

LAGraph: Linear Algebra, Network Analysis Libraries, and the Study of Graph Algorithms

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Abstract—Graphs and graph algorithms can be expressed in terms of linear algebra. The GraphBLAS are a library of low level building blocks for such algorithms. The GraphBLAS target algorithm *developers*. The LAGraph project targets graph algorithm *users* with high-level algorithms common in network analysis. In this paper, we describe the first release of the LAGraph library. We describe the design decisions behind the library, the contents of the library, and performance data using the GAP benchmark suite. LAGraph, however, is much more than a library development project. It is also a project to document and analyze the full range of algorithms enabled by the GraphBLAS. To that end we have developed a compact (and hopefully intuitive) notation for describing these algorithms. In this paper, we present that notation with examples from the GAP benchmark suite.

Index Terms—Graph Processing, Graph Algorithms, Graph Analytics, Linear Algebra, GraphBLAS

I. INTRODUCTION

Graphs represent networks of relationships. They play a key role in a wide range of applications. Consequently, numerous libraries of graph algorithms exist such as igraph [9], NetworkX [3], and SNAP [17]. These libraries let programmers work with graphs without the need to master the art of crafting graph algorithms.

There are multiple ways to build libraries of graph algorithms. One approach views graphs in terms of sparse matrices and graph algorithms in terms of linear algebra. This perspective led to the GraphBLAS [18], [20]; a community effort [1] to define low level building blocks for graph algorithms as linear algebra. The GraphBLAS, however, are for graph algorithm *developers*. They are too low level for graph algorithm *users*. To focus on users and the algorithms they require, researchers from the graphBLAS community launched the LAGraph project [19].

A major goal for the LAGraph project is to produce a library of high quality, production-worthy algorithms constructed on top the GraphBLAS library. In this paper, we describe the first release of the LAGraph library [2]. While LAGraph will eventually work with any implementation of the GraphBLAS, it is currently tied to the SuiteSparse GraphBLAS library [10].

In this first release of LAGraph, the range of algorithms is narrow (the algorithms found in the GAP benchmark suite).

With so few algorithms, we were able to focus our efforts on the key design decisions needed to establish a solid foundation we can build on for the future. Those design decisions and the rationale behind them are a key contribution of this paper. We also establish a performance baseline for our ongoing work with LAGraph with results for the GAP benchmark suite [6].

The LAGraph project is much more than a library project. Another goal for the LAGraph project is to create a repository of algorithms based on the GraphBLAS and study them in order to advance the state of the art in Graph algorithms expressed as Linear algebra. To support this goal, we created a concise notation for expressing graph algorithms in terms of the GraphBLAS. As an example of the use of this notation, we use it to describe the algorithms used in the GAP benchmark suite. We view this notation as a key contribution of this paper.

II. DESIGN DECISIONS

TODO: Jim

We investigate the following design questions:

- data structure for representing a graph
- basic/advanced mode
- algorithm calling conventions
- error handling
- user-contributed algorithms
- ———
- how to work around typecasting being expensive in GraphBLAS
- terminology: properties (parameters? features?)
- multi-threadable

A. Core data structure

The primary data structure in LAGraph is the `LAGraph_Graph` which consists of primary components and cached properties. The data structure is not opaque, providing the user with full access to access and modify all internal components, unlike the way GraphBLAS objects behave. This data structure is shown at the top of Listing 1 and defined ultimately on Line 14.

```

1 typedef struct LAGraph_Graph_struct
2 {
3     GrB_Matrix A;      // adjacency matrix of the graph
4     LAGraph_Kind kind; // kind of graph: directed, etc.
5
6     // cached properties
7     GrB_Matrix AT;     // transpose of A
8     GrB_Vector row_degree;
9     GrB_Vector col_degree;
10    LAGraph_BooleanProperty A_pattern_is_symmetric;
11    int64_t ndiag; // -1 if unknown
12 } *LAGraph_Graph;
13
14 typedef struct LAGraph_Graph_struct *LAGraph_Graph;
15
16 // creating a graph
17 GrB_Matrix M;
18 // ...construction of M omitted
19
20 LAGraph_Graph G;
21 LAGraph_New(&G, &M, LAGRAPH_DIRECTED_ADJACENCY);
22
23 // operating on properties
24 LAGraph_Property_AT(G, msg); // compute/cache

```

Listing 1: LAGraph_Graph data structure and methods.

The primary components of this struct are a GraphBLAS matrix named *A* and an enumeration *kind*. The *kind* indicates how the matrix should be interpreted. Currently, the only kinds defined are *LAGRAPH_ADJACENCY_UNDIRECTED* and *LAGRAPH_ADJACENCY_DIRECTED*, but more options will be added in the future. Creating the Graph object is performed on Line 21 of Listing 1. Following this call, *M* will be *NULL*. The matrix previously pointed to by *M* now lives at *G->A*. This “move” constructor helps avoid memory freeing errors.

Cached properties include the transpose of *A*, the row degrees, column degrees, etc. They can be computed from the primary components, but doing so repeatedly for each algorithm utilizing *A* would be wasteful. Having them live inside the Graph object helps simplify algorithm call signatures. Utility functions exist to compute each cached property. For example, Line 24 of Listing 1 will compute the transpose of *G->A* and store it as *G->AT*. Following this call, any algorithm which is given *G* will have access to both *A* and its transpose.

Because the Graph object is opaque, any piece of code may set the transpose as well. For instance, if an algorithm computes the transpose as part of its normal logic, it could directly set *G->AT*. The expectation is that the Graph object will always remain consistent. If *G->A* is modified, all cached properties must be either be set as unknown or modified to reflect the change. Properties which are not known are set to *NULL* or *LAGRAPH_BOOLEAN_UNKNOWN* in the case of Boolean properties. This expectation is a convention that all LAGraph algorithm implementers are expected to follow.

B. User modes

Algorithms in LAGraph target two types of user modes: Basic and Advanced. The Basic user mode is for those wanting things to “just work”, are less concerned about performance, and may be less experienced with the library. The Advanced

user mode is for those whose primary concern is performance and are willing to conform to stricter requirements to achieve that goal.

Algorithms targeting the Basic mode typically have limited options. Often, there will only be one function for a given algorithm. Under the hood, that single algorithm might take different paths depending on the shape or size of the input graph. The idea is that a basic user wants to compute PageRank or Betweenness Centrality, but doesn’t want to have to understand the five different ways to compute them. They simply want the correct answer.

Algorithms targeting the Advanced mode are often highly specialized implementations of an algorithm. The Advanced mode user is expected to understand details such as push-pull and batch mode and why different techniques are better for each graph. Advanced mode algorithms are very strict in their input. If the input doesn’t match the expected kind, an error will be raised.

