations can return all status codes. Any GraphBLAS method or operation can return an out-of-memory condition, GrB_OUT_OF_MEMORY, or a panic, GrB_PANIC. These two errors, and the GrB_INDEX_OUT_OF_BOUNDS error, are called execution errors. The other errors are called API errors. An API error is detecting immediately, regardless of the blocking mode. The detection of an execution error may be deferred until the pending operations complete.

In the discussions of each method and operation in this User Guide, most of the obvious error code returns are not discussed. For example, if a required input is a NULL pointer, then GrB_NULL_POINTER is returned. Only error codes specific to the method or that require elaboration are discussed here. For a full list of the status codes that each GraphBLAS function can return, refer to *The GraphBLAS C API Specification* [BMM+17b].

4.4 GrB_error: get more details on the last error

```
const char *GrB_error ( ) ; \hspace{0.4cm} // return a string describing the last error
```

Each GraphBLAS method and operation returns a GrB_Info error code. The GrB_error function returns additional information on the error in a thread-safe null-terminated string. The string returned by GrB_error is allocated in thread local storage and must not be freed or modified. Each user thread has its own error status. The simplest way to use it is just to print it out, such as:

```
info = GrB_some_method_here (...);
if (! (info == GrB_SUCCESS || info == GrB_NO_VALUE))
{
    printf ("info: %d error: %s\n", info, GrB_error ());
}
```

SuiteSparse:GraphBLAS reports many helpful details via GrB_error. For example, if a row or column index is out of bounds, the report will state what those bounds are. If a matrix dimension is incorrect, the mismatching dimensions will be provided. GrB_BinaryOp_new, GrB_UnaryOp_new, and GxB_SelectOp_new record the name the function passed to them, and GrB_Type_new records the name of its type parameter, and these are printed if the user-defined types and operators are used incorrectly. Refer to the output of the example programs in the Demo folder, which intentionally generate errors to illustrate the use of GrB_error.

Successful GraphBLAS methods do not modify the last error message recorded. If a GraphBLAS method fails and then subsequent GraphBLAS method succeeds, the error message is not modified from the last failure. Only a subsequent failure will cause GrB_error to return a different error message.

Note that GrB_NO_VALUE is an not error, but an informational status. $GrB_*=\text{extractElment(\&x,A,i,j)}$, which does x=A(i,j), returns this value to indicate that A(i,j) is not present in the matrix.

In SuiteSparse:GraphBLAS, some failures cannot be safely recorded for GrB_error to print. These include GrB_PANIC and errors in GrB_init and GxB_init.

4.5 GrB_finalize: finish GraphBLAS

```
GrB_Info GrB_finalize ( ) ; // finish GraphBLAS
```

GrB_finalize must be called as the last GraphBLAS operation, even after all calls to GrB_free. All GraphBLAS objects created by the user application should be freed first, before calling GrB_finalize since GrB_finalize will not free those objects. In non-blocking mode, GraphBLAS may leave some computations as pending. These computations can be safely abandoned if the user application frees all GraphBLAS objects it has created and then calls GrB_finalize. When the user application is finished, exactly one user thread must call GrB_finalize.

5 GraphBLAS Objects and their Methods

GraphBLAS defines eight different objects to represent matrices and vectors, their scalar data type (or domain), binary and unary operators on scalar types, monoids, semirings, and a *descriptor* object used to specify optional parameters that modify the behavior of a GraphBLAS operation. SuiteSparse:GraphBLAS adds two additional objects: a sparse scalar, and an operator for selecting entries from a matrix or vector.

The GraphBLAS API makes a distinction between *methods* and *operations*. A method is a function that works on a GraphBLAS object, creating it, destroying it, or querying its contents. An operation (not to be confused with an operator) acts on matrices and/or vectors in a semiring.

GrB_Type	a scalar data type
<pre>GrB_UnaryOp</pre>	a unary operator $z = f(x)$, where z and x are scalars
<pre>GrB_BinaryOp</pre>	a binary operator $z = f(x, y)$, where z, x, and y are scalars
<pre>GxB_SelectOp</pre>	a select operator
<pre>GrB_Monoid</pre>	an associative and commutative binary operator
	and its identity value
<pre>GrB_Semiring</pre>	a monoid that defines the "plus" and a binary operator
	that defines the "multiply" for an algebraic semiring
<pre>GrB_Matrix</pre>	a 2D sparse matrix of any type
<pre>GrB_Vector</pre>	a 1D sparse column vector of any type
<pre>GxB_Scalar</pre>	a sparse scalar of any type
GrB_Descriptor	a collection of parameters that modify an operation

Each of these objects is implemented in C as an opaque handle, which is a pointer to a data structure held by GraphBLAS. User applications may not examine the content of the object directly; instead, they can pass the handle back to GraphBLAS which will do the work. Assigning one handle to another is valid but it does not make a copy of the underlying object.

SPEC: SuiteSparse:GraphBLAS predefines all unique monoids and semirings that can be constructed from built-in types and operators, as an extension to the spec. They appear in GraphBLAS.h. The GxB_SelectOp object is an extension to GraphBLAS.

5.1 The GraphBLAS type: GrB_Type

A GraphBLAS GrB_Type defines the type of scalar values that a matrix or vector contains, and the type of scalar operands for a unary or binary operator. There are eleven built-in types, and a user application can define any types of its own as well. The built-in types correspond to built-in types in C (#include <stdbool.h> and #include <stdint.h>), and the classes in MATLAB, as listed in the following table.

$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$					
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	$\operatorname{GraphBLAS}$	C type	MATLAB	description	range
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	$_{\mathrm{type}}$		class		
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	GrB_BOOL	bool	logical	Boolean	true (1) , false (0)
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	GrB_INT8	int8_t	int8	8-bit signed integer	-128 to 127
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	GrB_UINT8	uint8_t	uint8	8-bit unsigned integer	· · · · - · ·
	GrB_INT16	int16_t	int16	16-bit integer	-2^{15} to $2^{15}-1$
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	GrB_UINT16	$uint16_t$	uint16	16-bit unsigned integer	
GrB_INT64 int64_t int64 64-bit integer -2^{63} to $2^{63}-1$ GrB_UINT64 uint64_t uint64 64-bit unsigned integer 0 to $2^{64}-1$	GrB_INT32	int32_t	int32	32-bit integer	-2^{31} to $2^{31}-1$
GrB_UINT64 uint64_t uint64 64-bit unsigned integer $0 \text{ to } 2^{64} - 1$	GrB_UINT32	uint32_t	uint32	32-bit unsigned integer	0 to $2^{32} - 1$
	GrB_INT64	int64_t	int64	64-bit integer	-2^{63} to $2^{63}-1$
GrB FP32 float single 32-bit IEEE 754 -Inf to +Inf	GrB_UINT64	$uint64_t$	uint64	64-bit unsigned integer	0 to $2^{64} - 1$
015_1102 11000 5116_10 02 510 1EEE 101 1111 00 1III	GrB_FP32	float	single	32-bit IEEE 754	-Inf to +Inf
GrB_FP64 double double 64-bit IEEE 754 -Inf to +Inf	GrB_FP64	double	double	64-bit IEEE 754	-Inf to +Inf

The user application can also define new types based on any typedef in the C language whose values are held in a contiguous region of memory. For example, a user-defined GrB_Type could be created to hold any C struct whose content is self-contained. A C struct containing pointers might be problematic because GraphBLAS would not know to dereference the pointers to traverse the entire "scalar" entry, but this can be done if the objects referenced by these pointers are not moved. A user-defined complex type with real and imaginary types can be defined, or even a "scalar" type containing a fixed-sized dense matrix (see Section 5.1.1). The possibilities are endless. GraphBLAS can create and operate on sparse matrices and vectors in any of these types, including any user-defined ones. For user-defined types, GraphBLAS simply moves the data around itself (via memcpy), and then passes the values back to user-defined functions when it needs to do any computations on the type. The next sections describe the methods for the GrB_Type object:

GrB_Type_new	create a user-defined type
<pre>GxB_Type_size</pre>	return the size of a type
<pre>GrB_Type_free</pre>	free a user-defined type

5.1.1 GrB_Type_new: create a user-defined type

GrB_Type_new creates a new user-defined type. The type is a handle, or a pointer to an opaque object. The handle itself must not be NULL on input, but the content of the handle can be undefined. On output, the handle contains a pointer to a newly created type. The ctype is the type in C that will be used to construct the new GraphBLAS type. It can be either a built-in C type, or defined by a typedef. The second parameter should be passed as sizeof(ctype). The only requirement on the C type is that sizeof(ctype) is valid in C, and that the type reside in a contiguous block of memory so that it can be moved with memcpy. For example, to create a user-defined type called Complex for double-precision complex values using the ANSI C11 double complex type, the following can be used. A complete example can be found in the usercomplex.c and usercomplex.h files in the Demo folder.

```
#include <math.h>
#include <complex.h>
GrB_Type Complex ;
GrB_Type_new (&Complex, sizeof (double complex)) ;
```

To demonstrate the flexibility of the GrB_Type, consider a "scalar" consisting of 4-by-4 floating-point matrix and a string. This type might be useful for the 4-by-4 translation/rotation/scaling matrices that arise in computer graphics, along with a string containing a description or even a regular expression that can be parsed and executed in a user-defined operator. All that is required is a fixed-size type, where sizeof(ctype) is a constant.

```
typedef struct
{
    float stuff [4][4];
    char whatstuff [64];
}
wildtype;
GrB_Type WildType;
GrB_Type_new (&WildType, sizeof (wildtype));
```

With this type a sparse matrix can be created in which each entry consists of a 4-by-4 dense matrix stuff and a 64-character string whatstuff. GraphBLAS treats this 4-by-4 as a "scalar." Any GraphBLAS method or operation that simply moves data can be used with this type without any further information from the user application. For example, entries of this type can be assigned to and extracted from a matrix or vector, and matrices containing this type can be transposed. A working example (wildtype.c in the Demo folder) creates matrices and multiplies them with a user-defined semiring with this type.

Performing arithmetic on matrices and vectors with user-defined types requires operators to be defined. For example, the user application can define its own type for complex numbers, but then transposing the matrix with GraphBLAS will not compute the complex conjugate transpose. This corresponds to the array transpose in MATLAB (C=A.') instead of the complex conjugate transpose (C=A'). To compute the complex conjugate transpose, the application would need to create a user-defined unary operator to conjugate a user-defined complex scalar, and then apply it to the matrix before or after the transpose, via GrB_apply. An extensive set of complex operators are provided in the usercomplex.c example in the Demo folder, along with an include file, usercomplex.h, that is suitable for inclusion in any user application. Thus, while GraphBLAS does not include any complex types or operators, SuiteSparse:GraphBLAS provides them in two simple "user" files in the Demo folder. Refer to Section 11.9 for more details on these two example user-defined types.

5.1.2 GxB_Type_size: return the size of a type

This function acts just like sizeof(type) in the C language. For example GxB_Type_size (&s, GrB_INT32) sets s to 4, the same as sizeof(int32_t).

```
SPEC: GxB_Type_size is an extension to the spec.
```

5.1.3 GrB_Type_free: free a user-defined type

GrB_Type_free frees a user-defined type. Either usage:

```
GrB_Type_free (&type) ;
GrB_free (&type) ;
```

frees the user-defined type and sets type to NULL. It safely does nothing if passed a NULL handle, or if type == NULL on input.

It is safe to attempt to free a built-in type. SuiteSparse:GraphBLAS silently ignores the request and returns GrB_SUCCESS. A user-defined type should not be freed until all operations using the type are completed. Suite-Sparse:GraphBLAS attempts to detect this condition but it must query a freed object in its attempt. This is hazardous and not recommended. Operations on such objects whose type has been freed leads to undefined behavior.

It is safe to first free a type, and then a matrix of that type, but after the type is freed the matrix can no longer be used. The only safe thing that can be done with such a matrix is to free it.

Note the function signature of GrB_Type_free, above. It is illustrated with the generic name, GrB_free. Any GraphBLAS object can be freed with the single function, GrB_free. Refer to Section 5.12 for more details.

GraphBLAS includes many such generic functions. When describing a specific variation, a function is described with its specific name in this User Guide (such as GrB_Type_free). When discussing features applicable to all specific forms, the generic name is used instead (such as GrB_free).

5.2 GraphBLAS unary operators: GrB_UnaryOp, z = f(x)

A unary operator is a scalar function of the form z = f(x). The domain (type) of z and x need not be the same.

There are six kinds of built-in unary operators: one, identity, additive inverse, absolute value, multiplicative inverse, and logical negation. In the notation in the table below, T is any of the 11 built-in types and is a place-holder for BOOL, INT8, UINT8, ... FP32, or FP64. For example, GrB_AINV_INT32 is a unary operator that computes z=-x for two values x and z of type GrB_INT32.

The logical negation operator $\mathtt{GrB_LNOT}$ only works on Boolean types. The $\mathtt{GxB_LNOT_}T$ functions operate on inputs of type T, implicitly typecasting their input to Boolean and returning result of type T, with a value 1 for true and 0 for false. The operators $\mathtt{GxB_LNOT_BOOL}$ and $\mathtt{GrB_LNOT}$ are identical. Considering all combinations, there are thus 67 built-in unary operators ((6 kinds of operators) \times (11 types), and $\mathtt{GrB_LNOT}$).

GraphBLAS name	types (domains)	expression	description
		z = f(x)	
$\mathtt{GxB_ONE_}T$	$T \to T$	z = 1	one
${\tt GrB_IDENTITY_}T$	$T \to T$	z = x	identity
${\tt GrB_AINV_}T$	$T \to T$	z = -x	additive inverse
${\tt GxB_ABS_}T$	$T \to T$	z = x	absolute value
${\tt GrB_MINV_}T$	$T \to T$	z = 1/x	multiplicative inverse
${\tt GxB_LNOT_}T$	$T \to T$	$z = \neg(x \neq 0)$	logical negation
GrB_LNOT	$\texttt{bool} \to \texttt{bool}$	$z = \neg x$	logical negation

SPEC: $\mathtt{GxB_ONE_}T$, $\mathtt{GxB_ABS_}T$ and $\mathtt{GxB_LNOT_}T$ are extensions to the spec.

Integer division by zero normally terminates an application, but this is avoided in SuiteSparse:GraphBLAS. For details, see the binary $\mathtt{GrB_DIV_}T$ operators.

SPEC: The definition of integer division by zero is an extension to the spec.

The next sections define the following methods for the GrB_UnaryOp object:

```
GrB_UnaryOp_new create a user-defined unary operator GxB_UnaryOp_ztype return the type of the output z for z=f(x) GxB_UnaryOp_stype return the type of the input x for z=f(x) free a user-defined unary operator
```

5.2.1 GrB_UnaryOp_new: create a user-defined unary operator

GrB_UnaryOp_new creates a new unary operator. The new operator is returned in the unaryop handle, which must not be NULL on input. On output, its contents contains a pointer to the new unary operator.

The two types xtype and ztype are the GraphBLAS types of the input x and output z of the user-defined function z = f(x). These types may be built-in types or user-defined types, in any combination. The two types need not be the same, but they must be previously defined before passing them to GrB_UnaryOp_new.

The function argument to GrB_UnaryOp_new is a pointer to a user-defined function with the following signature:

```
void (*f) (void *z, const void *x);
```

When the function f is called, the arguments z and x are passed as (void *) pointers, but they will be pointers to values of the correct type, defined by ztype and xtype, respectively, when the operator was created. **NOTE:** The pointers may not be unique. That is, the user function may be called with multiple pointers that point to the same space, such as when z=f(z,y) is to be computed by a binary operator, or z=f(z) for a unary operator. Any parameters passed to the user-callable function may be aliased to each other.

5.2.2 GxB_UnaryOp_ztype: return the type of z

GxB_UnaryOp_ztype returns the ztype of the unary operator, which is the type of z in the function z = f(x).

```
SPEC: GxB_UnaryOp_ztype is an extension to the spec.
```

5.2.3 GxB_UnaryOp_xtype: return the type of x

```
GrB_Info GxB_UnaryOp_xtype  // return the type of x
(
    GrB_Type *xtype,  // return type of input x
    GrB_UnaryOp unaryop  // unary operator
);
```

GxB_UnaryOp_xtype returns the xtype of the unary operator, which is the type of x in the function z = f(x).

```
SPEC: GxB_UnaryOp_xtype is an extension to the spec.
```

5.2.4 GrB_UnaryOp_free: free a user-defined unary operator

GrB_UnaryOp_free frees a user-defined unary operator. Either usage:

```
GrB_UnaryOp_free (&unaryop) ;
GrB_free (&unaryop) ;
```

frees the unaryop and sets unaryop to NULL. It safely does nothing if passed a NULL handle, or if unaryop == NULL on input. It does nothing at all if passed a built-in unary operator.

5.3 GraphBLAS binary operators: GrB_BinaryOp, z = f(x,y)

A binary operator is a scalar function of the form z = f(x, y). The types of z, x, and y need not be the same.

SuiteSparse:GraphBLAS has 21 kinds of built-in binary operators of the form $T \times T \to T$ that work on all 11 of the built-in types, T. These are listed in the table below. For each of these operators, all domains (types) of the three operands are the same. The six comparison operators and the logical operators all return a result one for true and zero for false, in the same domain T as their inputs. These six comparison operators are useful as "multiply" operators for creating semirings with non-Boolean monoids.

GraphBLAS	types (domains)	expression	description
name		z = f(x, y)	
${\tt GrB_FIRST_}T$	$T \times T \to T$	z = x	first argument
${\tt GrB_SECOND_}T$	$T\times T\to T$	z = y	second argument
${\tt GxB_ANY_}T$	$T\times T\to T$	z = x or y	pick x or y arbitrarily
${\tt GxB_PAIR_}T$	$T\times T\to T$	z = 1	one
${\tt GrB_MIN_}T$	$T\times T\to T$	$z = \min(x, y)$	minimum
${\tt GrB_MAX_}T$	$T\times T\to T$	$z = \max(x, y)$	maximum
${\tt GrB_PLUS_}T$	$T\times T\to T$	z = x + y	addition
${\tt GrB_MINUS_}T$	$T\times T\to T$	z = x - y	subtraction
${\tt GxB_RMINUS_}T$	$T\times T\to T$	z = y - x	reverse subtraction
${\tt GrB_TIMES_}T$	$T\times T\to T$	z = xy	multiplication
${\tt GrB_DIV_}T$	$T\times T\to T$	z = x/y	division
${\tt GxB_RDIV_}T$	$T\times T\to T$	z = y/x	reverse division
GxB_ISEQ_T	$T \times T \to T$	z = (x == y)	equal
${\tt GxB_ISNE_}T$	$T\times T\to T$	$z = (x \neq y)$	not equal
${\tt GxB_ISGT_}T$	$T\times T\to T$	z = (x > y)	greater than
${\tt GxB_ISLT_}T$	$T\times T\to T$	z = (x < y)	less than
${\tt GxB_ISGE_}T$	$T\times T\to T$	$z = (x \ge y)$	greater than or equal
${\tt GxB_ISLE_}T$	$T\times T\to T$	$z = (x \le y)$	less than or equal
GxB_LOR_T	$T \times T \to T$	$z = (x \neq 0) \lor (y \neq 0)$	logical OR
${\tt GxB_LAND_}T$	$T\times T\to T$	$z = (x \neq 0) \land (y \neq 0)$	logical AND
${\tt GxB_LXOR_}T$	$T\times T\to T$	$z = (x \neq 0) \lor (y \neq 0)$	logical XOR

SPEC: The $\texttt{GxB_IS*}_T$ $\texttt{GxB_RMINUS}_T$, $\texttt{GxB_RDIV}_T$ $\texttt{GxB_ANY}_T$, and $\texttt{GxB_PAIR}_T$, operators, and the Boolean $\texttt{GxB_L*}_T$ are extensions to the spec.

Another set of six kinds of built-in comparison operators have the form $T \times T \to \mathtt{bool}$. They are defined for all eleven built-in types, for a total of 66 binary operators. Note that when T is \mathtt{bool} , the six operators give the same results as the six $\mathtt{GxB_IS*_BOOL}$ operators in the table above. These six comparison operators are useful as "multiply" operators for creating semirings with Boolean monoids.

GraphBLAS	types (domains)	expression	description
name		z = f(x, y)	
${\tt GrB_EQ_}T$	$T \times T o \mathtt{bool}$	z = (x == y)	equal
${\tt GrB_NE_}T$	$T\times T\to \texttt{bool}$	$z = (x \neq y)$	not equal
${\tt GrB_GT_}T$	$T\times T\to \texttt{bool}$	z = (x > y)	greater than
${\tt GrB_LT_}T$	$T\times T\to \texttt{bool}$	z = (x < y)	less than
${\tt GrB_GE_}T$	$T\times T\to \texttt{bool}$	$z = (x \ge y)$	greater than or equal
${\tt GrB_LE_}T$	$T\times T\to \texttt{bool}$	$z = (x \le y)$	less than or equal

Finally, GraphBLAS has three built-in binary operators that operate purely in the Boolean domain. These three are identical to the GxB_L*_BOOL operators described above, just with a shorter name.

GraphBLAS	types (domains)	expression	description
name		z = f(x, y)	
GrB_LOR	$\texttt{bool} \times \texttt{bool} \to \texttt{bool}$	$z = x \vee y$	logical OR
GrB_LAND	$\texttt{bool} \times \texttt{bool} \to \texttt{bool}$	$z = x \wedge y$	logical AND
GrB_LXOR	$\mathtt{bool} \times \mathtt{bool} \to \mathtt{bool}$	$z = x \veebar y$	logical XOR

There are two sets of built-in comparison operators in SuiteSparse:Graph-BLAS, but they are not redundant. They are identical except for the type (domain) of their output, z. The $\mathtt{GrB_EQ_}T$ and related operators compare their inputs of type T and produce a Boolean result of true or false. The $\mathtt{GxB_ISEQ_}T$ and related operators do the same comparison and produce a result with same type T as their input operands, returning one for true or zero for false. The $\mathtt{IS*}$ comparison operators are useful when combining comparisons with other non-Boolean operators. For example, a PLUS-ISEQ semiring counts how many terms of the comparison are true. With this semiring, matrix multiplication $\mathbf{C} = \mathbf{AB}$ for two weighted undirected graphs \mathbf{A} and \mathbf{B} computes c_{ij} as the number of edges node i and j have in common that have identical edge weights. Since the output type of the "multiplier" operator in a semiring must match the type of its monoid, the Boolean \mathtt{EQ} cannot be combined with a non-Boolean PLUS monoid to perform this operation.

Likewise, SuiteSparse:GraphBLAS has two sets of logical OR, AND, and XOR operators. Without the $_T$ suffix, the three operators $\tt GrB_LOR$, $\tt GrB_LAND$, and $\tt GrB_LXOR$ operate purely in the Boolean domain, where all input and output types are $\tt GrB_BOOL$. The second set $\tt (GxB_LOR_T GxB_LAND_T$ and $\tt GxB_LXOR_T$) provides Boolean operators to all 11 real domains, implicitly typecasting their inputs from type T to Boolean and returning a value of type T that is 1 for true or zero for false. The set of $\tt GxB_L*_T$ operators are useful since they can be combined with non-Boolean monoids in a semiring.

SPEC: The definition of integer division by zero is an extension to the spec.

Floating-point operations follow the IEEE 754 standard. Thus, computing x/0 for a floating-point x results in +Inf if x is positive, -Inf if x is negative, and NaN if x is zero. The application is not terminated. However, integer division by zero normally terminates an application. Suite-Sparse:GraphBLAS avoids this by adopting the same rules as MATLAB, which are analogous to how the IEEE standard handles floating-point division by zero. For integers, when x is positive, x/0 is the largest positive integer, for negative x it is the minimum integer, and 0/0 results in zero. For example, for an integer x of type GrB_INT32 , 1/0 is $2^{31} - 1$ and (-1)/0 is -2^{31} . Refer to Section 5.1 for a list of integer ranges.

The next sections define the following methods for the <code>GrB_BinaryOp</code> object:

GrB_BinaryOp_new	create a user-defined binary operator
<pre>GxB_BinaryOp_ztype</pre>	return the type of the output z for $z = f(x, y)$
<pre>GxB_BinaryOp_xtype</pre>	return the type of the input x for $z = f(x, y)$
<pre>GxB_BinaryOp_ytype</pre>	return the type of the input y for $z = f(x, y)$
<pre>GrB_BinaryOp_free</pre>	free a user-defined binary operator

5.3.1 GrB_BinaryOp_new: create a user-defined binary operator

GrB_BinaryOp_new creates a new binary operator. The new operator is returned in the binaryop handle, which must not be NULL on input. On output, its contents contains a pointer to the new binary operator.

The three types xtype, ytype, and ztype are the GraphBLAS types of the inputs x and y, and output z of the user-defined function z = f(x, y). These types may be built-in types or user-defined types, in any combination. The three types need not be the same, but they must be previously defined before passing them to $GrB_BinaryOp_new$.

The final argument to GrB_BinaryOp_new is a pointer to a user-defined function with the following signature:

```
void (*f) (void *z, const void *x, const void *y);
```

When the function f is called, the arguments z, x, and y are passed as (void *) pointers, but they will be pointers to values of the correct type, defined by ztype, xtype, and ytype, respectively, when the operator was created. NOTE: SuiteSparse:GraphBLAS may call the function with the pointers z and x equal to one another, in which case z=f(z,y) should be computed. Future versions may use additional pointer aliasing.

5.3.2 GxB_BinaryOp_ztype: return the type of z

 $\texttt{GxB_BinaryOp_ztype}$ returns the ztype of the binary operator, which is the type of z in the function z = f(x, y).

SPEC: GxB_BinaryOp_ztype is an extension to the spec.

5.3.3 GxB_BinaryOp_xtype: return the type of x

GxB_BinaryOp_xtype returns the xtype of the binary operator, which is the type of x in the function z = f(x, y).

SPEC: GxB_BinaryOp_xtype is an extension to the spec.

5.3.4 GxB_BinaryOp_ytype: return the type of y

GxB_BinaryOp_ytype returns the ytype of the binary operator, which is the type of y in the function z = f(x, y).

SPEC: GxB_BinaryOp_ytype is an extension to the spec.

5.3.5 GrB_BinaryOp_free: free a user-defined binary operator

GrB_BinaryOp_free frees a user-defined binary operator. Either usage:

```
GrB_BinaryOp_free (&op) ;
GrB_free (&op) ;
```

frees the op and sets op to NULL. It safely does nothing if passed a NULL handle, or if op == NULL on input. It does nothing at all if passed a built-in binary operator.

5.3.6 ANY and PAIR operators

SuiteSparse:GraphBLAS v3.2.0 adds two new operators, ANY and PAIR.

The PAIR operator is simple to describe: just f(x,y) = 1. It is called the PAIR operator since it returns 1 in a semiring when a pair of entries a_{ik} and b_{kj} is found in the matrix multiply. This operator is simple yet very useful. It allows purely symbolic computations to be performed on matrices of any type, without having to typecast them to Boolean with all values being true. Typecasting need not be performed on the inputs to the PAIR operator, and the PAIR operator does not have to access the values of the matrix, so it is a very fast operator to use.

The ANY operator is very unusual, but very powerful. It is the function f(x,y) = x, or y, where GraphBLAS has to freedom to select either x, or y, at its own discretion. Do not confuse the ANY operator with the any function in MATLAB, which computes a reduction using the logical OR operator.

The ANY function is associative and commutative, and can thus serve as an operator for a monoid. The selection of x are y is not randomized. Instead, SuiteSparse:GraphBLAS uses this freedom to compute as fast a result as possible. When used in a dot product,

$$c_{ij} = \sum_{k} a_{ik} b_{kj}$$

for example, the computation can terminate as soon as any matching pair of entries is found. When used in a parallel saxpy-style computation, the ANY operator allows for a relaxed form of synchronization to be used, resulting in a fast benign race condition.

The result of the ANY monoid is non-deterministic, unless it is coupled with the PAIR multiplicative operator. In this case, the ANY_PAIR semiring will return a deterministic result, since f(1,1) is always 1, for the ANY operator.

When paired with a different operator, the results are non-deterministic. This gives a powerful method when computing results for which any value selected by the ANY operator is valid. One such example is the breadth-first-search tree. Suppose node j is at level v, and there are multiple nodes i at level v-1 for which edge (i,j) exists in the graph. Any of these nodes i can serve as a valid parent in the BFS tree. Using the ANY operator, GraphBLAS can quickly compute a valid BFS tree; if it used again on the same inputs, it might return a different, yet still valid, BFS tree, due to the non-deterministic nature of intra-thread synchronization.

5.4 SuiteSparse:GraphBLAS select operators: GxB_SelectOp

A select operator is a scalar function of the form $z = f(i, j, m, n, a_{ij}, \text{thunk})$ that is applied to the entries a_{ij} of an m-by-n matrix. The domain (type) of z is always boolean. The domain (type) of a_{ij} can be any built-in or user-defined type, or it can be $\texttt{GrB_NULL}$ if the operator is type-generic.

The GxB_SelectOp operator is used by GxB_select (see Section 8.13) to select entries from a matrix. Each entry A(i,j) is evaluated with the operator, which returns true if the entry is to be kept in the output, or false if it is not to appear in the output. The signature of the select function f is as follows:

Operators can be used on any type, including user-defined types, except that the comparisons GT, GE, LT, and LE can only be used with built-in types. User-defined select operators can also be created.

GraphBLAS name	MATLAB	description
	analog	
GxB_TRIL	C=tril(A,k)	true for $A(i,j)$ if $(j-i) \le k$
GxB_TRIU	<pre>C=triu(A,k)</pre>	true for $A(i,j)$ if $(j-i) >= k$
GxB_DIAG	<pre>C=diag(A,k)</pre>	true for $A(i,j)$ if $(j-i) == k$
GxB_OFFDIAG	C=A-diag(A,k)	true for $A(i,j)$ if $(j-i) != k$
GxB_NONZERO	C=A(A~=0)	true if A(i,j) is nonzero
GxB_EQ_ZERO	C=A(A==0)	true if A(i,j) is zero
GxB_GT_ZERO	C=A(A>0)	true if A(i,j) is greater than zero
GxB_GE_ZERO	C=A(A>=0)	true if A(i,j) is greater than or equal to zero
GxB_LT_ZERO	C=A(A<0)	true if A(i,j) is less than zero
GxB_LE_ZERO	$C=A(A \le 0)$	true if A(i,j) is less than or equal to zero
GxB_NE_THUNK	$C=A(A^{\sim}=k)$	true if A(i,j) is not equal to k
GxB_EQ_THUNK	C=A(A==k)	true if A(i,j) is equal to k
GxB_GT_THUNK	C=A(A>k)	true if A(i,j) is greater than k
GxB_GE_THUNK	C=A(A>=k)	true if A(i,j) is greater than or equal to k
GxB_LT_THUNK	C=A(A <k)< td=""><td>true if A(i,j) is less than k</td></k)<>	true if A(i,j) is less than k
GxB_LE_THUNK	$C=A(A\leq k)$	true if A(i,j) is less than or equal to k

SPEC: GxB_SelectOp and the table above are extensions to the spec.

The following methods operate on the GxB_SelectOp object:

5.4.1 GxB_SelectOp_new: create a user-defined select operator

GxB_SelectOp_new creates a new select operator. The new operator is returned in the selectop handle, which must not be NULL on input. On output, its contents contains a pointer to the new select operator.

The function argument to $GxB_SelectOp_new$ is a pointer to a user-defined function whose signature is given at the beginning of Section 5.4. Given the properties of an entry a_{ij} in an m-by-n matrix, the function should return true if the entry should be kept in the output of GxB_select , or false if it should not appear in the output.

The type xtype is the GraphBLAS type of the input x of the user-defined function z = f(i, j, m, n, x, thunk). The type may be built-in or user-defined, or it may even be $\texttt{GrB_NULL}$. If the xtype is $\texttt{GrB_NULL}$, then the selectop is type-generic.

The type is the GraphBLAS type of the input thunk of the user-defined function z = f(i, j, m, n, x, thunk). The type may be built-in or user-defined, or it may even be GrB_NULL . If the ttype is GrB_NULL , then the selectop does not access this parameter. The const void *thunk parameter on input to the user function will be passed as NULL.

5.4.2 GxB_SelectOp_xtype: return the type of x

 $\texttt{GxB_SelectOp_xtype}$ returns the xtype of the select operator, which is the type of x in the function z = f(i, j, m, n, x, thunk). If the select operator is type-generic, xtype is returned as $\texttt{GrB_NULL}$. This is not an error condition, but simply indicates that the $\texttt{GxB_SelectOp}$ is type-generic.

5.4.3 GxB_SelectOp_ttype: return the type of the thunk

 $\texttt{GxB_SelectOp_ttype}$ returns the ttype of the select operator, which is the type of thunk in the function z = f(i, j, m, n, x, thunk). If the select operator does not use this parameter, ttype is returned as $\texttt{GrB_NULL}$. This is not an error condition, but simply indicates that the $\texttt{GxB_SelectOp}$ does not use this parameter.

5.4.4 GxB_SelectOp_free: free a user-defined select operator

GxB_SelectOp_free frees a user-defined select operator. Either usage:

```
GxB_SelectOp_free (&selectop) ;
GrB_free (&selectop) ;
```

frees the selectop and sets selectop to NULL. It safely does nothing if passed a NULL handle, or if selectop == NULL on input. It does nothing at all if passed a built-in select operator.

5.5 GraphBLAS monoids: GrB_Monoid

A monoid is defined on a single domain (that is, a single type), T. It consists of an associative binary operator z = f(x, y) whose three operands x, y, and z are all in this same domain T (that is $T \times T \to T$). The associative operator must also have an identity element, or "zero" in this domain, such that f(x,0) = f(0,x) = 0. Recall that an associative operator f(x,y) is one for which the condition f(a,f(b,c)) = f(f(a,b),c) always holds. That is, operator can be applied in any order and the results remain the same.

Five kinds of built-in operators (MIN, MAX, PLUS, TIMES, and ANY) can be used to form monoids for each of the ten non-Boolean built-in types, and 12 can be used for Boolean monoids, all of which are listed in the table below. These are the valid monoids that can be constructed from built-in types and operators, although 8 of the 13 Boolean monoids are redundant (the four remaining being OR, AND, XOR, EQ, and ANY). There are a total of 55 unique monoids that can be constructed using built-in binary operators: five for each of the the 10 non-Boolean types, and 5 more for Boolean types. Since the built-in monoids are also commutative, all of them can be used to create a semiring. Recall that a commutative operator f(x,y) is one for which the condition f(a,b) = f(b,a) always holds. That is, the two operands can be swapped and the results remain the same. One of the components of a semiring is a commutative monoid.

GraphBLAS	types (domains)	expression	identity	terminal
operator		z = f(x, y)		
${\tt GrB_MIN_}T$	$T \times T \to T$	$z = \min(x, y)$	$+\infty$	$-\infty$
${\tt GrB_MAX_}T$	$T\times T\to T$	$z = \max(x, y)$	$-\infty$	$+\infty$
${\tt GrB_PLUS_}T$	$T \times T \to T$	z = x + y	0	none
${\tt GrB_TIMES_}T$	$T \times T \to T$	z = xy	1	0 (not fp)
${\tt GxB_ANY_}T$	$T\times T\to T$	z = x or y	any	any
GrB_LOR	$\texttt{bool} \times \texttt{bool} \to \texttt{bool}$	$z = x \vee y$	false	true
GrB_LAND	$\mathtt{bool} \times \mathtt{bool} \to \mathtt{bool}$	$z = x \wedge y$	true	false
GrB_LXOR	$\mathtt{bool} \times \mathtt{bool} \to \mathtt{bool}$	$z = x \veebar y$	false	none
GrB_EQ_BOOL	$\mathtt{bool} \times \mathtt{bool} \to \mathtt{bool}$	z = (x == y)	true	none

The above table lists the GraphBLAS operator, its type, expression, identity value, and terminal value (if any). For these built-in operators, the terminal values are the annihilators of the function, which is the value z so that z = f(z, y) regardless of the value of y. For example $\min(-\infty, y) = -\infty$ for any y. For integer domains, $+\infty$ and $-\infty$ are the largest and smallest

integer in their range. With unsigned integers, the smallest value is zero, and thus GrB_MIN_UINT8 has an identity of 255 and a terminal value of 0.

When computing with a monoid, the computation can terminate early if the terminal value arises. No further work is needed since the result will not change. This value is called the terminal value instead of the annihilator, since a user-defined operator can be created with a terminal value that is not an annihilator. See Section 5.5.2 for an example.

The GxB_ANY_* monoid can terminate as soon as it finds any value at all. The GrB_TIMES_FP* operators do not have a terminal value of zero, since they comply with the IEEE 754 standard, and 0*NaN is not zero, but NaN. Technically, their terminal value is NaN, but this value is rare in practice and thus the terminal condition is not worth checking.

SuiteSparse:GraphBLAS predefines each of the 55 unique monoids that can be constructed with built-in types and operators, with the naming convention GxB_op_type_MONOID. For the first 50, op is MIN, MAX, PLUS, TIMES, or ANY, and type is all but BOOL. The five Boolean monoids are GxB_LOR_BOOL_MONOID, GxB_LAND_BOOL_MONOID, GxB_LXOR_BOOL_MONOID, GxB_EQ_BOOL_MONOID, and GxB_ANY_BOOL_MONOID.

The next sections define the following methods for the GrB_Monoid object:

GrB_Monoid_new	create a monoid
<pre>GxB_Monoid_terminal_new</pre>	create a monoid that has a terminal value
<pre>GxB_Monoid_operator</pre>	return the monoid operator
<pre>GxB_Monoid_identity</pre>	return the monoid identity value
<pre>GxB_Monoid_terminal</pre>	return the monoid terminal value (if any)
<pre>GrB_Monoid_free</pre>	free a monoid

SPEC: The predefined monoids are an extension to the spec.

5.5.1 GrB Monoid new: create a monoid

GrB_Monoid_new creates a monoid. The operator, op, must be an associative binary operator, either built-in or user-defined.

In the definition above, <type> is a place-holder for the specific type of the monoid. For built-in types, it is the C type corresponding to the built-in type (see Section 5.1), such as bool, int32_t, float, or double. In this case, identity is a scalar value of the particular type, not a pointer. For user-defined types, <type> is void *, and thus identity is a not a scalar itself but a void * pointer to a memory location containing the identity value of the user-defined operator, op.

If op is a built-in operator with a known identity value, then the identity parameter is ignored, and its known identity value is used instead.

5.5.2 GxB_Monoid_terminal_new: create a monoid with terminal

GxB_Monoid_terminal_new is identical to GrB_Monoid_new, except that it allows for the specification of a *terminal value*. The <type> of the terminal value is the same as the identity parameter; see Section 5.5.1 for details.

