# The Knowledge Discovery Toolbox (Version 0.2) User Guide

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# 1 Overview

The Knowledge Discovery Toolbox (KDT) provides subject-matter experts with a simple interface to analyze very large graphs quickly and effectively without requiring knowledge of the underlying graph representation or algorithms. Domain experts are defined as experts in a field of study that is not graphs and graph algorithms, though they may have some familiarity with graph algorithms by using them. Because KDT is open-source, it can be customized or extended by interested (and intrepid) users.

#### Version 0.2

This early release provides a tiny selection of functions on directed graphs ranging from simple exploratory functions to complex algorithms, a set of constructors and conversion methods for hypergraphs, and the initial support for *semantic* graphs; *i.e.*, those with attributes on vertices and/or edges. The current version works on graphs contained in the memory of multiple computers in a cluster (while hiding data representation and partitioning from the user). Notes specific to this version will be denoted by "*Version 0.2*" and appear in blue text.

#### 1.1 Context

While graphs represent many real-world relationships in a mathematically robust way, their analysis with current methods does not scale to large graphs. The modern "data tsunami" has created graphs in critical scientific and societal domains that are large enough to be prohibitively time-consuming to analyze with well-known methods. This has led graph-analysis experts to create more efficient graph analysis algorithms, but has also led to a gap between those experts and the non-graph subject-matter experts who need to use them. KDT counters the trend by exposing an API through the Python language that is efficiently and scalably implemented on computer clusters, while remaining suitable for domain experts by hiding the underlying implementation.

KDT is intended to accelerate a virtuous cycle among (a) subject-matter experts who need to analyze graphs that don't fit in the memory of a single computer node; (b) researchers working on improved graph algorithms; and (c) developers of tool infrastructure. We envision that subject-matter experts will do more analysis of large graphs with the current algorithms in KDT and provide feedback on which algorithms are (not) most useful for the large graphs in their domains. This will spur algorithm researchers and tool developers to develop new variants to analyze the subject-matter experts' graphs better.

We believe that many of the subject-matter experts don't know exactly what analysis they need to perform on their data, so they need to explore different algorithms and analyses. The KDT's goal is for the subject-matter expert (or her graduate student) to be able to compose KDT building blocks at the Python level and explore interactively.

KDT's complex algorithms are difficult for even graph experts to verify, and so whenever possible KDT supports internal or user-driven verification of results. That variously consists of internal checks in KDT routines, companion routines that can validate a data structure (i.e., isBfsTree() can validate the results of bfsTree()), and synthetic inputs whose metrics can be analytically derived (e.g., each

vertex of the graph created by twoDTorus(nnodes) has an identical betweenness centrality value of 0.5 \* 2\*\*(3\*scale\*0.5) - 2\*\*scale + 1 for an even nnodes, where scale is equal to log2(nnodes\*\*2).

# 1.2 Intended use cases

#### Version 0.1

To tie KDT's development to the needs of potential users, this early release targets a small set of use cases. They all assume that the data about graph edges exists in "triplet" format, where the triplet is the source vertex, the destination vertex, and the attribute(s), such as weight and label, of the edge.

1.2.1 Creating a random power-law graph and calculating a breadth-first-search tree Following the <u>Graph500</u> benchmark, this use creates a random power-law graph in memory, calculates a breadth-first-search (BFS) tree from a starting vertex, and verifies the BFS tree.

# 1.2.2 Ranking the importance of Web pages

Analysis of the relative importance of Web pages led to the seminal PageRank algorithm, whose variations are the cornerstone of contemporary search engines.

# 1.2.3 Calculating the centrality of vertices of a graph

Centrality can be a step toward clustering that removes the most central vertices, but it is also an important metric in its own right for understanding which vertices most keep the graph connected. Exact betweenness centrality is simultaneously viewed by some researchers as not robust enough, because it provides information on only the single shortest path between two vertices, and by others as too computationally expensive, because its cost grows on the order of the number of vertices squared times the length of the longest path. For maximum flexibility, KDT provides a range of algorithms for calculating centrality from across the accuracy vs. execution-cost continuum.

#### 1.3 Deferred use cases

#### Version 0.2

KDT is envisioned to grow over time to support capabilities not included in this early release. The list of the deferred features includes: undirected graphs, bipartite graphs, multigraphs, and full support for hypergraphs; clustering; matching or alignment; general-purpose attributes, such as labels on vertices and edges; visualization of resulting graphs; support on other than x86-64 clusters; and a disk-based implementation, for problems that do not fit in the memory of a computational cluster.

#### 1.4 Performance

KDT's goal is to enable quick and effective analysis of very large graphs, and thus performance is a vital aspect of KDT. Its performance has not yet been fully characterized. Performance of specific methods, where available, is provided in their descriptions.

Much of the early performance work with KDT has focused on the Graph500 benchmark, whose units are traversed-edges per second (TEPS). On a 256-core cluster, the KDT implementation of Graph500 performs at over 1 GTEPS. See section 4 for details.

# 1.5 Legend and Naming Convention

In the function interface descriptions, required arguments are shown in black text, optional arguments in square brackets, and (optional) expert arguments are shown in grey.

```
cl = cluster('Markov',G[, nclus=k][, power=r])
```

For example, in the cluster function, 'Markov' and G are required arguments, nclus is an optional argument, and power is an optional expert argument.