Advanced mode algorithms will also raise an error if a cached property is needed by an algorithm, but is not currently available on the Graph object. While Basic mode algorithms are free to compute and cache properties on the Graph object, Advanced mode algorithms never will. The idea is to never surprise the user with unexpected additional computation. An Advanced mode user must opt-in to all computations.

Often, Basic mode algorithms will inspect the input, possibly compute properties or transform the data, and finally call one of the Advanced mode algorithms to do the actual work on the graph. Having these two user modes allows LAGraph to target a wider range of users who vary in their experience with graph algorithms.

C. Algorithm calling conventions

Algorithms in LAGraph follow a general calling convention.

```

1 int algorithm
2 (
3     // outputs:
4     TYPE *out1,
5     TYPE *out2,
6     ...
7     // input/output
8     TYPE inout,
9     ...
10    // inputs
11    TYPE input1,
12    TYPE input2,
13    ...
14    // error message holder
15    char *msg
16 )

```

The return value is always an *int* with the following meaning:

- *=0* -> success
- *<0* -> error
- *>0* -> warning

The meaning of a given error or warning value is algorithm-specific and should be listed in the documentation for the algorithm.

Outputs appear first and are passed by reference. A pointer should be created by the caller, but memory will be allocated by the algorithm. If the output is not needed, a NULL is passed and the algorithm will not return that output.

Input/Output arguments are passed by value. The expectation is that the object will be modified. This supports things like batch mode in which a frontier is updated and returned to the caller. It also supports Basic mode algorithms which may modify a Graph object by adding cached properties.

Inputs are passed by value and should never be changed by the algorithm.

The final argument of any LAGraph algorithm is the error message holder. This must be `char[]` of size `LAGRAPH_MSG_LEN`. When the algorithm returns an error or a warning, a message may be placed in this array as additional information. Because the caller creates this array, the caller must free the memory or reuse it as appropriate. If the algorithm is successful, it should fill the message array with an empty string to clear any previous message.

D. Error handling

Because every algorithm in LAGraph can return an error, the return value of every call should be checked before proceeding. To make this less burdensome for a C-based library, LAGraph provides a convenience macro which works similar to `try/catch` in other languages.

```

1 #define LAGraph_TRY(LAGraph_method)
2 {
3     int LAGraph_status = LAGraph_method;
4     if (LAGraph_status < 0)
5     {
6         LAGraph_CATCH (LAGraph_status);
7     }
8 }
```

`LAGraph_CATCH` can be defined before an algorithm and will be called in the event of an error. This allows for proper freeing of memory and other necessary tasks.

A similar macro, `GrB_TRY`, will call `GrB_CATCH` when making GraphBLAS calls which return a `GrB_Info` value other than `GrB_SUCCESS` or `GrB_NO_VALUE`.

`LAGraph_TRY` and `GrB_TRY` provide an easy to use and easy to read method for dealing with error checking while writing graph algorithms.

E. Contributing algorithms

The LAGraph project welcomes contributions from all graph practitioners who understand the GraphBLAS vision of using the language of linear algebra to express graph computations. However, as a matter of practical concern, many users want a stable experience when using LAGraph for doing real work. To balance these, the LAGraph repository will have both a stable and an experimental folder.

New algorithms or modifications of existing algorithms will first be added to the experimental folder. The release schedule of experimental algorithms will generally be much faster than the stable release and there is no expectation of a bug-free

experience. The goal is to generate lots of ideas and allow uninhibited contributions to push the boundary of what is possible with the GraphBLAS. The stable release will be fully tested and will move much slower, targeting the needs of those who want to use LAGraph as a complete, production-grade library rather than as a research project.

III. GRAPHBLAS THEORY AND NOTATION

In this section, we summarize the key concepts in GraphBLAS. Then, we present a concise notation for the operations and methods defined in the GraphBLAS standard. Additionally, we demonstrate how the operations can be interpreted as graph processing primitives if vectors encode nodes in the and the matrices are adjacency matrices. **TODO: adjacency matrices can be rectangular** Note that the standard does not have such restrictions, hence, users of the API are free to define other kinds of vectors and matrices (such as incidence matrices) as well.

A. Overview

Here, we give a brief overview of the theoretical aspects of the GraphBLAS. For more details, we refer the reader to tutorials [22] and the specification documents [7], [12].

a) *Data structures*: GraphBLAS builds on the duality between the graph and matrix data structures. Namely, a directed graph $G = (V, E)$ can be represented with an adjacency matrix $A \in \mathbb{B}^{|V| \times |V|}$ where $A_{i,j} = \text{TRUE}$ iff $(v_i, v_j) \in E$. Induced subgraphs where source nodes are selected from V_1 and target nodes are selected from V_2 ($V_1, V_2 \subseteq V$) can be represented with $A \in \mathbb{B}^{|V_1| \times |V_2|}$. Vectors are used to encode data for nodes, e.g., $u \in \mathbb{B}^{|V|}$ can be used to select a subset of nodes, while $u \in \mathbb{B}^{|V|}$. Vectors and matrices can be defined over multiple types, e.g., a nonnegative integer (\mathbb{N}) matrix can encode the number of paths between two nodes, while a floating point (FP64) matrix can encode edge weights.

b) *Semirings*: GraphBLAS uses matrix operations to express graph processing primitives, e.g., a matrix-vector multiplication $A \oplus \otimes u$ finds incoming neighbors of a given set of nodes. GraphBLAS allows users to use arbitrary semirings for multiplication operations. In general, the multiplication operator \otimes is used to define how to combine the values of matching input elements, while the addition operator \oplus defines how the results should be summarized. For example, the min.plus semiring uses \oplus as the multiplication operator to compute the path length and min as the addition operator to determine the length of the shortest path.

c) *Masks and accumulators*: All GraphBLAS operations whose output is a vector or a matrix, allow the use of masks to limit the scope of the computation. This implies that the computation should be performed on a given set of nodes (for vector masks) or a given set of edges (for matrix masks). The accumulator operator \odot is a binary operator that determines how the result of an operation should be applied to its output. The interplay of masks and the accumulators is discussed in the specifications [7], [12].