The terminal value of a monoid is the value z for which z = f(z, y) for any y, where z = f(x, y) is the binary operator of the monoid. This is also called the annihilator, but the term $terminal\ value$ is used here. This is because all annihilators are terminal values, but a terminal value need not be an annihilator, as described in the MIN example below.

If the terminal value is encountered during computation, the rest of the computations can be skipped. This can greatly improve the performance

of GrB_reduce , and matrix multiply in specific cases (when a dot product method is used). For example, using GrB_reduce to compute the sum of all entries in a GrB_FP32 matrix with e entries takes O(e) time, since a monoid based on GrB_PLUS_FP32 has no terminal value. By contrast, a reduction using GrB_LOR on a GrB_BOOL matrix can take as little as O(1) time, if a true value is found in the matrix very early.

Monoids based on the built-in GrB_MIN_* and GrB_MAX_* operators (for any type), the boolean GrB_LOR, and the boolean GrB_LAND operators all have terminal values. For example, the identity value of GrB_LOR is false, and its terminal value is true. When computing a reduction of a set of boolean values to a single value, once a true is seen, the computation can exit early since the result is now known.

If op is a built-in operator with known identity and terminal values, then the identity and terminal parameters are ignored, and its known identity and terminal values are used instead.

There may be cases in which the user application needs to use a non-standard terminal value for a built-in operator. For example, suppose the matrix has type GrB_FP32, but all values in the matrix are known to be non-negative. The annihilator value of MIN is -INFINITY, but this will never be seen. However, the computation could could terminate when finding the value zero. This is an example of using a terminal value that is not actually an annihilator, but it functions like one since the monoid will operate strictly on non-negative values. In this case, a monoid created with GrB_MIN_FP32 will not terminate early. To create a monoid that can terminate early, create a user-defined operator that computes the same thing as GrB_MIN_FP32, and then create a monoid based on this user-defined operator with a terminal value of zero and an identity of +INFINITY.

```
SPEC: GxB_Monoid_terminal_new is an extension to the spec.
```

5.5.3 GxB_Monoid_operator: return the monoid operator

```
GrB_Info GxB_Monoid_operator // return the monoid operator (

GrB_BinaryOp *op, // returns the binary op of the monoid GrB_Monoid monoid // monoid to query );
```

GxB_Monoid_operator returns the binary operator of the monoid.

SPEC: GxB_Monoid_operator is an extension to the spec.

5.5.4 GxB_Monoid_identity: return the monoid identity

```
GrB_Info GxB_Monoid_identity  // return the monoid identity (

void *identity,  // returns the identity of the monoid  
GrB_Monoid monoid  // monoid to query 
);
```

GxB_Monoid_identity returns the identity value of the monoid. The void * pointer, identity, must be non-NULL and must point to a memory space of size at least equal to the size of the type of the monoid. The type size can be obtained via GxB_Monoid_operator to return the monoid additive operator, then GxB_BinaryOp_ztype to obtain the ztype, followed by GxB_Type_size to get its size.

```
SPEC: GxB_Monoid_identity is an extension to the spec.
```

5.5.5 GxB_Monoid_terminal: return the monoid terminal value

GxB_Monoid_terminal returns the terminal value of the monoid (if any). The void * pointer, terminal, must be non-NULL and must point to a memory space of size at least equal to the size of the type of the monoid. The type size can be obtained via GxB_Monoid_operator to return the monoid additive operator, then GxB_BinaryOp_ztype to obtain the ztype, followed by GxB_Type_size to get its size.

If the monoid has a terminal value, then has_terminal is true, and its value is returned in the terminal parameter. If it has no terminal value, then has_terminal is false, and the terminal parameter is not modified.

SPEC: GxB_Monoid_terminal is an extension to the spec.

5.5.6 GrB_Monoid_free: free a monoid

GrB_Monoid_frees frees a monoid. Either usage:

```
GrB_Monoid_free (&monoid) ;
GrB_free (&monoid) ;
```

frees the monoid and sets monoid to NULL. It safely does nothing if passed a NULL handle, or if monoid == NULL on input. It does nothing at all if passed a built-in monoid.

5.6 GraphBLAS semirings: GrB_Semiring

A semiring defines all the operators required to define the multiplication of two sparse matrices in GraphBLAS, $\mathbf{C} = \mathbf{AB}$. The "add" operator is a commutative and associative monoid, and the binary "multiply" operator defines a function z = fmult(x,y) where the type of z matches the exactly with the monoid type. SuiteSparse:GraphBLAS includes 1,473 predefined built-in semirings, which are all those that can be constructed from built-in types and operators. The next sections define the following methods for the $\mathtt{GrB_Semiring}$ object:

```
GrB_Semiring_new create a semiring
GxB_Semiring_add return the additive monoid of a semiring
GxB_Semiring_multiply return the binary operator of a semiring
GrB_Semiring_free free a semiring
```

5.6.1 GrB_Semiring_new: create a semiring

```
GrB_Info GrB_Semiring_new  // create a semiring (

GrB_Semiring *semiring,  // handle of semiring to create  
GrB_Monoid add,  // add monoid of the semiring  
GrB_BinaryOp multiply  // multiply operator of the semiring  
);
```

GrB_Semiring_new creates a new semiring, with add being the additive monoid and multiply being the binary "multiply" operator. In addition to the standard error cases, the function returns GrB_DOMAIN_MISMATCH if the output (ztype) domain of multiply does not match the domain of the add monoid. Using built-in types and operators, 2,291 semirings can be built. This count excludes redundant Boolean operators (for example GrB_TIMES_BOOL and GrB_LAND are different operators but they are redundant since they always return the same result).

SuiteSparse:GraphBLAS pre-defines 1,473 of the 2,291 unique semirings that can be constructed from built-in types and operators, listed below, as an extension to the spec. The naming convention is GxB_add_mult_type, where add is the operator of the additive monoid, mult is the multiply operator, and type is the type of inputs to the multiply operator. The name of the domain for the additive monoid does not appear, since it always matches the type of the output of the mult operator. For example GxB_LAND_EQ_FP32 uses the

GxB_LAND_BOOL_MONOID as its additive monoid, and the GrB_EQ_FP32 as the binary multiplicative operator.

- 1000 semirings with a multiplier $T \times T \to T$ where T is non-Boolean, from the complete cross product of:
 - 5 add monoids (MIN, MAX, PLUS, TIMES, ANY)
 - 20 multiply operators (FIRST, SECOND, PAIR, MIN, MAX, PLUS, MINUS, RMINUS, TIMES, DIV, RDIV, ISEQ, ISNE, ISGT, ISLT, ISGE, ISLE, LOR, LAND, LXOR).
 - 10 non-Boolean types, T
- 300 semirings with a comparison operator $T \times T \to \mathsf{bool}$, where T is non-Boolean, from the complete cross product of:
 - 5 Boolean add monoids (LAND, LOR, LXOR, EQ, ANY)
 - 6 multiply operators (EQ, NE, GT, LT, GE, LE)
 - 10 non-Boolean types, T
- 55 semirings with purely Boolean types, bool × bool → bool, from the complete cross product of:
 - 5 Boolean add monoids (LAND, LOR, LXOR, EQ, ANY)
 - 11 multiply operators (FIRST, SECOND, PAIR, LOR, LAND, LXOR, EQ, GT, LT, GE, LE)
- 54 complex semirings, $T \times T \to T$ where T is float complex or double complex:
 - 3 complex monoids (PLUS, TIMES, ANY)
 - 9 complex multiply operators: (FIRST, SECOND, PAIR, PLUS, MINUS, TIMES, DIV, RDIV, RMINUS)
 - -2 complex types, T
- 64 bitwise semirings, $T \times T \to T$ where T is an unsigned integer.
 - 4 bitwise monoids (BOR, BAND, BXOR, BXNOR)
 - 4 bitwise multiply operators (the same list)
 - 4 unsigned integer types

SPEC: Predefined semirings are an extension to the spec.

5.6.2 GxB_Semiring_add: return the additive monoid of a semiring

GxB_Semiring_add returns the additive monoid of a semiring.

```
SPEC: GxB_Semiring_add is an extension to the spec.
```

5.6.3 GxB_Semiring_multiply: return multiply operator of a semiring

GxB_Semiring_multiply returns the binary multiplicative operator of a semiring.

```
SPEC: GxB_Semiring_multiply is an extension to the spec.
```

5.6.4 GrB_Semiring_free: free a semiring

GrB_Semiring_free frees a semiring. Either usage:

```
GrB_Semiring_free (&semiring) ;
GrB_free (&semiring) ;
```

frees the semiring and sets semiring to NULL. It safely does nothing if passed a NULL handle, or if semiring == NULL on input. It does nothing at all if passed a built-in semiring.

5.7 GraphBLAS scalars: GxB_Scalar

This section describes a set of methods that create, modify, query, and destroy a GraphBLAS sparse scalar, GxB_Scalar:

SPEC: GxB_Scalar is an extension to the spec.

```
GxB_Scalar_new
                               create a sparse scalar
GxB_Scalar_dup
                               copy a sparse scalar
GxB_Scalar_clear
                               clear a sparse scalar of its entry
GxB_Scalar_nvals
                               return the number of entries in a sparse scalar (0 or 1)
GxB_Scalar_type
                               return the type of a sparse scalar
                               set the single entry of a sparse scalar
GxB_Scalar_setElement
GxB_Scalar_extractElement
                               get the single entry from a sparse scalar
GxB_Scalar_free
                               free a sparse scalar
GxB_Scalar_wait
                               wait for pending work on a vector to finish
```

5.7.1 GxB_Scalar_new: create a sparse scalar

```
GrB_Info GxB_Scalar_new  // create a new GxB_Scalar with no entry
(
    GxB_Scalar *s,  // handle of GxB_Scalar to create
    GrB_Type type  // type of GxB_Scalar to create
);
```

GxB_Scalar_new creates a new sparse scalar with no entry in it, of the given type. This is analogous to MATLAB statement s = sparse (0), except that GraphBLAS can create sparse scalars any type. The pattern of the new scalar is empty.

5.7.2 GxB_Scalar_dup: copy a sparse scalar

GxB_Scalar_dup makes a deep copy of a sparse scalar, like s=t in MAT-LAB. In GraphBLAS, it is possible, and valid, to write the following:

Then s and t can be used interchangeably. However, only a pointer reference is made, and modifying one of them modifies both, and freeing one of them leaves the other as a dangling handle that should not be used. If two different sparse scalars are needed, then this should be used instead:

Then **s** and **t** are two different sparse scalars that currently have the same value, but they do not depend on each other. Modifying one has no effect on the other.

5.7.3 GxB_Scalar_clear: clear a sparse scalar of its entry

GxB_Scalar_clear clears the entry from a sparse scalar. The pattern of s is empty, just as if it were created fresh with GxB_Scalar_new. Analogous with s = sparse (0) in MATLAB. The type of s does not change. In SuiteSparse:GraphBLAS, any pending updates to the sparse scalar are discarded.

5.7.4 GxB_Scalar_nvals: return the number of entries in a sparse scalar

GxB_Scalar_nvals returns the number of entries in a sparse scalar, which is either 0 or 1. Roughly analogous to nvals = nnz(s) in MATLAB, except that the implicit value in GraphBLAS need not be zero and nnz (short for "number of nonzeros") in MATLAB is better described as "number of entries" in GraphBLAS.

5.7.5 GxB_Scalar_type: return the type of a sparse scalar

GxB_Scalar_type returns the type of a sparse scalar. Analogous to type = class (s) in MATLAB.

5.7.6 GxB_Scalar_setElement: set the single entry of a sparse scalar

GxB_Scalar_setElement sets the single entry in a sparse scalar, like s = sparse(x) in MATLAB notation. For further details of this function, see GxB_Matrix_setElement in Section 5.9.9.

5.7.7 GxB_Scalar_extractElement: get the single entry from a sparse scalar

GxB_Scalar_extractElement extracts the single entry from a sparse scalar, like x = full(s) in MATLAB. Further details of this method are discussed in Section 5.9.10, which discusses GrB_Matrix_extractElement. NOTE: if no entry is present in the sparse scalar s, then x is not modified, and the return value of GxB_Scalar_extractElement is GrB_NO_VALUE.

5.7.8 GxB_Scalar_free: free a sparse scalar

GxB_Scalar_free frees a sparse scalar. Either usage:

```
GxB_Scalar_free (&s) ;
GrB_free (&s) ;
```

frees the sparse scalar s and sets s to NULL. It safely does nothing if passed a NULL handle, or if s == NULL on input. In SuiteSparse:GraphBLAS, any pending updates to the sparse scalar are abandoned.

5.7.9 GxB_Scalar_wait: finish all pending work on a scalar

```
GrB_Info GxB_Scalar_wait
(
    GxB_Scalar *s
);
```

GxB_Scalar_wait forces all pending operations on a scalar to be completed.

5.8 GraphBLAS vectors: GrB_Vector

Many of the methods for GraphBLAS vectors require a row index or a size. Many methods for matrices require both a row and column index, or a row and column dimension. These are all integers of a specific type, GrB_Index, which is defined in GraphBLAS.h as

```
typedef uint64_t GrB_Index ;
```

Row and column indices of an nrows-by-ncols matrix range from zero to the nrows-1 for the rows, and zero to ncols-1 for the columns. Indices are zero-based, like C, and not one-based, like MATLAB. In Suite-Sparse:GraphBLAS, the largest size permitted for any integer of GrB_Index is 2^{60} . The largest GrB_Matrix that SuiteSparse:GraphBLAS can construct is thus 2^{60} -by- 2^{60} . An n-by-n matrix A that size can easily be constructed in practice with $O(|\mathbf{A}|)$ memory requirements, where $|\mathbf{A}|$ denotes the number of entries that explicitly appear in the pattern of \mathbf{A} . The time and memory required to construct a matrix that large does not depend on n, since Suite-Sparse:GraphBLAS can represent \mathbf{A} in hypersparse form (see Section 6.3). The largest GrB_Vector that can be constructed is 2^{60} -by-1.

This section describes a set of methods that create, modify, query, and destroy a GraphBLAS sparse vector, GrB_Vector:

GrB_Vector_new	create a vector
<pre>GrB_Vector_dup</pre>	copy a vector
<pre>GrB_Vector_clear</pre>	clear a vector of all entries
<pre>GrB_Vector_size</pre>	return the size of a vector
<pre>GrB_Vector_nvals</pre>	return the number of entries in a vector
<pre>GxB_Vector_type</pre>	return the type of a vector
<pre>GrB_Vector_build</pre>	build a vector from a set of tuples
<pre>GrB_Vector_setElement</pre>	add a single entry to a vector
<pre>GrB_Vector_extractElement</pre>	get a single entry from a vector
<pre>GrB_Vector_extractTuples</pre>	get all entries from a vector
<pre>GrB_Vector_resize</pre>	resize a vector
<pre>GrB_Vector_free</pre>	free a vector
<pre>GrB_Vector_wait</pre>	wait for all pending work to finish
GxB_Vector_import	import a vector (see Section 5.10)
<pre>GxB_Vector_export</pre>	export a vector (see Section 5.10)

5.8.1 GrB_Vector_new: create a vector

GrB_Vector_new creates a new n-by-1 sparse vector with no entries in it, of the given type. This is analogous to MATLAB statement v = sparse (n,1), except that GraphBLAS can create sparse vectors any type. The pattern of the new vector is empty.

5.8.2 GrB_Vector_dup: copy a vector

GrB_Vector_dup makes a deep copy of a sparse vector, like w=u in MAT-LAB. In GraphBLAS, it is possible, and valid, to write the following:

Then w and u can be used interchangeably. However, only a pointer reference is made, and modifying one of them modifies both, and freeing one of them leaves the other as a dangling handle that should not be used. If two different vectors are needed, then this should be used instead:

```
GrB_Vector u, w ;
GrB_Vector_new (&u, GrB_FP64, n) ;
GrB_Vector_dup (&w, u) ;  // like w = u, but making a deep copy
```

Then w and u are two different vectors that currently have the same set of values, but they do not depend on each other. Modifying one has no effect on the other.

5.8.3 GrB_Vector_clear: clear a vector of all entries

```
GrB_Info GrB_Vector_clear // clear a vector of all entries;
( // type and dimension remain unchanged.
GrB_Vector v // vector to clear
);
```

 GrB_Vector_clear clears all entries from a vector. All values v(i) are now equal to the implicit value, depending on what semiring ring is used to perform computations on the vector. The pattern of v is empty, just as if it were created fresh with GrB_Vector_new . Analogous with v (:) = 0 in MATLAB. The type and dimension of v do not change. In SuiteSparse:GraphBLAS, any pending updates to the vector are discarded.

5.8.4 GrB_Vector_size: return the size of a vector

 GrB_Vector_size returns the size of a vector (the number of rows). Analogous to n = length(v) or n = size(v, 1) in MATLAB.

5.8.5 GrB_Vector_nvals: return the number of entries in a vector

GrB_Vector_nvals returns the number of entries in a vector. Roughly analogous to nvals = nnz(v) in MATLAB, except that the implicit value in GraphBLAS need not be zero and nnz (short for "number of nonzeros") in MATLAB is better described as "number of entries" in GraphBLAS.

5.8.6 GxB_Vector_type: return the type of a vector

GxB_Vector_type returns the type of a vector. Analogous to type = class (v)
in MATLAB.

SPEC: GxB_Vector_type is an extension to the spec.

5.8.7 GrB_Vector_build: build a vector from a set of tuples

GrB_Vector_build constructs a sparse vector w from a set of tuples, I and X, each of length nvals. The vector w must have already been initialized with GrB_Vector_new, and it must have no entries in it before calling GrB_Vector_build.

This function is just like GrB_Matrix_build (see Section 5.9.8), except that it builds a sparse vector instead of a sparse matrix. For a description of what GrB_Vector_build does, refer to GrB_Matrix_build. For a vector, the list of column indices J in GrB_Matrix_build is implicitly a vector of length nvals all equal to zero. Otherwise the methods are identical.

SPEC: As an extension to the spec, results are defined even if dup is non-associative.

5.8.8 GrB_Vector_setElement: add a single entry to a vector

GrB_Vector_setElement sets a single entry in a vector, w(i) = x. The operation is exactly like setting a single entry in an n-by-1 matrix, A(i,0) = x, where the column index for a vector is implicitly j=0. For further details of this function, see GrB_Matrix_setElement in Section 5.9.9.

5.8.9 GrB_Vector_extractElement: get a single entry from a vector

GrB_Vector_extractElement extracts a single entry from a vector, $\mathbf{x} = \mathbf{v}(\mathbf{i})$. The method is identical to extracting a single entry $\mathbf{x} = \mathbf{A}(\mathbf{i}, \mathbf{0})$ from an n-by-1 matrix, so further details of this method are discussed in Section 5.9.10, which discusses GrB_Matrix_extractElement. In this case, the column index is implicitly $\mathbf{j} = \mathbf{0}$. NOTE: if no entry is present at $\mathbf{v}(\mathbf{i})$, then \mathbf{x} is not modified, and the return value of GrB_Vector_extractElement is GrB_NO_VALUE.

5.8.10 GrB_Vector_extractTuples: get all entries from a vector

GrB_Vector_extractTuples extracts all tuples from a sparse vector, analogous to [I,~,X] = find(v) in MATLAB. This function is identical to its GrB_Matrix_extractTuples counterpart, except that the array of column indices J does not appear in this function. Refer to Section 5.9.11 where further details of this function are described.

5.8.11 GrB_Vector_resize: resize a vector

```
GrB_Info GrB_Vector_resize // change the size of a vector

(
GrB_Vector u, // vector to modify
GrB_Index nrows_new // new number of rows in vector
);
```

GrB_Vector_resize changes the size of a vector. If the dimension decreases, entries that fall outside the resized vector are deleted.

5.8.12 GrB_Vector_free: free a vector

GrB_Vector_free frees a vector. Either usage:

```
GrB_Vector_free (&v) ;
GrB_free (&v) ;
```

frees the vector v and sets v to NULL. It safely does nothing if passed a NULL handle, or if v == NULL on input. In SuiteSparse:GraphBLAS, any pending updates to the vector are abandoned.

5.8.13 GrB_Vector_wait: finish all pending work on a vector

```
GrB_Info GrB_Vector_wait
(
        GrB_Vector *v
);
```

GrB_Vector_wait forces all pending operations on a vector to be completed.

5.9 GraphBLAS matrices: GrB_Matrix

This section describes a set of methods that create, modify, query, and destroy a GraphBLAS sparse matrix, GrB_Matrix:

```
GrB_Matrix_new
                               create a matrix
GrB_Matrix_dup
                               copy a matrix
GrB_Matrix_clear
                               clear a matrix of all entries
                               return the number of rows of a matrix
GrB_Matrix_nrows
GrB_Matrix_ncols
                               return the number of columns of a matrix
                               return the number of entries in a matrix
GrB_Matrix_nvals
GxB_Matrix_type
                               return the type of a matrix
                               build a matrix from a set of tuples
GrB_Matrix_build
GrB_Matrix_setElement
                               add a single entry to a matrix
                               get a single entry from a matrix
GrB_Matrix_extractElement
                               get all entries from a matrix
GrB_Matrix_extractTuples
GrB_Matrix_resize
                               resize a matrix
GrB_Matrix_free
                               free a matrix
                               wait for all pending work to finish
GrB_Matrix_wait
                               import a matrix in CSR form (see Section 5.10)
GxB_Matrix_import_CSR
GxB_Matrix_import_CSC
                               import a matrix in CSC form (see Section 5.10)
                               import a matrix in HyperCSR form (see Section 5.10)
GxB_Matrix_import_HyperCSR
GxB_Matrix_import_HyperCSC
                               import a matrix in HyperCSC form (see Section 5.10)
                               export a matrix in CSR form (see Section 5.10)
GxB_Matrix_export_CSR
GxB_Matrix_export_CSC
                               export a matrix in CSC form (see Section 5.10)
                               export a matrix in HyperCSR form (see Section 5.10)
GxB_Matrix_export_HyperCSR
GxB_Matrix_export_HyperCSC
                               export a matrix in HyperCSC form (see Section 5.10)
```

5.9.1 GrB_Matrix_new: create a matrix

GrB_Matrix_new creates a new nrows-by-ncols sparse matrix with no entries in it, of the given type. This is analogous to the MATLAB statement A = sparse (nrows, ncols), except that GraphBLAS can create sparse matrices of any type.

5.9.2 GrB_Matrix_dup: copy a matrix

```
GrB_Info GrB_Matrix_dup  // make an exact copy of a matrix

(
GrB_Matrix *C,  // handle of output matrix to create const GrB_Matrix A  // input matrix to copy
);
```

GrB_Matrix_dup makes a deep copy of a sparse matrix, like C=A in MAT-LAB. In GraphBLAS, it is possible, and valid, to write the following:

Then C and A can be used interchangeably. However, only a pointer reference is made, and modifying one of them modifies both, and freeing one of them leaves the other as a dangling handle that should not be used. If two different matrices are needed, then this should be used instead:

```
GrB_Matrix A, C ;
GrB_Matrix_new (&A, GrB_FP64, n) ;
GrB_Matrix_dup (&C, A) ;  // like C = A, but making a deep copy
```

Then C and A are two different matrices that currently have the same set of values, but they do not depend on each other. Modifying one has no effect on the other.

5.9.3 GrB_Matrix_clear: clear a matrix of all entries

GrB_Matrix_clear clears all entries from a matrix. All values A(i,j) are now equal to the implicit value, depending on what semiring ring is used to perform computations on the matrix. The pattern of A is empty, just as if it were created fresh with GrB_Matrix_new. Analogous with A (:,:) = 0 in MATLAB. The type and dimensions of A do not change. In SuiteSparse:Graph-BLAS, any pending updates to the matrix are discarded.

5.9.4 GrB_Matrix_nrows: return the number of rows of a matrix

GrB_Matrix_nrows returns the number of rows of a matrix (nrows=size(A,1) in MATLAB).

5.9.5 GrB_Matrix_ncols: return the number of columns of a matrix

```
GrB_Info GrB_Matrix_ncols // get the number of columns of a matrix (

GrB_Index *ncols, // matrix has ncols columns const GrB_Matrix A // matrix to query

);
```

GrB_Matrix_ncols returns the number of columns of a matrix (ncols=size(A,2) in MATLAB).

5.9.6 GrB_Matrix_nvals: return the number of entries in a matrix

```
GrB_Info GrB_Matrix_nvals // get the number of entries in a matrix (

GrB_Index *nvals, // matrix has nvals entries const GrB_Matrix A // matrix to query

);
```

GrB_Matrix_nvals returns the number of entries in a matrix. Roughly analogous to nvals = nnz(A) in MATLAB, except that the implicit value in GraphBLAS need not be zero and nnz (short for "number of nonzeros") in MATLAB is better described as "number of entries" in GraphBLAS.

5.9.7 GxB_Matrix_type: return the type of a matrix

GxB_Matrix_type returns the type of a matrix, like type=class(A) in MATLAB.

```
SPEC: GxB_Matrix_type is an extension to the spec.
```

5.9.8 GrB_Matrix_build: build a matrix from a set of tuples

GrB_Matrix_build constructs a sparse matrix C from a set of tuples, I, J, and X, each of length nvals. The matrix C must have already been initialized with GrB_Matrix_new, and it must have no entries in it before calling GrB_Matrix_build. Thus the dimensions and type of C are not changed by this function, but are inherited from the prior call to GrB_Matrix_new or GrB_matrix_dup.

An error is returned (GrB_INDEX_OUT_OF_BOUNDS) if any row index in I is greater than or equal to the number of rows of C, or if any column index in J is greater than or equal to the number of columns of C

Any duplicate entries with identical indices are assembled using the binary dup operator provided on input. All three types (x, y, z for z=dup(x,y)) must be identical. The types of dup, C and X must all be compatible. See Section 2.4 regarding typecasting and compatibility. The values in X are typecasted, if needed, into the type of dup. Duplicates are then assembled into a matrix T of the same type as dup, using $T(i,j) = dup \ (T \ (i,j), X \ (k))$. After T is constructed, it is typecasted into the result C. That is, typecasting does not occur at the same time as the assembly of duplicates.

SPEC: As an extension to the spec, results are defined even if dup is non-associative.

The GraphBLAS API requires dup to be associative so that entries can be assembled in any order, and states that the result is undefined if dup is not associative. However, SuiteSparse:GraphBLAS guarantees a well-defined order of assembly. Entries in the tuples $[\mathtt{I},\mathtt{J},\mathtt{X}]$ are first sorted in increasing order of row and column index, with ties broken by the position of the tuple in the $[\mathtt{I},\mathtt{J},\mathtt{X}]$ list. If duplicates appear, they are assembled in the order they appear in the $[\mathtt{I},\mathtt{J},\mathtt{X}]$ input. That is, if the same indices \mathtt{i} and \mathtt{j} appear in positions $\mathtt{k1}$, $\mathtt{k2}$, $\mathtt{k3}$, and $\mathtt{k4}$ in $[\mathtt{I},\mathtt{J},\mathtt{X}]$, where $\mathtt{k1} < \mathtt{k2} < \mathtt{k3} < \mathtt{k4}$, then the following operations will occur in order:

```
T (i,j) = X (k1);

T (i,j) = dup (T (i,j), X (k2));

T (i,j) = dup (T (i,j), X (k3));

T (i,j) = dup (T (i,j), X (k4));
```

This is a well-defined order but the user should not depend upon it when using other GraphBLAS implementations since the GraphBLAS API does not require this ordering.

However, SuiteSparse:GraphBLAS guarantees this ordering, even when it compute the result in parallel. With this well-defined order, several operators become very useful. In particular, the SECOND operator results in the last tuple overwriting the earlier ones. The FIRST operator means the value of the first tuple is used and the others are discarded.

The acronym dup is used here for the name of binary function used for assembling duplicates, but this should not be confused with the _dup suffix in the name of the function GrB_Matrix_dup. The latter function does not apply any operator at all, nor any typecasting, but simply makes a pure deep copy of a matrix.

The parameter X is a pointer to any C equivalent built-in type, or a void * pointer. The GrB_Matrix_build function uses the _Generic feature of ANSI C11 to detect the type of pointer passed as the parameter X. If X is a pointer to a built-in type, then the function can do the right typecasting. If X is a void * pointer, then it can only assume X to be a pointer to a user-defined type that is the same user-defined type of C and dup. This function has no way of checking this condition that the void * X pointer points to an array of the correct user-defined type, so behavior is undefined if the user breaks this condition.

The GrB_Matrix_build method is analogous to C = sparse (I,J,X) in MATLAB, with several important extensions that go beyond that which MATLAB can do. In particular, the MATLAB sparse function only provides one option for assembling duplicates (summation), and it can only build double, double complex, and logical sparse matrices.

5.9.9 GrB_Matrix_setElement: add a single entry to a matrix

GrB_Matrix_setElement sets a single entry in a matrix, C(i,j)=x. If the entry is already present in the pattern of C, it is overwritten with the new value. If the entry is not present, it is added to C. In either case, no entry is ever deleted by this function. Passing in a value of x=0 simply creates an explicit entry at position (i,j) whose value is zero, even if the implicit value is assumed to be zero.

An error is returned (Grb_INVALID_INDEX) if the row index i is greater than or equal to the number of rows of C, or if the column index j is greater than or equal to the number of columns of C. Note that this error code differs from the same kind of condition in Grb_Matrix_build, which returns Grb_INDEX_OUT_OF_BOUNDS. This is because Grb_INVALID_INDEX is an API error, and is caught immediately even in non-blocking mode, whereas Grb_INDEX_OUT_OF_BOUNDS is an execution error whose detection may wait until the computation completes sometime later.

The scalar **x** is typecasted into the type of **C**. Any value can be passed to this function and its type will be detected, via the _Generic feature of ANSI C11. For a user-defined type, **x** is a **void** * pointer that points to a memory space holding a single entry of this user-defined type. This user-defined type must exactly match the user-defined type of **C** since no typecasting is done between user-defined types.

Performance considerations: SuiteSparse:GraphBLAS exploits the non-blocking mode to greatly improve the performance of this method. Refer to the example shown in Section 2.2. If the entry exists in the pattern already, it is updated right away and the work is not left pending. Otherwise, it is placed in a list of pending updates, and the later on the updates are done all at once, using the same algorithm used for GrB_Matrix_build. In other words, setElement in SuiteSparse:GraphBLAS builds its own internal list of tuples [I,J,X], and then calls GrB_Matrix_build whenever the matrix is

needed in another computation, or whenever GrB_Matrix_wait is called.

As a result, if calls to setElement are mixed with calls to most other methods and operations (even extractElement) then the pending updates are assembled right away, which will be slow. Performance will be good if many setElement updates are left pending, and performance will be poor if the updates are assembled frequently.

A few methods and operations can be intermixed with setElement, in particular, some forms of the GrB_assign and GxB_subassign operations are compatible with the pending updates from setElement. Sections 8.11 gives more details on which GxB_subassign and GrB_assign operations can be interleaved with calls to setElement without forcing updates to be assembled. Other methods that do not access the existing entries may also be done without forcing the updates to be assembled, namely GrB_Matrix_clear (which erases all pending updates), GrB_Matrix_free, GrB_Matrix_ncols, GrB_Matrix_nrows, GxB_Matrix_type, and of course GrB_Matrix_setElement itself. All other methods and operations cause the updates to be assembled. Future versions of SuiteSparse:GraphBLAS may extend this list.

See Section 11.4 for an example of how to use GrB_Matrix_setElement.

5.9.10 GrB_Matrix_extractElement: get a single entry from a matrix

GrB_Matrix_extractElement extracts a single entry from a matrix x=A(i, j).

An error is returned (GrB_INVALID_INDEX) if the row index i is greater than or equal to the number of rows of C, or if column index j is greater than or equal to the number of columns of C.

NOTE: if no entry is present at A(i,j), then x is not modified, and the return value of GrB_Matrix_extractElement is GrB_NO_VALUE.

If the entry is not present then GraphBLAS does not know its value, since its value depends on the implicit value, which is the identity value of the additive monoid of the semiring. It is not a characteristic of the matrix itself, but of the semiring it is used in. A matrix can be used in any

compatible semiring, and even a mixture of semirings, so the implicit value can change as the semiring changes.

As a result, if the entry is present, x=A(i,j) is performed and the scalar x is returned with this value. The method returns $GrB_SUCCESS$. If the entry is not present, x is not modified, and GrB_NO_VALUE is returned to the caller. What this means is up to the caller.

The function knows the type of the pointer x, so it can do typecasting as needed, from the type of A into the type of x. User-defined types cannot be typecasted, so if A has a user-defined type then x must be a void * pointer that points to a memory space the same size as a single scalar of the type of A.

Currently, this method causes all pending updates from GrB_setElement, GrB_assign, or GxB_subassign to be assembled, so its use can have performance implications. Calls to this function should not be arbitrarily intermixed with calls to these other two functions. Everything will work correctly and results will be predictable, it will just be slow.

5.9.11 GrB_Matrix_extractTuples: get all entries from a matrix

GrB_Matrix_extractTuples extracts all the entries from the matrix A, returning them as a list of tuples, analogous to [I,J,X]=find(A) in MAT-LAB. Entries in the tuples [I,J,X] are unique. No pair of row and column indices (i,j) appears more than once.

The GraphBLAS API states the tuples can be returned in any order. SuiteSparse:GraphBLAS chooses to always return them in sorted order, depending on whether the matrix is stored by row or by column.

The number of tuples in the matrix A is given by GrB_Matrix_nvals(&anvals, A). If anvals is larger than the size of the arrays (nvals in the parameter list), an error GrB_INSUFFICIENT_SIZE is returned, and no tuples are extracted. If nvals is larger than anvals, then only the first anvals entries in the arrays

I J, and X are modified, containing all the tuples of A, and the rest of I J, and X are left unchanged. On output, nvals contains the number of tuples extracted.

5.9.12 GrB_Matrix_resize: resize a matrix

```
GrB_Info GrB_Matrix_resize  // change the size of a matrix

(

GrB_Matrix A,  // matrix to modify

const GrB_Index nrows_new,  // new number of rows in matrix

const GrB_Index ncols_new  // new number of columns in matrix
);
```

GrB_Matrix_resize changes the size of a matrix. If the dimensions decrease, entries that fall outside the resized matrix are deleted.

5.9.13 GrB_Matrix_free: free a matrix

GrB_Matrix_free frees a matrix. Either usage:

```
GrB_Matrix_free (&A) ;
GrB_free (&A) ;
```

frees the matrix A and sets A to NULL. It safely does nothing if passed a NULL handle, or if A == NULL on input. In SuiteSparse:GraphBLAS, any pending updates to the matrix are abandoned.

5.9.14 GrB_Matrix_wait: finish all pending work on a matrix

```
GrB_Info GrB_Matrix_wait
(
    GrB_Matrix *A
);
```

GrB_Matrix_wait forces all pending operations on a matrix to be completed.

5.10 GraphBLAS matrix and vector import/export

The import/export functions allow the user application to create a GrB_Matrix or GrB_Vector object, and to extract its contents, faster and with less memory overhead than the GrB_*_build and GrB_*_extractTuples functions.

The semantics of import/export are the same as the *move constructor* in C++. On import, the user provides a set of arrays that have been previously allocated via the ANSI C malloc, calloc, or realloc functions (by default), or by the corresponding functions passed to GxB_init. The arrays define the content of the matrix or vector. Unlike GrB_*_build, the GraphBLAS library then takes ownership of the user's input arrays and may either:

- 1. incorporate them into its internal data structure for the new GrB_Matrix or GrB_Vector, potentially creating the GrB_Matrix or GrB_Vector in constant time with no memory copying performed, or
- 2. if the library does not support the import format directly, then it may convert the input to its internal format, and then free the user's input arrays.
- 3. A GraphBLAS implementation may also choose to use a mix of the two strategies.

SuiteSparse:GraphBLAS takes the first approach, and so the import functions always take O(1) time, and require O(1) memory space to be allocated.

Regardless of the method chosen, as listed above, the input arrays are no longer owned by the user application. If A is a GrB_Matrix created by an import, the user input arrays are freed no later than GrB_free(&A), and may be freed earlier, at the discretion of the GraphBLAS library. The data structure of the GrB_Matrix and GrB_Vector remain opaque.

The export of a GrB_Matrix or GrB_Vector is symmetric with the import operation. The export changes the ownership of the arrays, where the GrB_Matrix or GrB_Vector no longer exists when the export completes, and instead the user is returned several arrays that contain the matrix or vector in the requested format. Ownership of these arrays is given to the user application, which is then responsible for freeing them via the ANSI C free function (by default), or by the free_function that was passed in to GxB_init. Alternatively, these arrays can be re-imported into a GrB_Matrix or GrB_Vector, at which point they again become the responsibility of GraphBLAS.

For a matrix export, if the output format matches the current internal format of the matrix then these arrays are returned to the user application in O(1) time and with no memory copying performed. Otherwise, the $\mathtt{GrB_Matrix}$ is first converted into the requested format, and then exported.

The vector import/export methods use a single format for a GrB_Vector. Four different formats are provided for the import/export of a GrB_Matrix. For each format, the Ax array has a C type corresponding to one of the 11 built-in types in GraphBLAS (bool, int*_t, uint*_t, float, and double), or that corresponds with the user-defined type. No typecasting is done on import or export.

The table below lists the methods presented in this section.

method	purpose	Section
GxB_Vector_import	import a vector	5.10.1
<pre>GxB_Vector_export</pre>	export a vector	5.10.2
GxB_Matrix_import_CSR	import a matrix in CSR form	5.10.3
<pre>GxB_Matrix_import_CSC</pre>	import a matrix in CSC form	5.10.4
<pre>GxB_Matrix_import_HyperCSR</pre>	import a matrix in HyperCSR form	5.10.5
<pre>GxB_Matrix_import_HyperCSC</pre>	import a matrix in HyperCSC form	5.10.6
<pre>GxB_Matrix_export_CSR</pre>	export a matrix in CSR form	5.10.7
<pre>GxB_Matrix_export_CSC</pre>	export a matrix in CSC form	5.10.8
<pre>GxB_Matrix_export_HyperCSR</pre>	export a matrix in HyperCSR form	5.10.9
<pre>GxB_Matrix_export_HyperCSC</pre>	export a matrix in HyperCSC form	5.10.10

SPEC: The import/export methods are extensions to the spec. However, they have been implemented in SuiteSparse:GraphBLAS at the request of the GraphBLAS C API Committee, as a prototype for future consideration for inclusion in a future specification. Their calling sequence may change if these functions are added to the specification as GrB_* functions. A GraphBLAS library need not implement these methods in constant time and memory. On import, a library may choose to copy the content of the user arrays into its internal data structure and then free the user arrays. On export, it may chose to malloc the output arrays, fill them with the requested data, and then GrB_free the GraphBLAS object being exported. The semantics of these options are the same as a move constructor; they just take more time and memory. The choice is up to the GraphBLAS implementation since the internal data structure is opaque to the user application.

