LAGraph: A Graph Algorithm and Network Analysis Library for GraphBLAS

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Abstract—design decisions, easy mode/expert mode, GAP algorithms, ...

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

I. INTRODUCTION

LAGraph is a library of Graph Algorithms based on the GraphBLAS

Key contributions:

- document design decisions for LAGraph
- present a concise notation for GraphBLAS algorihms
- algorithms of the GAP benchmark suite [4] used in the IISWC benchmark paper [2]
- improve data ingestion performance, e.g. using SIMD techniques [13]

Recently, numerous graph-specific have targeted GPUs such as Gunrock [18] and GraphBLAST [20], and FPGAs [5].

However, in the near future we expect even more heterogeneous hardware architectures including graph-specific hardware based on the Programmable Integrated Unified Memory Architecture (PIUMA) [1]. Additionally, graph processing workloads can be offloaded to machine learning accelerators, e.g., Tensor Processing Units (TPUs) [11], systolic arrays using reconfigurable dataflow architecture [9], sparse linear algebrabased deep learning accelerators [14].

Previous GraphBLAS design papers: theory [15], C API [17], C++ API [7], distributed API [6], LAGraph [16]

```
int main() {
   return 0; // return zero
}
```

Listing 1: Example



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II. DESIGN DECISIONS

We investigate the following design questions:

- easy/expert mode or standard/advanced more
- multi-threadable
- data structure for representing a graph/matrix
- error handling
- opacity of LAGraph
- how to work around typecasting being expensive in GraphBLAS
- terminology: properties (parameters? features?)

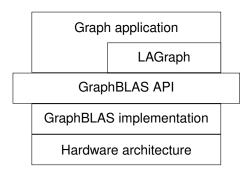


Fig. 1: Separation of concerns using the GraphBLAS API.

III. NOTATION

IV. ALGORITHMS

GAP algorithms: BFS, SSSP, TC, BC, PR. Not sure whether CC should be included.

A. BFS

push/pull [19]

GraphBLAST stores both directions and picks automatically [20]

B. Betweenness Centrality

also does push/pull

- C. PageRank
- D. SSSP
- E. Triangle Count

(Not too interesting from an API design point of view.)

F. Connected Components

Needs a few GxB extensions.

V. EVALUATION



A. SuiteSparse Extensions

In a prior paper ([2]), an early draft of Suite-Sparse:GraphBLAS v4.0.0 (Aug 2020) was compared with the GAP benchmark [3] 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

¹A non peer-reviewed comparison of 6 popular graph algorithms libraries is available at https://www.timlrx.com/blog/benchmark-of-popular-graph-network-packages-v2.

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 [3]. 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 [2] (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.

Direction-optimization is incredibly simple to add to an LA-Graph algorithm. For example, a batched direction-optimizing betweenness-centrality (BC) algorithm in LAGraph only requires a simple heuristic to determine which direction to use, followed by masked matrix-matrix multiplication with the matrix or its transpose: $F\langle \neg P \rangle = FB'$ (the pull) or $F\langle \neg P \rangle = FA$ (the push), where A is the adjacency matrix of the graph and B = A' is its explicit transpose, F is the frontier, and the complemented mask $\neg P$ is the set of unvisited nodes. The multiplication FB' relies on the descriptor to represent the transpose of B, which is not explicitly transposed. In the backward phase, the pull step is W = WA' while the push is W = WB, where W is the 4-by-n matrix in which centrality is accumulated.

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. 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.

Another useful addition to SS:GrB is the new positional binary operators. The BFS relies on the ANY-SECONDI semiring to compute a single step, $q\langle \neg \pi \rangle = q'A$, where q

Algorithm:	gı	aph, with	run time i	in seconds	
package	Kron	Urand	Twitter	Web	Road
BC : GAP	31.52	46.36	10.82	3.01	1.50
BC : SS	26.85	31.78	10.04	9.25	51.91
BFS : GAP	.31	.58	.22	.34	.25
BFS: SS	.52	1.31	.33	.67	3.33
PR : GAP	19.81	25.29	15.16	5.13	1.01
PR:SS	21.96	27.75	17.22	9.30	1.34
CC : GAP	.53	1.66	.23	.22	.05
CC : SS	3.42	4.59	1.48	1.97	1.00
SSSP : GAP	4.91	7.23	2.02	.81	.21
SSSP : SS	17.62	25.62	8.44	9.67	48.49
TC : GAP	374.08	21.83	79.58	22.18	.03
TC:SS	943.47	34.10	242.36	35.15	.29

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

is the current frontier π is the parent vector, and A is the adjacency matrix.

Consider a matrix multiply for conventional linear algebra, where the PLUS 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 C = 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 C=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.

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 (Jan 29, 2021) and SuiteSparse:GraphBLAS 4.0.3 (Jan 19, 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 I 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 II.

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 [2].

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 II: Benchmark matrices (https://sparse.tamu.edu/GAP)

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 (1.2x for Kron, 1.5x for Urand, and 1.08x for Twitter).

We expect the BC in LAGraph+SS:GrB to become faster still in the next release because we have not yet fully exploited the lazy sort. The frontier matrix F is left jumbled by the lazy sort, but it is sorted right away by a subsequent assignment. Most uses of GrB_assign are intolerant of jumbled input matrices, so it sorts them on input. However, GrB_assign includes about 40 internal variations, a few of which do not actually require sorted input matrices. The particular method used in GrB_assign in the LAGraph BC method is one of those methods, so this would be simple to exploit. With this change, the sort would be so lazy that it would *never* occur. The frontier would be computed, left jumbled, used in subsequent computations, and then recomputed (and thus discarded), just as we currently do in the LAGraph BFS.

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

- 1) The GAP assumes that the graph has fewer than 2³² 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 2³² nodes and edges [12].
- 2) In GraphBLAS, the BFS must be expressed as two calls. The first computes $q\langle \neg \pi \rangle = q'A$, and the second updates the parent vector, $\pi\langle q \rangle = q$:

 $\label{eq:grb_vxm} $$ $\operatorname{GrB_vxm} (q, pi, NULL, semiring, q, A, GrB_DESC_RSC) ; $$ $\operatorname{GrB_assign} (pi, 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 vector pi. 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 LAGraph+SS:GrB is quite slow for many algorithms (all but PR) on the Road graph. The primary reason for this is the high diameter of the Road graph (about 6000). This requires 6000 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 $C\langle L\rangle=LU'$, 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.

VI. UTILITY FUCTIONS

sort, diag/invdiag (?), equal

VII. CONCLUSION

Future work - ideas:

- Create a Python wrapper for LAGraph
- Implement the LDBC Graphalytics benchmark [10]
- Improve data ingestion performance using e.g., SIMD instructions [13]

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