| operation/method | description | notation |
|------------------------|--|--|
| mxm | matrix-matrix multiplication | $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A} \oplus \cdot \otimes \mathbf{B}$ |
| vxm | vector-matrix multiplication | $\mathbf{w}^\top \langle \mathbf{m}^\top \rangle \odot = \mathbf{u}^\top \oplus \cdot \otimes \mathbf{A}$ |
| mxv | matrix-vector multiplication | $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{A} \oplus \cdot \otimes \mathbf{u}$ |
| eWiseAdd | element-wise addition using operator op on elements in the set union of structures of \mathbf{A}/\mathbf{B} and \mathbf{u}/\mathbf{v} | $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A} \text{ op}_\cup \mathbf{B}$ $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{u} \text{ op}_\cup \mathbf{v}$ |
| eWiseMult | element-wise multiplication using operator op on elements in the set intersection of structures of \mathbf{A}/\mathbf{B} and \mathbf{u}/\mathbf{v} | $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A} \text{ op}_\cap \mathbf{B}$ $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{u} \text{ op}_\cap \mathbf{v}$ |
| extract | extract submatrix from matrix \mathbf{A} using indices i and indices j extract the j th column vector from matrix \mathbf{A} extract subvector from \mathbf{u} using indices i | $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A}(i, j)$ $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{A}(:, j)$ $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{u}(i)$ |
| assign | assign matrix to submatrix with mask for \mathbf{C} assign scalar to submatrix with mask for \mathbf{C} assign vector to subvector with mask for \mathbf{w} assign scalar to subvector with mask for \mathbf{w} | $\mathbf{C}\langle\mathbf{M}\rangle(i, j) \odot = \mathbf{A}$ $\mathbf{C}\langle\mathbf{M}\rangle(i, j) \odot = s$ $\mathbf{w}\langle\mathbf{m}\rangle(i) \odot = \mathbf{u}$ $\mathbf{w}\langle\mathbf{m}\rangle(i) \odot = s$ |
| subassign (GxB) | assign matrix to submatrix with submask for $\mathbf{C}(i, j)$ assign scalar to submatrix with submask for $\mathbf{C}(i, j)$ assign vector to subvector with submask for $\mathbf{w}(i)$ assign scalar to subvector with submask for $\mathbf{w}(i)$ | $\mathbf{C}(i, j)\langle\mathbf{M}\rangle \odot = \mathbf{A}$ $\mathbf{C}(i, j)\langle\mathbf{M}\rangle \odot = s$ $\mathbf{w}(i)\langle\mathbf{m}\rangle \odot = \mathbf{u}$ $\mathbf{w}(i)\langle\mathbf{m}\rangle \odot = s$ |
| apply | apply unary operator f with optional thunk k | $\mathbf{C}\langle\mathbf{M}\rangle \odot = f(\mathbf{A}, k)$ $\mathbf{w}\langle\mathbf{m}\rangle \odot = f(\mathbf{u}, k)$ |
| select | apply select operator f with optional thunk k | $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A}\langle f(\mathbf{A}, k) \rangle$ $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{u}\langle f(\mathbf{u}, k) \rangle$ |
| reduce | row-wise reduce matrix to column vector reduce matrix to scalar reduce vector to scalar | $\mathbf{w}\langle\mathbf{m}\rangle \odot = [\oplus_j \mathbf{A}(:, j)]$ $s \odot = [\oplus_{i,j} \mathbf{A}(i, j)]$ $s \odot = [\oplus_i \mathbf{u}(i)]$ |
| transpose | transpose | $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A}^\top$ |
| kroncker | Kronecker multiplication using operator op | $\mathbf{C}\langle\mathbf{M}\rangle \odot = \text{kron}(\mathbf{A}, \text{op}, \mathbf{B})$ |
| dup | duplicate matrix duplicate vector | $\mathbf{C} \leftarrow \mathbf{A}$ $\mathbf{w} \leftarrow \mathbf{u}$ |
| build | matrix from tuples vector from tuples | $\mathbf{C} \leftarrow \{i, j, x\}$ $\mathbf{w} \leftarrow \{i, x\}$ |
| extractTuples | extract index arrays (i, j) and value arrays (x) | $\{i, j, x\} \leftarrow \mathbf{A}$ $\{i, x\} \leftarrow \mathbf{u}$ |
| extractElement | extract element to scalar | $s = \mathbf{A}(i, j)$ $s = \mathbf{u}(i)$ |
| setElement | set element | $\mathbf{C}(i, j) = s$ $\mathbf{w}(i) = s$ |

TABLE I: GraphBLAS operations and methods based on [11]. *Notation:* Matrices and vectors are typeset in bold, starting with uppercase (\mathbf{A}) and lowercase (\mathbf{u}) letters, respectively. Scalars including indices are lowercase italic (k, i, j) while arrays are lowercase bold italic ($\mathbf{x}, \mathbf{i}, \mathbf{j}$). \oplus and \otimes are the addition and multiplication operators forming a semiring and default to conventional arithmetic $+$ and \times operators. \odot is the accumulator operator. Operations can be modified via a descriptor; matrices can be transposed (\mathbf{B}^\top), the mask can be complemented, and the mask can be valued (shown above) or structural ($\mathbf{C}\langle s(\mathbf{M}) \rangle$). **TODO: consider using $\mathbf{C}\langle \text{val}(\mathbf{M}) \rangle$ for valued** A structural mask can also be complemented ($\mathbf{C}\langle \neg s(\mathbf{M}) \rangle$). The result can be cleared after using it as input to the mask/accumulator step ($\mathbf{C}\langle \mathbf{M}, r \rangle$). Not all methods are listed (creating new ops, monoids, and semirings, clearing a matrix/vector, etc.). For the **assign** operation, the dimensions of the masks are the same as the output's: $\dim(\mathbf{M}) = \dim(\mathbf{C})$ and $\dim(\mathbf{m}) = \dim(\mathbf{w})$. For the **subassign** operation, the dimensions of the masks are the same as the submasks: $\text{nrows}(\mathbf{M}) = |\mathbf{i}|$, $\text{ncols}(\mathbf{M}) = |\mathbf{j}|$ for matrices and $\dim(\mathbf{m}) = |\mathbf{i}|$ for vectors.

d) *Notation*: To present our algorithms, we use the notation shown in Table I. Matrices and vectors are typeset in bold, starting with uppercase (**A**) and lowercase (**u**) letters, respectively. Scalars including indices are lowercase italic (*k*, *i*, *j*) while arrays are lowercase bold italic (***x***, ***i***, ***j***).