5.10.1 GxB_Vector_import: import a vector

```
GrB_Info GxB_Vector_import // import a vector in CSC format
    GrB_Vector *v,
                           // vector to create
    GrB_Type type,
                           // type of vector to create
    GrB_Index n,
                           // vector length
                           // number of entries in the vector
    GrB_Index nvals,
    GrB_Index **vi,
                           // indices, size nvals (in sorted order)
                           // values, size nvals
             **vx,
    void
    const GrB_Descriptor desc
                                   // currently unused
);
```

The GxB_Vector_import function is a fast way to construct a GrB_Vector , always taking just O(1) time. Calling GxB_Vector_import with:

```
GxB_Vector_import (&v, type, n, nvals, &vi, &vx, desc);
is identical to the following:
int64_t *Ap = calloc (2, sizeof (int64_t));
Ap [1] = nvals;
GxB_Matrix_import_CSC (&A, type, n, 1, nvals, -1, &Ap, &vi, &vx, desc);
```

except that the latter creates an n-by-1 matrix instead. For the vector import, described here, the first argument is a GrB_Vector. The arguments vi and vx take the place of Ai and Ax, and the Ap array for the CSC matrix import is not provided for a vector import. Refer to the description of GxB_Matrix_import_CSC for details (Section 5.10.4).

If successful, v is created as a n-by-1 vector. Its entries are the row indices given by vi, with the corresponding values in vx. The two pointers vi and vx are returned as NULL, which denotes that they are no longer owned by the user application. They have instead been moved into the new vector v. The row indices in vi must appear in sorted order, and no duplicates can appear. These conditions are not checked, so results are undefined if they are not met exactly. The user application can check the resulting vector v with GxB_print, if desired, which will determine if these conditions hold.

If not successful, v is returned as NULL and vi and vx are not modified.

```
SPEC: GxB_Vector_import is an extension to the spec.
```

5.10.2 GxB_Vector_export: export a vector

```
GrB_Info GxB_Vector_export // export and free a vector
   GrB_Vector *v,
                          // vector to export and free
                        // type of vector exported
   GrB_Type *type,
   GrB_Index *n,
                         // length of the vector
   GrB_Index *nvals,
                       // number of entries in the vector
   GrB_Index **vi,
                         // indices, size nvals
                         // values, size nvals
   void **vx,
   const GrB_Descriptor desc
                                  // currently unused
);
```

The GxB_Vector_export function is a fast way to extract the contents of a GrB_Vector , always taking just O(1) time. Using GxB_Vector_export with:

```
GxB_Vector_export (&v, &type, &n, &nvals, &vi, &vx, desc) ;
is analogous to:
GxB_Matrix_export_CSC (&A, &type, &n, &one, &nvals, &nonempty, &Ap, &Ai, &Ax, desc)
```

if A were an n-by-1 matrix. For the vector export, described here, the first argument is a GrB_Vector. The arguments vi and vx take the place of Ai and Ax, and the Ap array for the CSC matrix export is not returned from a vector export. Refer to the description of GxB_Matrix_export_CSC for details. (Section 5.10.8).

Exporting a vector forces completion of any pending operations on the vector.

If successful, v is returned as NULL, and its contents are returned to the user, with its type, dimension n, and number of entries nvals. A sorted list of row indices of entries that were in v is returned in vi, and the corresponding numerical values are returned in vx. If nvals is zero, the vi and vx arrays are returned as NULL; this is not an error condition.

If not successful, v is unmodified and vi and vx are not modified.

```
SPEC: GxB_Vector_export is an extension to the spec.
```

5.10.3 GxB_Matrix_import_CSR: import a CSR matrix

```
GrB_Info GxB_Matrix_import_CSR
                                    // import a CSR matrix
    GrB_Matrix *A,
                            // handle of matrix to create
    GrB_Type type,
                            // type of matrix to create
    GrB_Index nrows,
                            // matrix dimension is nrows-by-ncols
    GrB_Index ncols,
    GrB_Index nvals,
                            // number of entries in the matrix
    // CSR format:
    int64_t nonempty,
                            // number of rows with at least one entry:
                            // either < 0 if not known, or >= 0 if exact
                            // row "pointers", size nrows+1
    GrB_Index **Ap,
                            // column indices, size nvals
    GrB_Index **Aj,
    void
              **Ax,
                            // values, size nvals
    const GrB_Descriptor desc
                                    // currently unused
);
```

GxB_Matrix_import_CSR imports a matrix from 3 user arrays in CSR format. In the resulting GrB_Matrix A, the CSR format is a matrix with a format (GxB_FORMAT) of GxB_BY_ROW, in standard for instead of hypersparse form (See Section 6.3).

The first four arguments of <code>GxB_Matrix_import_CSR</code> are the same as all four arguments of <code>GrB_Matrix_new</code>, because this function is similar. It creates a new <code>GrB_Matrix A</code>, with the given type and dimensions. The <code>GrB_Matrix A</code> does not exist on input.

Unlike GrB_Matrix_new, this function also populates the new matrix A with the three arrays Ap, Aj and Ax, provided by the user, all of which must have been created with the ANSI C malloc, calloc, or realloc functions (by default), or by the corresponding malloc_function, calloc_function, or realloc_function provided to GxB_init. These arrays define the pattern and values of the new matrix A:

- GrB_Index Ap [nrows+1]; The Ap array is the row "pointer" array. It does not actual contain pointers. More precisely, it is an integer array that defines where the column indices and values appear in Aj and Ax, for each row. The number of entries in row i is given by the expression Ap [i+1] Ap [i].
- GrB_Index Aj [nvals]; The Aj array defines the column indices of entries in each row.

• ctype Aj [nvals]; The Ax array defines the values of entries in each row. It is passed in as a (void *) pointer, but it must point to an array of size nvals values, each of size sizeof(ctype), where ctype is the exact type in C that corresponds to the GrB_Type type parameter. That is, if type is GrB_INT32, then ctype is int32_t. User types may be used, just the same as built-in types.

The content of the three arrays Ap Aj, and Ax is very specific. This content is not checked, since this function takes only O(1) time. Results are undefined if the following specification is not followed exactly.

The column indices of entries in the ith row of the matrix are held in Aj [Ap [i] ... Ap[i+1]], and the corresponding values are held in the same positions in Ax. Column indices must be in the range 0 to ncols-1, and must appear in sorted order within each row. No duplicate column indices may appear in any row. Ap [0] must equal zero, and Ap [nrows] must equal nvals. The Ap array must be of size nrows+1 (or larger), and the Aj and Ax arrays must have size at least nvals.

If nvals is zero, then the content of the Aj and Ax arrays is not accessed and they may be NULL on input (if not NULL, they are still freed and returned as NULL, if the method is successful).

The nonempty parameter is optional. It states the number of rows that have at least one entry: if not known, use -1; if ≥ 0 , it must be exact.

An example of the CSR format is shown below. Consider the following matrix with 10 nonzero entries, and suppose the zeros are not stored.

$$A = \begin{bmatrix} 4.5 & 0 & 3.2 & 0 \\ 3.1 & 2.9 & 0 & 0.9 \\ 0 & 1.7 & 3.0 & 0 \\ 3.5 & 0.4 & 0 & 1.0 \end{bmatrix}$$
 (1)

The Ap array has length 5, since the matrix is 4-by-4. The first entry must always zero, and Ap [5] = 10 is the number of entries. The content of the arrays is shown below:

Spaces have been added to the Ap array, just for illustration. Row zero is in Aj [0..1] (column indices) and Ax [0..1] (values), starting at

Ap [0] = 0 and ending at Ap [0+1]-1 = 1. The list of column indices of row one is at Aj [2..4] and row two is in Aj [5..6]. The last row (three) appears Aj [7..9], because Ap [3] = 7 and Ap [4]-1 = 10-1 = 9. The corresponding numerical values appear in the same positions in Ax.

To iterate over the rows and entries of this matrix, the following code can be used:

On successful creation of A, the three pointers Ap, Aj, and Ax are set to NULL on output. This denotes to the user application that it is no longer responsible for freeing these arrays. Internally, GraphBLAS has moved these arrays into its internal data structure. They will eventually be freed no later than when the user does GrB_free(&A), but they may be freed or resized later, if the matrix changes.

If the matrix A is later exported in CSR form, and GraphBLAS has not yet reallocated these arrays, then these same three arrays are returned to the user by GxB_Matrix_export_CSR (see Section 5.10.7). If an export is performed, the freeing of these three arrays again becomes the responsibility of the user application.

The $GxB_Matrix_import_CSR$ function will rarely fail, since it allocates just O(1) space. If it does fail, it returns $GrB_0UT_0F_MEMORY$, and it leaves the three user arrays unmodified. They are still owned by the user application, which is eventually responsible for freeing them with free(Ap), etc.

SPEC: GxB_Matrix_import_CSR is an extension to the spec.

5.10.4 GxB_Matrix_import_CSC: import a CSC matrix

```
GrB_Info GxB_Matrix_import_CSC
                                    // import a CSC matrix
    GrB_Matrix *A,
                            // handle of matrix to create
    GrB_Type type,
                            // type of matrix to create
    GrB_Index nrows,
                            // matrix dimension is nrows-by-ncols
    GrB_Index ncols,
   GrB_Index nvals,
                            // number of entries in the matrix
    // CSC format:
    int64_t nonempty,
                            // number of columns with at least one entry:
                            // either < 0 if not known, or >= 0 if exact
                            // column "pointers", size ncols+1
    GrB_Index **Ap,
                            // row indices, size nvals
    GrB_Index **Ai,
    void
              **Ax.
                            // values, size nvals
                                    // currently unused
    const GrB_Descriptor desc
);
```

GxB_Matrix_import_CSC imports a matrix from 3 user arrays in CSC format. The GrB_Matrix A is created in the CSC format, which is a GxB_FORMAT of GxB_BY_COL. The arguments are identical to GxB_Matrix_import_CSR, except for how the 3 user arrays are interpreted. The column "pointer" array has size ncols+1. The row indices of the columns are in Ai, and must appear in ascending order in each column. The corresponding numerical values are held in Ax. The row indices of column j are held in Ai [Ap [j]...Ap [j+1]-1, and the corresponding numerical values are in the same locations in Ax.

The nonempty parameter is optional. It states the number of columns that have at least one entry: if not known, use -1; if ≥ 0 , it must be exact.

The same matrix from Equation 1 in the last section (repeated here):

$$A = \begin{bmatrix} 4.5 & 0 & 3.2 & 0 \\ 3.1 & 2.9 & 0 & 0.9 \\ 0 & 1.7 & 3.0 & 0 \\ 3.5 & 0.4 & 0 & 1.0 \end{bmatrix}$$
 (2)

is held in CSC form as follows:

```
int64_t Ap [] = { 0, 3, 6, 8, 10 };
int64_t Ai [] = { 0, 1, 3, 1, 2, 3, 0, 2, 1, 3 };
double Ax [] = { 4.5, 3.1, 3.5, 2.9, 1.7, 0.4, 3.2, 3.0, 0.9, 1.0 };
```

That is, the row indices of column 1 (the second column) are in Ai [3..5], and the values in the same place in Ax, since Ap [1] = 3 and Ap [2]-1 = 5.

To iterate over the columns and entries of this matrix, the following code can be used:

The method is identical to GxB_Matrix_import_CSR; just the format is different. That is, if the method is successful, the 3 user arrays are imported into the new GrB_Matrix A, with the given type and dimensions, and returned as NULL pointers to the user application.

If nvals is zero, then the content of the Ai and Ax arrays is not accessed and they may be NULL on input (if not NULL, they are still freed and returned as NULL, if the method is successful).

SPEC: GxB_Matrix_import_CSC is an extension to the spec.

5.10.5 GxB_Matrix_import_HyperCSR: import a HyperCSR matrix

```
GrB_Info GxB_Matrix_import_HyperCSR
                                        // import a hypersparse CSR matrix
    GrB_Matrix *A,
                            // handle of matrix to create
    GrB_Type type,
                            // type of matrix to create
    GrB_Index nrows,
                            // matrix dimension is nrows-by-ncols
    GrB_Index ncols,
    GrB_Index nvals,
                            // number of entries in the matrix
    // hypersparse CSR format:
    int64_t nonempty,
                            // number of rows in Ah with at least one entry,
                            // either < 0 if not known, or >= 0 if exact
                           // number of rows in Ah list
    GrB_Index nvec,
   GrB_Index **Ah,
                            // list of size nvec of rows that appear in A
   GrB_Index **Ap,
                            // row "pointers", size nvec+1
   GrB_Index **Aj,
                            // column indices, size nvals
    void
              **Ax,
                            // values, size nvals
    const GrB_Descriptor desc
                                    // currently unused
);
```

GxB_Matrix_import_HyperCSR imports a matrix in hypersparse CSR format in O(1) time. In the hypersparse format, the Ap array itself becomes sparse, if the matrix has rows that are completely empty. An array Ah of size nvec provides a list of rows that appear in the data structure. For example, consider Equation 3, which is a sparser version of the matrix in Equation 1. Row 2 and column 1 of this matrix are all zero.

$$A = \begin{bmatrix} 4.5 & 0 & 3.2 & 0 \\ 3.1 & 0 & 0 & 0.9 \\ 0 & 0 & 0 & 0 \\ 3.5 & 0 & 0 & 1.0 \end{bmatrix}$$
 (3)

The conventional CSR format would appear as follows. Since the third row (row 2) is all zero, accessing Ai [Ap [2] ... Ap [3]-1] gives an empty set ([2..1]), and the number of entries in this row is Ap [i+1] - Ap [i] = Ap [3] - Ap [2] = 0.

```
int64_t Ap [] = { 0, 2, 2, 4, 5 };
int64_t Aj [] = { 0, 2, 0, 3, 0 3 }
double Ax [] = { 4.5, 3.2, 3.1, 0.9, 3.5, 1.0 };
```

A hypersparse CSR format for this same matrix would discard these duplicate integers in Ap. Doing so requires another array, Ah, that keeps track of the rows that appear in the data structure.

Note that the Aj and Ax arrays are the same in the standard and hypersparse CSR formats. The row indices in Ah must appear in ascending order, and no duplicates can appear. To iterate over this data structure:

This is more complex than the standard CSR format, but it requires at most O(e) space, where A is m-by-n with $e = \mathtt{nvals}$ entries. The standard CSR format requires O(m+e) space. If e << m, then the size m+1 of Ap can dominate the memory required. In the hypersparse form, Ap takes on size $\mathtt{nvec+1}$, and Ah has size \mathtt{nvec} , where \mathtt{nvec} is the number of rows that appear in the data structure. The standard CSR format can be viewed as a dense array (of size \mathtt{nrows}) of sparse row vectors. By contrast, the hypersparse CSR format is a sparse array (of size \mathtt{nvec}) of sparse row vectors.

The import takes O(1) time. If successful, the four arrays Ah, Ap, Aj, and Ax are returned as NULL, and the hypersparse GrB_Matrix A is created.

If nvals is zero, then the content of the Aj and Ax arrays is not accessed and they may be NULL on input (if not NULL, they are still freed and returned as NULL, if the method is successful). The nonempty parameter is optional. It states the number of rows that have at least one entry: if not known, use -1; if ≥ 0 , it must be exact.

SPEC: GxB_Matrix_import_HyperCSR is an extension to the spec.

5.10.6 GxB_Matrix_import_HyperCSC: import a HyperCSC matrix

```
GrB_Info GxB_Matrix_import_HyperCSC
                                        // import a hypersparse CSC matrix
    GrB_Matrix *A,
                           // handle of matrix to create
                           // type of matrix to create
    GrB_Type type,
    GrB_Index nrows,
                           // matrix dimension is nrows-by-ncols
    GrB_Index ncols,
    GrB_Index nvals,
                           // number of entries in the matrix
    // hypersparse CSC format:
    int64_t nonempty,
                           // number of columns in Ah with at least one entry,
                           // either < 0 if not known, or >= 0 if exact
                           // number of columns in Ah list
    GrB_Index nvec,
                           // list of size nvec of columns that appear in A
    GrB_Index **Ah,
    GrB_Index **Ap,
                           // column "pointers", size nvec+1
   GrB_Index **Ai,
                           // row indices, size nvals
                           // values, size nvals
    void
             **Ax,
    const GrB_Descriptor desc
                                    // currently unused
);
```

 $GxB_Matrix_import_HyperCSC$ imports a matrix in hypersparse CSC format in O(1) time. It is identical to $GxB_Matrix_import_HyperCSR$, except for the data structure defined by the four arrays Ah, Ap, Ai, and Ax. It is a sparse array of size nvec of sparse column vectors. The following code iterates over the matrix:

The nonempty parameter is optional. It states the number of columns that have at least one entry: if not known, use -1; if ≥ 0 , it must be exact.

SPEC: GxB_Matrix_import_HyperCSC is an extension to the spec.

5.10.7 GxB_Matrix_export_CSR: export a CSR matrix

```
GrB_Info GxB_Matrix_export_CSR // export and free a CSR matrix
    GrB_Matrix *A,
                            // handle of matrix to export and free
    GrB_Type *type,
                            // type of matrix exported
    GrB_Index *nrows,
                            // matrix dimension is nrows-by-ncols
    GrB_Index *ncols,
    GrB_Index *nvals,
                            // number of entries in the matrix
    // CSR format:
    int64_t *nonempty,
                            // number of rows with at least one entry
   GrB_Index **Ap,
                            // row "pointers", size nrows+1
    GrB_Index **Aj,
                            // column indices, size nvals
                            // values, size nvals
    void
              **Ax,
    const GrB_Descriptor desc
                                    // currently unused
);
```

GxB_Matrix_export_CSR exports a matrix in CSR form:

On successful output, the GrB_Matrix A is freed, and A is returned as NULL. Its type is returned in the type parameter, its dimensions in nrows and ncols, its number of entries in nvals, and the CSR format is in the three arrays Ap, Aj, and Ax. If nvals is zero, the Aj and Ax arrays are returned as NULL; this is not an error, and GxB_Matrix_import_CSR also allows these two arrays to be NULL on input when nvals is zero. After a successful export, the user application is responsible for freeing these three arrays via free (or the free function passed to GxB_init). The CSR format is described in Section 5.10.3.

This method takes O(1) time if the matrix is already in standard (non-hypersparse) CSR format internally. If it is in hypersparse CSR form, the export must first convert the matrix to standard CSR form, taking O(m) time and memory, where m = nrows. If the matrix is in CSC format, it is first transposed to convert it to CSR format, and then exported. This takes O(m+n+e) or $O(m+e\log e)$ time and memory, whichever is less, where n = ncols and e = nvals.

SPEC: GxB_Matrix_export_CSR is an extension to the spec.

5.10.8 GxB_Matrix_export_CSC: export a CSC matrix

```
GrB_Info GxB_Matrix_export_CSC // export and free a CSC matrix
    GrB_Matrix *A,
                            // handle of matrix to export and free
    GrB_Type *type,
                            // type of matrix exported
    GrB_Index *nrows,
                            // matrix dimension is nrows-by-ncols
    GrB_Index *ncols,
   GrB_Index *nvals,
                           // number of entries in the matrix
    // CSC format:
                           // number of columns with at least one entry
    int64_t *nonempty,
    GrB_Index **Ap,
                           // column "pointers", size ncols+1
   GrB_Index **Ai,
                           // row indices, size nvals
                            // values, size nvals
    void
              **Ax,
    const GrB_Descriptor desc
                                    // currently unused
);
```

GxB_Matrix_export_CSC exports a matrix in CSC form:

On successful output, the GrB_Matrix A is freed, and A is returned as NULL. Its type is returned in the type parameter, its dimensions in nrows and ncols, its number of entries in nvals, and the CSC format is in the three arrays Ap, Ai, and Ax. If nvals is zero, the Ai and Ax arrays are returned as NULL; this is not an error, and GxB_Matrix_import_CSC also allows these two arrays to be NULL on input when nvals is zero. After a successful export, the user application is responsible for freeing these three arrays via free (or the free function passed to GxB_init). The CSC format is described in Section 5.10.4.

This method takes O(1) time if the matrix is already in standard (non-hypersparse) CSC format internally. If it is in hypersparse CSC form, the export must first convert the matrix to standard CSC form, taking O(n) time and memory, where n = ncols. If the matrix is in CSR format, it is first transposed to convert it to CSC format, and then exported. This takes O(m+n+e) or $O(n+e\log e)$ time and memory, whichever is less, where m = nrows and e = nvals.

SPEC: GxB_Matrix_export_CSC is an extension to the spec.

5.10.9 GxB_Matrix_export_HyperCSR: export a HyperCSR matrix

```
GrB_Info GxB_Matrix_export_HyperCSR // export and free a hypersparse CSR matrix
    GrB_Matrix *A,
                            // handle of matrix to export and free
    GrB_Type *type,
                            // type of matrix exported
    GrB_Index *nrows,
                            // matrix dimension is nrows-by-ncols
    GrB_Index *ncols,
    GrB_Index *nvals,
                            // number of entries in the matrix
    // hypersparse CSR format:
    int64_t *nonempty,
                            // number of rows in Ah with at least one entry
   GrB_Index *nvec,
                            // number of rows in Ah list
    GrB_Index **Ah,
                            // list of size nvec of rows that appear in A
                            // row "pointers", size nvec+1
   GrB_Index **Ap,
    GrB_Index **Aj,
                            // column indices, size nvals
    void
              **Ax,
                            // values, size nvals
    const GrB_Descriptor desc
                                    // currently unused
);
```

GxB_Matrix_export_HyperCSR exports a matrix in CSR form:

On successful output, the GrB_Matrix A is freed, and A is returned as NULL. Its type is returned in the type parameter, its dimensions in nrows and ncols, its number of entries in nvals, and the number of non-empty rows in nvec. The hypersparse CSR format is in the four arrays Ah, Ap, Aj, and Ax. If nvals is zero, the Aj and Ax arrays are returned as NULL; this is not an error. After a successful export, the user application is responsible for freeing these three arrays via free (or the free function passed to GxB_init). The hypersparse CSR format is described in Section 5.10.5.

This method takes O(1) time if the matrix is already in hypersparse CSR format internally. If it is in standard CSR form, the export must first convert the matrix to hypersparse CSR form, taking O(m) time and memory, where m = nrows. If the matrix is in CSC format, it is first transposed to convert it to hypersparse CSR format, and then exported. If in standard CSC form, the transpose takes O(m + n + e) or $O(n + e \log e)$ time and memory, whichever is less. If in hypersparse CSC format, it takes $O(e \log e)$ time.

SPEC: GxB_Matrix_export_HyperCSR is an extension to the spec.

5.10.10 GxB_Matrix_export_HyperCSC: export a HyperCSC matrix

```
GrB_Info GxB_Matrix_export_HyperCSC // export and free a hypersparse CSC matrix
    GrB_Matrix *A,
                            // handle of matrix to export and free
    GrB_Type *type,
                            // type of matrix exported
    GrB_Index *nrows,
                            // matrix dimension is nrows-by-ncols
    GrB_Index *ncols,
    GrB_Index *nvals,
                            // number of entries in the matrix
    // hypersparse CSC format:
    int64_t *nonempty,
                            // number of columns in Ah with at least one entry
   GrB_Index *nvec,
                            // number of columns in Ah list
    GrB_Index **Ah,
                            // list of size nvec of columns that appear in A
                            // columns "pointers", size nvec+1
    GrB_Index **Ap,
    GrB_Index **Ai,
                            // row indices, size nvals
                            // values, size nvals
    void
              **Ax,
    const GrB_Descriptor desc
                                    // currently unused
);
```

GxB_Matrix_export_HyperCSC exports a matrix in CSC form:

```
GxB_Matrix_export_HyperCSC (&A, &type, &nrows, &ncols, &nvals, &nonempty, &nvec, &Ah, &Ap, &Ai, &Ax, desc);
```

On successful output, the GrB_Matrix A is freed, and A is returned as NULL. Its type is returned in the type parameter, its dimensions in nrows and ncols, its number of entries in nvals, and the number of non-empty rows in nvec. The hypersparse CSC format is in the four arrays Ah, Ap, Ai, and Ax. If nvals is zero, the Ai and Ax arrays are returned as NULL; this is not an error. After a successful export, the user application is responsible for freeing these three arrays via free (or the free function passed to GxB_init). The hypersparse CSC format is described in Section 5.10.6.

This method takes O(1) time if the matrix is already in hypersparse CSR format internally. If it is in standard CSR form, the export must first convert the matrix to hypersparse CSR form, taking O(m) time and memory, where m = nrows. If the matrix is in CSC format, it is first transposed to convert it to hypersparse CSR format, and then exported. If in standard CSC form, the transpose takes O(m + n + e) or $O(n + e \log e)$ time and memory, whichever is less. If in hypersparse CSC format, it takes $O(e \log e)$ time.

SPEC: GxB_Matrix_export_HyperCSC is an extension to the spec.

5.11 GraphBLAS descriptors: GrB_Descriptor

A GraphBLAS descriptor modifies the behavior of a GraphBLAS operation. If the descriptor is GrB_NULL, defaults are used.

The access to these parameters and their values is governed by two enum types, GrB_Desc_Field and GrB_Desc_Value:

```
#define GxB_NTHREADS 5 // for both GrB_Desc_field and GxB_Option_field
#define GxB_CHUNK 7
typedef enum
    GrB_OUTP = 0, // descriptor for output of a method
                  // descriptor for the mask input of a method
    GrB\_MASK = 1,
    GrB_INPO = 2, // descriptor for the first input of a method
    GrB_INP1 = 3, // descriptor for the second input of a method
    GxB_DESCRIPTOR_NTHREADS = GxB_NTHREADS,
                                              // number of threads to use
    GxB_DESCRIPTOR_CHUNK = GxB_CHUNK,
                                       // chunk size for small problems
    GxB_AxB_METHOD = 1000, // descriptor for selecting C=A*B algorithm
}
GrB_Desc_Field ;
typedef enum
{
    // for all GrB_Descriptor fields:
                       // default behavior of the method
   GxB_DEFAULT = 0,
   // for GrB_OUTP only:
    GrB_REPLACE = 1,
                       // clear the output before assigning new values to it
    // for GrB_MASK only:
   GrB\_COMP = 2,
                       // use the complement of the mask
    GrB_STRUCTURE = 4, // use the structure of the mask
    // for GrB_INPO and GrB_INP1 only:
    GrB_TRAN = 3,
                        // use the transpose of the input
    // for GxB_AxB_METHOD only:
    GxB_AxB_GUSTAVSON = 1001,
                               // gather-scatter saxpy method
    GxB_AxB_HEAP
                     = 1002,
                               // heap-based saxpy method
    GxB_AxB_DOT
                     = 1003, // dot product
                     = 1004, // hash-based saxpy method
    GxB_AxB_HASH
    GxB_AxB_SAXPY
                     = 1005
                               // saxpy method (any kind)
GrB_Desc_Value ;
```

SPEC: GxB_DEFAULT, GxB_NTHRADS, GxB_CHUNK, GxB_AxB_METHOD, and GxB_AxB_* are extensions to the spec.

The internal representation is opaque to the user, but in this User Guide the five descriptor fields of a descriptor desc are illustrated as an array of five items, as described in the list below. The underlying implementation need not be an array:

• desc [GrB_0UTP] is a parameter that modifies the output of a Graph-BLAS operation. Currently, there are two possible settings. In the default case, the output is not cleared, and $\mathbf{C}\langle\mathbf{M}\rangle=\mathbf{Z}=\mathbf{C}\odot\mathbf{T}$ is computed as-is, where \mathbf{T} is the results of the particular GraphBLAS operation.

In the non-default case, $\mathbf{Z} = \mathbf{C} \odot \mathbf{T}$ is first computed, using the results of \mathbf{T} and the accumulator \odot . After this is done, if the $\mathtt{GrB_OUTP}$ descriptor field is set to $\mathtt{GrB_REPLACE}$, then the output is cleared of its entries. Next, the assignment $\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{Z}$ is performed.

• desc [GrB_MASK] is a parameter that modifies the Mask, even if the mask is not present.

If this parameter is set to its default value, and if the mask is not present (Mask==NULL) then implicitly Mask(i,j)=1 for all i and j. If the mask is present then Mask(i,j)=1 means that C(i,j) is to be modified by the $C\langle M\rangle=Z$ update. Otherwise, if Mask(i,j)=0, then C(i,j) is not modified, even if Z(i,j) is an entry with a different value; that value is simply discarded.

If the desc [GrB_MASK] parameter is set to GrB_COMP, then the use of the mask is complemented. In this case, if the mask is not present (Mask==NULL) then implicitly Mask(i,j)=0 for all i and j. This means that none of C is modified and the entire computation of Z might as well have been skipped. That is, a complemented empty mask means no modifications are made to the output object at all, except perhaps to clear it in accordance with the GrB_OUTP descriptor. With a complemented mask, if the mask is present then Mask(i,j)=0 means that C(i,j) is to be modified by the $C\langle M \rangle = Z$ update. Otherwise, if Mask(i,j)=1, then C(i,j) is not modified, even if Z(i,j) is an entry with a different value; that value is simply discarded.

If the desc [GrB_MASK] parameter is set to GrB_STRUCTURE, then the values of the mask are ignored, and just the pattern of the entries is used. Any entry M(i,j) in the pattern is treated as if it were true.

The GrB_COMP and GrB_STRUCTURE settings can be combined, either by setting the mask option twice (once with each value), or by setting the mask option to GrB_COMP+GrB_STRUCTURE (the latter is an extension to the spec).

Using a parameter to complement the Mask is very useful because constructing the actual complement of a very sparse mask is impossible since it has too many entries. If the number of places in C that should be modified is very small, then use a sparse mask without complementing it. If the number of places in C that should be protected from modification is very small, then use a sparse mask to indicate those places, and use a descriptor GrB_MASK that complements the use of the mask.

• desc [GrB_INP0] and desc [GrB_INP1] modify the use of the first and second input matrices A and B of the GraphBLAS operation.

If the desc [GrB_INPO] is set to GrB_TRAN, then A is transposed before using it in the operation. Likewise, if desc [GrB_INP1] is set to GrB_TRAN, then the second input, typically called B, is transposed.

Vectors are never transposed via the descriptor. If a method's first parameter is a matrix and the second a vector, then desc [GrB_INP0] modifies the matrix parameter and desc [GrB_INP1] is ignored. If a method's first parameter is a vector and the second a matrix, then desc [GrB_INP1] modifies the matrix parameter and desc [GrB_INP0] is ignored.

To clarify this in each function, the inputs are labeled as first input: and second input: in the function signatures.

- desc [GxB_AxB_METHOD] suggests the method that should be used to compute C=A*B. All the methods compute the same result, except they may have different floating-point roundoff errors. This descriptor should be considered as a hint; SuiteSparse:GraphBLAS is free to ignore it. The current version always follows the hint, however.
 - GxB_DEFAULT means that a method is selected automatically.

- GxB_AxB_SAXPY: select any saxpy-based method: GxB_AxB_GUSTAVSON,
 GxB_AxB_HEAP, and/or GxB_AxB_HASH, or any mix of the three, in contrast to the dot-product method.
- GxB_AxB_GUSTAVSON: an extended version of Gustavson's method [Gus78], which is a very good general-purpose method, but sometimes the workspace can be too large. Assuming all matrices are stored by column, it computes C(:,j)=A*B(:,j) with a sequence of saxpy operations (C(:,j)+=A(:,k)*B(k:,j) for each nonzero B(k,j)). Each internal thread requires workspace of size m, to the number of rows of C, which is not suitable if the matrices are extremely sparse or if there are many threads. If all matrices are stored by row, then it computes C(i,:)=A(i,:)*B in a sequence of sparse saxpy operations, and using workspace of size n per thread, corresponding to the number of columns of C.
- GxB_AxB_HEAP: no longer appears in v3.2.0, but will likely be reintroduced in a future version. This is silently replaced with GxB_AxB_HASH.
- GxB_AxB_HASH: a hash-based method, based on [NMAB18]. Very efficient for hypersparse matrices, matrix-vector-multiply, and when |B| is small.
- GxB_AxB_DOT: computes C(i,j)=A(i,:)*B(j,:)', for each entry C(i,j). If the mask is present and not complemented, only entries for which M(i,j)=1 are computed. This is a very specialized method that works well only if the mask is present, very sparse, and not complemented, or when C is tiny. For example, it works very well when A and B are tall and thin, and C<M>=A*B' or C=A*B' are computed. These expressions assume all matrices are in CSR format. If in CSC format, then the dot-product method used for A'*B. The method is impossibly slow if C is large and the mask is not present, since it takes Ω(mn) time if C is m-by-n in that case. It does not use any workspace at all. Since it uses no workspace, it can work very well for extremely sparse or hypersparse matrices, when the mask is present and not complemented.

5.11.1 GrB_Descriptor_new: create a new descriptor

```
GrB_Info GrB_Descriptor_new  // create a new descriptor
(
    GrB_Descriptor *descriptor // handle of descriptor to create
);
```

GrB_Descriptor_new creates a new descriptor, with all fields set to their defaults (output is not replaced, mask is not complemented, neither input matrix is transposed, and the method used in C=A*B is selected automatically).

5.11.2 GrB_Descriptor_set: set a parameter in a descriptor

GrB_Descriptor_set sets a descriptor field (GrB_OUTP, GrB_MASK, GrB_INPO, GrB_INP1, or GxB_AxB_METHOD) to a particular value (GxB_DEFAULT, GrB_COMP, GrB_STRUCTURE, GrB_COMP+GrB_STRUCTURE, GrB_TRAN, GrB_REPLACE, GxB_AxB_GUSTAVSON, GxB_AxB_HEAP, GxB_AxB_HASH, GxB_AxB_SAXPY, or GxB_AxB_DOT).

Descriptor	Default	Non-default		
field				
GrB_OUTP	GxB_DEFAULT: The output matrix is	GrB_REPLACE: After computing		
	not cleared. The operation computes	$\mathbf{Z} = \mathbf{C} \odot \mathbf{T}$, the output \mathbf{C} is		
	$\mathbf{C}\langle\mathbf{M} angle = \mathbf{C}\odot\mathbf{T}.$	cleared of all entries. Then		
		$ \mathbf{C}\langle\mathbf{M}\rangle = \mathbf{Z}$ is performed.		
GrB_MASK	GxB_DEFAULT: The Mask is not com-	GrB_COMP: The Mask is comple-		
	plemented. Mask(i,j)=1 means the	mented. Mask(i,j)=0 means the		
	value C_{ij} can be modified by the op-	value C_{ij} can be modified by the		
	eration, while Mask(i,j)=0 means	operation, while Mask(i,j)=1		
the value C_{ij} shall not be mo		means the value C_{ij} shall not be		
	by the operation.	modified by the operation.		
		GrB_STRUCTURE: The values of		
		the Mask are ignored. If		
		Mask(i,j) is an entry in the		
		Mask matrix, it is treated as if		
		Mask(i,j)=1. The two options		
		GrB_COMP and GrB_STRUCTURE		
		can be combined.		
GrB_INPO	GxB_DEFAULT: The first input is not	GrB_TRAN: The first input is		
	transposed prior to using it in the	transposed prior to using it in		
	operation.	the operation. Only matrices are		
		transposed, never vectors.		
GrB_INP1	GxB_DEFAULT: The second input is	GrB_TRAN: The second input is		
	not transposed prior to using it in	transposed prior to using it in		
	the operation.	the operation. Only matrices are		
		transposed, never vectors.		
GrB_AxB_METHOD	GxB_DEFAULT: The method used for	GxB_AxB_method: The selected		
	computing C=A*B is selected auto-	method is used to compute		
	matically.	C=A*B.		

5.11.3 GxB_Desc_set: set a parameter in a descriptor

GxB_Desc_set is identical to GrB_Descriptor_set, except that the type of the third parameter can vary with the field. All descriptor fields are currently of type GrB_Desc_Value, so currently this function is identical in all ways to GrB_Descriptor_set, except for the name of the function. Future versions of this function will allow for arbitrary types of the third parameter, depending on the field. For a simpler-to-use alternative, see GxB_set described in Section 6.

```
SPEC: GxB_Desc_set is an extension to the spec.
```

5.11.4 GxB_Desc_get: get a parameter from a descriptor

GxB_Desc_get returns the value of a single field in a descriptor. The type of the third parameter is a pointer to a variable type, whose type depends on the field. Currently, all descriptor values are of type GrB_Desc_Value, so this third parameter is a pointer to a scalar value of type GrB_Desc_Value. For a simpler-to-use alternative, see GxB_get described in Section 6.

```
SPEC: GxB_Desc_get is an extension to the spec.
```

5.11.5 GrB_Descriptor_free: free a descriptor

 ${\tt GrB_Descriptor_free}$ frees a descriptor. Either usage:

```
GrB_Descriptor_free (&descriptor) ;
GrB_free (&descriptor) ;
```

frees the descriptor and sets descriptor to NULL. It safely does nothing if passed a NULL handle, or if descriptor == NULL on input.

5.11.6 GrB_DESC_*: predefined descriptors

Version 1.3 of the GraphBLAS C API Specification adds predefined descriptors, and these have been added as of v3.2.0 of SuiteSparse:GraphBLAS. They are listed in the table below. These descriptors may not be modified or freed. Attempts to modify them result in an error (GrB_INVALID_VALUE); attempts to free them are silently ignored. GrB_NULL is the default descriptor, with all settings at their defaults: OUTP: do not replace the output, MASK: mask is valued and not complemented, INPO: first input not transposed, and INP1: second input not transposed.

Descriptor	OUTP	MASK	MASK	INPO	INP1
		structural	complement		
GrB_NULL	-	-	-	-	-
GrB_DESC_T1	-	-	-	-	GrB_TRAN
GrB_DESC_TO	-	-	-	GrB_TRAN	-
GrB_DESC_TOT1	-	-	-	GrB_TRAN	GrB_TRAN
GrB_DESC_C	-	-	GrB_COMP	-	-
GrB_DESC_CT1	-	-	GrB_COMP	-	GrB_TRAN
GrB_DESC_CTO	-	-	GrB_COMP	GrB_TRAN	-
GrB_DESC_CTOT1	-	-	GrB_COMP	GrB_TRAN	GrB_TRAN
GrB_DESC_S	-	GrB_STRUCTURE	_	-	-
GrB_DESC_ST1	-	GrB_STRUCTURE	_	-	GrB_TRAN
GrB_DESC_STO	-	GrB_STRUCTURE	_	GrB_TRAN	_
GrB_DESC_STOT1	-	GrB_STRUCTURE	-	GrB_TRAN	GrB_TRAN
GrB_DESC_SC	-	GrB_STRUCTURE	GrB_COMP	-	-
GrB_DESC_SCT1	-	GrB_STRUCTURE	GrB_COMP	-	GrB_TRAN
GrB_DESC_SCTO	-	GrB_STRUCTURE	GrB_COMP	GrB_TRAN	-
GrB_DESC_SCTOT1	-	<pre>Grb_STRUCTURE</pre>	GrB_COMP	GrB_TRAN	GrB_TRAN
GrB_DESC_R	GrB_REPLACE	-	-	-	
GrB_DESC_RT1	GrB_REPLACE	-	-	-	GrB_TRAN
GrB_DESC_RTO	GrB_REPLACE	-	-	GrB_TRAN	-
GrB_DESC_RTOT1	GrB_REPLACE	-	-	GrB_TRAN	GrB_TRAN
GrB_DESC_RC	GrB_REPLACE	-	GrB_COMP	-	-
GrB_DESC_RCT1	GrB_REPLACE	-	GrB_COMP	-	GrB_TRAN
GrB_DESC_RCTO	GrB_REPLACE	-	GrB_COMP	GrB_TRAN	-
GrB_DESC_RCTOT1	GrB_REPLACE	-	GrB_COMP	GrB_TRAN	GrB_TRAN
GrB_DESC_RS	GrB_REPLACE	GrB_STRUCTURE	-	-	-
GrB_DESC_RST1	GrB_REPLACE	GrB_STRUCTURE	-	-	GrB_TRAN
GrB_DESC_RSTO	GrB_REPLACE	GrB_STRUCTURE	-	GrB_TRAN	-
GrB_DESC_RSTOT1	GrB_REPLACE	GrB_STRUCTURE	-	GrB_TRAN	GrB_TRAN
GrB_DESC_RSC	GrB_REPLACE	GrB_STRUCTURE	GrB_COMP	_	_
GrB_DESC_RSCT1	GrB_REPLACE	GrB_STRUCTURE	GrB_COMP	-	GrB_TRAN
GrB_DESC_RSCTO	GrB_REPLACE	GrB_STRUCTURE	GrB_COMP	GrB_TRAN	-
GrB_DESC_RSCTOT1	GrB_REPLACE	GrB_STRUCTURE	GrB_COMP	GrB_TRAN	GrB_TRAN

5.12 GrB_free: free any GraphBLAS object

Each of the ten objects has GrB_*_new and GrB_*_free methods that are specific to each object. They can also be accessed by a generic function, GrB_free, that works for all ten objects. If G is any of the ten objects, the statement

```
GrB_free (&G) ;
```

frees the object and sets the variable G to NULL. It is safe to pass in a NULL handle, or to free an object twice:

```
GrB_free (NULL); // SuiteSparse:GraphBLAS safely does nothing GrB_free (&G); // the object G is freed and G set to NULL GrB_free (&G); // SuiteSparse:GraphBLAS safely does nothing
```

However, the following sequence of operations is not safe. The first two are valid but the last statement will lead to undefined behavior.

```
H = G;  // valid; creates a 2nd handle of the same object GrB\_free (&G);  // valid; G is freed and set to NULL; H now undefined GrB\_some\_method (H);  // not valid; H is undefined
```

Some objects are predefined, such as the built-in types. If a user application attempts to free a built-in object, SuiteSparse:GraphBLAS will safely do nothing. In all cases, the GrB_free function in SuiteSparse:GraphBLAS always returns GrB_SUCCESS.

6 SuiteSparse:GraphBLAS Options

SPEC: GxB_set and GxB_get are extensions to the specification.

SuiteSparse:GraphBLAS includes two type-generic methods, GxB_set and GxB_get, that set and query various options and parameters settings, including a generic way to set values in the GrB_Descriptor object. Using these methods, the user application can provide hints to SuiteSparse:GraphBLAS on how it should store and operate on its matrices. These hints have no effect on the results of any GraphBLAS operation (except perhaps floating-point roundoff differences), but they can have a great impact on the amount of time or memory taken.

- GxB_set (field, value) provides hints to SuiteSparse:GraphBLAS on how it should store all matrices created after calling this function: by row, by column, and whether or not to use a *hypersparse* format [BG08, BG12]. These are global options that modify all matrices created after calling this method.
- GxB_set (GrB_Matrix A, field, value) provides hints to SuiteSparse: GraphBLAS on how to store a particular matrix. This method allows SuiteSparse:GraphBLAS to transform a specific matrix from one format to another. The format has no effect on the result computed by GraphBLAS; it only affects the time and memory taken to do the computations.
- GxB_set (GrB_Descriptor desc, field, value) is another way to set the value of a field in a GrB_Descriptor. It is identical to GrB_Descriptor_set, just with a generic name.

The GxB_get method queries a GrB_Descriptor, a GrB_Matrix, or the global options.

- GxB_get (field, &value) retrieves the current value of a global option.
- GxB_get (GrB_Matrix A, field, &value) retrieves the current value of an option from a particular matrix A.
- GxB_get (GrB_Descriptor desc, field, &value) retrieves the value of a field in a descriptor.

6.1 OpenMP parallelism

SuiteSparse:GraphBLAS Version 3 is a parallel library, based on OpenMP. By default, all GraphBLAS operations will use up to the maximum number of threads specified by the omp_get_max_threads OpenMP function. For small problems, GraphBLAS may choose to use fewer threads, using two parameters: the maximum number of threads to use (which may differ from the omp_get_max_threads value), and a parameter called the chunk. Suppose work is a measure of the work an operation needs to perform (say the number of entries in the two input matrices for GrB_eWiseAdd). No more than floor(work/chunk) threads will be used (or one thread if the ratio is less than 1).

The default **chunk** value is 65,536, but this may change in future versions, or it may be modified when GraphBLAS is installed on a particular machine. Both parameters can be set in two ways:

• Globally: If the following methods are used, then all subsequent Graph-BLAS operations will use these settings. Note the typecast, (double) chunk. This is necessary if a literal constant such as 20000 is passed as this argument. The type of the constant must be double.

```
int nthreads_max = 40 ;
GxB_set (GxB_NTHREADS, nthreads_max) ;
GxB_set (GxB_CHUNK, (double) 20000) ;
```

• Per operation: Most GraphBLAS operations take a GrB_Descriptor input, and this can be modified to set the number of threads and chunk size for the operation that uses this descriptor. Note that chunk is a double.

```
GrB_Descriptor desc ;
GrB_Descriptor_new (&desc)
int nthreads_max = 40 ;
GxB_set (desc, GxB_NTHREADS, nthreads_max) ;
double chunk = 20000 ;
GxB_set (desc, GxB_CHUNK, chunk) ;
```

The smaller of nthreads_max and floor(work/chunk) is used for any given GraphBLAS operation, except that a single thread is used if this value is zero or less.

If either parameter is set to GxB_DEFAULT, then default values are used. The default for nthreads_max is the return value from omp_get_max_threads, and the default chunk size is currently 65,536.

If a descriptor value for either parameter is left at its default, or set to GxB_DEFAULT, then the global setting is used. This global setting may have been modified from its default, and this modified value will be used.

For example, suppose omp_get_max_threads reports 8 threads. If GxB_set (GxB_NTHREADS, 4) is used, then the global setting is four threads, not eight. If a descriptor is used but its GxB_NTHREADS is not set, or set to GxB_DEFAULT, then any operation that uses this descriptor

6.2 Storing a matrix by row or by column

The GraphBLAS GrB_Matrix is entirely opaque to the user application, and the GraphBLAS API does not specify how the matrix should be stored. However, choices made in how the matrix is represented in a particular implementation, such as SuiteSparse:GraphBLAS, can have a large impact on performance.

Many graph algorithms are just as fast in any format, but some algorithms are much faster in one format or the other. For example, suppose the user application stores a directed graph as a matrix A, with the edge (i, j) represented as the value A(i,j), and the application makes many accesses to the *i*th row of the matrix, with $GrB_Col_extract(w, ..., A, GrB_ALL, ..., i, desc)$ with the transposed descriptor (GrB_INPO set to GrB_TRAN). If the matrix is stored by column this can be extremely slow, just like the expression w=A(i,:) in MATLAB, where i is a scalar. Since this is a typical usecase in graph algorithms, the default format in SuiteSparse:GraphBLAS is to store its matrices by row, in Compressed Sparse Row format (CSR).

MATLAB stores its sparse matrices by column, in "non-hypersparse" format, in what is called the Compressed Sparse Column format, or CSC for short. An m-by-n matrix in MATLAB is represented as a set of n column vectors, each with a sorted list of row indices and values of the nonzero entries in that column. As a result, w=A(:,j) is very fast in MATLAB, since the result is already held in the data structure a single list, the jth column vector. However, w=A(i,:) is very slow in MATLAB, since every column in the matrix has to be searched to see if it contains row i. In MATLAB, if many such accesses are made, it is much better to transpose the matrix (say AT=A') and then use w=AT(:,i) instead. This can have a dramatic impact

on the performance of MATLAB.

Likewise, if u is a very sparse column vector and A is stored by column, then $w=u^**A$ (via GrB_vxm) is slower than w=A*u (via GrB_mxv). The opposite is true if the matrix is stored by row.

An example of this can be found in Section B.1 of Version 1.2 of the GraphBLAS API Specification, where the breadth-first search BFS uses GrB_vxm to compute q'=q'*A. This method is not fast if the matrix A is stored by column. The bfs5 and bfs6 examples in the Demo/ folder of Suite-Sparse:GraphBLAS use GrB_vxm, which is fast since the matrices are assumed to be stored in their default format, by row.

SuiteSparse:GraphBLAS stores its sparse matrices by row, by default. In Versions 2.1 and earlier, the matrices were stored by column, by default. However, it can also be instructed to store any selected matrices, or all matrices, by column instead (just like MATLAB), so that w=A(:,j) (via GrB_Col_extract) is very fast. The change in data format has no effect on the result, just the time and memory usage. To use a column-oriented format by default, the following can be done in a user application that tends to access its matrices by column.

```
GrB_init (...);
// just after GrB_init: do the following:
#ifdef GxB_SUITESPARSE_GRAPHBLAS
GxB_set (GxB_FORMAT, GxB_BY_COL);
#endif
```

If this is done, and no other GxB_set calls are made with GxB_FORMAT, all matrices will be stored by column. Alternatively, SuiteSparse:GraphBLAS can be compiled with -DBYCOL, which changes the default format to GxB_BY_COL, with no calls to any GxB_* function. The default format is now GxB_BY_ROW.

6.3 Hypersparse matrices

MATLAB can store an m-by-n matrix with a very large value of m, since a CSC data structure takes $O(n + |\mathbf{A}|)$ memory, independent of m, where $|\mathbf{A}|$ is the number of nonzeros in the matrix. It cannot store a matrix with a huge n, and this structure is also inefficient when $|\mathbf{A}|$ is much smaller than n. In contrast, SuiteSparse:GraphBLAS can store its matrices in *hypersparse* format, taking only $O(|\mathbf{A}|)$ memory, independent of how it is stored (by row or by column) and independent of both m and n [BG08, BG12].

In both the CSR and CSC formats, the matrix is held as a set of sparse vectors. In non-hypersparse format, the set of sparse vectors is itself dense; all vectors are present, even if they are empty. For example, an m-by-n matrix in non-hypersparse CSC format contains n sparse vectors. Each column vector takes at least one integer to represent, even for a column with no entries. This allows for quick lookup for a particular vector, but the memory required is $O(n+|\mathbf{A}|)$. With a hypersparse CSC format, the set of vectors itself is sparse, and columns with no entries take no memory at all. The drawback of the hypersparse format is that finding an arbitrary column vector \mathbf{j} , such as for the computation $\mathbf{C}=\mathbf{A}(:,\mathbf{j})$, takes $O(\log k)$ time if there $k\leq n$ vectors in the data structure. One advantage of the hypersparse structure is the memory required for an m-by-n hypersparse CSC matrix is only $O(|\mathbf{A}|)$, independent of m and n. Algorithms that must visit all non-empty columns of a matrix are much faster when working with hypersparse matrices, since empty columns can be skipped.

The hyper_ratio parameter controls the hypersparsity of the internal data structure for a matrix. The parameter is typically in the range 0 to 1. The default is hyper_ratio = GxB_HYPER_DEFAULT, which is an extern const double value, currently set to 0.0625, or 1/16. This default ratio may change in the future.

The hyper_ratio determines how the matrix is converted between the hypersparse and non-hypersparse formats. Let n be the number of columns of a CSC matrix, or the number of rows of a CSR matrix. The matrix can have at most n non-empty vectors.

Let k be the actual number of non-empty vectors. That is, for the CSC format, $k \leq n$ is the number of columns that have at least one entry. Let h be the value of hyper_ratio.

If a matrix is currently hypersparse, it can be converted to non-hypersparse if the either condition $n \leq 1$ or k > 2nh holds, or both. Otherwise, it stays hypersparse. Note that if $n \leq 1$ the matrix is always stored as non-hypersparse.

If currently non-hypersparse, it can be converted to hypersparse if both conditions n > 1 and $k \le nh$ hold. Otherwise, it stays non-hypersparse. Note that if $n \le 1$ the matrix always remains non-hypersparse.

The default value of hyper_ratio is assigned at startup by GrB_init, and can then be modified globally with GxB_set. All new matrices are created with the same hyper_ratio, determined by the global value. Once a particular matrix A has been constructed, its hypersparsity ratio can be

modified from the default with:

```
double hyper_ratio = 0.2 ;
GxB_set (A, GxB_HYPER, hyper_ratio) ;
```

To force a matrix to always be non-hypersparse, use hyper_ratio equal to GxB_NEVER_HYPER. To force a matrix to always stay hypersparse, set hyper_ratio to GxB_ALWAYS_HYPER.

A GrB_Matrix can thus be held in one of four formats: any combination of hyper/non-hyper and CSR/CSC. All GrB_Vector objects are always stored in non-hypersparse CSC format.

A new matrix created via $\mathtt{GrB_Matrix_new}$ starts with k=0 and is created in hypersparse form by default unless $n\leq 1$ or if h<0, where h is the global hyper_ratio value. The matrix is created in in either $\mathtt{GxB_BY_ROW}$ or $\mathtt{GxB_BY_COL}$ format, as determined by the last call to $\mathtt{GxB_set}(\mathtt{GxB_FORMAT},\ldots)$ or $\mathtt{GrB_init}$.

A new matrix C created via GrB_dup (&C,A) inherits the CSR/CSC format, hypersparsity format, and hyper_ratio from A.

Parameter types: The GxB_Option_Field enumerated type gives the type of the field parameter for the second argument of GxB_set and GxB_get, for setting global options or matrix options.

```
typedef enum
{
   GxB_HYPER = 0,
                       // defines switch to hypersparse format (double value)
   GxB_FORMAT = 1,
                      // defines CSR/CSC format: GxB_BY_ROW or GxB_BY_COL
   GxB\_MODE = 2,
                       // mode passed to GrB_init (blocking or non-blocking)
   GxB_THREAD_SAFETY = 3, // thread library for thread safety
    GxB\_THREADING = 4,
                           // currently none (in progress)
   GxB_GLOBAL_NTHREADS = GxB_NTHREADS, // max number of threads to use
                                       // chunk size for small problems
   GxB_GLOBAL_CHUNK = GxB_CHUNK,
    GxB_IS_HYPER = 6
                        // query a matrix to see if it hypersparse or not
                         // (GxB_Matrix_Option_get only)
GxB_Option_Field ;
```

The GxB_FORMAT field can be by row or by column, set to a value with the type GxB_Format_Value:

The default format (in SuiteSparse:GraphBLAS Version 2.2 and later) is by row. The format in SuiteSparse:GraphBLAS Version 2.1 and earlier was by column, just like MATLAB.

The default format is given by the predefined value GxB_FORMAT_DEFAULT, which is equal to GxB_BY_ROW if default compile-time options are used. To change the default at compile time to GxB_BY_COL, compile the SuiteSparse: GraphBLAS library with -DBYCOL. This changes GxB_FORMAT_DEFAULT to GxB_BY_COL. The default hypersparsity ratio is 0.0625 (1/16), but this value may change in the future.

Setting the GxB_HYPER field to GxB_ALWAYS_HYPER ensures a matrix always stays hypersparse. If set to GxB_NEVER_HYPER, it always stays non-hypersparse. At startup, GrB_init defines the following initial settings:

```
GxB_set (GxB_HYPER, GxB_HYPER_DEFAULT) ;
GxB_set (GxB_FORMAT, GxB_FORMAT_DEFAULT) ;
```

That is, by default, all new matrices are held by column in CSR format, unless -DBYCOL is used at compile time, in which case the default is to store all new matrices by row in CSC format. If a matrix has fewer than n/16 columns, it can be converted to hypersparse format. If it has more than n/8 columns, it can be converted to non-hypersparse format. These options can be changed for all future matrices with GxB_set . For example, to change all future matrices to be in non-hypersparse CSC when created, use:

```
GxB_set (GxB_HYPER, GxB_NEVER_HYPER) ;
GxB_set (GxB_FORMAT, GxB_BY_COL) ;
Then if a particular matrix needs a different format, then (as an example):
GxB_set (A, GxB_HYPER, 0.1) ;
GxB_set (A, GxB_FORMAT, GxB_BY_ROW) ;
```

This changes the matrix A so that it is stored by row, and it is converted from non-hypersparse to hypersparse format if it has fewer than 10% non-empty columns. If it is hypersparse, it is a candidate for conversion to non-hypersparse if has 20% or more non-empty columns. If it has between 10% and 20% non-empty columns, it remains in its current format. MATLAB only supports a non-hypersparse CSC format. The format in SuiteSparse:GraphBLAS that is equivalent to the MATLAB format is:

```
GrB_init (...);
GxB_set (GxB_HYPER, GxB_NEVER_HYPER);
GxB_set (GxB_FORMAT, GxB_BY_COL);
// no subsequent use of GxB_HYPER or GxB_FORMAT
```

The GxB_HYPER and GxB_FORMAT options should be considered as suggestions from the user application as to how SuiteSparse:GraphBLAS can obtain the best performance for a particular application. SuiteSparse:GraphBLAS is free to ignore any of these suggestions, both now and in the future, and the available options and formats may be augmented in the future. Any prior options no longer needed in future versions of SuiteSparse:GraphBLAS will be silently ignored, so the use these options is safe for future updates.

The hypersparse status of a matrix can be queried with the following usage:

```
bool is_hyper ;
GxB_get (A, GxB_IS_HYPER, &is_hyper) ;
printf (is_hyper ? "A is hypersparse" : "A is standard sparse") ;
```

6.4 Other global options

GxB_MODE, GxB_THREAD_SAFETY, and GxB_THREADING can only be queried by GxB_get; they cannot be modified by GxB_set. The mode is the value passed to GrB_init (blocking or non-blocking). The GxB_THREAD* options are returned as an enum type with one of the following options:

SuiteSparse:GraphBLAS multi-threaded, using only OpenMP for its internal parallelism. It is also thread-safe if it is compiled with OpenMP or POSIX pthreads, and if the user application threads do not operate on the same matrices at the same time. The user threads may use OpenMP or POSIX pthreads. If multiple user threads make simultaneous calls to

GraphBLAS, then output matrices and vectors used by different threads must be different, and input matrices and vectors can be safely used only if any pending computations on them have finished, via GrB_Matrix_wait, GrB_Vector_wait, or GxB_Scalar_wait.

The GxB_THREAD_SAFETY option returns the threading model used internally to synchronize user threads, solely for the now-deprected GrB_wait(). This is determined during installation (see Section 12.4). Since GxB_THREAD_NONE is zero, the following can be used:

```
GxB_Thread_Model thread_safety ;
GxB_get (GxB_THREAD_SAFETY, &thread_safety) ;
if (thread_safety)
{
    printf ("GraphBLAS is thread-safe\n") ;
}
else
{
    // neither OpenMP, POSIX pthreads, nor any other threading model
    // was available at compile-time
    printf ("GraphBLAS is not thread-safe!\n") ;
}
```

The GxB_THREADING option returns the internal parallelism used inside SuiteSparse:GraphBLAS, depending on how the library was compiled:

```
GxB_Thread_Model threading ;
GxB_get (GxB_THREADING, &threading) ;
if (threading == GxB_THREAD_NONE)
{
    printf ("GraphBLAS is single-threaded, internally.\n") ;
} else
{
    printf ("GraphBLAS is multi-threaded, internally, using OpenMP.\n") ;
}
```

All threads in the same user application share the same global options, including hypersparsity and CSR/CSC format determined by GxB_set, the blocking mode determined by GrB_init, and the threading options. Specific format and hypersparsity parameters of each matrix are specific to that matrix and can be independently changed.

6.5 GxB_Global_Option_set: set a global option

This usage of GxB_set sets the value of a global option. The field parameter can be GxB_HYPER, GxB_FORMAT, GxB_NTHREADS, or GxB_CHUNK.

For example, the following usage sets the global hypersparsity ratio to 0.2, the format of future matrices to GxB_BY_COL, the maximum number of threads to 4, and the chunk size to 10000. No existing matrices are changed.

```
GxB_set (GxB_HYPER, 0.2) ;
GxB_set (GxB_FORMAT, GxB_BY_COL) ;
GxB_set (GxB_NTHREADS, 4) ;
GxB_set (GxB_CHUNK, (double) 10000) ;
```

6.6 GxB_Matrix_Option_set: set a matrix option

This usage of GxB_set sets the value of a matrix option, for a particular matrix. The field parameter can be GxB_HYPER or GxB_FORMAT.

For example, the following usage sets the hypersparsity ratio to 0.2, and the format of GxB_BY_COL, for a particular matrix A. SuiteSparse:GraphBLAS currently applies these changes immediately, but since they are simply hints, future versions of SuiteSparse:GraphBLAS may delay the change in format if it can obtain better performance.

For performance, the matrix option should be set as soon as it is created with GrB_Matrix_new, so the internal transformation takes less time.

```
GxB_set (A, GxB_HYPER, 0.2) ;
GxB_set (A, GxB_FORMAT, GxB_BY_COL) ;
```

6.7 GxB_Desc_set: set a GrB_Descriptor value

This usage is similar to GrB_Descriptor_set, just with a name that is consistent with the other usages of this generic function. Unlike GrB_Descriptor_set, the field may also be GxB_NTHREADS, or GxB_CHUNK. Refer to Sections 5.11.2 and 5.11.3 for details.

6.8 GxB_Global_Option_get: retrieve a global option

This usage of GxB_get retrieves the value of a global option. The field parameter can be GxB_HYPER, GxB_FORMAT. GxB_MODE, GxB_THREAD_SAFETY, GxB_THREADING, GxB_NTHREADS, or GxB_CHUNK. For example:

```
double h ;
GxB_get (GxB_HYPER, &h) ;
printf ("hyper_ratio = %g for all new matrices\n", h) ;

GxB_Format_Value s ;
GxB_get (GxB_FORMAT, &s) ;
if (s == GxB_BY_COL) printf ("all new matrices are stored by column\n") :
else printf ("all new matrices are stored by row\n") ;

GrB_mode mode ;
GxB_get (GxB_MODE, &mode) ;
if (mode == GrB_BLOCKING) printf ("GrB_init(GrB_BLOCKING) was called.\n") :
else printf ("GrB_init(GrB_NONBLOCK) was called.\n") ;
int nthreads_max ;
GxB_get (GxB_NTHREADS, &nthreads_max) ;
printf ("max # of threads to use: %d\n", nthreads_max) ;
```

```
double chunk ;
GxB_get (GxB_CHUNK, &chunk) ;
printf ("chunk size: %g\n", chunk) ;

// see Demo/Program/pthread_demo.c and openmp_demo.c for examples:
GxB_Threading_Model thread_safety, threading ;
GxB_get (GxB_THREAD_SAFETY, &thread_safey) ;
GxB_get (GxB_THREADING, &threading) ;
```

6.9 GxB_Matrix_Option_get: retrieve a matrix option

```
GrB_Info GxB_get  // gets the current option of a matrix (

GrB_Matrix A,  // matrix to query  // option to query  // return value of the matrix option

);
```

This usage of GxB_get retrieves the value of a matrix option. The field parameter can be GxB_HYPER or GxB_FORMAT. For example:

```
double h ;
GxB_get (A, GxB_HYPER, &h) ;
printf ("matrix A has hyper_ratio = %g\n", h) ;

GxB_Format_Value s ;
GxB_get (A, GxB_FORMAT, &s) ;
if (s == GxB_BY_COL) printf ("matrix A is stored by column\n") :
else printf ("matrix A is stored by row\n") ;
```

6.10 GxB_Desc_get: retrieve a GrB_Descriptor value

This usage is the same as GxB_Desc_get. The field parameter can be GrB_OUTP, GrB_MASK, GrB_INPO, GrB_INP1, GxB_AxB_METHOD, GxB_NTHREADS, or GxB_CHUNK. Refer to Section 5.11.4 for details.

6.11 Summary of usage of GxB_set and GxB_get

The different usages of GxB_set and GxB_get are summarized below. To set/get the global options:

```
GxB_set (GxB_HYPER, double h) ;
   GxB_set (GxB_HYPER, GxB_ALWAYS_HYPER) ;
   GxB_set (GxB_HYPER, GxB_NEVER_HYPER) ;
   GxB_get (GxB_HYPER, double *h) ;
   GxB_set (GxB_FORMAT, GxB_BY_ROW) ;
   GxB_set (GxB_FORMAT, GxB_BY_COL) ;
   GxB_get (GxB_FORMAT, GxB_Format_Value *s) ;
   GxB_set (GxB_THREADS, int nthreads_max) ;
   GxB_get (GxB_THREADS, int *nthreads_max) ;
   GxB_set (GxB_CHUNK, double chunk) ;
   GxB_get (GxB_CHUNK, double *chunk) ;
To get global options that can be queried but not modified:
   GxB_get (GxB_MODE,
                                GrB_Mode *mode) ;
   GxB_get (GxB_THREAD_SAFETY, GxB_Thread_Model *thread_safety) ;
   GxB_get (GxB_THREADING,
                                GxB_Thread_Model *threading) ;
To set/get a matrix option:
   GxB_set (GrB_Matrix A, GxB_HYPER, double h) ;
   GxB_set (GrB_Matrix A, GxB_HYPER, GxB_ALWAYS_HYPER) ;
   GxB_set (GrB_Matrix A, GxB_HYPER, GxB_NEVER_HYPER) ;
   GxB_get (GrB_Matrix A, GxB_HYPER, double *h) ;
   GxB_set (GrB_Matrix A, GxB_FORMAT, GxB_BY_ROW) ;
   GxB_set (GrB_Matrix A, GxB_FORMAT, GxB_BY_COL) ;
   GxB_get (GrB_Matrix A, GxB_FORMAT, GxB_Format_Value *s) ;
To get the hypersparse status of a matrix:
   GxB_get (GrB_Matrix A, GxB_IS_HYPER, bool *is_hyper) ;
```

To set/get a descriptor field:

```
GxB_set (GrB_Descriptor d, GrB_OUTP, GxB_DEFAULT) ;
GxB_set (GrB_Descriptor d, GrB_OUTP, GrB_REPLACE) ;
GxB_get (GrB_Descriptor d, GrB_OUTP, GrB_Desc_Value *v) ;
GxB_set (GrB_Descriptor d, GrB_MASK, GxB_DEFAULT) ;
GxB_set (GrB_Descriptor d, GrB_MASK, GrB_COMP) ;
GxB_set (GrB_Descriptor d, GrB_MASK, GrB_STRUCTURE) ;
GxB_set (GrB_Descriptor d, GrB_MASK, GrB_COMP+GrB_STRUCTURE) ;
GxB_get (GrB_Descriptor d, GrB_MASK, GrB_Desc_Value *v) ;
GxB_set (GrB_Descriptor d, GrB_INPO, GxB_DEFAULT) ;
GxB_set (GrB_Descriptor d, GrB_INPO, GrB_TRAN) ;
GxB_get (GrB_Descriptor d, GrB_INPO, GrB_Desc_Value *v) ;
GxB_set (GrB_Descriptor d, GrB_INP1, GxB_DEFAULT) ;
GxB_set (GrB_Descriptor d, GrB_INP1, GrB_TRAN) ;
GxB_get (GrB_Descriptor d, GrB_INP1, GrB_Desc_Value *v) ;
GxB_set (GrB_Descriptor d, GxB_AxB_METHOD, GxB_DEFAULT) ;
GxB_set (GrB_Descriptor d, GxB_AxB_METHOD, GxB_AxB_GUSTAVSON) ;
GxB_set (GrB_Descriptor d, GxB_AxB_METHOD, GxB_AxB_HEAP) ;
GxB_set (GrB_Descriptor d, GxB_AxB_METHOD, GxB_AxB_HASH) ;
GxB_set (GrB_Descriptor d, GxB_AxB_METHOD, GxB_AxB_SAXPY) ;
GxB_set (GrB_Descriptor d, GxB_AxB_METHOD, GxB_AxB_DOT) ;
GxB_get (GrB_Descriptor d, GrB_AxB_METHOD, GrB_Desc_Value *v) ;
GxB_set (GrB_Descriptor d, GxB_NTHREADS, int nthreads) ;
GxB_get (GrB_Descriptor d, GxB_NTHREADS, int *nthreads) ;
GxB_set (GrB_Descriptor d, GxB_CHUNK, double chunk) ;
GxB_get (GrB_Descriptor d, GxB_CHUNK, double *chunk) ;
```

7 SuiteSparse:GraphBLAS Colon and Index Notation

MATLAB uses a colon notation to index into matrices, such as C=A(2:4,3:8), which extracts C as 3-by-6 submatrix from A, from rows 2 through 4 and columns 3 to 8 of the matrix A. A single colon is used to denote all rows, C=A(:,9), or all columns, C=A(12,:), which refers to the 9th column and 12th row of A, respectively. An arbitrary integer list can be given as well, such as the MATLAB statements:

```
I = [2 1 4];

J = [3 5];

C = A (I,J);
```

which creates the 3-by-2 matrix C as follows:

$$C = \left[\begin{array}{cc} a_{2,3} & a_{2,5} \\ a_{1,3} & a_{1,5} \\ a_{4,3} & a_{4,5} \end{array} \right]$$

The GraphBLAS API can do the equivalent of C=A(I,J), C=A(:,J), C=A(I,:), and C=A(:,:), by passing a parameter const GrB_Index *I as either an array of size ni, or as the special value GrB_ALL, which corresponds to the stand-alone colon C=A(:,J), and the same can be done for J.. To compute C=A(2:4,3:8) in GraphBLAS requires the user application to create two explicit integer arrays I and J of size 3 and 5, respectively, and then fill them with the explicit values [2,3,4] and [3,4,5,6,7,8]. This works well if the lists are small, or if the matrix has more entries than rows or columns.

However, particularly with hypersparse matrices, the size of the explicit arrays I and J can vastly exceed the number of entries in the matrix. When using its hypersparse format, SuiteSparse:GraphBLAS allows the user application to create a GrB_Matrix with dimensions up to 2^{60} , with no memory constraints. The only constraint on memory usage in a hypersparse matrix is the number of entries in the matrix.

For example, creating a n-by-n matrix A of type GrB_FP64 with $n=2^{60}$ and one million entries is trivial to do in Version 2.1 (and later) of Suite-Sparse:GraphBLAS, taking at most 24MB of space. SuiteSparse:GraphBLAS

Version 2.1 (or later) could do this on an old smartphone. However, using just the pure GraphBLAS API, constructing C=A(0:(n/2),0:(n/2)) in SuiteSparse Version 2.0 would require the creation of an integer array I of size 2⁵⁹, containing the sequence 0, 1, 2, 3,, requiring about 4 ExaBytes of memory (4 million terabytes). This is roughly 1000 times larger than the memory size of the world's largest computer in 2018.

SuiteSparse:GraphBLAS Version 2.1 and later extends the GraphBLAS API with a full implementation of the MATLAB colon notation for integers, I=begin:inc:end. This extension allows the construction of the matrix C=A(0:(n/2),0:(n/2)) in this example, with dimension 2⁵⁹, probably taking just milliseconds on an old smartphone.

The GrB_extract, GrB_assign, and GxB_subassign operations (described in the Section 8) each have parameters that define a list of integer indices, using two parameters:

These two parameters define five kinds of index lists, which can be used to specify either an explicit or implicit list of row indices and/or column indices. The length of the list of indices is denoted |I|. This discussion applies equally to the row indices I and the column indices J. The five kinds are listed below.

- 1. An explicit list of indices, such as I = [2 1 4 7 2] in MATLAB notation, is handled by passing in I as a pointer to an array of size 5, and passing ni=5 as the size of the list. The length of the explicit list is ni=|I|. Duplicates may appear, except that for some uses of GrB_assign and GxB_subassign, duplicates lead to undefined behavior according to the GraphBLAS C API Specification. Suite-Sparse:GraphBLAS specifies how duplicates are handled in all cases, as an addition to the specification. See Section 8.10 for details.
- 2. To specify all rows of a matrix, use I = GrB_ALL. The parameter ni is ignored. This is equivalent to C=A(:,J) in MATLAB. In GraphBLAS, this is the sequence 0:(m-1) if A has m rows, with length |I|=m. If J is used the columns of an m-by-n matrix, then J=GrB_ALL refers to all columns, and is the sequence 0:(n-1), of length |J|=n.

3. To specify a contiguous range of indices, such as I=10:20 in MATLAB, the array I has size 2, and ni is passed to SuiteSparse:GraphBLAS as the special value ni = GxB_RANGE. The beginning index is I[GxB_BEGIN] and the ending index is I[GxB_END]. Both values must be non-negative since GrB_Index is an unsigned integer (uint64_t). The value of I[GxB_INC] is ignored.

```
// to specify I = 10:20
GrB_Index I [2], ni = GxB_RANGE;
I [GxB_BEGIN] = 10;  // the start of the sequence
I [GxB_END] = 20;  // the end of the sequence
```

Let $b = I[GxB_BEGIN]$, let $e = I[GxB_END]$, The sequence has length zero if b > e; otherwise the length is |I| = (e - b) + 1.

4. To specify a strided range of indices with a non-negative stride, such as I=3:2:10, the array I has size 3, and ni has the special value GxB_STRIDE. This is the sequence 3, 5, 7, 9, of length 4. Note that 10 does not appear in the list. The end point need not appear if the increment goes past it.

The GxB_STRIDE sequence is the same as the List generated by the following for loop:

```
int64_t k = 0 ;
GrB_Index *List = (a pointer to an array of large enough size)
for (int64_t i = I [GxB_BEGIN] ; i <= I [GxB_END] ; i += I [GxB_INC])
{
    // i is the kth entry in the sequence
    List [k++] = i ;
}</pre>
```

Then passing the explicit array List and its length ni=k has the same effect as passing in the array I of size 3, with ni=GxB_STRIDE. The

latter is simply much faster to produce, and much more efficient for SuiteSparse:GraphBLAS to process.

Let $b = I[GxB_BEGIN]$, let $e = I[GxB_END]$, and let $\Delta = I[GxB_INC]$. The sequence has length zero if b > e or $\Delta = 0$. Otherwise, the length of the sequence is

$$|I| = \left| \frac{e - b}{\Lambda} \right| + 1$$

5. In MATLAB notation, if the stride is negative, the sequence is decreasing. For example, 10:-2:1 is the sequence 10, 8, 6, 4, 2, in that order. In SuiteSparse:GraphBLAS, use ni = GxB_BACKWARDS, with an array I of size 3. The following example specifies defines the equivalent of the MATLAB expression 10:-2:1 in SuiteSparse:GraphBLAS:

The value -2 cannot be assigned to the GrB_Index array I, since that is an unsigned type. The signed increment is represented instead with the special value ni = GxB_BACKWARDS. The GxB_BACKWARDS sequence is the same as generated by the following for loop:

```
int64_t k = 0 ;
GrB_Index *List = (a pointer to an array of large enough size)
for (int64_t i = I [GxB_BEGIN] ; i >= I [GxB_END] ; i -= I [GxB_INC])
{
    // i is the kth entry in the sequence
    List [k++] = i ;
}
```

Let $b = I[GxB_BEGIN]$, let $e = I[GxB_END]$, and let $\Delta = I[GxB_INC]$ (note that Δ is not negative). The sequence has length zero if b < e or $\Delta = 0$. Otherwise, the length of the sequence is

$$|I| = \left\lfloor \frac{b - e}{\Delta} \right\rfloor + 1$$

Since GrB_Index is an unsigned integer, all three values I[GxB_BEGIN], I[GxB_INC], and I[GxB_END] must be non-negative.

Just as in MATLAB, it is valid to specify an empty sequence of length zero. For example, I = 5:3 has length zero in MATLAB and the same is true for a GxB_RANGE sequence in SuiteSparse:GraphBLAS, with $I[GxB_BEGIN]=5$ and $I[GxB_END]=3$. This has the same effect as array I with ni=0.

 $\mathbf{SPEC}\textsc{:}\ \mathtt{GxB_RANGE},\ \mathtt{GxB_STRIDE},\ \mathrm{and}\ \mathtt{GxB_BACKWARDS}$ are extensions to the specification.

8 GraphBLAS Operations

The next sections define each of the GraphBLAS operations, also listed in the table below. SuiteSparse:GraphBLAS extensions (GxB_subassign, GxB_select) are included in the table.