B. Operations

a) *Matrix multiplication*: The *matrix-matrix multiplication* operation $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A} \oplus \otimes \mathbf{B}$ expresses navigation step starting from edges of **A** and uses the edges of **B**. The result matrix **C** contains paths with $\mathbf{C}_{i,k}$ containing the summarized paths with start node *i* in the graph of **A** and end node *k* in the graph of **B**. The *vector-matrix multiplication* operation $\mathbf{w}^T\langle\mathbf{m}^T\rangle \odot = \mathbf{u}^T \oplus \otimes \mathbf{A}$ performs navigation starting from the nodes selected in vector **u** among the edges encoded in matrix **A**, while the *matrix-vector multiplication* operation $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{A} \oplus \otimes \mathbf{u}$ performs navigation in the reverse direction of the edges encoded in **A**. The result vector **w** contains the set of reached nodes with the values computed on the semiring (combining the initial node values with the outgoing edge values using \otimes and summarizing the results in the same end node using \oplus).

b) *Element-wise addition*: The *element-wise addition* operation applies the operator op on the elements selected by the *union of the structures of its inputs*, i.e., nodes/edges which are present in at least one of the inputs.

c) *Element-wise multiplication*: The *element-wise multiplication* operation applies the operator op on the elements selected by the *intersection of the structures of its inputs*, i.e., nodes/edges which are present in both inputs. The notation is the same as that of the element-wise addition operation except using op_\cap .

d) *Extract*: For adjacency matrix **A**, the *extract submatrix* operation $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A}(\mathbf{i}, \mathbf{j})$ selects the induced subgraph where the source nodes of the edges are in array **i** and the target nodes of the edges are in array **j**. The *extract vector* operation $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{A}(\mathbf{i}, :)$ selects a column vector containing node *i*'s neighbors along incoming edges. Finally, the *extract subvector* operation $\mathbf{w}\langle\mathbf{m}\rangle \odot = \mathbf{u}(\mathbf{i})$ selects the nodes with indices in array **i**.

e) *Assign*: **TODO: edge values, vertex values, constant value**

f) *Apply*: **TODO: apply**

g) *Select*: **TODO: selectively keep edges/nodes based on condition, including the indices of ops**

h) *Reduce*: For adjacency matrix **A**, the *row-wise reduction* $\mathbf{w}\langle\mathbf{m}\rangle \odot = [\oplus_j \mathbf{A}(:, j)]$ represents a summarization of the values on outgoing edges for each node (represented by row vector $\mathbf{A}(:, j)$) to vector **w** with an optional mask $\langle\mathbf{m}\rangle$ to limit the computation to certain nodes. For matrix **A**, the *reduction to scalar* $s \odot = [\oplus_{i,j} \mathbf{A}(i, j)]$ represents a summarization of all edge values. For vector **u**, the *reduction to scalar* $s \odot = [\oplus_i \mathbf{u}(i)]$ represents a summarization of all node values.

i) *Transposition*: The transposed variant of matrix **A** is denoted with \mathbf{A}^T where $\mathbf{A}^T(i, j) = \mathbf{A}(j, i)$. For adjacency matrix **A**, matrix \mathbf{A}^T contains the edges in the reverse direction.

Transposition can be applied as a standalone GraphBLAS operation $\mathbf{C}\langle\mathbf{M}\rangle \odot = \mathbf{A}^T$ with an optional mask $\langle\mathbf{M}\rangle$ and also to the input/output matrices of operations, for example:

$$\mathbf{C}^{[T]}\langle\mathbf{M}\rangle \odot = \mathbf{A}^{[T]} \oplus \otimes \mathbf{B}^{[T]}$$

C. Masks

TODO: explain masks

TODO: explain replace/merge as per Scott's email

TODO: revise Scott, Tim D, and Tim M

TODO: consider adding semiring table

what do masks mean?

the mask restricts the scope of the computation to the elements selected by the mask (**m**) or the complement of these elements ($\neg\mathbf{m}$)

there are variations based on (1) how the elements are selected (*valued* masks take the element values into consideration, while *structural* masks only use the structure (pattern) of the mask (2) how the elements outside the selected ones are treated (they are *replaced* with implicit zeros or they are kept intact and *merged* with the results of the computation)

by default, masks should use *merge* semantics, i.e. the computation can only effect elements selected by the mask, elements outside the mask are unaffected $\mathbf{w}\langle\mathbf{m}\rangle$ and $\mathbf{w}\langle\neg\mathbf{m}\rangle$

replace semantics, i.e. they annihilate all elements outside the mask. this is denoted with $\mathbf{w}\langle\mathbf{m}, r\rangle$ and $\mathbf{w}\langle\neg\mathbf{m}, r\rangle$

by default, element values in the mask are checked and elements with zero value are not used to avoid this, we use *structural masks* $\mathbf{w}\langle s(\mathbf{m})\rangle$ and $\mathbf{w}\langle\neg s(\mathbf{m})\rangle$

Combining *replace semantics* and *structural masks* is possible: $\mathbf{w}\langle s(\mathbf{m}), r\rangle$ $\mathbf{w}\langle\neg s(\mathbf{m}), r\rangle$

Initializing scalars, vectors, and matrices (GraphBLAS methods):

- let: $s \in \mathbb{Q}_{64}$
- let: $\mathbf{w} \in \mathbb{Q}_{32}^n$
- let: $\mathbf{A} \in \mathbb{N}_{16}^{m \times n}$
- let: $\mathbf{A} \in \mathbb{Z}_{64}^{k \times m}$

separate operator + descriptor (T)

vectors can be interpreted as row/column vectors, we do not transpose them manually
extensions

D. Methods

TODO: conversion table?

IV. ALGORITHMS

A. Breadth-First Search (BFS)

The breadth-first search (BFS) builds on the observation that a vector-matrix multiplication $\mathbf{f}^T \mathbf{A}$ expresses the navigation from the nodes selected by vector **f** in the graph represented by **A**. The direction-optimizing push/pull BFS [5] is simple to express in GraphBLAS [24]. If **A** is held by row, then $\mathbf{f}^T \mathbf{A}$ is a push step, while $\mathbf{B} \mathbf{f}$ is a pull step, where $\mathbf{B} = \mathbf{A}^T$ is the explicit transpose of **A**, also held by row. Other GraphBLAS libraries, e.g., GraphBLAST, store both directions and perform

direction-optimization automatically [25]. The push-only BFS is shown in Alg. 1, while the push/pull BFS is Alg. 2.

The GraphBLAS BFS relies on the any.secondi semiring to compute a single step, $\mathbf{q}\langle\neg s(\mathbf{p})\rangle = \mathbf{q}^\top \mathbf{A}$, where \mathbf{q} is the current frontier (using \mathbf{q} as short for queue), \mathbf{p} is the parent vector, and \mathbf{A} is the adjacency matrix. This step assigns the parents of newly nodes, which do not yet have a parent, using the complemented structural mask $\langle\neg s(\mathbf{p})\rangle$.