GrB_mxm	matrix-matrix multiply	$\mathbf{C}\langle\mathbf{M} angle=\mathbf{C}\odot\mathbf{AB}$
${\tt GrB_vxm}$	vector-matrix multiply	$\mathbf{w}^{T} \langle \mathbf{m}^{T} \rangle = \mathbf{w}^{T} \odot \mathbf{u}^{T} \mathbf{A}$
GrB_mxv	matrix-vector multiply	$\mathbf{w}\langle\mathbf{m}\rangle=\mathbf{w}\odot\mathbf{A}\mathbf{u}$
GrB_eWiseMult	element-wise,	$\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{C} \odot (\mathbf{A} \otimes \mathbf{B})$
	set union	$\mathbf{w}\langle\mathbf{m}\rangle=\mathbf{w}\odot(\mathbf{u}\otimes\mathbf{v})$
GrB_eWiseAdd	element-wise,	$\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{C} \odot (\mathbf{A} \oplus \mathbf{B})$
	set intersection	$\mathbf{w}\langle\mathbf{m}\rangle=\mathbf{w}\odot(\mathbf{u}\oplus\mathbf{v})$
GrB_extract	extract submatrix	$\mathbf{C}\langle\mathbf{M} angle=\mathbf{C}\odot\mathbf{A}(\mathbf{I},\mathbf{J})$
		$\mathbf{w}\langle\mathbf{m} angle=\mathbf{w}\odot\mathbf{u}(\mathbf{i})$
GxB_subassign	assign submatrix,	$\mathbf{C}(\mathbf{I},\mathbf{J})\langle\mathbf{M} angle = \mathbf{C}(\mathbf{I},\mathbf{J})\odot\mathbf{A}$
	with submask for $\mathbf{C}(\mathbf{I}, \mathbf{J})$	$\mathbf{w}(\mathbf{i})\langle\mathbf{m} angle = \mathbf{w}(\mathbf{i})\odot\mathbf{u}$
GrB_assign	assign submatrix	$\mathbf{C}\langle\mathbf{M} angle(\mathbf{I},\mathbf{J})=\mathbf{C}(\mathbf{I},\mathbf{J})\odot\mathbf{A}$
	with submask for ${f C}$	$\mathbf{w}\langle\mathbf{m} angle(\mathbf{i})=\mathbf{w}(\mathbf{i})\odot\mathbf{u}$
GrB_apply	apply unary operator	$\mathbf{C}\langle\mathbf{M} angle = \mathbf{C}\odot f(\mathbf{A})$
		$\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot f(\mathbf{u})$
GxB_select	apply select operator	$\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot f(\mathbf{A}, \mathbf{k})$
		$\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot f(\mathbf{u},\mathbf{k})$
GrB_reduce	reduce to vector	$\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot[\oplus_{j}\mathbf{A}(:,j)]$
	reduce to scalar	$s = s \odot [\oplus_{ij} \mathbf{A}(I,J)]$
GrB_transpose	transpose	$\mathbf{C}\langle\mathbf{M} angle = \mathbf{C}\odot\mathbf{A}^T$
GrB_kronecker	Kronecker product	$\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{C} \odot \operatorname{kron}(\mathbf{A}, \mathbf{B})$
-		

8.1 The GraphBLAS specification in MATLAB

SuiteSparse:GraphBLAS includes a MATLAB implementation of nearly the entire GraphBLAS specification, including all built-in types and operators. The typecasting rules and integer operator rules from GraphBLAS are implemented in MATLAB via mexFunctions that call the GraphBLAS routines in C. All other functions are written purely in MATLAB M-files, and are given names of the form GB_spec_*. All of these MATLAB interfaces and M-file functions they are provided in the software distribution of Suite-Sparse:GraphBLAS. The purpose of this is two-fold:

- Illustration and documentation: MATLAB is so expressive, and so beautiful to read and write, that the GB_spec_* functions read almost like the exact specifications from the GraphBLAS API. Excerpts and condensed versions of these functions have already been used to this point in the User Guide, such as Figure 1, and the subsequent sections rely on them as well. This is why the discussion here is not just relegated to an Appendix on testing; the reader can benefit from studying the GB_spec_* functions to understand what a GraphBLAS operation is computing. For example, GrB_mxm (Section 8.2) includes a condensed and simplified version of GB_spec_mxm.
- Testing: Testing the C interface to SuiteSparse:GraphBLAS is a significant challenge since it supports so many different kinds of operations on a vast range of semirings. It is difficult to tell from looking at the result from a C function in GraphBLAS if the result is correct. Thus, each function has been written twice: once in a highly-optimized function in C, and again in a simple and elegant MATLAB function. The latter is almost a direct translation of all the mathematics behind the GraphBLAS API, so it is much easier to visually inspect the GB_spec_* version in MATLAB to ensure the correct mathematics are being computed.

The following functions are included in the SuiteSparse:GraphBLAS software distribution. Each has a name of the form GB_spec_*, and each of them is a "mimic" of a corresponding C function in GraphBLAS. Not all functions in the C API have a corresponding mimic; in particular, many of the vector functions can be computed directly with the corresponding matrix version in the MATLAB implementations. A list of these files is shown below:

MATLAB GB_spec function	${\it corresponding GraphBLAS}$	Section
	function or method	
GB_spec_accum.m	$\mathbf{Z} = \mathbf{C} \odot \mathbf{T}$	2.3
GB_spec_mask.m	$\mathbf{C}\langle\mathbf{M} angle=\mathbf{Z}$	2.3
GB_spec_accum_mask.m	$\mathbf{C}\langle\mathbf{M} angle=\mathbf{C}\odot\mathbf{T}$	2.3
GB_spec_Vector_extractElement.m	<pre>GrB_Vector_extractElement</pre>	5.8.9
GB_spec_build.m	GrB_Matrix_build	5.9.8
<pre>GB_spec_Matrix_extractElement.m</pre>	<pre>GrB_Matrix_extractElement</pre>	5.9.10
${\tt GB_spec_extractTuples.m}$	${\tt GrB_Matrix_extractTuples}$	5.9.11
GB_spec_mxm.m	GrB_mxm	8.2
GB_spec_vxm.m	GrB_vxm	8.3
GB_spec_mxv.m	GrB_mxv	8.4
GB_spec_Vector_eWiseMult.m	<pre>GrB_Vector_eWiseMult</pre>	8.5
<pre>GB_spec_Matrix_eWiseMult.m</pre>	<pre>GrB_Matrix_eWiseMult</pre>	8.5
${\tt GB_spec_Vector_eWiseAdd.m}$	<pre>GrB_Vector_eWiseAdd</pre>	8.6
${\tt GB_spec_Matrix_eWiseAdd.m}$	<pre>GrB_Matrix_eWiseAdd</pre>	8.6
GB_spec_Vector_extract.m	<pre>GrB_Vector_extract</pre>	8.7.1
<pre>GB_spec_Matrix_extract.m</pre>	<pre>GrB_Matrix_extract</pre>	8.7.2
<pre>GB_spec_Col_extract.m</pre>	<pre>GrB_Col_extract</pre>	8.7.3
GB_spec_subassign.m	GxB_subassign	8.8
GB_spec_assign.m	GrB_assign	8.9
GB_spec_apply.m	GrB_apply	8.12
GB_spec_select.m	GxB_select	8.13
GB_spec_reduce_to_vector.m	GrB_reduce (to vector)	8.14.1
GB_spec_reduce_to_scalar.m	GrB_reduce (to scalar)	8.14.3
GB_spec_transpose.m	GrB_transpose	8.15
GB_spec_kron.m	GrB_kronecker	8.16

Additional files are included for creating test problems and providing inputs to the above files, or supporting functions:

MATLAB GB_spec function	purpose
GB_spec_compare.m	Compares output of C and MATLAB functions
GB_spec_random.m	Generates a random matrix
GB_spec_op.m	MATLAB mimic of built-in operators
GB_spec_operator.m	Like GrB_*Op_new
GB_spec_opsall.m	List operators, types, and semirings
GB_spec_semiring.m	Like GrB_Semiring_new
GB_spec_descriptor.m	mimics a GraphBLAS descriptor
GB_spec_identity.m	returns the identity of a monoid
<pre>GB_spec_matrix.m</pre>	conforms a MATLAB sparse matrix to GraphBLAS
GB_define.m	creates draft of GraphBLAS.h

An intensive test suite has been written that generates test graphs in MATLAB, then computes the result in both the C version of the Suite-Sparse:GraphBLAS and in the MATLAB GB_spec_* functions. Each C function in GraphBLAS has a direct mexFunction interface that allow the test suite in MATLAB to call both functions.

This approach has its limitations:

- matrix classes: MATLAB only supports sparse double, sparse double complex, and sparse logical matrices. MATLAB can represent dense matrices in all eleven built-in GraphBLAS data types, so in all these specification M-files, the matrices are either in dense format in the corresponding MATLAB class, or they are held as sparse double or sparse logical, and the actual GraphBLAS type is held with it as a string member of a MATLAB struct. To ensure the correct typecasting is computed, most of the MATLAB scripts work on dense matrices, not sparse ones. As a result, the MATLAB GB_spec_* function are not meant for production use, but just for testing and illustration.
- integer operations: MATLAB and GraphBLAS handle integer operations differently. In MATLAB, an integer result outside the range of the integer is set to maximum or minimum integer. For example, int8(127)+1 is 127. This is useful for many computations such as image processing, but GraphBLAS follows the C rules instead, where integer values wrap, modulo style. For example, in GraphBLAS and in C, incrementing (int8_t) 127 by one results in -128. Of course, an alternative would be for a MATLAB interface to create its own integer operators, each of which would follow the MATLAB integer rules of arithmetic. However, this would obscure the purpose of these GB_spec_* and GB_mex_* test functions, which is to test the C API of GraphBLAS. When the GB_spec_* functions need to perform integer computations and typecasting, they call GraphBLAS to do the work, instead doing the work in MATLAB. This ensures that the GB_spec_* functions obtain the same results as their GraphBLAS counterparts.
- elegance: to simplify testing, each MATLAB mexFunction interface a GraphBLAS function is a direct translation of the C API. For example, GB_mex_mxm is a direct interface to the GraphBLAS GrB_mxm, even down the order of parameters. This approach abandons some of the potential features of MATLAB for creating elegant M-file interfaces in a

highly usable form, such as the ability to provide fewer parameters when optional parameters are not in use. These mexFunctions, as written, are not meant to be usable in a user application. They are not highly documented. They are meant to be fast, and direct, to accomplish the goal of testing SuiteSparse:GraphBLAS in MATLAB and comparing their results with the corresponding GB_spec_* function. They are not recommended for use in general applications in MATLAB.

• generality: the MATLAB mexFunction interface needs to test the C API directly, so it must access content of SuiteSparse:GraphBLAS objects that are normally opaque to an end user application. As a result, these mexFunctions do not serve as a general interface to any conforming GraphBLAS implementation, but only to SuiteSparse:GraphBLAS.

In the MATLAB mimic functions, GB_spec_*, a GraphBLAS matrix A is represented as a MATLAB struct with the following components:

- A.matrix: the values of the matrix. If A.matrix is a sparse double matrix, it holds a typecasted copy of the values of a GraphBLAS matrix, unless the GraphBLAS matrix is also double (GrB_FP64).
- A.pattern: a logical matrix holding the pattern; A.pattern(i,j)=true if (i,j) is in the pattern of A, and false otherwise.
- A.class: the MATLAB class of the matrix corresponding to one of the eleven built-in types. Normally this is simply class(A.matrix).
- A.values: most of the GraphBLAS test mexFunctions return their result as a MATLAB sparse matrix, in the double class. This works well for all types except for the 64-bit integer types, since a double has about 54 bits of mantissa which is less than the 64 bits available in a long integer. To ensure no bits are lots, these values are also returned as a vector. This enables GB_spec_compare to ensure the test results are identical down to the very last bit, and not just to within roundoff error. Nearly all tests, even in double precision, check for perfect equality, not just for results accurate to within round-off error.

8.2 GrB_mxm: matrix-matrix multiply

```
GrB_Info GrB_mxm
                                     // C<Mask> = accum (C, A*B)
                                    // input/output matrix for results
    GrB_Matrix C,
   const GrB_Matrix Mask,
                                    // optional mask for C, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C,T)
   const GrB_Semiring semiring,
                                    // defines '+' and '*' for A*B
    const GrB_Matrix A,
                                    // first input: matrix A
    const GrB_Matrix B,
                                    // second input: matrix B
    const GrB_Descriptor desc
                                    // descriptor for C, Mask, A, and B
);
```

GrB_mxm multiplies two sparse matrices A and B using the semiring. The input matrices A and B may be transposed according to the descriptor, desc (which may be NULL) and then typecasted to match the multiply operator of the semiring. Next, T=A*B is computed on the semiring, precisely defined in the GB_spec_mxm.m script. The actual algorithm exploits sparsity and does not take $O(n^3)$ time, but what computes is the following:

```
[m s] = size (A.matrix) ;
[s n] = size (B.matrix) ;
T.matrix = zeros (m, n, multiply.ztype) ;
T.pattern = zeros (m, n, 'logical') ;
T.matrix (:,:) = identity ;
                                        % the identity of the semiring's monoid
T.class = multiply.ztype ;
                                        % the ztype of the semiring's multiply op
A = cast (A.matrix, multiply.xtype);
                                       % the xtype of the semiring's multiply op
                                        % the ytype of the semiring's multiply op
B = cast (B.matrix, multiply.ytype);
for j = 1:n
   for i = 1:m
       for k = 1:s
            % T (i,j) += A (i,k) * B (k,j), using the semiring
            if (A.pattern (i,k) && B.pattern (k,j))
                z = multiply (A (i,k), B (k,j));
                T.matrix (i,j) = add (T.matrix (i,j), z);
                T.pattern(i,j) = true;
            end
        end
    end
end
```

Finally, T is typecasted into the type of C, and the results are written back into C via the accum and Mask, $\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot\mathbf{T}$. The latter step is reflected in the MATLAB function $\mathtt{GB_spec_accum_mask.m}$, discussed in Section 2.3.

Performance considerations: Suppose all matrices are in GxB_BY_COL format, and B is extremely sparse but A is not as sparse. Then computing C=A*B is very fast, and much faster than when A is extremely sparse. For example, if A is square and B is a column vector that is all nonzero except for one entry B(j,0)=1, then C=A*B is the same as extracting column A(:,j). This is very fast if A is stored by column but slow if A is stored by row. If A is a sparse row with a single entry A(0,i)=1, then C=A*B is the same as extracting row B(i,:). This is fast if B is stored by row but slow if B is stored by column.

If the user application needs to repeatedly extract rows and columns from a matrix, whether by matrix multiplication or by <code>GrB_extract</code>, then keep two copies: one stored by row, and other by column, and use the copy that results in the fastest computation.

8.3 GrB_vxm: vector-matrix multiply

```
GrB_Info GrB_vxm
                                    // w'<mask> = accum (w, u'*A)
   GrB_Vector w,
                                    // input/output vector for results
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
                                    // optional accum for z=accum(w,t)
    const GrB_BinaryOp accum,
                                    // defines '+' and '*' for u'*A
    const GrB_Semiring semiring,
    const GrB_Vector u,
                                    // first input: vector u
    const GrB_Matrix A,
                                    // second input: matrix A
    const GrB_Descriptor desc
                                    // descriptor for w, mask, and A
);
```

GrB_vxm multiplies a row vector u' times a matrix A. The matrix A may be first transposed according to desc (as the second input, GrB_INP1); the column vector u is never transposed via the descriptor. The inputs u and A are typecasted to match the xtype and ytype inputs, respectively, of the multiply operator of the semiring. Next, an intermediate column vector t=A'*u is computed on the semiring using the same method as GrB_mxm. Finally, the column vector t is typecasted from the ztype of the multiply operator of the semiring into the type of w, and the results are written back into w using the optional accumulator accum and mask.

The last step is $\mathbf{w}\langle \mathbf{m} \rangle = \mathbf{w} \odot \mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

Performance considerations: If the GxB_FORMAT of A is GxB_BY_ROW, and the default descriptor is used (A is not transposed), then GrB_vxm is faster than than GrB_mxv with its default descriptor, when the vector u is very sparse. However, if the GxB_FORMAT of A is GxB_BY_COL, then GrB_mxv with its default descriptor is faster than GrB_vxm with its default descriptor, when the vector u is very sparse. Using the non-default GrB_TRAN descriptor for A makes the GrB_vxm operation equivalent to GrB_mxv with its default descriptor (with the operands reversed in the multiplier, as well). The reverse is true as well; GrB_mxv with GrB_TRAN is the same as GrB_vxm with a default descriptor.

The breadth-first search presented in Section 11.2 of this User Guide uses GrB_vxm instead of GrB_mxv, since the default format in SuiteSparse:GraphBLAS is GxB_BY_ROW. If the matrix is stored by column, then use GrB_mxv instead.

8.4 GrB_mxv: matrix-vector multiply

```
GrB_Info GrB_mxv
                                    // w<mask> = accum (w, A*u)
   GrB_Vector w,
                                    // input/output vector for results
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w,t)
                                    // defines '+' and '*' for A*B
    const GrB_Semiring semiring,
    const GrB_Matrix A,
                                    // first input: matrix A
    const GrB_Vector u,
                                    // second input: vector u
    const GrB_Descriptor desc
                                    // descriptor for w, mask, and A
);
```

GrB_mxv multiplies a matrix A times a column vector u. The matrix A may be first transposed according to desc (as the first input); the column vector u is never transposed via the descriptor. The inputs A and u are typecasted to match the xtype and ytype inputs, respectively, of the multiply operator of the semiring. Next, an intermediate column vector t=A*u is computed on the semiring using the same method as GrB_mxm. Finally, the column vector t is typecasted from the ztype of the multiply operator of the semiring into the type of w, and the results are written back into w using the optional accumulator accum and mask.

The last step is $\mathbf{w}\langle \mathbf{m} \rangle = \mathbf{w} \odot \mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

Performance considerations: Refer to the discussion of GrB_vxm. In SuiteSparse:GraphBLAS, GrB_mxv is very efficient when u is sparse or dense, when the default descriptor is used, and when the matrix is GxB_BY_COL. When u is very sparse and GrB_INPO is set to its non-default GrB_TRAN, then this method is not efficient if the matrix is in GxB_BY_COL format. If an application needs to perform A'*u repeatedly where u is very sparse, then use the GxB_BY_ROW format for A instead.

8.5 GrB_eWiseMult: element-wise operations, set intersection

Element-wise "multiplication" is shorthand for applying a binary operator element-wise on two matrices or vectors A and B, for all entries that appear in the set intersection of the patterns of A and B. This is like A.*B for two sparse matrices in MATLAB, except that in GraphBLAS any binary operator can be used, not just multiplication.

The pattern of the result of the element-wise "multiplication" is exactly this set intersection. Entries in A but not B, or visa versa, do not appear in the result.

Let \otimes denote the binary operator to be used. The computation $\mathbf{T} = \mathbf{A} \otimes \mathbf{B}$ is given below. Entries not in the intersection of \mathbf{A} and \mathbf{B} do not appear in the pattern of \mathbf{T} . That is:

for all entries
$$(i, j)$$
 in $\mathbf{A} \cap \mathbf{B}$
 $t_{ij} = a_{ij} \otimes b_{ij}$

Depending on what kind of operator is used and what the implicit value is assumed to be, this can give the Hadamard product. This is the case for A.*B in MATLAB since the implicit value is zero. However, computing a Hadamard product is not necessarily the goal of the eWiseMult operation. It simply applies any binary operator, built-in or user-defined, to the set intersection of A and B, and discards any entry outside this intersection. Its usefulness in a user's application does not depend upon it computing a Hadamard product in all cases. The operator need not be associative, commutative, nor have any particular property except for type compatibility with A and B, and the output matrix C.

The generic name for this operation is **GrB_eWiseMult**, which can be used for both matrices and vectors.

8.5.1 GrB_eWiseMult_Vector: element-wise vector multiply

```
GrB_Info GrB_eWiseMult
                                    // w<mask> = accum (w, u.*v)
    GrB_Vector w,
                                    // input/output vector for results
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w,t)
    const <operator> multiply,
                                    // defines '.*' for t=u.*v
    const GrB_Vector u,
                                    // first input: vector u
    const GrB_Vector v,
                                    // second input: vector v
    const GrB_Descriptor desc
                                    // descriptor for w and mask
);
```

GrB_Vector_eWiseMult computes the element-wise "multiplication" of two vectors \mathbf{u} and \mathbf{v} , element-wise using any binary operator (not just times). The vectors are not transposed via the descriptor. The vectors \mathbf{u} and \mathbf{v} are first typecasted into the first and second inputs of the multiply operator. Next, a column vector \mathbf{t} is computed, denoted $\mathbf{t} = \mathbf{u} \otimes \mathbf{v}$. The pattern of \mathbf{t} is the set intersection of \mathbf{u} and \mathbf{v} . The result \mathbf{t} has the type of the output ztype of the multiply operator.

The operator is typically a GrB_BinaryOp, but the method is type-generic for this parameter. If given a monoid (GrB_Monoid), the additive operator of the monoid is used as the multiply binary operator. If given a semiring (GrB_Semiring), the multiply operator of the semiring is used as the multiply binary operator.

The next and final step is $\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot\mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices. Note for all GraphBLAS operations, including this one, the accumulator $\mathbf{w}\odot\mathbf{t}$ is always applied in a set union manner, even though $\mathbf{t} = \mathbf{u}\otimes\mathbf{v}$ for this operation is applied in a set intersection manner.

8.5.2 GrB_eWiseMult_Matrix: element-wise matrix multiply

```
GrB_Info GrB_eWiseMult
                                    // C<Mask> = accum (C, A.*B)
    GrB_Matrix C,
                                    // input/output matrix for results
    const GrB_Matrix Mask,
                                    // optional mask for C, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C,T)
    const <operator> multiply,
                                    // defines '.*' for T=A.*B
    const GrB_Matrix A,
                                    // first input: matrix A
    const GrB_Matrix B,
                                    // second input: matrix B
    const GrB_Descriptor desc
                                    // descriptor for C, Mask, A, and B
);
```

GrB_Matrix_eWiseMult computes the element-wise "multiplication" of two matrices A and B, element-wise using any binary operator (not just times). The input matrices may be transposed first, according to the descriptor desc. They are then typecasted into the first and second inputs of the multiply operator. Next, a matrix T is computed, denoted $\mathbf{T} = \mathbf{A} \otimes \mathbf{B}$. The pattern of T is the set intersection of A and B. The result T has the type of the output ztype of the multiply operator.

The multiply operator is typically a GrB_BinaryOp, but the method is type-generic for this parameter. If given a monoid (GrB_Monoid), the additive operator of the monoid is used as the multiply binary operator. If given a semiring (GrB_Semiring), the multiply operator of the semiring is used as the multiply binary operator.

The operation can be expressed in MATLAB notation as:

```
[nrows, ncols] = size (A.matrix);
T.matrix = zeros (nrows, ncols, multiply.ztype);
T.class = multiply.ztype;
p = A.pattern & B.pattern;
A = cast (A.matrix (p), multiply.xtype);
B = cast (B.matrix (p), multiply.ytype);
T.matrix (p) = multiply (A, B);
T.pattern = p;
```

The final step is $\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{C} \odot \mathbf{T}$, as described in Section 2.3. Note for all GraphBLAS operations, including this one, the accumulator $\mathbf{C} \odot \mathbf{T}$ is always applied in a set union manner, even though $\mathbf{T} = \mathbf{A} \otimes \mathbf{B}$ for this operation is applied in a set intersection manner.

8.6 GrB_eWiseAdd: element-wise operations, set union

Element-wise "addition" is shorthand for applying a binary operator element-wise on two matrices or vectors A and B, for all entries that appear in the set intersection of the patterns of A and B. This is like A+B for two sparse matrices in MATLAB, except that in GraphBLAS any binary operator can be used, not just addition. The pattern of the result of the element-wise "addition" is the set union of the pattern of A and B. Entries in neither in A nor in B do not appear in the result.

Let \oplus denote the binary operator to be used. The computation $\mathbf{T} = \mathbf{A} \oplus \mathbf{B}$ is exactly the same as the computation with accumulator operator as described in Section 2.3. It acts like a sparse matrix addition, except that any operator can be used. The pattern of $\mathbf{A} \oplus \mathbf{B}$ is the set union of the patterns of \mathbf{A} and \mathbf{B} , and the operator is applied only on the set intersection of \mathbf{A} and \mathbf{B} . Entries not in either the pattern of \mathbf{A} or \mathbf{B} do not appear in the pattern of \mathbf{T} . That is:

```
for all entries (i, j) in \mathbf{A} \cap \mathbf{B}

t_{ij} = a_{ij} \oplus b_{ij}

for all entries (i, j) in \mathbf{A} \setminus \mathbf{B}

t_{ij} = a_{ij}

for all entries (i, j) in \mathbf{B} \setminus \mathbf{A}

t_{ij} = b_{ij}
```

The only difference between element-wise "multiplication" ($\mathbf{T} = \mathbf{A} \otimes \mathbf{B}$) and "addition" ($\mathbf{T} = \mathbf{A} \oplus \mathbf{B}$) is the pattern of the result, and what happens to entries outside the intersection. With \otimes the pattern of \mathbf{T} is the intersection; with \oplus it is the set union. Entries outside the set intersection are dropped for \otimes , and kept for \oplus ; in both cases the operator is only applied to those (and only those) entries in the intersection. Any binary operator can be used interchangeably for either operation.

Element-wise operations do not operate on the implicit values, even implicitly, since the operations make no assumption about the semiring. As a result, the results can be different from MATLAB, which can always assume the implicit value is zero. For example, C=A-B is the conventional matrix subtraction in MATLAB. Computing A-B in GraphBLAS with eWiseAdd will apply the MINUS operator to the intersection, entries in A but not B will be unchanged and appear in C, and entries in neither A nor B do not appear in C. For these cases, the results matches the MATLAB C=A-B. Entries in B but not A do appear in C but they are not negated; they cannot be subtracted

from an implicit value in A. This is by design. If conventional matrix subtraction of two sparse matrices is required, and the implicit value is known to be zero, use GrB_apply to negate the values in B, and then use eWiseAdd with the PLUS operator, to compute A+(-B).

The generic name for this operation is GrB_eWiseAdd, which can be used for both matrices and vectors.

There is another minor difference in two variants of the element-wise functions. If given a semiring, the eWiseAdd functions use the binary operator of the semiring's monoid, while the eWiseMult functions use the multiplicative operator of the semiring.

8.6.1 GrB_eWiseAdd_Vector: element-wise vector addition

```
// w<mask> = accum (w, u+v)
GrB_Info GrB_eWiseAdd
    GrB_Vector w,
                                    // input/output vector for results
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w,t)
    const <operator> add,
                                    // defines '+' for t=u+v
    const GrB_Vector u,
                                    // first input: vector u
    const GrB_Vector v,
                                    // second input: vector v
    const GrB_Descriptor desc
                                    // descriptor for w and mask
);
```

GrB_Vector_eWiseAdd computes the element-wise "addition" of two vectors u and v, element-wise using any binary operator (not just plus). The vectors are not transposed via the descriptor. Entries in the intersection of u and v are first typecasted into the first and second inputs of the add operator. Next, a column vector t is computed, denoted $t = u \oplus v$. The pattern of t is the set union of u and v. The result t has the type of the output ztype of the add operator.

The add operator is typically a GrB_BinaryOp, but the method is type-generic for this parameter. If given a monoid (GrB_Monoid), the additive operator of the monoid is used as the add binary operator. If given a semiring (GrB_Semiring), the additive operator of the monoid of the semiring is used as the add binary operator.

The final step is $\mathbf{w}\langle \mathbf{m} \rangle = \mathbf{w} \odot \mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

8.6.2 GrB eWiseAdd Matrix: element-wise matrix addition

```
GrB_Info GrB_eWiseAdd
                                    // C<Mask> = accum (C, A+B)
    GrB_Matrix C,
                                    // input/output matrix for results
    const GrB_Matrix Mask,
                                    // optional mask for C, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C,T)
                                    // defines '+' for T=A+B
    const <operator> add,
    const GrB_Matrix A,
                                    // first input: matrix A
    const GrB_Matrix B,
                                    // second input: matrix B
    const GrB_Descriptor desc
                                    // descriptor for C, Mask, A, and B
);
```

GrB_Matrix_eWiseAdd computes the element-wise "addition" of two matrices A and B, element-wise using any binary operator (not just plus). The input matrices may be transposed first, according to the descriptor desc. Entries in the intersection then typecasted into the first and second inputs of the add operator. Next, a matrix T is computed, denoted $\mathbf{T} = \mathbf{A} \oplus \mathbf{B}$. The pattern of T is the set union of A and B. The result T has the type of the output ztype of the add operator.

The add operator is typically a GrB_BinaryOp, but the method is type-generic for this parameter. If given a monoid (GrB_Monoid), the additive operator of the monoid is used as the add binary operator. If given a semiring (GrB_Semiring), the additive operator of the monoid of the semiring is used as the add binary operator.

The operation can be expressed in MATLAB notation as:

```
[nrows, ncols] = size (A.matrix);
T.matrix = zeros (nrows, ncols, add.ztype);
p = A.pattern & B.pattern;
A = GB_mex_cast (A.matrix (p), add.xtype);
B = GB_mex_cast (B.matrix (p), add.ytype);
T.matrix (p) = add (A, B);
p = A.pattern & ~B.pattern; T.matrix (p) = cast (A.matrix (p), add.ztype);
p = ~A.pattern & B.pattern; T.matrix (p) = cast (B.matrix (p), add.ztype);
T.pattern = A.pattern | B.pattern;
T.class = add.ztype;
```

Except for when typecasting is performed, this is identical to how the accum operator is applied in Figure 1.

The final step is $C(M) = C \odot T$, as described in Section 2.3.

8.7 GrB_extract: submatrix extraction

The GrB_extract function is a generic name for three specific functions: GrB_Vector_extract, GrB_Col_extract, and GrB_Matrix_extract. The generic name appears in the function signature, but the specific function name is used when describing what each variation does.

8.7.1 GrB_Vector_extract: extract subvector from vector

```
GrB_Info GrB_extract
                                    // w<mask> = accum (w, u(I))
                                    // input/output vector for results
    GrB_Vector w,
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
                                    // optional accum for z=accum(w,t)
    const GrB_BinaryOp accum,
    const GrB_Vector u,
                                    // first input: vector u
    const GrB_Index *I,
                                    // row indices
    const GrB_Index ni,
                                    // number of row indices
    const GrB_Descriptor desc
                                    // descriptor for w and mask
) ;
```

GrB_Vector_extract extracts a subvector from another vector, identical to $\mathbf{t} = \mathbf{u}$ (I) in MATLAB where I is an integer vector of row indices. Refer to GrB_Matrix_extract for further details; vector extraction is the same as matrix extraction with n-by-1 matrices. See Section 7 for a description of I and ni. The final step is $\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot\mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

8.7.2 GrB Matrix extract: extract submatrix from matrix

```
GrB_Info GrB_extract
                                    // C<Mask> = accum (C, A(I,J))
    GrB_Matrix C,
                                    // input/output matrix for results
    const GrB_Matrix Mask,
                                    // optional mask for C, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C,T)
    const GrB_Matrix A,
                                    // first input: matrix A
    const GrB_Index *I,
                                    // row indices
    const GrB_Index ni,
                                    // number of row indices
                                    // column indices
    const GrB_Index *J,
    const GrB_Index nj,
                                    // number of column indices
    const GrB_Descriptor desc
                                    // descriptor for C, Mask, and A
);
```

GrB_Matrix_extract extracts a submatrix from another matrix, identical to T = A(I,J) in MATLAB where I and J are integer vectors of row and column indices, respectively, except that indices are zero-based in Graph-BLAS and one-based in MATLAB. The input matrix A may be transposed first, via the descriptor. The type of T and A are the same. The size of C is |I|-by-|J|. Entries outside A(I,J) are not accessed and do not take part in the computation. More precisely, assuming the matrix A is not transposed, the matrix T is defined as follows:

If duplicate indices are present in I or J, the above method defines the result in T. Duplicates result in the same values of A being copied into different places in T. See Section 7 for a description of the row indices I and ni, and the column indices J and nj. The final step is $\mathbf{C}\langle\mathbf{M}\rangle=\mathbf{C}\odot\mathbf{T}$, as described in Section 2.3.

Performance considerations: If A is not transposed via input descriptor: if |I| is small, then it is fastest if A is GxB_BY_ROW; if |J| is small, then it is fastest if A is GxB_BY_COL. The opposite is true if A is transposed.

8.7.3 GrB_Col_extract: extract column vector from matrix

```
GrB_Info GrB_extract
                                    // w<mask> = accum (w, A(I,j))
    GrB_Vector w,
                                    // input/output matrix for results
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w,t)
    const GrB_Matrix A,
                                    // first input: matrix A
    const GrB_Index *I,
                                    // row indices
    const GrB_Index ni,
                                    // number of row indices
    const GrB_Index j,
                                    // column index
    const GrB_Descriptor desc
                                    // descriptor for w, mask, and A
);
```

GrB_Col_extract extracts a subvector from a matrix, identical to t = A (I,j) in MATLAB where I is an integer vector of row indices and where j is a single column index. The input matrix A may be transposed first, via the descriptor, which results in the extraction of a single row j from the matrix A, the result of which is a column vector w. The type of t and A are the same. The size of w is |I|-by-1.

See Section 7 for a description of the row indices I and ni. The final step is $\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot\mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

Performance considerations: If A is not transposed: it is fastest if the format of A is GxB_BY_COL. The opposite is true if A is transposed.

8.8 GxB_subassign: submatrix assignment

The methods described in this section are all variations of the form C(I,J)=A, which modifies a submatrix of the matrix C. All methods can be used in their generic form with the single name GxB_subassign. This is reflected in the prototypes. However, to avoid confusion between the different kinds of assignment, the name of the specific function is used when describing each variation. If the discussion applies to all variations, the simple name GxB_subassign is used.

See Section 7 for a description of the row indices I and ni, and the column indices J and nj.

GxB_subassign is very similar to GrB_assign, described in Section 8.9. The two operations are compared and contrasted in Section 8.11. For a discussion of how duplicate indices are handled in I and J, see Section 8.10.

SPEC: All variants of GxB_subassign are extensions to the spec.

8.8.1 GxB_Vector_subassign: assign to a subvector

```
GrB_Info GxB_subassign
                                    // w(I) < mask > = accum (w(I),u)
    GrB_Vector w,
                                    // input/output matrix for results
    const GrB_Vector mask,
                                    // optional mask for w(I), unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w(I),t)
    const GrB_Vector u,
                                    // first input: vector u
    const GrB_Index *I,
                                    // row indices
                                    // number of row indices
    const GrB_Index ni,
    const GrB_Descriptor desc
                                    // descriptor for w(I) and mask
);
```

GxB_Vector_subassign operates on a subvector w(I) of w, modifying it with the vector u. The method is identical to GxB_Matrix_subassign described in Section 8.8.2, where all matrices have a single column each. The mask has the same size as w(I) and u. The only other difference is that the input u in this method is not transposed via the GrB_INPO descriptor.

8.8.2 GxB_Matrix_subassign: assign to a submatrix

```
GrB_Info GxB_subassign
                                    // C(I,J) < Mask > = accum (C(I,J),A)
    GrB_Matrix C,
                                    // input/output matrix for results
    const GrB_Matrix Mask,
                                    // optional mask for C(I,J), unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C(I,J),T)
                                    // first input: matrix A
    const GrB_Matrix A,
    const GrB_Index *I,
                                    // row indices
                                    // number of row indices
    const GrB_Index ni,
    const GrB_Index *J,
                                    // column indices
    const GrB_Index nj,
                                    // number of column indices
    const GrB_Descriptor desc
                                    // descriptor for C(I,J), Mask, and A
);
```

GxB_Matrix_subassign operates only on a submatrix S of C, modifying it with the matrix A. For this operation, the result is not the entire matrix C, but a submatrix S=C(I,J) of C. The steps taken are as follows, except that A may be optionally transposed via the GrB_INPO descriptor option.

Step	GraphBLAS	description
	notation	
1	S = C(I, J)	extract the $C(I, J)$ submatrix
2	$\mathbf{S}\langle\mathbf{M} angle=\mathbf{S}\odot\mathbf{A}$	apply the accumulator/mask to the submatrix ${f S}$
3	$\mathbf{C}(\mathbf{I},\mathbf{J}) = \mathbf{S}$	put the submatrix S back into $C(I, J)$

The accumulator/mask step in Step 2 is the same as for all other Graph-BLAS operations, described in Section 2.3, except that for $GxB_subassign$, it is applied to just the submatrix S = C(I, J), and thus the Mask has the same size as A, S, and C(I, J).

The GxB_subassign operation is the reverse of matrix extraction:

- For submatrix extraction, GrB_Matrix_extract, the submatrix A(I,J) appears on the right-hand side of the assignment, C=A(I,J), and entries outside of the submatrix are not accessed and do not take part in the computation.
- For submatrix assignment, GxB_Matrix_subassign, the submatrix C(I, J) appears on the left-hand-side of the assignment, C(I, J)=A, and entries outside of the submatrix are not accessed and do not take part in the computation.

In both methods, the accumulator and mask modify the submatrix of the assignment; they simply differ on which side of the assignment the submatrix resides on. In both cases, if the Mask matrix is present it is the same size as the submatrix:

- For submatrix extraction, $\mathbf{C}\langle\mathbf{M}\rangle=\mathbf{C}\odot\mathbf{A}(\mathbf{I},\mathbf{J})$ is computed, where the submatrix is on the right. The mask \mathbf{M} has the same size as the submatrix $\mathbf{A}(\mathbf{I},\mathbf{J})$.
- For submatrix assignment, $\mathbf{C}(\mathbf{I}, \mathbf{J})\langle \mathbf{M} \rangle = \mathbf{C}(\mathbf{I}, \mathbf{J}) \odot \mathbf{A}$ is computed, where the submatrix is on the left. The mask \mathbf{M} has the same size as the submatrix $\mathbf{C}(\mathbf{I}, \mathbf{J})$.

In Step 1, the submatrix S is first computed by the GrB_Matrix_extract operation, S=C(I,J).

Step 2 accumulates the results $S\langle M \rangle = S \odot T$, exactly as described in Section 2.3, but operating on the submatrix S, not C, using the optional Mask and accum operator. The matrix T is simply T = A, or $T = A^T$ if A is transposed via the desc descriptor, GrB_INPO . The $GrB_REPLACE$ option in the descriptor clears S after computing C = T or $C = C \odot T$, not all of C since this operation can only modify the specified submatrix of C.

Finally, Step 3 writes the result (which is the modified submatrix S and not all of C) back into the C matrix that contains it, via the assignment C(I,J)=S, using the reverse operation from the method described for matrix extraction:

Performance considerations: If A is not transposed: if |I| is small, then it is fastest if the format of C is GxB_BY_ROW; if |J| is small, then it is fastest if the format of C is GxB_BY_COL. The opposite is true if A is transposed.

8.8.3 GxB_Col_subassign: assign to a sub-column of a matrix

```
GrB_Info GxB_subassign
                                    // C(I,j) < mask > = accum (C(I,j),u)
    GrB_Matrix C,
                                    // input/output matrix for results
                                    // optional mask for C(I,j), unused if NULL
    const GrB_Vector mask,
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(C(I,j),t)
   const GrB_Vector u,
                                    // input vector
    const GrB_Index *I,
                                    // row indices
    const GrB_Index ni,
                                    // number of row indices
    const GrB_Index j,
                                    // column index
    const GrB_Descriptor desc
                                    // descriptor for C(I,j) and mask
);
```

GxB_Col_subassign modifies a single sub-column of a matrix C. It is the same as GxB_Matrix_subassign where the index vector J[0]=j is a single column index (and thus nj=1), and where all matrices in GxB_Matrix_subassign (except C) consist of a single column. The mask vector has the same size as u and the sub-column C(I,j). The input descriptor GrB_INPO is ignored; the input vector u is not transposed. Refer to GxB_Matrix_subassign for further details.

Performance considerations: GxB_Col_subassign is much faster than GxB_Row_subassign if the format of C is GxB_BY_COL. GxB_Row_subassign is much faster than GxB_Col_subassign if the format of C is GxB_BY_ROW.

8.8.4 GxB_Row_subassign: assign to a sub-row of a matrix

```
// C(i,J) < mask' > = accum (C(i,J),u')
GrB_Info GxB_subassign
                                    // input/output matrix for results
    GrB_Matrix C,
    const GrB_Vector mask,
                                    // optional mask for C(i,J), unused if NULL
                                    // optional accum for z=accum(C(i,J),t)
    const GrB_BinaryOp accum,
    const GrB_Vector u,
                                    // input vector
    const GrB_Index i,
                                    // row index
    const GrB_Index *J,
                                    // column indices
                                    // number of column indices
    const GrB_Index nj,
    const GrB_Descriptor desc
                                    // descriptor for C(i,J) and mask
);
```

GxB_Row_subassign modifies a single sub-row of a matrix C. It is the same as GxB_Matrix_subassign where the index vector I[0]=i is a single

row index (and thus ni=1), and where all matrices in GxB_Matrix_subassign (except C) consist of a single row. The mask vector has the same size as u and the sub-column C(I,j). The input descriptor GrB_INPO is ignored; the input vector u is not transposed. Refer to GxB_Matrix_subassign for further details.

Performance considerations: GxB_Col_subassign is much faster than GxB_Row_subassign if the format of C is GxB_BY_COL. GxB_Row_subassign is much faster than GxB_Col_subassign if the format of C is GxB_BY_ROW.

8.8.5 GxB_Vector_subassign_<type>: assign a scalar to a subvector

```
GrB_Info GxB_subassign
                                       // w(I) < mask > = accum (w(I),x)
   GrB_Vector w,
                                    // input/output vector for results
                                    // optional mask for w(I), unused if NULL
    const GrB_Vector mask,
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w(I),x)
    const <type> x,
                                    // scalar to assign to w(I)
    const GrB_Index *I,
                                    // row indices
    const GrB_Index ni,
                                    // number of row indices
   const GrB_Descriptor desc
                                    // descriptor for w(I) and mask
);
```

GxB_Vector_subassign_<type> assigns a single scalar to an entire subvector of the vector w. The operation is exactly like setting a single entry in an n-by-1 matrix, A(I,0) = x, where the column index for a vector is implicitly j=0. For further details of this function, see GxB_Matrix_subassign_<type> in Section 8.8.6.

8.8.6 GxB_Matrix_subassign_<type>: assign a scalar to a submatrix

```
GrB_Info GxB_subassign
                                        // C(I,J) < Mask > = accum (C(I,J),x)
    GrB_Matrix C,
                                     // input/output matrix for results
    const GrB_Matrix Mask,
                                     // optional mask for C(I,J), unused if NULL
    const GrB_BinaryOp accum,
                                     // optional accum for Z=accum(C(I,J),x)
                                     // scalar to assign to C(I,J)
    const <type> x,
    const GrB_Index *I,
                                     // row indices
    const GrB_Index ni,
                                     // number of row indices
    const GrB_Index *J,
                                     // column indices
    const GrB_Index nj,
                                     // number of column indices
    const GrB_Descriptor desc
                                     // descriptor for C(I,J) and Mask
) ;
```

GxB_Matrix_subassign_<type> assigns a single scalar to an entire submatrix of C, like the scalar expansion C(I,J)=x in MATLAB. The scalar x is implicitly expanded into a matrix A of size ni by nj, and then the matrix A is assigned to C(I,J) using the same method as in GxB_Matrix_subassign. Refer to that function in Section 8.8.2 for further details. For the accumulation step, the scalar x is typecasted directly into the type of C when the accum operator is not applied to it, or into the ytype of the accum operator, if accum is not NULL, for entries that are already present in C.

The <type> x notation is otherwise the same as GrB_Matrix_setElement (see Section 5.9.9). Any value can be passed to this function and its type will be detected, via the _Generic feature of ANSI C11. For a user-defined type, x is a void * pointer that points to a memory space holding a single entry of a scalar that has exactly the same user-defined type as the matrix C. This user-defined type must exactly match the user-defined type of C since no typecasting is done between user-defined types.

If a void * pointer is passed in and the type of the underlying scalar does not exactly match the user-defined type of C, then results are undefined. No error status will be returned since GraphBLAS has no way of catching this error.

Performance considerations: If A is not transposed: if |I| is small, then it is fastest if the format of C is GxB_BY_ROW; if |J| is small, then it is fastest if the format of C is GxB_BY_COL. The opposite is true if A is transposed.

8.9 GrB_assign: submatrix assignment

The methods described in this section are all variations of the form C(I, J)=A, which modifies a submatrix of the matrix C. All methods can be used in their generic form with the single name GrB_assign. These methods are very similar to their GxB_subassign counterparts in Section 8.8. They differ primarily in the size of the Mask, and how the GrB_REPLACE option works. Refer to Section 8.11 for a complete comparison of GxB_subassign and GrB_assign. See Section 7 for a description of I, ni, J, and nj.

8.9.1 GrB_Vector_assign: assign to a subvector

```
GrB_Info GrB_assign
                                    // w<mask>(I) = accum (w(I),u)
    GrB_Vector w,
                                    // input/output matrix for results
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
   const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w(I),t)
    const GrB_Vector u,
                                    // first input: vector u
    const GrB_Index *I,
                                    // row indices
    const GrB_Index ni,
                                    // number of row indices
    const GrB_Descriptor desc
                                    // descriptor for w and mask
);
```

GrB_Vector_assign operates on a subvector w(I) of w, modifying it with the vector u. The mask vector has the same size as w. The method is identical to GrB_Matrix_assign described in Section 8.9.2, where all matrices have a single column each. The only other difference is that the input u in this method is not transposed via the GrB_INPO descriptor.

8.9.2 GrB_Matrix_assign: assign to a submatrix

```
GrB_Info GrB_assign
                                    // C<Mask>(I,J) = accum (C(I,J),A)
    GrB_Matrix C,
                                    // input/output matrix for results
                                    // optional mask for C, unused if NULL
    const GrB_Matrix Mask,
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C(I,J),T)
                                    // first input: matrix A
    const GrB_Matrix A,
    const GrB_Index *I,
                                    // row indices
                                    // number of row indices
    const GrB_Index ni,
    const GrB_Index *J,
                                    // column indices
    const GrB_Index nj,
                                    // number of column indices
    const GrB_Descriptor desc
                                    // descriptor for C, Mask, and A
);
```

GrB_Matrix_assign operates on a submatrix S of C, modifying it with the matrix A. It may also modify all of C, depending on the input descriptor desc and the Mask.

Step	GraphBLAS	description
	notation	
1	$\mathbf{S} = \mathbf{C}(\mathbf{I}, \mathbf{J})$	extract $\mathbf{C}(\mathbf{I}, \mathbf{J})$ submatrix
2	$\mathbf{S} = \mathbf{S} \odot \mathbf{A}$	apply the accumulator (but not the mask) to ${\bf S}$
3	$\mathbf{Z} = \mathbf{C}$	make a copy of \mathbf{C}
4	$\mathbf{Z}(\mathbf{I},\mathbf{J}) = \mathbf{S}$	put the submatrix into $\mathbf{Z}(\mathbf{I}, \mathbf{J})$
5	$\mathbf{C}\langle\mathbf{M} angle=\mathbf{Z}$	apply the mask/replace phase to all of ${\bf C}$

In contrast to GxB_subassign, the Mask has the same as C.

Step 1 extracts the submatrix and then Step 2 applies the accumulator (or S = A if accum is NULL). The Mask is not yet applied.

Step 3 makes a copy of the C matrix, and then Step 4 writes the submatrix S into Z. This is the same as Step 3 of GxB_subassign, except that it operates on a temporary matrix Z.

Finally, Step 5 writes \mathbf{Z} back into \mathbf{C} via the Mask, using the Mask/Replace Phase described in Section 2.3. If $\mathtt{GrB_REPLACE}$ is enabled, then all of \mathbf{C} is cleared prior to writing \mathbf{Z} via the mask. As a result, the $\mathtt{GrB_REPLACE}$ option can delete entries outside the $\mathbf{C}(\mathbf{I},\mathbf{J})$ submatrix.

Performance considerations: If A is not transposed: if |I| is small, then it is fastest if the format of C is GxB_BY_ROW; if |J| is small, then it is fastest if the format of C is GxB_BY_COL. The opposite is true if A is transposed.

8.9.3 GrB_Col_assign: assign to a sub-column of a matrix

```
GrB_Info GrB_assign
                                    // C < mask > (I,j) = accum (C(I,j),u)
    GrB_Matrix C,
                                    // input/output matrix for results
                                    // optional mask for C(:,j), unused if NULL
    const GrB_Vector mask,
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(C(I,j),t)
    const GrB_Vector u,
                                    // input vector
    const GrB_Index *I,
                                    // row indices
                                    // number of row indices
    const GrB_Index ni,
    const GrB_Index j,
                                    // column index
    const GrB_Descriptor desc
                                    // descriptor for C(:,j) and mask
);
```

GrB_Col_assign modifies a single sub-column of a matrix C. It is the same as GrB_Matrix_assign where the index vector J[0]=j is a single column index, and where all matrices in GrB_Matrix_assign (except C) consist of a single column.

Unlike GrB_Matrix_assign, the mask is a vector with the same size as a single column of C.

The input descriptor GrB_INPO is ignored; the input vector **u** is not transposed. Refer to GrB_Matrix_assign for further details.

Performance considerations: GrB_Col_assign is much faster than GrB_Row_assign if the format of C is GxB_BY_COL. GrB_Row_assign is much faster than GrB_Col_assign if the format of C is GxB_BY_ROW.

8.9.4 GrB_Row_assign: assign to a sub-row of a matrix

```
GrB_Info GrB_assign
                                    // C < mask' > (i,J) = accum (C(i,J),u')
    GrB_Matrix C,
                                    // input/output matrix for results
   const GrB_Vector mask,
                                    // optional mask for C(i,:), unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(C(i,J),t)
   const GrB_Vector u,
                                    // input vector
    const GrB_Index i,
                                    // row index
                                    // column indices
    const GrB_Index *J,
    const GrB_Index nj,
                                    // number of column indices
    const GrB_Descriptor desc
                                    // descriptor for C(i,:) and mask
);
```

GxB_Row_subassign modifies a single sub-row of a matrix C. It is the same as GxB_Matrix_subassign where the index vector I[0]=i is a single row index, and where all matrices in GxB_Matrix_subassign (except C) consist of a single row.

Unlike GrB_Matrix_assign, the mask is a vector with the same size as a single row of C.

The input descriptor GrB_INPO is ignored; the input vector u is not transposed. Refer to GxB_Matrix_subassign for further details.

Performance considerations: GrB_Col_assign is much faster than GrB_Row_assign if the format of C is GxB_BY_COL. GrB_Row_assign is much faster than GrB_Col_assign if the format of C is GxB_BY_ROW.

8.9.5 GrB_Vector_assign_<type>: assign a scalar to a subvector

```
GrB_Info GrB_assign
                                    // w<mask>(I) = accum (w(I),x)
    GrB_Vector w,
                                    // input/output vector for results
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w(I),x)
    const <type> x,
                                    // scalar to assign to w(I)
    const GrB_Index *I,
                                    // row indices
                                    // number of row indices
    const GrB_Index ni,
    const GrB_Descriptor desc
                                    // descriptor for w and mask
);
```

GrB_Vector_assign_<type> assigns a single scalar to an entire subvector of the vector w. The operation is exactly like setting a single entry in an n-by-1 matrix, A(I,0) = x, where the column index for a vector is implicitly j=0. The mask vector has the same size as w. For further details of this function, see GrB_Matrix_assign_<type> in the next section.

Following the C API Specification, results are well-defined if I contains duplicate indices. Duplicate indices are simply ignored. See Section 8.10 for more details.

8.9.6 GrB_Matrix_assign_<type>: assign a scalar to a submatrix

```
// C<Mask>(I,J) = accum (C(I,J),x)
GrB_Info GrB_assign
    GrB_Matrix C,
                                    // input/output matrix for results
    const GrB_Matrix Mask,
                                    // optional mask for C, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C(I,J),x)
    const <type> x,
                                    // scalar to assign to C(I,J)
                                    // row indices
    const GrB_Index *I,
    const GrB_Index ni,
                                    // number of row indices
    const GrB_Index *J,
                                    // column indices
    const GrB_Index nj,
                                    // number of column indices
    const GrB_Descriptor desc
                                    // descriptor for C and Mask
);
```

GrB_Matrix_assign_<type> assigns a single scalar to an entire submatrix of C, like the scalar expansion C(I,J)=x in MATLAB. The scalar x is implicitly expanded into a matrix A of size ni by nj, and then the matrix A is assigned to C(I,J) using the same method as in GrB_Matrix_assign. Refer to that function in Section 8.9.2 for further details.

The Mask has the same size as C.

For the accumulation step, the scalar \mathbf{x} is typecasted directly into the type of C when the accum operator is not applied to it, or into the ytype of the accum operator, if accum is not NULL, for entries that are already present in C.

The <type> x notation is otherwise the same as GrB_Matrix_setElement (see Section 5.9.9). Any value can be passed to this function and its type will be detected, via the _Generic feature of ANSI C11. For a user-defined type, x is a void * pointer that points to a memory space holding a single entry of a scalar that has exactly the same user-defined type as the matrix C. This user-defined type must exactly match the user-defined type of C since no typecasting is done between user-defined types.

If a void * pointer is passed in and the type of the underlying scalar does not exactly match the user-defined type of C, then results are undefined. No error status will be returned since GraphBLAS has no way of catching this error.

Following the C API Specification, results are well-defined if I or J contain duplicate indices. Duplicate indices are simply ignored. See Section 8.10 for more details.

Performance considerations: If A is not transposed: if |I| is small, then it is fastest if the format of C is GxB_BY_ROW; if |J| is small, then it is fastest if the format of C is GxB_BY_COL. The opposite is true if A is transposed.

8.10 Duplicate indices in GrB_assign and GxB_subassign

According to the GraphBLAS C API Specification if the index vectors I or J contain duplicate indices, the results are undefined for GrB_Matrix_assign GrB_Matrix_assign and GrB_Row_assign. Only the scalar assignment operations (GrB_Matrix_assign_TYPE and GrB_Matrix_assign_TYPE) are well-defined when duplicates appear in I and J. In those two functions, duplicate indices are ignored.

As an extension to the specification, SuiteSparse:GraphBLAS provides a definition of how duplicate indices are handled in all cases. If I has duplicate indices, they are ignored and the last unique entry in the list is used. When no mask and no accumulator is present, the results are identical to how MAT-LAB handles duplicate indices in the built-in expression C(I,J)=A. Details of how this is done is shown below.

```
function C = subassign (C, I, J, A)
% submatrix assignment with pre-sort of I and J; and remove duplicates
% delete duplicates from I, keeping the last one seen
[I2 I2k] = sort(I);
Idupl = [(I2 (1:end-1) == I2 (2:end)), false];
I2 = I2 ( (Idupl) ;
I2k = I2k (~Idupl);
assert (isequal (I2, unique (I)))
% delete duplicates from J, keeping the last one seen
[J2 \ J2k] = sort(J);
Jdupl = [(J2 (1:end-1) == J2 (2:end)), false];
J2 = J2 (~Jdupl);
J2k = J2k (~Jdupl);
assert (isequal (J2, unique (J)))
% do the submatrix assignment, with no duplicates in I2 or J2
C (I2,J2) = A (I2k,J2k);
```

If a mask is present, then it is replaced with M = M (I2k, J2k) for $GxB_subassign$, or with M = M (I2, J2) for GrB_assign . If an accumulator operator is present, it is applied after the duplicates are removed, as (for example):

```
C(I2,J2) = C(I2,J2) + A(I2k,J2k);
```

These definitions allow the MATLAB interface to GraphBLAS to return the same results for C(I,J)=A for a Grb object as they do for built-in MATLAB matrices. They also allow the assignment to be done in parallel.

Results are always well-defined in SuiteSparse:GraphBLAS, but they might not be what you expect. For example, suppose the MIN operator is being used the following assignment to the vector \mathbf{x} , and suppose I contains the entries [0 0]. Suppose \mathbf{x} is initially empty, of length 1, and suppose \mathbf{y} is a vector of length 2 with the values [5 7].

```
#include "GraphBLAS.h"
#include <stdio.h>
int main (void)
{
    GrB_init (GrB_NONBLOCKING) ;
    GrB_Vector x, y ;
    GrB_Vector_new (&x, GrB_INT32, 1);
    GrB_Vector_new (&y, GrB_INT32, 2);
    GrB_Index I [2] = \{0, 0\};
    GrB_Vector_setElement (y, 5, 0);
    GrB_Vector_setElement (y, 7, 1);
    GrB_Vector_wait (&y) ;
    GxB_print (x, 3);
    GxB_print (y, 3);
    GrB_assign (x, NULL, GrB_MIN_INT32, y, I, 2, NULL) ;
    GrB_Vector_wait (&y) ;
    GxB_print (x, 3);
    GrB_finalize ( ) ;
}
```

You might (wrongly) expect the result to be the vector $\mathbf{x}(0)=5$, since two entries seem to be assigned, and the min operator might be expected to take the minimum of the two. This is not how SuiteSparse:GraphBLAS handles duplicates.

Instead, the first duplicate index of I is discarded (I [0] = 0, and y(0)=5). and only the second entry is used (I [1] = 0, and y(1)=7). The output of the above program is:

```
1x1 GraphBLAS int32_t vector, sparse by col:
x, no entries
```

```
2x1 GraphBLAS int32_t vector, sparse by col:
y, 2 entries

(0,0) 5
(1,0) 7

1x1 GraphBLAS int32_t vector, sparse by col:
x, 1 entry

(0,0) 7
```

You see that the result is x(0)=7, since the y(0)=5 entry has been ignored because of the duplicate indices in I.

SPEC: Providing a well-defined behavior for duplicate indices with matrix and vector assignment is an extension to the spec. The spec only defines the behavior when assigning a scalar into a matrix or vector, and states that duplicate indices otherwise lead to undefined results.

8.11 Comparing GrB_assign and GxB_subassign

The GxB_subassign and GrB_assign operations are very similar, but they differ in two ways:

- 1. The Mask has a different size: The mask in GxB_subassign has the same dimensions as w(I) for vectors and C(I,J) for matrices. In GrB_assign, the mask is the same size as w or C, respectively (except for the row/col variants). The two masks are related. If M is the mask for GrB_assign, then M(I,J) is the mask for GxB_subassign. If there is no mask, or if I and J are both GrB_ALL, the two masks are the same. For GrB_Row_assign and GrB_Col_assign, the mask vector is the same size as a row or column of C, respectively. For the corresponding GxB_Row_subassign and GxB_Col_subassign operations, the mask is the same size as the sub-row C(i,J) or subcolumn C(I,j), respectively.
- 2. GrB_REPLACE is different: They differ in how C is affected in areas outside the C(I,J) submatrix. In GxB_subassign, the C(I,J) submatrix is the only part of C that can be modified, and no part of C outside the submatrix is ever modified. In GrB_assign, it is possible to delete entries in C outside the submatrix, but only in one specific manner. Suppose the mask M is present (or, suppose it is not present but GrB_COMP is true). After (optionally) complementing the mask, the value of M(i,j) can be 0 for some entry outside the C(I,J) submatrix. If the GrB_REPLACE descriptor is true, GrB_assign deletes this entry.

GxB_subassign and GrB_assign are identical if GrB_REPLACE is set to its default value of false, and if the masks happen to be the same. The two masks can be the same in two cases: either the Mask input is NULL (and it is not complemented via GrB_COMP), or I and J are both GrB_ALL. If all these conditions hold, the two algorithms are identical and have the same performance. Otherwise, GxB_subassign is much faster than GrB_assign when the latter must examine the entire matrix C to delete entries (when GrB_REPLACE is true), and if it must deal with a much larger Mask matrix. However, both methods have specific uses.

Consider using C(I,J)+=F for many submatrices F (for example, when assembling a finite-element matrix). If the Mask is meant as a specification for which entries of C should appear in the final result, then use GrB_assign.

If instead the Mask is meant to control which entries of the submatrix C(I,J) are modified by the finite-element F, then use GxB_subassign. This is particularly useful is the Mask is a template that follows along with the finite-element F, independent of where it is applied to C. Using GrB_assign would be very difficult in this case since a new Mask, the same size as C, would need to be constructed for each finite-element F.

In GraphBLAS notation, the two methods can be described as follows:

matrix and vector subassign	$\mathbf{C}(\mathbf{I},\mathbf{J})\langle\mathbf{M} angle=\mathbf{C}(\mathbf{I},\mathbf{J})\odot\mathbf{A}$
matrix and vector assign	$\mathbf{C}\langle\mathbf{M} angle(\mathbf{I},\mathbf{J})=\mathbf{C}(\mathbf{I},\mathbf{J})\odot\mathbf{A}$

This notation does not include the details of the GrB_COMP and GrB_REPLACE descriptors, but it does illustrate the difference in the Mask. In the subassign, Mask is the same size as C(I,J) and A. If I[0]=i and J[0]=j, Then Mask(0,0) controls how C(i,j) is modified by the subassign, from the value A(0,0). In the assign, Mask is the same size as C, and Mask(i,j) controls how C(i,j) is modified.

The GxB_subassign and GrB_assign functions have the same signatures; they differ only in how they consider the Mask and the GrB_REPLACE descriptor

Details of each step of the two operations are listed below:

Step	<pre>GrB_Matrix_assign</pre>	GxB_Matrix_subassign
1	$\mathbf{S} = \mathbf{C}(\mathbf{I}, \mathbf{J})$	$\mathbf{S} = \mathbf{C}(\mathbf{I}, \mathbf{J})$
2	$\mathbf{S} = \mathbf{S} \odot \mathbf{A}$	$\mathbf{S}\langle \mathbf{M} \rangle = \mathbf{S} \odot \mathbf{A}$
3	$\mathbf{Z} = \mathbf{C}$	$\mathbf{C}(\mathbf{I},\mathbf{J})=\mathbf{S}$
4	$\mathbf{Z}(\mathbf{I},\mathbf{J}) = \mathbf{S}$	
5	$\mathbf{C}\langle \mathbf{M} angle = \mathbf{Z}$	

Step 1 is the same. In the Accumulator Phase (Step 2), the expression $\mathbf{S} \odot \mathbf{A}$, described in Section 2.3, is the same in both operations. The result is simply \mathbf{A} if accum is NULL. It only applies to the submatrix \mathbf{S} , not the whole matrix. The result $\mathbf{S} \odot \mathbf{A}$ is used differently in the Mask/Replace phase.

The Mask/Replace Phase, described in Section 2.3 is different:

• For Grb_assign (Step 5), the mask is applied to all of C. The mask has the same size as C. Just prior to making the assignment via the mask, the Grb_REPLACE option can be used to clear all of C first. This is the only way in which entries in C that are outside the C(I, J) submatrix can be modified by this operation.

• For GxB_subassign (Step 2b), the mask is applied to just S. The mask has the same size as C(I, J), S, and A. Just prior to making the assignment via the mask, the GrB_REPLACE option can be used to clear S first. No entries in C that are outside the C(I, J) can be modified by this operation. Thus, GrB_REPLACE has no effect on entries in C outside the C(I, J) submatrix.

The differences between GrB_assign and $GxB_subassign$ can be seen in Tables 1 and 2. The first table considers the case when the entry c_{ij} is in the C(I, J) submatrix, and it describes what is computed for both GrB_assign and $GxB_subassign$. They perform the exact same computation; the only difference is how the value of the mask is specified.

The first column of the table is yes if $GrB_REPLACE$ is enabled, and a dash otherwise. The second column is yes if an accumulator operator is given, and a dash otherwise. The third column is c_{ij} if the entry is present in \mathbf{C} , and a dash otherwise. The fourth column is $a_{i'j'}$ if the corresponding entry is present in \mathbf{A} , where $i = \mathbf{I}(i')$ and $j = \mathbf{J}(i')$.

The mask column is 1 if the mask allows C to be modified, and 0 otherwise. This is m_{ij} for GrB_assign , and $m_{i'j'}$ for $GxB_subassign$, to reflect the difference in the mask, but this difference is not reflected in the table. The value 1 or 0 is the value of the entry in the mask after it is optionally complemented via the GrB_COMP option.

Finally, the last column is the action taken in this case. It is left blank if no action is taken, in which case c_{ij} is not modified if present, or not inserted into \mathbf{C} if not present.

repl	accum	\mathbf{C}	A	mask	action taken by GrB_assign and GxB_subassign
-	_	c_{ij}	$a_{i'j'}$	1	$c_{ij} = a_{i'j'}$, update
-	-	-	$a_{i'j'}$	1	$c_{ij} = a_{i'j'}$, insert
-	-	c_{ij}	-	1	delete c_{ij} because $a_{i'j'}$ not present
-	-	-	-	1	
-	-	c_{ij}	$a_{i'j'}$	0	
-	-	-	$a_{i'j'}$	0	
-	-	c_{ij}	-	0	
-	-	-	-	0	
yes	-	c_{ij}	$a_{i'j'}$	1	$c_{ij} = a_{i'j'}$, update
yes	-	-	$a_{i'j'}$	1	$c_{ij} = a_{i'j'}$, insert
yes	-	c_{ij}	-	1	delete c_{ij} because $a_{i'j'}$ not present
yes	-	-	-	1	
yes	-	c_{ij}	$a_{i'j'}$	0	delete c_{ij} (because of GrB_REPLACE)
yes	-	-	$a_{i'j'}$	0	
yes	-	c_{ij}	-	0	delete c_{ij} (because of GrB_REPLACE)
yes	-	-	-	0	
-	yes	c_{ij}	$a_{i'j'}$	1	$c_{ij} = c_{ij} \odot a_{i'j'}$, apply accumulator
-	yes	-	$a_{i'j'}$	1	$c_{ij} = a_{i'j'}$, insert
-	yes	c_{ij}	-	1	
_	yes	-	-	1	
-	yes	c_{ij}	$a_{i'j'}$	0	
-	yes	-	$a_{i'j'}$	0	
-	yes	c_{ij}	-	0	
-	yes	-	-	0	
yes	yes	c_{ij}	$a_{i'j'}$	1	$c_{ij} = c_{ij} \odot a_{i'j'}$, apply accumulator
yes	yes	-	$a_{i'j'}$	1	$c_{ij} = a_{i'j'}$, insert
yes	yes	c_{ij}	-	1	
yes	yes	-	-	1	
yes	yes	c_{ij}	$a_{i'j'}$	0	delete c_{ij} (because of $GrB_REPLACE$)
yes	yes	-	$a_{i'j'}$	0	
yes	yes	c_{ij}	-	0	delete c_{ij} (because of $GrB_REPLACE$)
yes	yes	_	_	0	

Table 1: Results of assign and subassign for entries in the $\mathbf{C}(\mathbf{I},\mathbf{J})$ submatrix

repl	accum	\mathbf{C}	$\mathbf{C} = \mathbf{Z}$	mask	action taken by GrB_assign
-	-	c_{ij}	c_{ij}	1	
-	-	-	-	1	
-	-	c_{ij}	c_{ij}	0	
-	-	-	-	0	
yes	-	c_{ij}	c_{ij}	1	
yes	-	-	-	1	
yes	-	c_{ij}	c_{ij}	0	delete c_{ij} (because of $GrB_REPLACE$)
yes	-	-	-	0	
-	yes	c_{ij}	c_{ij}	1	
-	yes	-	-	1	
-	yes	c_{ij}	c_{ij}	0	
-	yes	-	-	0	
yes	yes	c_{ij}	c_{ij}	1	
yes	yes	-	-	1	
yes	yes	c_{ij}	c_{ij}	0	delete c_{ij} (because of $GrB_REPLACE$)
yes	yes	-	-	0	

Table 2: Results of assign for entries outside the C(I, J) submatrix. Subassign has no effect on these entries.

Table 2 illustrates how GrB_assign and $GxB_subassign$ differ for entries outside the submatrix. $GxB_subassign$ never modifies any entry outside the C(I, J) submatrix, but GrB_assign can modify them in two cases listed in Table 2. When the $GrB_REPLACE$ option is selected, and when the Mask(i,j) for an entry c_{ij} is false (or if the Mask(i,j) is true and GrB_COMP is enabled via the descriptor), then the entry is deleted by GrB_assign .

The fourth column of Table 2 differs from Table 1, since entries in **A** never affect these entries. Instead, for all index pairs outside the $I \times J$ submatrix, **C** and **Z** are identical (see Step 3 above). As a result, each section of the table includes just two cases: either c_{ij} is present, or not. This in contrast to Table 1, where each section must consider four different cases.

The GrB_Row_assign and GrB_Col_assign operations are slightly different. They only affect a single row or column of C. For GrB_Row_assign, Table 2 only applies to entries in the single row C(i, J) that are outside the list of indices, J. For GrB_Col_assign, Table 2 only applies to entries in the single column C(I,j) that are outside the list of indices, I.

8.11.1 Example

The difference between GxB_subassign and GrB_assign is illustrated in the following example. Consider the 2-by-2 matrix C where all entries are present.

$$\mathbf{C} = \left[\begin{array}{cc} 11 & 12 \\ 21 & 22 \end{array} \right]$$

Suppose GrB_REPLACE is true, and GrB_COMP is false. Let the Mask be:

$$\mathbf{M} = \left[\begin{array}{cc} 1 & 1 \\ 0 & 1 \end{array} \right].$$

Let $\mathbf{A} = 100$, and let the index sets be $\mathbf{I} = 0$ and $\mathbf{J} = 1$. Consider the computation $\mathbf{C}\langle\mathbf{M}\rangle(0,1) = \mathbf{C}(0,1) + \mathbf{A}$, using the GrB_assign operation. The result is:

$$\mathbf{C} = \left[\begin{array}{cc} 11 & 112 \\ - & 22 \end{array} \right].$$

The (0,1) entry is updated and the (1,0) entry is deleted because its Mask is zero. The other two entries are not modified since $\mathbf{Z} = \mathbf{C}$ outside the submatrix, and those two values are written back into \mathbf{C} because their Mask values are 1. The (1,0) entry is deleted because the entry $\mathbf{Z}(1,0) = 21$ is prevented from being written back into \mathbf{C} since Mask(1,0)=0.

Now consider the analogous $GxB_subassign$ operation. The Mask has the same size as A, namely:

$$\mathbf{M} = [1].$$

After computing $\mathbf{C}(0,1)\langle \mathbf{M} \rangle = \mathbf{C}(0,1) + \mathbf{A}$, the result is

$$\mathbf{C} = \left[\begin{array}{cc} 11 & 112 \\ 21 & 22 \end{array} \right].$$

Only the $\mathbf{C}(\mathbf{I}, \mathbf{J})$ submatrix, the single entry $\mathbf{C}(0, 1)$, is modified by $\mathtt{GxB_subassign}$. The entry $\mathbf{C}(1, 0) = 21$ is unaffected by $\mathtt{GxB_subassign}$, but it is deleted by $\mathtt{GrB_assign}$.

8.11.2 Performance of GxB_subassign, GrB_assign and GrB_*_setElement

When SuiteSparse:GraphBLAS uses non-blocking mode, the modifications to a matrix by GxB_subassign, GrB_assign, and GrB_*_setElement can postponed, and computed all at once later on. This has a huge impact on performance.

A sequence of assignments is fast if their completion can be postponed for as long as possible, or if they do not modify the pattern at all. Modifying the pattern can be costly, but it is fast if non-blocking mode can be fully exploited.

Consider a sequence of t submatrix assignments C(I,J)=C(I,J)+A to an n-by-n matrix C where each submatrix A has size a-by-a with s entries, and where C starts with s entries. Assume the matrices are all stored in non-hypersparse form, by row (GxB_BY_ROW) .

If blocking mode is enabled, or if the sequence requires the matrix to be completed after each assignment, each of the t assignments takes $O(a + s \log n)$ time to process the A matrix and then $O(n + c + s \log s)$ time to complete C. The latter step uses GrB_*build to build an update matrix and then merge it with C. This step does not occur if the sequence of assignments does not add new entries to the pattern of C, however. Assuming in the worst case that the pattern does change, the total time is $O(t [a + s \log n + n + c + s \log s])$.

If the sequence can be computed with all updates postponed until the end of the sequence, then the total time is no worse than $O(a+s\log n)$ to process each A matrix, for t assignments, and then a single build at the end, taking $O(n+c+st\log st)$ time. The total time is $O(t\left[a+s\log n\right]+(n+c+st\log st))$. If no new entries appear in C the time drops to $O(t\left[a+s\log n\right])$, and in this case, the time for both methods is the same; both are equally efficient.

A few simplifying assumptions are useful to compare these times. Consider a graph of n nodes with O(n) edges, and with a constant bound on the degree of each node. The asymptotic bounds assume a worst-case scenario where C has a least some dense rows (thus the $\log n$ terms). If these are not present, if both t and c are O(n), and if a and s are constants, then the total time with blocking mode becomes $O(n^2)$, assuming the pattern of C changes at each assignment. This very high for a sparse graph problem. In contrast, the non-blocking time becomes $O(n \log n)$ under these same assumptions, which is asymptotically much faster.

The difference in practice can be very dramatic, since n can be many millions for sparse graphs with n nodes and O(n), which can be handled on a commodity laptop.

The following guidelines should be considered when using GxB_subassign, GrB_assign and GrB_*_setElement.

- 1. A sequence of assignments that does not modify the pattern at all is fast, taking as little as $\Omega(1)$ time per entry modified. The worst case time complexity is $O(\log n)$ per entry, assuming they all modify a dense row of C with n entries, which can occur in practice. It is more common, however, that most rows of C have a constant number of entries, independent of n. No work is ever left pending when the pattern of C does not change.
- 2. A sequence of assignments that modifies the entries that already exist in the pattern of a matrix, or adds new entries to the pattern (using the same accum operator), but does not delete any entries, is fast. The matrix is not completed until the end of the sequence.
- 3. Similarly, a sequence that modifies existing entries, or deletes them, but does not add new ones, is also fast. This sequence can also repeatedly delete pre-existing entries and then reinstate them and still be fast. The matrix is not completed until the end of the sequence.
- 4. A sequence that mixes assignments of types (2) and (3) above can be costly, since the matrix may need to be completed after each assignment. The time complexity can become quadratic in the worst case.
- 5. However, any single assignment takes no more than $O(a+s\log n+n+c+s\log s)$ time, even including the time for a matrix completion, where C is n-by-n with c entries and A is a-by-a with s entries. This time is essentially linear in the size of the matrix C, if A is relatively small and sparse compared with C. In this case, n+c are the two dominant terms.
- 6. In general, GxB_subassign is faster than GrB_assign. If GrB_REPLACE is used with GrB_assign, the entire matrix C must be traversed. This is much slower than GxB_subassign, which only needs to examine the C(I,J) submatrix. Furthermore, GrB_assign must deal with a much larger Mask matrix, whereas GxB_subassign has a smaller mask. Since

its mask is smaller, GxB_subassign takes less time than GrB_assign to access the mask.

Submatrix assignment in SuiteSparse:GraphBLAS is extremely efficient, even without considering the advantages of non-blocking mode discussed in Section 8.11. It can be up to 500x faster than MATLAB R2019b, or even higher depending on the kind of matrix assignment. MATLAB logical indexing (the mask of GraphBLAS) is much faster with GraphBLAS than in MATLAB R2019b.

All of the 28 variants (each with their own source code) are either asymptotically optimal, or to within a log factor of being asymptotically optimal. The methods are also fully parallel. For hypersparse matrices, the term n in the expressions in the above discussion is dropped, and is replaced with $h \log h$, at the worst case, where h << n is the number of non-empty columns of a hypersparse matrix stored by column, or the number of non-empty rows of a hypersparse matrix stored by row. In many methods, n is replaced with h, not $h \log h$.

8.12 GrB_apply: apply a unary operator

The GrB_apply function is the generic name for two specific functions: GrB_Vector_apply and GrB_Matrix_apply. The generic name appears in the function prototypes, but the specific function name is used when describing each variation. When discussing features that apply to both versions, the simple name GrB_apply is used.

8.12.1 GrB_Vector_apply: apply a unary operator to a vector

GrB_Vector_apply applies a unary operator to the entries of a vector, analogous to $\mathbf{t} = op(\mathbf{u})$ in MATLAB except the operator op is only applied to entries in the pattern of \mathbf{u} . Implicit values outside the pattern of \mathbf{u} are not affected. The entries in \mathbf{u} are typecasted into the xtype of the unary operator. The vector \mathbf{t} has the same type as the ztype of the unary operator. The final step is $\mathbf{w}\langle \mathbf{m} \rangle = \mathbf{w} \odot \mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

8.12.2 GrB_Matrix_apply: apply a unary operator to a matrix

GrB_Matrix_apply applies a unary operator to the entries of a matrix, analogous to T = op(A) in MATLAB except the operator op is only applied to entries in the pattern of A. Implicit values outside the pattern of A are not affected. The input matrix A may be transposed first. The entries in A are typecasted into the xtype of the unary operator. The matrix T has the same type as the ztype of the unary operator. The final step is $C\langle M \rangle = C \odot T$, as described in Section 2.3.

The built-in $GrB_IDENTITY_T$ operators (one for each built-in type T) are very useful when combined with this function, enabling it to compute $C\langle M \rangle = C \odot A$. This makes GrB_apply a direct interface to the accumulator/mask function for both matrices and vectors.

To compute $\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{A}$ or $\mathbf{C}\langle \mathbf{M} \rangle = \mathbf{C} \odot \mathbf{A}$ for user-defined types, the user application would need to define an identity operator for the type. Since GraphBLAS cannot detect that it is an identity operator, it must call the operator to make the full copy T=A and apply the operator to each entry of the matrix or vector.

The other GraphBLAS operation that provides a direct interface to the accumulator/mask function is GrB_transpose, which does not require an operator to perform this task. As a result, GrB_transpose can be used as an efficient and direct interface to the accumulator/mask function for both built-in and user-defined types. However, it is only available for matrices, not vectors.

8.13 GxB_select: apply a select operator

The GxB_select function is the generic name for two specific functions: GxB_Vector_select and GxB_Matrix_select. The generic name appears in the function prototypes, but the specific function name is used when describing each variation. When discussing features that apply to both versions, the simple name GxB_select is used.

SPEC: The GxB_select operation and GxB_SelectOp operator are extensions to the spec.