Consider a matrix multiply for conventional linear algebra, where the \oplus monoid sums a set of t entries to obtain a single scalar for computing $c_{ij} = \sum a_{ik}b_{kj}$ in the matrix multiply $\mathbf{C} = \mathbf{AB}$. The any monoid performs the reduction of t entries to a single number by merely selecting any one of the t entries as the result c_{ij} . The selection is done non-deterministically, allowing for a benign race condition. In the BFS, this corresponds to selecting any valid parent of a newly discovered node. Indeed, the creation of the any operator was inspired by Scott Beamer’s `bfs.cc` method in the GAP benchmark, which has the same benign race condition. The any monoid translates the concept of this benign race condition to construct a valid BFS tree into a linear algebraic operation, suitable for implementation in GraphBLAS.

The secondi operator is the multiplicative operator in the any.secondi semiring, where the result of $a_{ik}b_{kj}$ is simply the index k in the semiring for $\mathbf{C} = \mathbf{AB}$. This gives the id of the parent node for a newly discovered node in the next frontier. The any monoid then selects any valid parent k .

Algorithm 1: Parents BFS.

Input: \mathbf{A} , $startVertex$

```

1 Function ParentsBFS
2    $\mathbf{p}(startVertex) = startVertex$ 
3    $\mathbf{q}(startVertex) = startVertex$ 
4   for  $level = 1$  to  $nrows(\mathbf{A}) - 1$  do
5      $\mathbf{q}^\top \langle \neg s(\mathbf{p}^\top), r \rangle = \mathbf{q}^\top \text{any.secondi } \mathbf{A}$ 
6      $\mathbf{p}\langle s(\mathbf{q}) \rangle = \mathbf{q}$ 
7     if  $nvals(\mathbf{q}) = 0$  then
8       return
```

B. Betweenness Centrality (BC)

The vertex betweenness-centrality metric is a weighted measure of the number of shorted paths that go through any given node,

$$\sum_{s \neq i \neq t} \frac{\sigma(s, t|i)}{\sigma(s, t)},$$

where $\sigma(s, t)$ is the total number of shortest paths from node s to t , and $\sigma(s, t|i)$ is the total number of shortest paths from node s to t that pass through node i . This is expensive to compute, so in practice, a subset of source nodes are chosen at random (a *batch*), of size ns .

Like the BFS, direction-optimization is incredibly simple to add to the LAGraph algorithm for batched betweenness-centrality (BC). It only requires a simple heuristic to determine

Algorithm 2: Direction Optimizing Parent BFS.

Input: \mathbf{A} , \mathbf{A}^\top , $startVertex$

```

1 Function DirectionOptimizingBFS
2    $\mathbf{q}(startVertex) = 0$ 
3   for  $level = 1$  to  $nrows(\mathbf{A}) - 1$  do
4     if  $Push(\mathbf{A}, \mathbf{q})$  then
5        $\mathbf{q}^\top \langle \neg s(\mathbf{p}^\top), r \rangle = \mathbf{q}^\top \text{any.secondi } \mathbf{A}$ 
6     else
7        $\mathbf{q} \langle \neg s(\mathbf{p}), r \rangle = \mathbf{A}^\top \text{any.secondi } \mathbf{q}$ 
8      $\mathbf{p}\langle s(\mathbf{q}) \rangle = \mathbf{q}$ 
9     if  $nvals(\mathbf{q}) = 0$  then
10      return
```

Algorithm 3: Betweenness centrality.

```

1 Function BrandesBC
2   //  $\mathbf{P}(k, j) = \#$  paths from  $k$ th source to node  $j$ 
3    $\mathbf{P} \leftarrow \{s, [1, 1, \dots, 1]\}$ 
4   //  $\mathbf{F}$ : # paths in the current frontier
5    $\mathbf{F} \leftarrow \{s, [1, 1, \dots, 1]\}$ 
6    $\mathbf{F} \langle \neg s(\mathbf{P}), r \rangle = \mathbf{F} \text{ plus.first } \mathbf{A}$ 
7   // BFS phase
8   for  $d = 0$  to  $nrows(\mathbf{A})$  do
9     let:  $\mathbf{S}[d] \in \mathbb{B}^{ns \times n}$ 
10     $\mathbf{S}[d] \langle s(\mathbf{F}), r \rangle = 1$  //  $\mathbf{S}[d] = \text{pattern of } \mathbf{F}$ 
11     $\mathbf{P} += \mathbf{F}$ 
12     $\mathbf{F} \langle \neg s(\mathbf{P}), r \rangle = \mathbf{F} \text{ plus.first } \mathbf{A}$ 
13    if  $nvals(\mathbf{F}) = 0$  then
14      break
15  // Backtrack phase
16  let:  $\mathbf{B} \in \mathbb{Q}_{64}^{ns \times n}$ 
17   $\mathbf{B}(\cdot) = 1.0$ 
18  let:  $\mathbf{W} \in \mathbb{Q}_{64}^{ns \times n}$ 
19  for  $i = d - 1$  downto  $0$  do
20     $\mathbf{W} \langle s(\mathbf{S}[i]), r \rangle = \mathbf{B} \text{div}_\cap \mathbf{P}$ 
21     $\mathbf{W} \langle s(\mathbf{S}[i - 1]), r \rangle = \mathbf{W} \text{ plus.first } \mathbf{A}^\top$ 
22     $\mathbf{B} += \mathbf{W} \times_\cap \mathbf{P}$ 
23  //  $\text{centrality}(j) = \sum_i (\mathbf{B}(i, j) - 1)$ 
24   $\text{centrality}(\cdot) = -ns$ 
25   $\text{centrality} += [+_i \mathbf{B}(i, :)]$ 
```

which direction to use, followed by masked matrix-matrix multiplication with the matrix or its transpose: $\mathbf{F} \langle \neg s(\mathbf{P}) \rangle = \mathbf{FB}^\top$ (the pull) or $\mathbf{F} \langle \neg s(\mathbf{P}) \rangle = \mathbf{FA}$ (the push), where \mathbf{A} is the adjacency matrix of the graph and $\mathbf{B} = \mathbf{A}^\top$ is its explicit transpose, \mathbf{F} is the frontier, and the complemented structural mask $\neg s(\mathbf{P})$ is the set of unvisited nodes. The multiplication \mathbf{FB}^\top relies on the descriptor to represent the transpose of \mathbf{B} , which is not explicitly transposed. In the backward phase, the pull step is $\mathbf{W} = \mathbf{WA}^\top$ while the push is $\mathbf{W} = \mathbf{WB}$, where \mathbf{W} is the ns -by- n matrix in which centrality is accumulated (where $ns = 4$ is a typical batch size).

To simplify the presentation of the entire BC algorithm, Alg. 3 does not show the direction-optimization. It is the same transformation as the pair of BFS algorithms, where the push-only step (line 5 of Alg. 1), is expanded to a push/pull heuristic (lines 4-7 of Alg. 2).