8.13.1 GxB_Vector_select: apply a select operator to a vector

```
GrB_Info GxB_select
                                    // w<mask> = accum (w, op(u,k))
                                    // input/output vector for results
    GrB_Vector w,
    const GrB_Vector mask,
                                    // optional mask for w, unused if NULL
    const GrB_BinaryOp accum,
                                    // optional accum for z=accum(w,t)
                                    // operator to apply to the entries
    const GxB_SelectOp op,
    const GrB_Vector u,
                                    // first input: vector u
    const GxB_Scalar Thunk,
                                    // optional input for the select operator
    const GrB_Descriptor desc
                                    // descriptor for w and mask
);
```

GxB_Vector_select applies a select operator to the entries of a vector, analogous to t = u.*op(u) in MATLAB except the operator op is only applied to entries in the pattern of u. Implicit values outside the pattern of u are not affected. If the operator is not type-generic, the entries in u are type-casted into the xtype of the select operator. The vector t has the same type and size as u. The final step is $\mathbf{w}\langle\mathbf{m}\rangle = \mathbf{w}\odot \mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

This operation operates on vectors just as if they were m-by-1 matrices, except that GraphBLAS never transposes a vector via the descriptor. The op is passed n=1 as the number of columns. Refer to the next section on GxB_Matrix_select for more details.

8.13.2 GxB_Matrix_select: apply a select operator to a matrix

```
// C<Mask> = accum (C, op(A,k)) or op(A',k)
GrB_Info GxB_select
    GrB_Matrix C,
                                    // input/output matrix for results
                                    // optional mask for C, unused if NULL
    const GrB_Matrix Mask,
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C,T)
                                    // operator to apply to the entries
    const GxB_SelectOp op,
    const GrB_Matrix A,
                                    // first input: matrix A
    const GxB_Scalar Thunk,
                                    // optional input for the select operator
    const GrB_Descriptor desc
                                    // descriptor for C, mask, and A
);
```

GxB_Matrix_select applies a select operator to the entries of a matrix, analogous to T = A .* op(A) in MATLAB except the operator op is only applied to entries in the pattern of A. Implicit values outside the pattern of A are not affected. The input matrix A may be transposed first. If the operator is not type-generic, the entries in A are typecasted into the xtype of the select operator. The final step is $C\langle M \rangle = C \odot T$, as described in Section 2.3.

The matrix T has the same size and type as A (or the transpose of A if the input is transposed via the descriptor). The entries of T are a subset of those of A. Each entry A(i,j) of A is passed to the op, as $z = f(i,j,m,n,a_{ij},\text{thunk})$, where A is m-by-n. If A is transposed first then the operator is applied to entries in the transposed matrix, A'. If z is returned as true, then the entry is copied into T, unchanged. If it returns false, the entry does not appear in T.

If Thunk is not NULL, it must be a valid GxB_Scalar. If it has no entry, it is treated as if it had a single entry equal to zero, for built-in types (not user-defined types).

For user-defined select operators, the entry is passed to the user-defined select operator, with no typecasting. Its type must be identical to ttype of the select operator.

For the GxB_TRIL , GxB_TRIU , GxB_DIAG , and GxB_DIAG , the Thunk parameter may be NULL, or it may be present but contain no entry. In this case, these operators use the value of k=0, the main diagonal. If present, the Thunk can be any built-in type. The value of this entry is typecasted: $k = (int64_t)$ Thunk. The value k=0 specifies the main diagonal of the matrix, k=1 is the +1 diagonal (the entries just above the main diagonal), k=-1 is the -1 diagonal, and so on.

For the GxB_*ZERO select operators, Thunk is ignored, and may be NULL.

For built-in types, with the GxB_*THUNK operators, the value of Thunk is typecasted to the same type as the A matrix. For user-defined types, Thunk is passed to the select operator without typecasting.

The action of GxB_select with the built-in select operators is described in the table below. The MATLAB analogs are precise for tril and triu, but shorthand for the other operations. The MATLAB diag function returns a column with the diagonal, if A is a matrix, whereas the matrix T in GxB_select always has the same size as A (or its transpose if the GrB_INPO is set to GrB_TRAN). In the MATLAB analog column, diag is as if it operates like GxB_select, where T is a matrix.

The following operators may be used on matrices with a user-defined type: GxB_TRIL, GxB_TRIU, GxB_DIAG, GxB_OFFIAG, GxB_NONZERO, GxB_EQ_ZERO, GxB_NE_THUNK, and GxB_EQ_THUNK.

The comparators <code>GxB_GT_*GxB_GE_*GxB_LT_*</code>, and <code>GxB_LE_*</code> only work for built-in types. All other built-in select operators can be used for any type, both built-in and any user-defined type.

NOTE: For floating-point values, comparisons with NaN always return false. The built-in select operators should not be used with a scalar thunk that is equal to NaN. For this case, create a user-defined select operator that performs the test with the ANSI C isnan function instead.

GraphBLAS	MATLAB	
name	analog	
GxB_TRIL	T=tril(A,k)	Entries in T are the entries on and below the kth diagonal of A.
GxB_TRIU	T=triu(A,k)	Entries in T are the entries on and above the
		kth diagonal of A.
GxB_DIAG	T=diag(A,k)	Entries in T are the entries on the kth diagonal
		of A.
GxB_OFFDIAG	T=A-diag(A,k)	Entries in T are all entries not on the kth di-
		agonal of A.
GxB_NONZERO	T=A(A~=0)	Entries in T are all entries in A that have
		nonzero value.
GxB_EQ_ZERO	T=A(A==0)	Entries in T are all entries in A that are equal
		to zero.
GxB_GT_ZERO	T=A(A>0)	Entries in T are all entries in A that are greater
		than zero.
GxB_GE_ZERO	$T=A(A \le 0)$	Entries in T are all entries in A that are greater
		than or equal to zero.
GxB_LT_ZERO	T=A(A<0)	Entries in T are all entries in A that are less
		than zero.
GxB_LE_ZERO	$T=A(A \le 0)$	Entries in T are all entries in A that are less
		than or equal to zero.
GxB_NE_THUNK	$T=A(A^{\sim}=k)$	Entries in T are all entries in A that are not
		equal to k.
GxB_EQ_THUNK	T=A(A==k)	Entries in T are all entries in A that are equal
		to k.
GxB_GT_THUNK	T=A(A>k)	Entries in T are all entries in A that are greater
		than k.
GxB_GE_THUNK	T=A(A>=k)	Entries in T are all entries in A that are greater
		than or equal to k.
GxB_LT_THUNK	T=A(A <k)< td=""><td>Entries in T are all entries in A that are less</td></k)<>	Entries in T are all entries in A that are less
		than k.
GxB_LE_THUNK	$T=A(A\leq k)$	Entries in T are all entries in A that are less
		than or equal to k.

8.14 GrB reduce: reduce to a vector or scalar

The generic function name <code>GrB_reduce</code> may be used for all specific functions discussed in this section. When the details of a specific function are discussed, the specific name is used for clarity.

8.14.1 GrB_Matrix_reduce_<op>: reduce a matrix to a vector

GrB_Matrix_reduce_<op> is a generic name for two specific methods. Both methods reduce a matrix to a column vector using an operator, roughly analogous to t = sum (A') in MATLAB, in the default case, where t is a column vector. By default, the method reduces across the rows to obtain a column vector; use GrB_TRAN to reduce down the columns.

GrB_Matrix_reduce_BinaryOp relies on a binary operator for the reduction: the fourth argument reduce, a GrB_BinaryOp. All three domains of the operator must be the same. GrB_Matrix_reduce_Monoid performs the same reduction using a GrB_Monoid as its fourth argument. In both cases the reduction operator must be commutative and associative. Otherwise the results are undefined.

The input matrix A may be transposed first. Its entries are then typecast into the type of the reduce operator or monoid. The reduction is applied to all entries in A (i,:) to produce the scalar t (i). This is done without the use of the identity value of the monoid. If the ith row A (i,:) has no entries, then (i) is not an entry in t and its value is implicit. If A (i,:) has a single entry, then that is the result t (i) and reduce is not applied at all for the ith row. Otherwise, multiple entries in row A (i,:) are reduced via the reduce operator or monoid to obtain a single scalar, the result t (i).

The final step is $\mathbf{w}\langle \mathbf{m} \rangle = \mathbf{w} \odot \mathbf{t}$, as described in Section 2.3, except that all the terms are column vectors instead of matrices.

8.14.2 GrB_Vector_reduce_<type>: reduce a vector to a scalar

GrB_Vector_reduce_<type> reduces a vector to a scalar, analogous to t = sum (u) in MATLAB, except that in GraphBLAS any commutative and associative monoid can be used in the reduction.

The reduction operator is a commutative and associative monoid with an identity value. Results are undefined if the monoid does not have these properties. This function differs from <code>GrB_Matrix_reduce_BinaryOp</code> (which reduces a matrix to a vector) in that it requires a valid monoid additive identity value. If the vector u has no entries, that identity value is copied into the scalar t. Otherwise, all of the entries in the vector are reduced to a single scalar using the <code>reduce</code> operator.

The scalar type is any of the built-in types, or a user-defined type. In the function signature it is a C type: bool, int8_t, ... float, double, or void * for a user-defined type. The user-defined type must be identical to the type of the vector u. This cannot be checked by GraphBLAS and thus results are undefined if the types are not the same.

The descriptor is unused, but it appears in case it is needed in future versions of the GraphBLAS API. This function has no mask so its accumulator/mask step differs from the other GraphBLAS operations. It does not use the methods described in Section 2.3, but uses the following method instead.

If accum is NULL, then the scalar t is typecast into the type of c, and c = t is the final result. Otherwise, the scalar t is typecast into the ytype of the accum operator, and the value of c (on input) is typecast into the xtype of the accum operator. Next, the scalar z = accum (c,t) is computed, of the ztype of the accum operator. Finally, z is typecast into the final result, c.

8.14.3 GrB_Matrix_reduce_<type>: reduce a matrix to a scalar

GrB_Matrix_reduce_<type> reduces a matrix A to a scalar, roughly analogous to t = sum (A (:)) in MATLAB. This function is identical to reducing a vector to a scalar, since the positions of the entries in a matrix or vector have no effect on the result. Refer to the reduction to scalar described in the previous Section 8.14.2.

8.15 GrB_transpose: transpose a matrix

GrB_transpose transposes a matrix A, just like the array transpose T = A.' in MATLAB. The internal result matrix T = A' (or merely T = A if A is transposed via the descriptor) has the same type as A. The final step is $\mathbf{C}\langle\mathbf{M}\rangle = \mathbf{C}\odot\mathbf{T}$, as described in Section 2.3, which typecasts T as needed and applies the mask and accumulator.

To be consistent with the rest of the GraphBLAS API regarding the descriptor, the input matrix A may be transposed first. It may seem counterintuitive, but this has the effect of not doing any transpose at all. As a result, $GrB_transpose$ is useful for more than just transposing a matrix. It can be used as a direct interface to the accumulator/mask operation, $C\langle M \rangle = C \odot A$. This step also does any typecasting needed, so $GrB_transpose$ can be used to typecast a matrix A into another matrix C. To do this, simply use NULL for the Mask and accum, and provide a non-default descriptor desc that sets the transpose option:

```
// C = typecasted copy of A
GrB_Descriptor_set (desc, GrB_INPO, GrB_TRAN) ;
GrB_transpose (C, NULL, NULL, A, desc) ;
```

If the types of C and A match, then the above two lines of code are the same as GrB_Matrix_dup (&C, A), except that for GrB_transpose the matrix C must already exist and be the right size. If C does not exist, the work of GrB_Matrix_dup can be replicated with this:

```
// C = create an exact copy of A, just like GrB_Matrix_dup
GrB_Matrix C;
GrB_Type type;
GrB_Index nrows, ncols;
GrB_Descriptor desc;
GxB_Matrix_type (&type, A);
GrB_Matrix_nrows (&nrows, A);
```

```
GrB_Matrix_ncols (&ncols, A) ;
GrB_Matrix_new (&C, type, nrows, ncols) ;
GrB_Descriptor_new (&desc) ;
GrB_Descriptor_set (desc, GrB_INPO, GrB_TRAN) ;
GrB_transpose (C, NULL, NULL, A, desc) ;
```

Since the input matrix A is transposed by the descriptor, SuiteSparse:Graph-BLAS does the right thing and does not transpose the matrix at all. Since T = A is not typecasted, SuiteSparse:GraphBLAS can construct T internally in O(1) time and using no memory at all. This makes $Grb_transpose$ a fast and direct interface to the accumulator/mask function in GraphBLAS.

This example is of course overkill, since the work can all be done by a single call to the <code>GrB_Matrix_dup</code> function. However, the <code>GrB_Matrix_dup</code> function can only create <code>C</code> as an exact copy of <code>A</code>, whereas variants of the code above can do many more things with these two matrices. For example, the <code>type</code> in the example can be replaced with any other type, perhaps selected from another matrix or from an operator.

Consider the following code excerpt, which uses $GrB_transpose$ to remove all diagonal entries from a square matrix. It first creates a diagonal Mask, which is complemented so that $\mathbf{C}\langle \neg \mathbf{M}\rangle = \mathbf{A}$ does not modify the diagonal of \mathbf{C} . The REPLACE ensures that \mathbf{C} is cleared first, and then $\mathbf{C}\langle \neg \mathbf{M}\rangle = \mathbf{A}$ modifies all entries in \mathbf{C} where the mask \mathbf{M} is false. These correspond to all the off-diagonal entries. The descriptor ensures that \mathbf{A} is not transposed at all. The Mask can have any pattern, of course, and wherever it is set true, the corresponding entries in \mathbf{A} are deleted from the copy \mathbf{C} .

```
// remove all diagonal entries from the matrix A
// Mask = speye (n);
GrB_Matrix_new (&Mask, GrB_BOOL, n, n);
for (int64_t i = 0 ; i < n ; i++)
{
        GrB_Matrix_setElement (Mask, (bool) true, i, i);
}
// C<~Mask> = A, clearing C first. No transpose.
GrB_Descriptor_new (&desc);
GrB_Descriptor_set (desc, GrB_INPO, GrB_TRAN);
GrB_Descriptor_set (desc, GrB_MASK, GrB_COMP);
GrB_Descriptor_set (desc, GrB_OUTP, GrB_REPLACE);
GrB_transpose (A, Mask, NULL, A, desc);
```

8.16 GrB_kronecker: Kronecker product

```
GrB_Info GrB_kronecker
                                    // C<Mask> = accum (C, kron(A,B))
    GrB_Matrix C,
                                    // input/output matrix for results
                                    // optional mask for C, unused if NULL
    const GrB_Matrix Mask,
    const GrB_BinaryOp accum,
                                    // optional accum for Z=accum(C,T)
   const GrB_BinaryOp op,
                                    // defines '*' for T=kron(A,B)
    const GrB_Matrix A,
                                    // first input: matrix A
    const GrB_Matrix B,
                                    // second input: matrix B
    const GrB_Descriptor desc
                                    // descriptor for C, Mask, A, and B
);
```

GrB_kronecker computes the Kronecker product, $\mathbf{C}\langle\mathbf{M}\rangle=\mathbf{C}\odot\mathrm{kron}(\mathbf{A},\mathbf{B})$ where

$$\operatorname{kron}(\mathbf{A}, \mathbf{B}) = \begin{bmatrix} a_{00} \otimes \mathbf{B} & \dots & a_{0,n-1} \otimes \mathbf{B} \\ \vdots & \ddots & \vdots \\ a_{m-1,0} \otimes \mathbf{B} & \dots & a_{m-1,n-1} \otimes \mathbf{B} \end{bmatrix}$$

The \otimes operator is defined by the op parameter. It is applied in an element-wise fashion (like $GrB_eWiseMult$), where the pattern of the submatrix $a_{ij}\otimes B$ is the same as the pattern of B if a_{ij} is an entry in the matrix A, or empty otherwise. The input matrices A and B can be of any dimension, and both matrices may be transposed first via the descriptor, desc. Entries in A and B are typecast into the input types of the op. The matrix T=kron(A,B) has the same type as the ztype of the binary operator, op. The final step is $C\langle M \rangle = C \odot T$, as described in Section 2.3.

9 Printing GraphBLAS objects

SPEC: The GraphBLAS API has no mechanism for printing the contents of GraphBLAS objects. This entire section is an extension to the specification.

The ten different objects handled by SuiteSparse:GraphBLAS are all opaque, although nearly all of their contents can be extracted via methods such as GrB_Matrix_extractTuples, GrB_Matrix_extractElement, GxB_Matrix_type, and so on. The GraphBLAS C API has no mechanism for printing all the contents of GraphBLAS objects, but this is helpful for debugging. Ten type-specific methods and two type-generic methods are provided:

GxB_Type_fprint	print and check a GrB_Type
<pre>GxB_UnaryOp_fprint</pre>	print and check a GrB_UnaryOp
<pre>GxB_BinaryOp_fprint</pre>	print and check a GrB_BinaryOp
<pre>GxB_SelectOp_fprint</pre>	print and check a GxB_SelectOp
<pre>GxB_Monoid_fprint</pre>	print and check a GrB_Monoid
<pre>GxB_Semiring_fprint</pre>	print and check a GrB_Semiring
<pre>GxB_Descriptor_fprint</pre>	print and check a GrB_Descriptor
<pre>GxB_Matrix_fprint</pre>	print and check a GrB_Matrix
<pre>GxB_Vector_fprint</pre>	print and check a GrB_Vector
<pre>GxB_Scalar_fprint</pre>	print and check a GxB_Scalar
GxB_fprint	print/check any object to a file
GxB_print	print/check any object to stdout

These methods do not modify the status of any object. If a matrix or vector has not been completed, the pending computations are guaranteed to *not* be performed. The reason is simple. It is possible for a bug in the user application (such as accessing memory outside the bounds of an array) to mangle the internal content of a GraphBLAS object, and the <code>GxB_*print</code> methods can be helpful tools to track down this bug. If <code>GxB_*print</code> attempted to complete any computations prior to printing or checking the contents of the matrix or vector, then further errors could occur, including a segfault.

By contrast, all GraphBLAS methods and operations that return values into user-provided arrays or variables force the completion of pending operations (GrB_*_nvals, GrB_*_extractElement, GrB_*_extractTuples, and GrB_reduce (to scalar)). The GxB_*print methods provide a useful alternative for debugging, and for a quick understanding of what GraphBLAS is computing while developing a user application.

Each of the methods has a parameter of type GxB_Print_Level that specifies the amount to print:

The ten type-specific functions include an additional argument, the name string. The name is printed at the beginning of the display (assuming the print level is not GxB_SILENT) so that the object can be more easily identified in the output. For the type-generic methods GxB_fprint and GxB_print, the name string is the variable name of the object itself.

If the file f is NULL, nothing is printed (pr is effectively GxB_SILENT). If name is NULL, it is treated as the empty string. These are not error conditions.

The methods check their input objects carefully and extensively, even when pr is equal to GxB_SILENT. The following error codes can be returned:

- Grb_Success: object is valid
- Grb_UNINITIALIZED_OBJECT: object is not initialized
- Grb_INVALID_OBJECT: object is not valid
- Grb_NULL_POINTER: object is a NULL pointer
- GrB_INVALID_VALUE: fprintf returned an I/O error; see the ANSI C error or GrB_error() for details.

The content of any GraphBLAS object is opaque, and subject to change. As a result, the exact content and format of what is printed is implementation-dependent, and will change from version to version of SuiteSparse:GraphBLAS. Do not attempt to rely on the exact content or format by trying to parse the resulting output via another program. The intent of these functions is to produce a report of an object for visual inspection. If the user application needs to extract content from a GraphBLAS matrix or vector, use GrB_*extractTuples instead.

9.1 GxB_fprint: Print a GraphBLAS object to a file

The GxB_fprint function prints the contents of any of the ten Graph-BLAS objects to the file f. If f is NULL, the results are printed to stdout. For example, to print the entire contents of a matrix A to the file f, use GxB_fprint (A, GxB_COMPLETE, f).

9.2 GxB_print: Print a GraphBLAS object to stdout

GxB_print is the same as GxB_fprint, except that it prints the contents of the object to stdout instead of a file f. For example, to print the entire contents of a matrix A, use GxB_print (A, GxB_COMPLETE).

9.3 GxB_Type_fprint: Print a GrB_Type

For example, GxB_Type_fprint (GrB_BOOL, "boolean type", GxB_COMPLETE, f) prints the contents of the GrB_BOOL object to the file f.

9.4 GxB_UnaryOp_fprint: Print a GrB_UnaryOp

For example, GxB_UnaryOp_fprint (GrB_LNOT, "not", GxB_COMPLETE, f) prints the GrB_LNOT unary operator to the file f.

9.5 GxB_BinaryOp_fprint: Print a GrB_BinaryOp

For example, GxB_BinaryOp_fprint (GrB_PLUS_FP64, "plus", GxB_COMPLETE, f) prints the GrB_PLUS_FP64 binary operator to the file f.

9.6 GxB_SelectOp_fprint: Print a GxB_SelectOp

For example, GxB_SelectOp_fprint (GxB_TRIL, "tril", GxB_COMPLETE, f) prints the GxB_TRIL select operator to the file f.

9.7 GxB_Monoid_fprint: Print a GrB_Monoid

For example, GxB_Monoid_fprint (GxB_PLUS_FP64_MONOID, "plus monoid", GxB_COMPLETE, f) prints the predefined GxB_PLUS_FP64_MONOID (based on the binary operator GrB_PLUS_FP64) to the file f.

9.8 GxB_Semiring_fprint: Print a GrB_Semiring

For example, GxB_Semiring_fprint (GxB_PLUS_TIMES_FP64, "standard", GxB_COMPLETE, f) prints the predefined GxB_PLUS_TIMES_FP64 semiring to the file f.

9.9 GxB_Descriptor_fprint: Print a GrB_Descriptor

For example, GxB_Descriptor_fprint (d, "descriptor", GxB_COMPLETE, f) prints the descriptor d to the file f.

9.10 GxB_Matrix_fprint: Print a GrB_Matrix

For example, GxB_Matrix_fprint (A, "my matrix", GxB_SHORT, f) prints about 30 entries from the matrix A to the file f.

9.11 GxB_Vector_fprint: Print a GrB_Vector

For example, GxB_Vector_fprint (v, "my vector", GxB_SHORT, f) prints about 30 entries from the vector v to the file f.

9.12 GxB_Scalar_fprint: Print a GxB_Scalar

```
GrB_Info GxB_Scalar_fprint  // print and check a GrB_Scalar (

GxB_Sclarr s,  // object to print and check const char *name,  // name of the object GxB_Print_Level pr,  // print level FILE *f  // file for output 
);
```

For example, GxB_Scalar_fprint (s, "my scalar", GxB_SHORT, f) prints a short description of the sparse scalar s to the file f.

9.13 Performance and portability considerations

Even when the print level is GxB_SILENT, these methods extensively check the contents of the objects passed to them, which can take some time. They should be considered debugging tools only, not for final use in production.

The return value of the <code>GxB_*print</code> methods can be relied upon, but the output to the file (or <code>stdout</code>) can change from version to version. If these methods are eventually added to the GraphBLAS C API Specification, a conforming implementation might never print anything at all, regardless of the <code>pr</code> value. This may be essential if the GraphBLAS library is installed in a dedicated device, with no file output, for example.

Some implementations may wish to print nothing at all if the matrix is not yet completed, or just an indication that the matrix has pending operations and cannot be printed, when non-blocking mode is employed. In this case, use <code>GrB_Matrix_wait</code>, <code>GrB_Vector_wait</code>, or <code>GxB_Scalar_wait</code> to finish all pending computations first. If a matrix or vector has pending operations, SuiteSparse:GraphBLAS prints a list of the *pending tuples*, which are the entries not yet inserted into the primary data structure. It can also print out entries that remain in the data structure but are awaiting deletion; these are called *zombies* in the output report.

Most of the rest of the report is self-explanatory.

10 Creating user-defined objects at compiletime (feature removed)

This feature has been removed as of SuiteSparse:GraphBLAS v3.2.0. User objects defined with GxB_*_define when GraphBLAS is compiled are not compatible with the faster matrix operations. Use the run-time definitions instead, from the GraphBLAS C API Specification (GrB_Type_new instead of GxB_Type_define in the GraphBLAS/User/*.m4 script).

11 Examples

Several examples of how to use GraphBLAS are listed below. They all appear in the Demo folder of SuiteSparse:GraphBLAS.

- 1. performing a breadth-first search,
- 2. finding a maximal independent set,
- 3. creating a random matrix,
- 4. creating a finite-element matrix,
- 5. reading a matrix from a file, and
- 6. complex numbers as a user-defined type.
- 7. triangle counting
- 8. PageRank
- 9. matrix import/export

Additional examples appear in the newly created LAGraph project, currently in progress. Finally, the Extras folder includes triangle counting and k-truss algorithms in GraphBLAS, and methods that do not GraphBLAS (both simple sequential methods, and methods that use OpenMP).

11.1 LAGraph

The LAGraph project is a community-wide effort to create graph algorithms based on GraphBLAS (any implementation of the API, not just SuiteSparse: GraphBLAS). As of Oct, 2019, the library includes the algorithms and utilities listed in the table below. Many additional algorithms are planned, such as betweenness centrality, PageRank, single-source shortest path (via delta stepping), minimum spanning trees, connected components, and many more. Refer to https://github.com/GraphBLAS/LAGraph for a current list of algorithms (the one below will soon be out of date). Most of the functions in the Demo/ and the Extras folder in SuiteSparse:GraphBLAS will eventually be translated into algorithms or utilities for LAGraph.

To use LAGraph with SuiteSparse:GraphBLAS, place the two folders LAGraph and GraphBLAS in the same parent directory. This allows the cmake script in LAGraph to find the copy of GraphBLAS. Alternatively, the GraphBLAS source could be placed anywhere, as long as sudo make install is performed.

Build GraphBLAS first, then the LAGraph library, and then the tests in LAGraph/Test.

Many of these algorithms are described in $[ACD^{+}20]$.

Algorithms	description
LAGraph_bfs_pushpull	a direction-optimized BFS [BAP12, YBO18],
	typically 2x faster than bfs5m
LAGraph_bfs_simple	a simple BFS (about the same as bfs5m)
LAGraph_bc_batch	batched betweenness-centrality
LAGraph_bc	betweenness-centrality
LAGraph_cdlp	community detection via label propagation
LAGraph_cc	connected components
LAGraph_BF_*	three variants of Bellman-Ford
LAGraph_allktruss	construct all k -trusses
LAGraph_dnn	sparse deep neural network [DAK19]
LAGraph_ktruss	construct a k -trusses
LAGraph_lcc	local clustering coefficient
LAGraph_pagerank	PageRank
LAGraph_pagerank2	PageRank variant
LAGraph_tricount	triangle counting

Utilities	description
LAGraph_Vector_isall	tests 2 vectors with a binary operator
${\tt LAGraph_Vector_isequal}$	tests if 2 vectors are equal
LAGraph_Vector_to_dense	converts a vector to dense
LAGraph_alloc_global	types, operators, monoids, and semirings
LAGraph_finalize	ends LAGraph
LAGraph_free	wrapper for free
LAGraph_free_global	frees objects created by _alloc_global
LAGraph_get_nthreads	get $\#$ of threads used
LAGraph_grread	read a binary matrix in Galois format
LAGraph_init	starts LAGraph
LAGraph_isall	tests 2 matrices with a binary operator
LAGraph_isequal	tests if 2 matrices are equal
LAGraph_ispattern	tests if all entries in a matrix are 1
LAGraph_malloc	wrapper for malloc
LAGraph_mmread	read a Matrix Market file
LAGraph_mmwrite	write a Matrix Market file
LAGraph_pattern	extracts the pattern of a matrix
LAGraph_prune_diag	diagonal entries from a matrix
LAGraph_rand	simple random number generator
LAGraph_rand64	int64_t random number generator
LAGraph_random	random matrix generator
LAGraph_randx	double random number generator
LAGraph_set_nthreads	set # of threads to use
LAGraph_tic	start a timer
LAGraph_toc	end a timer
LAGraph_tsvread	read a TSV file
LAGraph_xinit	starts LAGraph, with different malloc
LAgraph_1_to_n	construct the vector 1:n
GB_*sort*	sorting for LAGraph_cdlp

11.2 Breadth-first search

The bfs examples in the Demo folder provide several examples of how to compute a breadth-first search (BFS) in GraphBLAS. Additional BFS examples are in LAGraph, shown below. The LAGraph_bfs_simple function starts at a given source node s of an undirected graph with n nodes. The graph is represented as an n-by-n matrix, A, where A(i,j) is the edge (i,j). The matrix A can have any type (even a user-defined type), since the PAIR operator does not access its values. No typecasting will be done.

The vector \mathbf{v} of size \mathbf{n} holds the level of each node in the BFS, where $\mathbf{v}(\mathtt{i})=0$ if the node has not yet been seen. This particular value makes \mathbf{v} useful for another role. It can be used as a Boolean mask, since 0 is \mathtt{false} and nonzero is \mathtt{true} . Initially the entire \mathbf{v} vector is zero. It is initialized as a dense vector, with all entries present, to improve performance (otherwise, it will slowly grow, incrementally, and this will take a lot of time if the number of BFS levels is high).

The vector **q** is the set of nodes just discovered at the current level, where **q(i)=true** if node **i** is in the current level. It starts out with just a single entry set to true, **q(s)**, the starting node.

Each iteration of the BFS consists of three calls to GraphBLAS. The first one uses q as a mask. It modifies all positions in v where q has an entry, setting them all to the current level.

```
// v < q > = level, using vector assign with q as the mask GrB_assign (v, q, NULL, level, GrB_ALL, n, GrB_DESC_S);
```

The next call to GraphBLAS is the heart of the algorithm:

```
// q<!v> = q ||.&& A ; finds all the unvisited
// successors from current q, using !v as the mask
GrB_vxm (q, v, NULL, GxB_ANY_PAIR_BOOL, q, A, GrB_DESC_RC) ;
```

The vector \mathbf{q} is all the set of nodes at the current level. Suppose $\mathbf{q}(\mathbf{j})$ is true, and it has a neighbor \mathbf{i} . Then $\mathbf{A}(\mathbf{i},\mathbf{j})=1$, and the dot product of $\mathbf{A}(\mathbf{i},\mathbf{i})*\mathbf{q}$ using the ANY_PAIR semiring will use the PAIR multiplier on these two terms, \mathbf{f} ($\mathbf{A}(\mathbf{i},\mathbf{j})$, $\mathbf{q}(\mathbf{j})$), resulting in a value 1. The ANY monoid will "sum" up all the results in this single row \mathbf{i} ; note that the OR monoid would compute the same thing. If the result is a column vector $\mathbf{t}=\mathbf{A}*\mathbf{q}$, then this $\mathbf{t}(\mathbf{i})$ will be true. The vector \mathbf{t} will be true for any node adjacent to any node in the set \mathbf{q} .

Some of these neighbors of the nodes in q have already been visited by the BFS, either in the current level or in a prior level. These results must be discarded; what is desired is the set of all nodes i for which t(i) is true, and yet v(i) is still zero.

Enter the mask. The vector \mathbf{v} is complemented for use a mask, via the desc descriptor. This means that wherever the vector is true, that position in the result is protected and will not be modified by the assignment. Only where \mathbf{v} is false will the result be modified. This is exactly the desired result, since these represent newly seen nodes for the next level of the BFS. A node \mathbf{k} already visited will have a nonzero $\mathbf{v}(\mathbf{k})$, and thus $\mathbf{q}(\mathbf{k})$ will not be modified by the assignment.

The result t is written back into the vector q, through the mask, but to do this correctly, another descriptor parameter is used: GrB_REPLACE. The vector q was used to compute t=A*q, and after using it to compute t, the entire q vector needs to be cleared. Only new nodes are desired, for the next level. This is exactly what the REPLACE option does.

As a result, the vector **q** now contains the set of nodes at the new level of the BFS. It contains all those nodes (and only those nodes) that are neighbors of the prior set and that have not already been seen in any prior level.

A single call to GrB_Vector_nvals finds how many entries are in the current level. If this is zero, the BFS can terminate.

```
#include "LAGraph_internal.h"
#define LAGRAPH_FREE_ALL { GrB_free (&v) ; GrB_free (&q) ; }
GrB_Info LAGraph_bfs_simple
                              // push-only BFS
   GrB_Vector *v_output,
                         // v(i) is the BFS level of node i in the graph
   GrB_Matrix A,
                           // input graph, treated as if boolean in semiring
   GrB_Index source
                          // starting node of the BFS
)
{
   GrB_Info info ;
   GrB_Vector q = NULL ;
                                   // nodes visited at each level
                                   // result vector
   GrB_Vector v = NULL ;
   if (v_output == NULL) LAGRAPH_ERROR ("argument missing", GrB_NULL_POINTER) ;
   GrB_Index n, nvals ;
   GrB_Matrix_nrows (&n, A) ;
   // create an empty vector v, and make it dense
   GrB_Vector_new (&v, (n > INT32_MAX) ? GrB_INT64 : GrB_INT32, n) ;
   GrB_assign (v, NULL, NULL, 0, GrB_ALL, n, NULL) ;
   // create a boolean vector q, and set q(source) to true
   GrB_Vector_new (&q, GrB_BOOL, n) ;
   GrB_Vector_setElement (q, true, source) ;
   // BFS traversal and label the nodes
   for (int64_t level = 1 ; level <= n ; level++)</pre>
       // v < q > = level
       GrB_assign (v, q, NULL, level, GrB_ALL, n, GrB_DESC_S) ;
       // break if q is empty
       GrB_Vector_nvals (&nvals, q) ;
       if (nvals == 0) break ;
       // q' <! v> = q' *A
       // free workspace and return result
    (*v_output) = v;
                          // return result
                           // set to NULL so LAGRAPH_FREE_ALL doesn't free it
   v = NULL;
   LAGRAPH_FREE_ALL ;
                          // free all workspace (except for result v)
   return (GrB_SUCCESS) ;
}
```

11.3 Maximal independent set

The maximal independent set problem is to find a set of nodes S such that no two nodes in S are adjacent to each other (an independent set), and all nodes not in S are adjacent to at least one node in S (and thus S is maximal since it cannot be augmented by any node while remaining an independent set). The mis function in the Demo folder solves this problem using Luby's method [Lub86]. The key operations in the method are replicated on the next page.

The gist of the algorithm is this. In each phase, all candidate nodes are given a random score. If a node has a score higher than all its neighbors, then it is added to the independent set. All new nodes added to the set cause their neighbors to be removed from the set of candidates. The process must be repeated for multiple phases until no new nodes can be added. This is because in one phase, a node \mathbf{i} might not be added because one of its neighbors \mathbf{j} has a higher score, yet that neighbor \mathbf{j} might not be added because one of its neighbors \mathbf{k} is added to the independent set instead. The node \mathbf{j} is no longer a candidate and can never be added to the independent set, but node \mathbf{i} could be added to S in a subsequent phase.

The initialization step, before the while loop, computes the degree of each node with a PLUS reduction. The set of candidates is Boolean vector, the ith component is true if node i is a candidate. A node with no neighbors causes the algorithm to stall, so these nodes are not candidates. Instead, they are immediately added to the independent set, represented by another Boolean vector iset. Both steps are done with an assign, using the degree as a mask, except the assignment to iset uses the complement of the mask, via the sr_desc descriptor. Finally, the GrB_Vector_nvals statement counts how many candidates remain.

Each phase of Luby's algorithm consists of 11 calls to GraphBLAS operations, all of which are either parallel, or take O(1) time. Not all of them are described here since they are commented in the code itself. The two matrix-vector multiplications are the important parts and also take the most time. They also make interesting use of semirings and masks. The first one computes the largest score of all the neighbors of each node in the candidate set:

```
// compute the max probability of all neighbors
GrB_vxm (neighbor_max, candidates, NULL, maxFirst, prob, A, r_desc);
```

```
// compute the degree of each node
GrB_reduce (degrees, NULL, NULL, GrB_PLUS_FP64, A, NULL) ;
// singletons are not candidates; they are added to iset first instead
// candidates[degree != 0] = 1
GrB_assign (candidates, degrees, NULL, true, GrB_ALL, n, NULL);
// add all singletons to iset
// iset[degree == 0] = 1
GrB_assign (iset, degrees, NULL, true, GrB_ALL, n, sr_desc) ;
// Iterate while there are candidates to check.
GrB_Index nvals ;
GrB_Vector_nvals (&nvals, candidates) ;
while (nvals > 0)
    // sparsify the random number seeds (just keep it for each candidate)
    GrB_assign (Seed, candidates, NULL, Seed, GrB_ALL, n, r_desc) ;
    // compute a random probability scaled by inverse of degree
    prand_xget (X, Seed) ; // two calls to GrB_apply
    GrB_eWiseMult (prob, candidates, NULL, set_random, degrees, X, r_desc) ;
    // compute the max probability of all neighbors
    GrB_vxm (neighbor_max, candidates, NULL, maxFirst, prob, A, r_desc) ;
    // select node if its probability is > than all its active neighbors
    GrB_eWiseAdd (new_members, NULL, NULL, GrB_GT_FP64, prob, neighbor_max,0)
    // add new members to independent set.
    GrB_eWiseAdd (iset, NULL, NULL, GrB_LOR, iset, new_members, NULL) ;
    // remove new members from set of candidates c = c & !new
    GrB_apply (candidates, new_members, NULL, GrB_IDENTITY_BOOL,
        candidates, sr_desc);
    GrB_Vector_nvals (&nvals, candidates);
    if (nvals == 0) { break ; }
                                                 // early exit condition
    // Neighbors of new members can also be removed from candidates
    GrB_vxm (new_neighbors, candidates, NULL, Boolean,
        new_members, A, NULL);
    GrB_apply (candidates, new_neighbors, NULL, GrB_IDENTITY_BOOL,
        candidates, sr_desc);
    GrB_Vector_nvals (&nvals, candidates) ;
```

A is a symmetric Boolean matrix and prob is a sparse real vector (of type FP32), where prob(i) is nonzero only if node i is a candidate. The prob vector is computed from a random vector computed by a utility function prand_xget, in the Demo folder. It uses two calls to GrB_apply to construct

n random numbers in parallel, using a repeatable pseudo-random number generator.

The maxFirst semiring uses z=FIRST(x,y) as the multiplier operator. The column A(:,j) is the adjacency of node j, and the dot product prob'*A(:,j) applies the FIRST operator on all entries that appear in the intersection of prob and A(:,j), where z=FIRST(prob(i),A(i,j)) which is just prob(i) if A(i,j) is present. If A(i,j) not an explicit entry in the matrix, then this term is not computed and does not take part in the reduction by the MAX monoid.

Thus, each term z=FIRST(prob(i),A(i,j)) is the score, prob(i), of all neighbors i of node j that have a score. Node i does not have a score if it is not also a candidate and so this is skipped. These terms are then "summed" up by taking the maximum score, using MAX as the additive monoid.

Finally, the results of this matrix-vector multiply are written to the result, neighbor_max. The r_desc descriptor has the REPLACE option enabled. Since neighbor_max does not also take part in the computation prob'*A, it is simply cleared first. Next, is it modified only in those positions i where candidates(i) is true, using candidates as a mask. This sets the neighbor_max only for candidate nodes, and leaves the other components of neighbor_max as zero (implicit values not in the pattern of the vector).