C. PageRank (PR)

PageRank (PR) computes the importance of each node as a recursively-defined metric: a web page is important if important pages link to it. Alg. 4 shows the GraphBLAS implementation of PR as specified in the GAP Benchmark. It uses the plus.second semiring, where $\text{second}(x, y) = y$, so it can ignore any edge weights in the input matrix. The PR in GAP does not properly handle dangling vertices in the graph. The Graphalytics benchmark has a PageRank variant which avoids this problem [14]. We have included this version to compare its performance with the GAP benchmark algorithm `pr.cc`.

Algorithm 4: PageRank (as specified in the GAPBS).

Data: $\mathbf{A} \in \mathbb{B}^{n \times n}$ // adjacency matrix
 $damping$ // damping factor
 tol // stopping tolerance
 $itermax$ // maximum number of iterations

Result: $\mathbf{r} \in \mathbb{Q}^n$

```

1 Function PageRank
2    $teleport = \frac{1-\alpha}{n}$ 
3    $\mathbf{r}(0 : n-1) = \frac{1}{n}, \mathbf{t} = \mathbb{Q}^n$ 
4    $\mathbf{d}_{out} = [+_j \mathbf{A}(:, j)]$  // precomputed rowdegree
5    $\mathbf{d} = \mathbf{d}_{out} \text{div}_{\cap} damping$  // prescale with damping
6   for  $k = 1$  to  $numIterations$  do
7     swap  $\mathbf{t}$  and  $\mathbf{r}$  //  $\mathbf{t}$  is now the prior rank
8      $\mathbf{w} = \mathbf{t} \text{div}_{\cap} \mathbf{d}$ 
9      $\mathbf{r}(0 : n-1) = teleport$ 
10     $\mathbf{r} += \mathbf{A}^T \text{plus.second } \mathbf{w}$ 
11     $\mathbf{t} -= \mathbf{r}$ 
12     $\mathbf{t} = abs(\mathbf{t})$ 
13    if  $[+_j \mathbf{t}(:)] < tol$  then
14      return
```

D. Single-Source Shortest-Paths (SSSP)

A Delta-Stepping Single-Source-Shortest-Path algorithm in GraphBLAS is shown in Alg. 5. It relies on the min.plus semiring. Since it is a fairly complex algorithm, refer to [21] for a description of the method.

E. Triangle Counting (TC)

The triangle counting (TC) problem is to compute the number of unique cliques of size 3 in a graph. The TC algorithm is shown in Alg. 6, based on [23]. It starts with a heuristic that decides when to sort the input graph by ascending degree. Next, it constructs the lower and upper triangular part and computes a masked matrix multiply using the plus.pair semiring. Internally, a dot product method is used in SS:GrB,

Algorithm 5: SSSP (delta-stepping).

Data:
 $\mathbf{A}, \mathbf{A}_H, \mathbf{A}_L \in \mathbb{Q}^{|V| \times |V|}$
 $s, i \in \mathbb{N}$
 $\Delta \in \mathbb{Q}$
 $\mathbf{t}, \mathbf{t}_{Req} \in \mathbb{Q}^{|V|}$
 $\mathbf{t}_{B_i}, \mathbf{e} \in \mathbb{N}^{|V|}$

```

1 Function DeltaStepping
2    $\mathbf{A}_L = \mathbf{A} \langle 0 < \mathbf{A} \leq \Delta \rangle$ 
3    $\mathbf{A}_H = \mathbf{A} \langle \Delta < \mathbf{A} \rangle$ 
4    $\mathbf{t}(:) = \infty$ 
5    $\mathbf{t}(s) = 0$ 
6   while  $nvals(\mathbf{t} \langle i\Delta \leq \mathbf{t} \rangle) \neq 0$  do
7      $s = 0$ 
8      $\mathbf{t}_{B_i} = \mathbf{t} \langle i\Delta \leq \mathbf{t} < (i+1)\Delta \rangle$ 
9     while  $\mathbf{t}_{B_i} \neq 0$  do
10       $\mathbf{t}_{Req} = \mathbf{t} \times_{\cap} \mathbf{t}_{B_i}$ 
11       $\mathbf{t}_{Req} = \mathbf{A}_L^T \text{min.plus } \mathbf{t}_{Req}$ 
12       $\mathbf{e} = \mathbf{t} \langle 0 < \mathbf{e} \oplus \mathbf{t}_{B_i} \rangle$ 
13       $\mathbf{t}_{B_i} = \mathbf{t} \langle i\Delta \leq \mathbf{t}_{Req} < (i+1)\Delta \rangle$ 
14       $\mathbf{t}_{B_i} = \mathbf{t}_{B_i} \langle \mathbf{t}_{Req} < \mathbf{t} \rangle$ 
15       $\mathbf{t} = \mathbf{t} \text{min}_{\cup} \mathbf{t}_{Req}$ 
16       $\mathbf{t}_{Req} = \mathbf{A}_H^T \text{min.plus } (\mathbf{t} \times_{\cap} \mathbf{e})$ 
17       $\mathbf{t} = \mathbf{t} \text{min}_{\cup} \mathbf{t}_{Req}$ 
18       $i = i + 1$ 
```

Algorithm 6: Triangle counting

Data: $\mathbf{A} \in \mathbb{B}^{n \times n}$
Result: $t \in \mathbb{N}$

```

1 Function TriangleCount
2   sample the mean and median degree of  $\mathbf{A}$ 
3   if  $mean > 4 \times median$  then
4      $\mathbf{p}$  = permutation to sort degree, ascending order
5      $\mathbf{A} = \mathbf{A}(\mathbf{p}, \mathbf{p})$ 
6    $\mathbf{L} = \text{tril}(\mathbf{A})$ 
7    $\mathbf{U} = \text{triu}(\mathbf{A})$ 
8    $\mathbf{C} \langle s(\mathbf{L}) \rangle = \mathbf{L} \text{plus.pair } \mathbf{U}^T$ 
9    $t = [+_{ij} \mathbf{C}(i, j)]$ 
```

because \mathbf{U} is transposed via the descriptor. The pair is the simple function $\text{pair}(x, y) = 1$. When used in a semiring, it acts like the times operator of the conventional semiring, except that it can ignore the values of its inputs and treat them both as 1. This semiring is useful for structural computations, such as triangle counting, when the edge weights of a graph may be present but should be ignored in a particular algorithm.

F. Connected Components

The connected components algorithm in LAGraph (Alg. 7) is written by Zhang, Azad, and Buluç [26], [27]. The method maintains a forest of trees represented by a parent vector, and

iteratively merges trees until no more merging is possible. The method as shown in Alg. 7 is a simplified variant that it operates on the entire graph. In the LAGraph version, a subgraph is constructed first, and the method finds the connected components of the subgraph, and then operates on the entire graph.

Algorithm 7: Connected components (FastSV).

```

1 Function FastSV
2    $n = \text{nrows}(\mathbf{A})$ 
3    $\mathbf{gf} = \mathbf{f}$ 
4    $\mathbf{dup} = \mathbf{gf}$ 
5    $\mathbf{mngf} = \mathbf{gf}$ 
6    $\{i, x\} \leftarrow \mathbf{f}$ 
7   repeat
8     // Step 1: Stochastic hooking
9      $\mathbf{mngf} = \mathbf{mngf} \min \mathbf{A}$ 
10     $\mathbf{mngf} = \mathbf{mngf} \text{ second.min } \mathbf{gf}$ 
11     $\mathbf{f}(x) = \mathbf{f} \min \mathbf{mngf}$ 
12    // Step 2: Aggressive hooking
13     $\mathbf{f} = \mathbf{f} \min \mathbf{mngf}$ 
14    // Step 3: Shortcutting
15     $\mathbf{f} = \mathbf{f} \min \mathbf{gf}$ 
16    // Step 4: Calculate grandparents
17     $\{i, x\} \leftarrow \mathbf{f}$ 
18     $\mathbf{gf} = \mathbf{f}(x)$ 
19    // Step 5: Check termination
20     $\mathbf{diff} = \mathbf{dup} \neq \mathbf{gf}$ 
21     $\mathbf{sum} = [\oplus_i \mathbf{diff}(i)]$ 
22     $\mathbf{dup} = \mathbf{gf}$ 
23  until  $\mathbf{sum} == 0$ 

```

V. UTILITY FUCTIONS

LAGraph includes a set of utility functions that operate on a graph. All function names are prefixed with `LAGraph_` so we exclude that prefix in the names below, for brevity.

- **Graph Properties:** An `LAGraph_Graph` includes cached properties which can be assigned by Basic methods, or which are required by Advanced methods. `DeleteProperties` clear all properies, `Property_AT` computes the transpose of the adjacency matrix $G \rightarrow A$, `Property_RowDegree` computes the row degrees of $G \rightarrow A$, `Property_ColDegree` computes the column degrees of $G \rightarrow A$, and `Property_ASymmetricPattern` determines if the pattern of $G \rightarrow A$ is symmetric or unsymmetric.
- **Display and debug:** `CheckGraph` checks the validity of a graph. Since the graph is not opaque, a user application is able to change a graph arbitrarily and thus might make it an invalid object. `DisplayGraph` displays a graph and its properties.
- **Memory management:** Wrappers for `malloc`, `calloc`, `realloc`, and `free`, allowing a user application to select the memory manager to be used. These default to the ANSI C11 library functions.

- **Graph I/O:** `BinRead` and `BinWrite` read/write a `GrB_Matrix` in binary form. `MMRead` and `MMWrite` read/write a `GrB_Matrix` in Matrix Market form.
- **Matrix operations:** `Pattern` returns a boolean matrix containing the pattern of a matrix. `IsEqual` determines if two matrices are equal. It selects the appropriate `GrB_EQ_T` operator that matches the matrix type, and then calls `IsAll`. `IsAll` compares two matrices and returns false if the pattern of the two matrices differ. It then uses a given comparator operator to compare all pairs of entries, and returns true if all comparisons return true.
- **Degree operations:** `SortByDegree` returns a permutation that sorts a graph by its row/column degrees, and `SampleDegree` computes a quick estimate of the mean and median row/column degrees.
- **Error handling:** `LAGraph_TRY` and `GrB_TRY` are helper macros for a simple try/catch mechanism. They require the user application to define `LAGraph_CATCH` and `GrB_CATCH`.
- **Other:** `TypeName` returns a string with the name of a `GrB_Type`. `KindName` returns a string with of graph kind (directed or undirected). `Tic` and `Toc` provide a portable timer. `Sort1`, `Sort2`, and `Sort3` sort 1, 2, or 3 integer arrays.

VI. EVALUATION

The performance of LAGraph can only be considered in context of an implementation of the underlying GraphBLAS library. This is discussed in Section VI-A, followed by performance results of the new LAGraph API on the 6 algorithms in the GAP Benchmark [5].

A. SuiteSparse Extensions

In a prior paper ([4]), an early draft of SuiteSparse:GraphBLAS v4.0.0 (Aug 2020) was compared with the GAP benchmark [5] and four other graph libraries. This prior version of SS:GrB included two primary data structures for its sparse matrices: compressed sparse vector, and a hypersparse variant [8], both held by row or by column. It included a draft implementation of a bitmap data structure that could only be used in a prototype breadth-first search. Since then, SuiteSparse:GraphBLAS v4.0.3 has been released, with full support for bitmap and full matrices for all its operations. In an m -by- n bitmap matrix, the values are held in a full array of size mn , and another `int8_t` array of size mn holds the sparsity pattern of the matrix. A full matrix is a simple dense array of size mn .

The bitmap format is particularly important for the “pull” phase of an algorithm, as used in direction-optimizing breadth-first-search [5], [24]. The GAP benchmark suite uses this method by holding its frontier as a bitmap in the pull step and as a list in the push step. The GAP BFS was shown to be typically the fastest BFS amongst the 6 graph libraries compared in [4] (for 4 of the 5 benchmark graphs). With the addition of the bitmap format to SS:GrB, LAGraph+SS:GrB is able to come within a factor of 2 or so of the performance of

the highly-tuned BFS GAP benchmark (see the results in the next section), for those 4 graphs. At the same time, however, the BFS is very easily expressed in LAGraph as an easy-to-read and easy-to-write code. This enables non-experts to obtain a reasonably high level of performance with modest programming effort when writing their own graph algorithms.

Additional optimizations added to SS:GrB in the past year include a *lazy sort*. Normally, SS:GrB keeps its vectors sorted (row vectors in a CSR matrix, or column vectors if the matrix is held by column), with entries sorted in ascending order of column or row index, respectively. This simplifies the many algorithms that operate on a `GrB_Matrix`. However, some algorithms naturally produce a jumbled result (matrix multiply in particular), while many algorithms are tolerant of jumbled input matrices. We thus allow the sort to be left pending. The lazy sort joins two other kinds of pending work in SS:GrB: *pending tuples* and *zombies* [11]. A pending tuple is an entry that is held inside a matrix in an unsorted list, awaiting insertion into the CSR/CSC format of a `GrB_Matrix`. A zombie is the opposite: it is an entry in the CSR/CSC format that has been marked for deletion, but has not yet been deleted from the matrix. With the lazy sort, the sort is postponed until another algorithm requires sorted input matrices. If the sort is lazy enough, it might never occur, which is the case for the LAGraph BFS and BC.