All of the above work is done in a single matrix-vector multiply, with an elegant use of the maxFirst semiring coupled with a mask. The matrix-vector multiplication is described above as if it uses dot products of rows of A with the column vector prob, but SuiteSparse:GraphBLAS does not compute it that way. Sparse dot products are much slower the optimal method for multiplying a sparse matrix times a sparse vector. The result is the same, however.

The second matrix-vector multiplication is more straight-forward. Once the set of new members in the independent is found, it is used to remove all neighbors of those new members from the set of candidates.

The resulting method is very efficient. For the Freescale2 matrix, the algorithm finds an independent set of size 1.6 million in 1.7 seconds (on the same MacBook Pro referred to in Section 11.2, using a single core), taking four iterations of the while loop. For comparison, removing its diagonal entries (required for the algorithm to work) takes 0.3 seconds in GraphBLAS (see Section 8.15), and simply transposing the matrix takes 0.24 seconds in both MATLAB and GraphBLAS.

11.4 Creating a random matrix

The random_matrix function in the Demo folder generates a random matrix with a specified dimension and number of entries, either symmetric or unsymmetric, and with or without self-edges (diagonal entries in the matrix). It relies on simple_rand* functions in the Demo folder to provide a portable random number generator that creates the same sequence on any computer and operating system.

random_matrix can use one of two methods: GrB_Matrix_setElement
and GrB_Matrix_build. The former method is very simple to use:

```
GrB_Matrix_new (&A, GrB_FP64, nrows, ncols);
for (int64_t k = 0; k < ntuples; k++)
{
    GrB_Index i = simple_rand_i () % nrows;
    GrB_Index j = simple_rand_i () % ncols;
    if (no_self_edges && (i == j)) continue;
    double x = simple_rand_x ();
    // A (i,j) = x
    GrB_Matrix_setElement (A, x, i, j);
    if (make_symmetric)
    {
        // A (j,i) = x
        GrB_Matrix_setElement (A, x, j, i);
    }
}</pre>
```

The above code can generate a million-by-million sparse double matrix with 200 million entries in 66 seconds (6 seconds of which is the time to generate the random i, j, and x), including the time to finish all pending computations. The user application does not need to create a list of all the tuples, nor does it need to know how many entries will appear in the matrix. It just starts from an empty matrix and adds them one at a time in arbitrary order. GraphBLAS handles the rest. This method is not feasible in MATLAB.

The next method uses GrB_Matrix_build. It is more complex to use than setElement since it requires the user application to allocate and fill the tuple lists, and it requires knowledge of how many entries will appear in the matrix, or at least a good upper bound, before the matrix is constructed. It is slightly faster, creating the same matrix in 60 seconds, 51 seconds of which is spent in GrB_Matrix_build.

```
GrB_Index *I, *J;
double *X ;
int64_t s = ((make_symmetric) ? 2 : 1) * nedges + 1 ;
I = malloc (s * sizeof (GrB_Index)) ;
J = malloc (s * sizeof (GrB_Index));
X = malloc (s * sizeof (double
                                 ));
if (I == NULL || J == NULL || X == NULL)
{
    // out of memory
    if (I != NULL) free (I) :
    if (J != NULL) free (J) :
    if (X != NULL) free (X) :
   return (GrB_OUT_OF_MEMORY) ;
int64_t ntuples = 0 ;
for (int64_t k = 0 ; k < nedges ; k++)
    GrB_Index i = simple_rand_i ( ) % nrows ;
    GrB_Index j = simple_rand_i ( ) % ncols ;
    if (no_self_edges && (i == j)) continue ;
    double x = simple_rand_x ( ) ;
    // A (i,j) = x
    I [ntuples] = i ;
    J [ntuples] = j ;
    X [ntuples] = x ;
   ntuples++;
    if (make_symmetric)
        // A (j,i) = x
        I [ntuples] = j ;
        J [ntuples] = i ;
        X [ntuples] = x ;
        ntuples++;
    }
GrB_Matrix_build (A, I, J, X, ntuples, GrB_SECOND_FP64) ;
```

The equivalent sprandsym function in MATLAB takes 150 seconds, but sprandsym uses a much higher-quality random number generator to create the tuples [I,J,X]. Considering just the time for sparse(I,J,X,n,n) in sprandsym (equivalent to GrB_Matrix_build), the time is 70 seconds. That is, each of these three methods, setElement and build in Suite-Sparse:GraphBLAS, and sparse in MATLAB, are equally fast.

11.5 Creating a finite-element matrix

Suppose a finite-element matrix is being constructed, with k=40,000 finite-element matrices, each of size 8-by-8. The following operations (in pseudo-MATLAB notation) are very efficient in SuiteSparse:GraphBLAS.

```
A = sparse (m,n) ; % create an empty n-by-n sparse GraphBLAS matrix
for i = 1:k
    construct a 8-by-8 sparse or dense finite-element F
    I and J define where the matrix F is to be added:
    I = a list of 8 row indices
    J = a list of 8 column indices
    % using GrB_assign, with the 'plus' accum operator:
    A (I,J) = A (I,J) + F
end
```

If this were done in MATLAB or in GraphBLAS with blocking mode enabled, the computations would be extremely slow. This example is taken from Loren Shure's blog on MATLAB Central, Loren on the Art of MATLAB [Dav07], which discusses the built-in wathen function. In MATLAB, a far better approach is to construct a list of tuples [I,J,X] and to use sparse(I,J,X,n,n). This is identical to creating the same list of tuples in GraphBLAS and using the GrB_Matrix_build, which is equally fast. The difference in time between using sparse or GrB_Matrix_build, and using submatrix assignment with blocking mode (or in MATLAB which does not have a nonblocking mode) can be extreme. For the example matrix discussed in [Dav07], using sparse instead of submatrix assignment in MATLAB cut the run time of wathen from 305 seconds down to 1.6 seconds.

In SuiteSparse:GraphBLAS, the performance of both methods is essentially identical, and roughly as fast as sparse in MATLAB. Inside SuiteSparse:GraphBLAS, GrB_assign is doing the same thing. When performing A(I,J)=A(I,J)+F, if it finds that it cannot quickly insert an update into the A matrix, it creates a list of pending tuples to be assembled later on. When the matrix is ready for use in a subsequent GraphBLAS operation (one that normally cannot use a matrix with pending computations), the tuples are assembled all at once via GrB_Matrix_build.

GraphBLAS operations on other matrices have no effect on when the pending updates of a matrix are completed. Thus, any GraphBLAS method or operation can be used to construct the F matrix in the example above, without affecting when the pending updates to A are completed.

The MATLAB wathen.m script is part of Higham's gallery of matrices [Hig02]. It creates a finite-element matrix with random coefficients for a 2D mesh of size nx-by-ny, a matrix formulation by Wathen [Wat87]. The pattern of the matrix is fixed; just the values are randomized. The GraphBLAS equivalent can use either GrB_Matrix_build, or GrB_assign. Both methods have good performance. The GrB_Matrix_build version below is about 15% to 20% faster than the MATLAB wathen.m function, regardless of the problem size. It uses the identical algorithm as wathen.m.

```
int64_t ntriplets = nx*ny*64 ;
I = malloc (ntriplets * sizeof (int64_t));
J = malloc (ntriplets * sizeof (int64_t));
X = malloc (ntriplets * sizeof (double )) ;
if (I == NULL || J == NULL || X == NULL)
    FREE_ALL ;
    return (GrB_OUT_OF_MEMORY) ;
}
ntriplets = 0 ;
for (int j = 1; j \le ny; j++)
   for (int i = 1; i <= nx; i++)
        nn [0] = 3*j*nx + 2*i + 2*j + 1;
        nn [1] = nn [0] - 1;
       nn [2] = nn [1] - 1;
        nn [3] = (3*j-1)*nx + 2*j + i - 1;
        nn [4] = 3*(j-1)*nx + 2*i + 2*j - 3;
        nn [5] = nn [4] + 1;
        nn [6] = nn [5] + 1;
        nn [7] = nn [3] + 1;
        for (int krow = 0; krow < 8; krow++) nn [krow]--;
        for (int krow = 0; krow < 8; krow++)
        {
            for (int kcol = 0; kcol < 8; kcol++)
            {
               I [ntriplets] = nn [krow] ;
                J [ntriplets] = nn [kcol] ;
               X [ntriplets] = em (krow,kcol) ;
               ntriplets++ ;
            }
       }
   }
}
```

```
// A = sparse (I,J,X,n,n) ;
GrB_Matrix_build (A, I, J, X, ntriplets, GrB_PLUS_FP64) ;
```

The GrB_assign version has the advantage of not requiring the user application to construct the tuple list, and is almost as fast as using GrB_Matrix_build. The code is more elegant than either the MATLAB wathen.m function or its GraphBLAS equivalent above. Its performance is comparable with the other two methods, but slightly slower, being about 5% slower than the MATLAB wathen, and 20% slower than the GraphBLAS method above.

```
GrB_Matrix_new (&F, GrB_FP64, 8, 8);
for (int j = 1; j \le ny; j++)
    for (int i = 1; i <= nx; i++)
        nn [0] = 3*j*nx + 2*i + 2*j + 1;
        nn [1] = nn [0] - 1;
        nn [2] = nn [1] - 1;
        nn [3] = (3*j-1)*nx + 2*j + i - 1;
        nn [4] = 3*(j-1)*nx + 2*i + 2*j - 3;
        nn [5] = nn [4] + 1;
       nn [6] = nn [5] + 1;
       nn [7] = nn [3] + 1;
        for (int krow = 0; krow < 8; krow++) nn [krow]--;</pre>
        for (int krow = 0; krow < 8; krow++)
        {
            for (int kcol = 0; kcol < 8; kcol++)
                // F (krow,kcol) = em (krow, kcol)
               GrB_Matrix_setElement (F, em (krow,kcol), krow, kcol);
            }
        // A (nn,nn) += F
        GrB_assign (A, NULL, GrB_PLUS_FP64, F, nn, 8, nn, 8, NULL) ;
   }
}
```

Since there is no Mask, and since GrB_REPLACE is not used, the call to GrB_assign in the example above is identical to GxB_subassign. Either one can be used, and their performance would be identical.

Refer to the wathen.c function in the Demo folder, which uses GraphBLAS to implement the two methods above, and two additional ones.

11.6 Reading a matrix from a file

NOTE: see also LAGraph_mmread and LAGraph_mmwrite, which can read and write any matrix in Matrix Market format, and LAGraph_binread and LAGraph_binwrite, which read/write a matrix from a binary file. The binary file I/O functions are much faster than the read_matrix function described here, and also much faster than LAGraph_mmread and LAGraph_mmwrite.

The read_matrix function in the Demo reads in a triplet matrix from a file, one line per entry, and then uses GrB_Matrix_build to create the matrix. It creates a second copy with GrB_Matrix_setElement, just to test that method and compare the run times. A comparison of build versus setElement has already been discussed in Section 11.4.

The function can return the matrix as-is, which may be rectangular or unsymmetric. If an input parameter is set to make the matrix symmetric, read_matrix computes A=(A+A')/2 if A is square (turning all directed edges into undirected ones. If A is rectangular, it creates a bipartite graph, which is the same as the augmented matrix, $A = [0 \ A \ ; A' \ 0]$. If C is an n-by-n matrix, then C=(C+C')/2 can be computed as follows in GraphBLAS, (the scale2 function divides an entry by 2):

```
GrB_Descriptor_new (&dt2) ;
GrB_Descriptor_set (dt2, GrB_INP1, GrB_TRAN) ;
GrB_Matrix_new (&A, GrB_FP64, n, n) ;
GrB_eWiseAdd (A, NULL, NULL, GrB_PLUS_FP64, C, C, dt2) ; // A=C+C'
GrB_free (&C) ;
GrB_Matrix_new (&C, GrB_FP64, n, n) ;
GrB_UnaryOp_new (&scale2_op, scale2, GrB_FP64, GrB_FP64) ;
GrB_apply (C, NULL, NULL, scale2_op, A, NULL) ; // C=A/2
GrB_free (&A) ;
GrB_free (&scale2_op) ;
```

This is of course not nearly as elegant as A=(A+A')/2 in MATLAB, but with minor changes it can work on any type and use any built-in operators instead of PLUS, or it can use any user-defined operators and types. The above code in SuiteSparse:GraphBLAS takes 0.60 seconds for the Freescale2 matrix, slightly slower than MATLAB (0.55 seconds).

Constructing the augmented system is more complicated using the Graph-BLAS C API Specification since it does not yet have a simple way of specifying a range of row and column indices, as in A(10:20,30:50) in MATLAB (GxB_RANGE is a SuiteSparse:GraphBLAS extension that is not in the Speci-

fication). Using the C API in the Specification, the application must instead build a list of indices first, I=[10, 11 ... 20].

Thus, to compute the MATLAB equivalent of $A = [0 \ A \ ; \ A' \ 0]$, index lists I and J must first be constructed:

```
int64_t n = nrows + ncols ;
I = malloc (nrows * sizeof (int64_t)) ;
J = malloc (ncols * sizeof (int64_t)) ;
// I = 0:nrows-1
// J = nrows:n-1
if (I == NULL || J == NULL)
{
    if (I != NULL) free (I) ;
    if (J != NULL) free (J) ;
    return (GrB_OUT_OF_MEMORY) ;
}
for (int64_t k = 0 ; k < nrows ; k++) I [k] = k ;
for (int64_t k = 0 ; k < ncols ; k++) J [k] = k + nrows ;</pre>
```

Once the index lists are generated, however, the resulting GraphBLAS operations are fairly straightforward, computing A=[0 C; C'0].

```
GrB_Descriptor_new (&dt1) ;
GrB_Descriptor_set (dt1, GrB_INPO, GrB_TRAN) ;
GrB_Matrix_new (&A, GrB_FP64, n, n) ;
// A (nrows:n-1, 0:nrows-1) = C'
GrB_assign (A, NULL, NULL, C, J, ncols, I, nrows, dt1) ;
// A (0:nrows-1, nrows:n-1) = C
GrB_assign (A, NULL, NULL, C, I, nrows, J, ncols, NULL) ;
```

This takes 1.38 seconds for the Freescale2 matrix, almost as fast as A=[sparse(m,m) C; C' sparse(n,n)] in MATLAB (1.25 seconds).

Both calls to GrB_assign use no accumulator, so the second one causes the partial matrix A=[0 0; C'0] to be built first, followed by the final build of A=[0 C; C'0]. A better method, but not an obvious one, is to use the GrB_FIRST_FP64 accumulator for both assignments. An accumulator enables SuiteSparse:GraphBLAS to determine that that entries created by the first assignment cannot be deleted by the second, and thus it need not force completion of the pending updates prior to the second assignment.

SuiteSparse:GraphBLAS also adds a GxB_RANGE mechanism that mimics the MATLAB colon notation. This speeds up the method and simplifies the code the user needs to write to compute $A=[0\ C\ ;\ C'\ 0]$:

Any operator will suffice because it is not actually applied. An operator is only applied to the set intersection, and the two assignments do not overlap. If an accum operator is used, only the final matrix is built, and the time in GraphBLAS drops slightly to 1.25 seconds. This is a very small improvement because in this particular case, SuiteSparse:GraphBLAS is able to detect that no sorting is required for the first build, and the second one is a simple concatenation. In general, however, allowing GraphBLAS to postpone pending updates can lead to significant reductions in run time.

11.7 PageRank

The Demo folder contains three methods for computing the PageRank of the nodes of a graph. One uses floating-point arithmetic (GrB_FP64) and two user-defined unary operators (dpagerank.c). The second (ipagerank.c) is very similar, relying on integer arithmetic instead (GrB_UINT64). Neither method include a stopping condition. They simply compute a fixed number of iterations. The third example is more extensive (dpagerank2.c), and serves as an example of the power and flexibility of user-defined types, operators, monoids, and semirings. It creates a semiring for the entire PageRank computation. It terminates if the 2-norm of the change in the rank vector r is below a threshold.

11.8 Triangle counting

A triangle in an undirected graph is a clique of size three: three nodes i, j, and k that are all pairwise connected. There are many ways of counting the number of triangles in a graph. Let A be a symmetric matrix with values 0 and 1, and no diagonal entries; this matrix is the adjacency matrix of the graph. Let E be the edge incidence matrix with exactly two 1's per column. A column of E with entries in rows i and j represents the edge (i, j) in the graph, A(i,j)=1 where i<j. Let L and U be the strictly lower and upper triangular parts of A, respectively.

The methods are listed in the table below. Most of them use a form of masked matrix-matrix multiplication. The methods are implemented in MATLAB in the tricount.m file, and in GraphBLAS in the tricount.c file, both in the GraphBLAS/Demo folder. Refer to the comments in those two files for details and derivations on how these methods work.

When the matrix is stored by row, and a mask is present and not complemented, GrB_INP1 is GrB_TRAN , and GrB_INP0 is $GxB_DEFAULT$, the Suite-Sparse:GraphBLAS implementation of GrB_mxm always uses a dot-product formulation. Thus, the $C\langle L\rangle = LU^T$ method uses dot products. This provides a mechanism for the end-user to select a masked dot product matrix multiplication method in SuiteSparse:GraphBLAS, which is occasionally faster than the outer product method. The MATLAB form assumes the matrices are stored by column (the only option in MATLAB).

Each method is followed by a reduction to a scalar, via GrB_reduce in GraphBLAS or by nnz or sum(sum(...)) in MATLAB.

method and	in MATLAB	in GraphBLAS
citation		
minitri [WBS15]	nnz(A*E==2)/3	$\mathbf{C} = \mathbf{AE}, ext{ then GrB_apply}$
Burkhardt [Bur16]	$sum(sum((A^2).*A))/6$	$\mathbf{C}\langle\mathbf{A} angle=\mathbf{A}^2$
Cohen [ABG15, Coh09]	sum(sum((L*U).*A))/2	$\mathbf{C}\langle\mathbf{A} angle=\mathbf{L}\mathbf{U}$
Sandia [WDB ⁺ 17]	sum(sum((U*U).*U))	$\mathbf{C}\langle\mathbf{L}\rangle = \mathbf{L}\mathbf{L}$ (outer product)
SandiaDot	sum(sum((U'*L).*L))	$\mathbf{C}\langle\mathbf{U}\rangle = \mathbf{L}\mathbf{U}^{T} \text{ (dot product)}$
Sandia2	sum(sum((L*L).*L))	$\mathbf{C}\langle\mathbf{U}\rangle = \mathbf{U}\mathbf{U}$ (outer product)

In general, the Sandia methods are the fastest of the 6 methods when implemented in GraphBLAS. For full details on the triangle counting and k-truss algorithms, and performance results, see [Dav18], a copy of which appears in the SuiteSparse/GraphBLAS/Doc folder. The code appears in Extras. That paper uses an earlier version of SuiteSparse:GraphBLAS in which all matrices are stored by column.

11.9 User-defined types and operators: double complex and struct-based

The Demo folder contains two working examples of user-defined types, first discussed in Section 5.1.1: double complex, and a user-defined typedef called wildtype with a struct containing a string and a 4-by-4 float matrix.

Double Complex: Prior to v3.3, GraphBLAS did not have a native complex type. It now appears as the GxB_FC64 predefined type, but a complex type can also easily added as a user-defined type. The Complex_init function in the usercomplex.c file in the Demo folder creates the Complex type based on the ANSI C11 double complex type.

```
GrB_Type_new (&Complex, sizeof (double complex));
```

Next, it creates a full suite of operators that correspond to every built-in GraphBLAS operator, both binary and unary. In addition, it creates the operators listed in the following table, where D is double and C is Complex.

name	types	MATLAB equivalent	description
Complex_complex	$D \times D \to C$	z=complex(x,y)	complex from real and imag.
Complex_conj	$C \to C$	z=conj(x)	complex conjugate
Complex_real	$C \to D$	z=real(x)	real part
Complex_imag	$C \to D$	z=imag(x)	imaginary part
Complex_angle	$C \to D$	z=angle(x)	phase angle
Complex_complex_real	$D \to C$	z=complex(x,0)	real to complex real
Complex_complex_imag	$D \to C$	z=complex(0,x)	real to complex imag.

The Complex_init function creates two monoids (Complex_add_monoid and Complex_times_monoid) and a semiring Complex_plus_times that corresponds to the conventional linear algebra for complex matrices. The include file usercomplex.h in the Demo folder is available so that this user-defined Complex type can easily be imported into any other user application. When the user application is done, the Complex_finalize function frees the Complex type and its operators, monoids, and semiring.

NOTE: the Complex type is not supported in this Demo in Microsoft Visual Studio.

Struct-based: In addition, the wildtype.c program creates a user-defined typedef of a struct containing a dense 4-by-4 float matrix, and a 64-character string. It constructs an additive monoid that adds two 4-by-4 dense matrices, and a multiplier operator that multiplies two 4-by-4 matrices.

Each of these 4-by-4 matrices is treated by GraphBLAS as a "scalar" value, and they can be manipulated in the same way any other GraphBLAS type can be manipulated. The purpose of this type is illustrate the endless possibilities of user-defined types and their use in GraphBLAS.

11.10 User applications using OpenMP or POSIX pthreads

Two example demo programs are included that illustrate how a multi-threaded user application can use GraphBLAS: openmp_demo uses OpenMP for its user threads and pthread_demo uses POSIX pthreads. To be thread-safe, SuiteSparse:GraphBLAS must be compiled with a threading library, either OpenMP or POSIX. Either option used inside GraphBLAS can typically be combined with any user threading model. See Section 12.

The openmp_demo can be compiled without OpenMP, in which case it becomes single-threaded. GraphBLAS can be compiled with OpenMP, POSIX pthreads, or no threading support (and is not thread-safe in this latter case). This gives 9 different combinations:

User	GraphBLAS	Demo/Output file	comments
applic.			
none	none	user_none_grb_none.out	OK
none	OpenMP	user_none_grb_openmp.out	OK
none	pthread	${\tt user_none_grb_pthread.out}$	OK
OpenMP	none	user_openmp_grb_none.out	fail
OpenMP	OpenMP	user_openmp_grb_openmp.out	OK, random
OpenMP	pthread	${\tt user_openmp_grb_pthread.out}$	OK, random
pthread	none	user_pthread_grb_none.out	fail
pthread	OpenMP	user_pthread_grb_openmp.out	OK, random
pthread	pthread	${\tt user_pthread_grb_pthread.out}$	OK, random

When the user application is multithreaded, GraphBLAS must be compiled with a threading library to be thread-safe. The results listed above as OK, random mean that the output of the program will appear out of order. This is by design, simply to show that the user application is running in parallel. The output of each thread should be the same. In particular, each thread generates an intentional error, and later on prints it with GrB_error . It will print its own error, not an error from another thread. When all the threads finish, the master thread prints out each matrix generated by each thread, and these results are identical for all 7 cases listed above as OK.

The GraphBLAS C API requires GraphBLAS to be thread-safe. If Suite-Sparse:GraphBLAS is not compiled with a threading library it will not be thread-safe (the two *fail* cases above). For these cases, a thread will not retrieve its own error, but the last error of any thread. In addition, since there is no critical section that SuiteSparse:GraphBLAS can use, the output will include errors about an invalid state of the global matrix queue. These errors are to be expected if SuiteSparse:GraphBLAS is not thread-safe.

12 Compiling and Installing SuiteSparse:GraphBLAS

12.1 On Linux and Mac

GraphBLAS makes extensive use of features in the ANSI C11 standard, and thus a C compiler supporting this version of the C standard is required to use all features of GraphBLAS. On the Mac (OS X), clang 8.0.0 in Xcode version 8.2.1 is sufficient, although earlier versions of Xcode may work as well. For the GNU gcc compiler, version 4.9 or later is required. For the Intel icc compiler, version 18.0 or later is required. Version 2.8.12 or later of cmake is required; version 3.0.0 is preferred.

If you are using a pre-C11 ANSI C compiler, or Microsoft Visual Studio, then the _Generic keyword is not available. SuiteSparse:GraphBLAS will still compile, but you will not have access to polymorphic functions such as GrB_assign. You will need to use the non-polymorphic functions instead.

NOTE: icc is generally an excellent compiler, but it will generate slower code than gcc for v3.2.0. This is merely because of how the two compilers treat #pragma omp atomic read and #pragma omp atomic write. The use of gcc for SuiteSparse:GraphBLAS v3.2.0 is recommended. This difference in performance should be resolved in a future version.

To compile SuiteSparse:GraphBLAS and the demo programs, simply type make in the main GraphBLAS folder, which compiles the library. To use a non-default compiler:

```
make CC=icc CXX=icc JOBS=4
```

After compiling the library, you can run the demos by typing ./demo in the Demo folder.

If cmake or make fail, it might be that your default compiler does not support ANSI C11. Try another compiler. For example, try one of these options. Go into the build directory and type one of these:

```
CC=gcc cmake ..

CC=gcc-6 cmake ..

CC=xlc cmake ..

CC=icc cmake ..
```

You can also do the following in the top-level GraphBLAS folder instead:

```
CC=gcc make
CC=gcc-6 cmake
CC=xlc cmake
CC=icc cmake
```

For faster compilation, you can specify a parallel make. For example, to use 32 parallel jobs and the gcc compiler, do the following:

```
JOBS=32 CC=gcc make
```

12.2 On Microsoft Windows

SuiteSparse:GraphBLAS is now ported to Microsoft Visual Studio. However, that compiler is not ANSI C11 compliant and does not support OpenMP v4.0. As a result, GraphBLAS on Windows will have a few limitations.

- The MS Visual Studio compiler does not support the _Generic keyword, required for the polymorphic GraphBLAS functions. So for example, you will need to use GrB_Matrix_free instead of just GrB_free.
- Another limitation is the lack of support for OpenMP tasking, used in the parallel sort inside GraphBLAS. With Microsoft Visual Studio, the sort is compiled to use just a single thread. The sort is used for GrB_Matrix_build and GrB_Vector_build, and for GrB_assign and GxB_subassign when the index lists are unsorted on input. The internal sort still works as specified; it will just be single-threaded and thus these GraphBLAS functions will be slower on Windows as compared to Linux or MacOS.
- In addition, variable-length arrays are not supported, so user-defined types are limited to 128 bytes in size.

If you use a recent gcc or icc compiler on Windows other than the Microsoft Compiler (c1), these limitations can be avoided.

The following instructions apply to Windows 10, CMake 3.16, and Visual Studio 2019, but may work for earlier versions.

1. Install CMake 3.16 or later, if not already installed. See https://cmake.org/ for details.

- 2. Install Microsoft Visual Studio, if not already installed. See https://visualstudio.microsoft.com/ for details. Version 2019 is preferred, but earlier versions may also work.
- 3. Open a terminal window and type the following in the SuiteSparse/GraphBLAS/build folder.

cmake ..

- 4. The cmake command generates many files in SuiteSparse/GraphBLAS/build, and the file graphblas.sln in particular. Open the generated graphblas.sln file in Visual Studio.
- 5. Optionally: right-click graphblas in the left panel (Solution Explorer) and select properties; then navigate to Configuration Properties, C/C++, General and change the parameter Multiprocessor Compilation to Yes (/MP). Click OK. This will significantly speed up the compilation of GraphBLAS.
- 6. Select the Build menu item at the top of the window and select Build Solution. This should create a folder called Release and place the compiled graphblas.dll, graphblas.lib, and graphblas.exp files there. Please be patient; some files may take a while to compile and sometimes may appear to be stalled. Just wait.
- 7. Add the GraphBLAS/build/Release folder to the Windows System path:
 - Open the Start Menu and type Control Panel.
 - Select the Control Panel app.
 - When the app opens, select System.
 - From the top left side of the System window, select Advanced System Settings. You may have to authenticate at this step.
 - The Systems Properties window should appear with the Advanced tab selected; select Environment Variables.

- The Environment Variables window displays 2 sections, one for User variables and the other for System variables. Under the Systems variable section, scroll to and select Path, then select Edit. A editor window appears allowing to add, modify, delete or re-order the parts of the Path.
- Add the full path of the GraphBLAS\build\Release folder (typically starting with C:\Users\you\..., where you is your Windows username) to the Path.
- If the above steps do not work, you can instead copy the graphblas.* files from GraphBLAS\build\Release into any existing folder listed in your Path.
- 8. The GraphBLAS/Include/GraphBLAS.h file must be included in user applications via #include "GraphBLAS.h". This is already done for you in the MATLAB interface discussed in the next section.

12.3 Compiling the MATLAB interface

First, compile the SuiteSparse:GraphBLAS dynamic library (libgraphblas.so for Linux, libgraphblas.dylib for Mac, or graphblas.dll for Windows), as described in the prior two subsections. Next:

1. In the MATLAB command window:

cd GraphBLAS/GraphBLAS/@GrB/private
gbmake

- 2. Follow the remaining instructions in the GraphBLAS/GraphBLAS/README.md file, to revise your MATLAB path and startup.m file.
- 3. As a quick test, try the MATLAB command GrB(1), which creates and displays a 1-by-1 GraphBLAS matrix. For a longer test, do the following:

cd GraphBLAS/GraphBLAS/test
gbtest

4. In Windows, if the tests fail with an error stating that the mex file is invalid because the module could not be found, it means that MAT-LAB could not find the compiled graphblas.lib, *.dll or *.exp files in the build/Release folder. This can happen if your Windows System path is not set properly, or if Windows is not recognizing the GraphBLAS/build/Release folder (see Section 12.2) Or, you might p have permission to change your Windows System path. In this case, do the following in the MATLAB Command Window:

cd GraphBLAS/build/Release
GrB(1)

After this step, the GraphBLAS library will be loaded into MATLAB. You may need to add the above lines in your Documents/MATLAB/startup.m file, so that they are done each time MATLAB starts. You will also need to do this after clear all or clear mex, since those MATLAB commands remove all loaded libraries from MATLAB.

12.4 Thread-safety in multithreaded user applications

SuiteSparse:GraphBLAS is parallel, based on OpenMP. It is thread-safe if multiple simultaneous calls are made to GraphBLAS functions, from user threads that rely on either OpenMP or POSIX pthreads. The output variables of those calls to GraphBLAS must be unique; you cannot safely modify one GraphBLAS object in parallel, with two or more simultaneous GraphBLAS functions operating on the same output object. In addition, all pending operations of objects that appear in parallel calls to GraphBLAS must be complete. This can be done for all objects via GrB_Matrix_wait, GrB_Vector_wait, and GxB_Scalar_wait, which force completion of a particular object. If multiple parallel calls to GraphBLAS functions operate on unique inputs, then those input objects can safely have pending operations.

To use GraphBLAS from a multithreaded user application, GraphBLAS requires access to a critical section for the (now deprecated) GrB_wait() queue of matrices with pending operations, and to a thread-local storage space so that each user thread can safely retrieve its own error message with GrB_error. In v4.0, the no-argument GrB_wait() function will be removed, and thus the user-thread-based critical section will be no longer needed.

SuiteSparse:GraphBLAS supports the following user threading models. By default, the cmake script detects the presence of OpenMP and POSIX pthreads. If OpenMP is present, it uses OpenMP critical sections for GrB_wait() and OpenMP threadprivate(...) for thread-local storage for GrB_error. Otherwise, if POSIX pthreads are available, it uses a POSIX mutex, and POSIX thread-local storage via pthread_key_create.

These methods used inside GraphBLAS can typically inter-operate with any user threading model. That is, a user application that relies on POSIX threads, OpenMP, ANSI C11 threads, or Microsoft Windows threads will find GraphBLAS thread-safe, even though GraphBLAS uses OpenMP or POSIX internally to synchronize the user threads. However, for the most reliable results, the preferred approach is to use the same threading model in GraphBLAS as is used in the user application.

You can modify the automatic selection of a user thread synchronization model by adding the following settings for cmake. This setting does not determine how SuiteSparse:GraphBLAS creates and exploits multiple threads *inside* any given GraphBLAS operation. Rather, it determines which threading library it will use to synchronize multiple calls to GraphBLAS from more than one user thread.

- OpenMP: this is the default if your compiler supports OpenMP. It can also be specified with cmake -DUSER_OPENMP=1 in the cmake command line. Internal parallelism in SuiteSparse:GraphBLAS version is based on OpenMP. This is typically safe to use with any user threading models.
- POSIX: this is used if OpenMP is not available. If OpenMP is available but you still want GraphBLAS to use POSIX synchronization, compile with cmake -DUSER_POSIX=1
- no user threading: compile with cmake -DUSER_NONE=1. **GraphBLAS** will not be thread-safe.

12.5 Default matrix format

By default, SuiteSparse:GraphBLAS stores its matrices by row, using the GxB_BY_ROW format. You can change the default at compile time to GxB_BY_COL using cmake -DBYCOL=1. For example:

```
cmake -DBYCOL=1 ...
```

The user application can also use GxB_get and GxB_set to set and query the global option (see also Sections 6.7 and 6.8):

12.6 Setting the C flags and using CMake

The above options can also be combined. For example, to use the gcc compiler, to change the default format GxB_FORMAT_DEFAULT to GxB_BY_COL, and to use a POSIX mutex inside GraphBLAS to synchronize user threads, use the following cmake command while in the GraphBLAS/build directory:

```
CC=gcc cmake -DBYCOL=1 -DUSER_POSIX=1 ..
```

Then do make in the build directory. If this still fails, see the CMakeLists.txt file. You can edit that file to pass compiler-specific options to your compiler. Locate this section in the CMakeLists.txt file. Use the set command in cmake, as in the example below, to set the compiler flags you need.

```
# check which compiler is being used. If you need to make
# compiler-specific modifications, here is the place to do it.
if ("${CMAKE_C_COMPILER_ID}" STREQUAL "GNU")
    # cmake 2.8 workaround: gcc needs to be told to do ANSI C11.
    # cmake 3.0 doesn't have this problem.
    set ( CMAKE_C_FLAGS "${CMAKE_C_FLAGS} -std=c11 -lm " )
    ...
elseif ("${CMAKE_C_COMPILER_ID}" STREQUAL "Intel")
    ...
elseif ("${CMAKE_C_COMPILER_ID}" STREQUAL "Clang")
    ...
elseif ("${CMAKE_C_COMPILER_ID}" STREQUAL "MSVC")
    ...
endif ( )
```

To compile SuiteSparse:GraphBLAS without running the demos, use make library in the top-level directory, or make in the build directory.

Several compile-time options can be selected by editing the Source/GB.h file, but these are meant only for code development of SuiteSparse:GraphBLAS itself, not for end-users of SuiteSparse:GraphBLAS.

12.7 Using a plain makefile

The GraphBLAS/alternative directory contains a simple Makefile that can be used to compile SuiteSparse:GraphBLAS. This is a useful option if you do not have the required version of cmake.

12.8 Running the Demos

By default, make in the top-level directory compiles the library and runs the demos. You can also run the demos after compiling:

```
cd Demo
```

The ./demo command is a script that runs the demos with various input matrices in the Demo/Matrix folder. The output of the demos will be compared with expected output files in Demo/Output.

12.9 Installing SuiteSparse:GraphBLAS

To install the library (typically in /usr/local/lib and /usr/local/include for Linux systems), go to the top-level GraphBLAS folder and type:

sudo make install

12.10 Running the tests

To perform the extensive tests in the Test folder, and the statement coverage tests in Tcov, MATLAB R2017A is required. See the README.txt files in those two folders for instructions on how to run the tests. The tests in the Test folder have been ported to MATLAB on Linux, MacOS, and Windows. The Tcov tests do not work on Windows.

12.11 Cleaning up

To remove all compiled files, type make distclean in the top-level Graph-BLAS folder.

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https://redislabs.com/blog/new-redisgraph-1-0-achieves-600x-faster-performance-graph-databases/).

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14 Additional Resources

See http://graphblas.org for the GraphBLAS community page. See https://github.com/GraphBLAS/GraphBLAS-Pointers for an up-to-date list of additional resources on GraphBLAS, maintained by Gábor Szárnyas.

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Provides a basic overview of many sparse matrix algorithms and a simple sparse matrix data structure. The sparse data structure used in the book is much like the one in both MATLAB and SuiteSparse:GraphBLAS. A series of 42 lectures are available on YouTube; see the link at http://faculty.cse.tamu.edu/davis/publications.html For the book, see https://dx.doi.org/10.1137/1.9780898718881

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Abstract: Wilkinson defined a sparse matrix as one with enough zeros that it pays to take advantage of them. This informal yet practical definition captures the essence of the goal of direct methods for solving sparse matrix problems. They exploit the sparsity of a matrix to solve problems economically: much faster and using far less memory than if all the entries of a matrix were stored and took part in explicit computations. These methods

form the backbone of a wide range of problems in computational science. A glimpse of the breadth of applications relying on sparse solvers can be seen in the origins of matrices in published matrix benchmark collections (Duff and Reid 1979a, Duff, Grimes and Lewis 1989a, Davis and Hu 2011). The goal of this survey article is to impart a working knowledge of the underlying theory and practice of sparse direct methods for solving linear systems and least-squares problems, and to provide an overview of the algorithms, data structures, and software available to solve these problems, so that the reader can both understand the methods and know how best to use them. DOI: https://dx.doi.org/10.1017/S0962492916000076

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From the preface: Graphs are among the most important abstract data types in computer science, and the algorithms that operate on them are critical to modern life. Graphs have been shown to be powerful tools for modeling complex problems because of their simplicity and generality. Graph algorithms are one of the pillars of mathematics, informing research in such diverse areas as combinatorial optimization, complexity theory, and topology. Algorithms on graphs are applied in many ways in today's worldfrom Web rankings to metabolic networks, from finite element meshes to semantic graphs. The current exponential growth in graph data has forced a shift to parallel computing for executing graph algorithms. Implementing parallel graph algorithms and achieving good parallel performance have proven difficult. This book addresses these challenges by exploiting the well-known duality between a canonical representation of graphs as abstract collections of vertices and edges and a sparse adjacency matrix representation. This linear algebraic approach is widely accessible to

scientists and engineers who may not be formally trained in computer science. The authors show how to leverage existing parallel matrix computation techniques and the large amount of software infrastructure that exists for these computations to implement efficient and scalable parallel graph algorithms. The benefits of this approach are reduced algorithmic complexity, ease of implementation, and improved performance. DOI: https://dx.doi.org/10.1137/1.9780898719918

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Triangle counting serves as a key building block for a set of important graph algorithms in network science. In this paper, we address the IEEE HPEC Static Graph Challenge problem of triangle counting, focusing on obtaining the best parallel performance on a single multicore node. Our implementation uses a linear algebra-based approach to triangle counting that has grown out of work related to our miniTri data analytics miniapplication and our efforts to pose graph algorithms in the language of linear algebra. We leverage KokkosKernels to implement this approach efficiently on multicore architectures. Our performance results are competitive with the fastest known graph traversal-based approaches and are significantly faster than the Graph Challenge reference implementations, up to 670,000 times faster than the C++ reference and 10,000 times faster than the Python reference on a single Intel Haswell node.

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