Another useful addition to SS:GrB is the new positional binary operators, such as the `any.secondi` used in the BFS.

B. Performance Results

Our benchmark environment is an NVIDIA DGX Station (donated to Texas A&M by NVIDIA in support of this research). It includes a 20-core Intel(R) Xeon(R) CPU E5-2698 v4 @ 2.20GHz, with 40 threads. All codes were compiled with gcc 5.4.0 (-O3). All default settings were used, which means that hyperthreading was enabled. The system has 256GB of RAM in a single socket (no NUMA effects). LAGraph (Feb 7, 2021) and SuiteSparse:GraphBLAS 4.0.4-draft (Feb 7, 2021) were used. The NVIDIA DGX Station includes four P100 GPUs, but no GPUs were used by this experiment (a GPU-accelerated SS:GrB is in progress). Table II lists the run time (in seconds) for the GAP benchmark and LAGraph+SS:GrB for the 6 algorithms on the 5 benchmark matrices. The benchmark matrices are listed in Table III.

With the simple addition of the bitmap (needed for the pull step), the push/pull optimization in BC resulted in a nearly 2x performance gain in the GraphBLAS method for the largest matrices, as compared to the SS:GrB version used for the results presented in [4].

With this change the BC method in LAGraph+SS:GrB is not only expressible in a simple, elegant code, but it is also faster than the highly-tuned GAP benchmark method, `bc.cc`, for the three largest matrices (1.3x for Kron, 1.5x for Urand, and 1.2x for Twitter).

With the addition of bitmap format (which makes push/pull optimization very simple to express, and very fast) and the `any.secondi` semiring, the BFS of a directed or undirected graph

| Algorithm : package | graph, with run time in seconds | | | | |
|------------------------|---------------------------------|-------|--------|-------|-------|
| BC : GAP | 31.52 | 46.36 | 10.82 | 3.01 | 1.50 |
| BC : SS | 23.61 | 32.69 | 9.25 | 8.20 | 34.40 |
| BFS : GAP | .31 | .58 | .22 | .34 | .25 |
| BFS : SS | .52 | 1.22 | .33 | .66 | 3.32 |
| PR : GAP | 19.81 | 25.29 | 15.16 | 5.13 | 1.01 |
| PR : SS | 22.17 | 27.71 | 17.21 | 9.30 | 1.34 |
| CC : GAP | .53 | 1.66 | .23 | .22 | .05 |
| CC : SS | 3.36 | 4.47 | 1.47 | 1.97 | .98 |
| SSSP : GAP | 4.91 | 7.23 | 2.02 | .81 | .21 |
| SSSP : SS | 17.37 | 25.54 | 8.54 | 9.61 | 46.79 |
| TC : GAP | 374.08 | 21.83 | 79.58 | 22.18 | .03 |
| TC : SS | 917.99 | 34.01 | 239.58 | 34.65 | .23 |

TABLE II: Run time of GAP and LAGraph+SS:GrB

| graph | nodes | entries in A | graph kind |
|---------|-------------|----------------|------------|
| Kron | 134,217,726 | 4,223,264,644 | undirected |
| Urand | 134,217,728 | 4,294,966,740 | undirected |
| Twitter | 61,578,415 | 1,468,364,884 | directed |
| Web | 50,636,151 | 1,930,292,948 | directed |
| Road | 23,947,347 | 57,708,624 | directed |

TABLE III: Benchmark matrices (<https://sparse.tamu.edu/GAP>)

is easily expressed in GraphBLAS, and has a performance that is only about 1.5x to 2x slower than the GAP benchmark. We expect the remaining performance gap arises from two issues:

- 1) The GAP assumes that the graph has fewer than 2^{32} nodes and edges, and thus uses 32-bit integers throughout. GraphBLAS is written for larger problems, and thus relies solely on 64-bit integers. This cannot be easily changed in GraphBLAS, but rather than “fixing” GraphBLAS to use smaller integers, it is the GAP algorithms that would need to be updated for larger graphs in the future. In the current GAP benchmark graphs, two graphs are chosen with almost exactly 4 billion edges. Graphs of current interest in large data science can easily exceed 2^{32} nodes and edges [15].
- 2) In GraphBLAS, the BFS must be expressed as two calls. The first computes $\mathbf{q}(\neg \mathbf{p}) = \mathbf{q}^T \mathbf{A}$, and the second updates the parent vector, $\mathbf{p}(s(\mathbf{q})) = \mathbf{q}$:

```
GrB_vxm (q, p, NULL, semiring, q, A, GrB_DESC_RSC) ;
GrB_assign (p, q, NULL, q, GrB_ALL, n, GrB_DESC_S) ;
```

In Scott Beamer’s `bfs.cc`, these two steps are fused, and the matrix-vector multiplication can write its result directly into the parent vector `p`. This could be implemented in a future GraphBLAS library, since the GraphBLAS API allows for a non-blocking mode where work is queued and done later, thus enabling a fusion of these two steps. SS:GrB exploits the non-blocking mode (for its lazy sort, pending tuples, and zombies) but does not *yet* exploit the fusion of `GrB_vxm` and `GrB_assign`. We intend to exploit this in the future.

Note that for the Road graph, LAGraph+SS:GrB is quite slow for all but PageRank (PR). The primary reason for this is the high diameter of the Road graph (about 6980). This requires 6980 iterations of GraphBLAS in the BFS, each with a tiny amount of work. Each call to GraphBLAS does several

malloc and frees, and in some cases the workspace must be initialized. A future version of SS:GrB is planned that will eliminate this work entirely, by implementing an internal memory pool. There may be other overheads, but we hope that a memory pool, fusion to fully exploit non-blocking mode, and other optimization will address this large performance gap for the Road graph for these algorithms.

LAGraph+SS:GrB is also up to 3x slower than the GAP for the triangle counting problem (for all but the Road graph, where it is even slower). This performance gap can be eliminated entirely in the future, if the GrB_mxm and GrB_reduce are combined in a single fused step, by a full exploitation of the GraphBLAS non-blocking mode. The current method computes $C\langle s(L) \rangle = LU^T$, followed by the reduction of C to a single scalar. The matrix C is then discarded. All that GraphBLAS needs is a fused kernel that does not explicitly instantiate the temporary matrix C . This is permitted by the GraphBLAS C API Specification, but not yet implemented in SS:GrB.

VII. CONCLUSION

Future work - ideas:

- Create a Python wrapper for LAGraph
- Implement the LDBC Graphalytics benchmark [13]
- Improve data ingestion performance using e.g., SIMD instructions [16]

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TODO: add acks

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