Names follow the Python convention of generally using lower case and capitalizing the first letter of a class name (DiGraph, e.g.), sometimes referred to as Pascal case. Multi-word member names follow the so-called camel case, where the first letter is lower case but the first letter of subsequent words is capitalized, such as sendFeedback.

# 1.6 Sending feedback to the KDT developers

KDT includes <code>sendFeedback</code>, a built-in feedback method that enables users to type in code that they wish KDT could execute and then send it to the developers. It uses IPython's <code>%logstart</code> facility to capture the code snippet. In Python code interpreters other than IPython the feedback mechanism will not work but will not otherwise obstruct program execution.

For sites that cannot directly send email onto the Internet, the default email address (in feedback.py, variable name \_kdt\_Alias) can be changed to an internal collection point.

The feedback mechanism can be used as follows:

In the course of solving your problem, when you need a function not implemented by KDT, type the code that you want KDT to support (which will evoke an error) and then invoke its sendFeedback method. It will capture the most recent lines you've typed, create a file from them, and give you an opportunity to edit the file and add supporting comments. If you have feedback that is not directly related to a desired method, that can also be edited into the file. It will then prompt you for confirmation to send the file to the the KDT developers.

With a legend of user input / system responses / user annotation, an IPython session might look like:

```
In [43]: G = kdt.Graph500Edges(scale=32)
In [44]: bc = G.centrality('approxBC',sample=0.01)
In [45]: # delete the top 10 most central vertices; topK and
In [46]: # deleteverts() method doesn't exist
In [47]: discG = G.deleteverts(bc.topK(10))
In [48]: kdt.sendFeedback()
The code example you want to send to the KDT developers is
in /home/sam/graphdev/KDT_email.
If you wish, edit it with your favorite editor. Type 'Send'
when you are ready to send it or 'Cancel' to cancel sending it.
```

```
>>>
```

[... If desired, edit the file with an editor ...]

Send In [49]:

# 2 Graph500 example

The Graph500 benchmark has replaced SSCA #2 [SSCA] as the primary benchmark for a segment of the graph-analysis research community. This new benchmark is "needed in order to guide the design of hardware architectures and software systems intended to support such applications and to help procurements. Graph algorithms are a core part of many analytics workloads." The specification currently describes two kernels that, respectively, create the graph from the input edge tuples and perform a breadth-first search of the graph from a start vertex. (Future kernels are expected to calculate single-source shortest path and the maximal independent set.) This section illustrates the implementation of kernel 2 of the benchmark with KDT.

This section assumes that Python, IPython, and KDT have already been installed in their default locations and that the environment variable KDTINSTALL has been set to the location of the unzipped/untarred package (see section 5.3.3 for details). You can type the following to start a serial Python session to run the Graph500 script and understand how it works. For instructions on how to run KDT and Graph500 in parallel with Python, see section 5.3.3.

```
[sam@neumann ~]$ which ipython
/usr/bin/python
[sam@neumann ~] $ ipython -debug $KDTINSTALL/examples/Graph500.py
Activating auto-logging. Current session state plus future input
saved.
               : .KDT_log #This output is from the startup of the IPython logstart
Filename
                            # mechanism used for the sendFeedback function
Mode
               : over
Output logging : False
                           # from Section 1.6
Raw input log : False
                                   w #
Timestamping
               : False
                            #
                                   w //
State
               : active
Generating a Graph500 RMAT graph with 2^15 vertices...
Duplicates removed (or summed): 78842 and self-loops removed: 0
Generation took 0.328309s.
iteration 1: start=28166, BFS took 0.027983s, verification took
2.482376s and succeeded, TEPS=31,832,419
iteration 2: start=27666, BFS took 0.028234s, verification took
2.441357s and succeeded, TEPS=31,549,367
iteration 3: start=27587, BFS took 0.028846s, verification took
2.421630s and succeeded, TEPS=30,879,990
iteration 4: start=2117, BFS took 0.029042s, verification took
2.409176s and succeeded, TEPS=30,671,607
     [...]
```

```
iteration 61: start=18643, BFS took 0.028643s, verification took
2.409366s and succeeded, TEPS=31,098,729
iteration 62: start=26284, BFS took 0.029493s, verification took
2.411485s and succeeded, TEPS=30,202,494
iteration 63: start=7687, BFS took 0.030046s, verification took
2.421176s and succeeded, TEPS=29,646,722
iteration 64: start=30285, BFS took 0.029399s, verification took
2.412161s and succeeded, TEPS=30,299,244
Graph500 benchmark run for scale = 15
Kernel 1 time = 0.3283 seconds
Kernel 2 BFS execution times
            min time: 2.79829502105712891e-02
  firstquartile_time: 2.86501049995422363e-02
         median_time: 2.90445089340209961e-02
  thirdquartile time: 2.93260216712951660e-02
           max time: 3.00459861755371094e-02
           mean time: 2.89925746619701385e-02
         stddev_time: 4.45555267017642357e-04
Kernel 2 number of edges traversed
            min_nedge: 8.90765000000000000e+05
  firstquartile_nedge: 8.90765000000000000e+05
         median_nedge: 8.90765000000000000e+05
  thirdquartile nedge: 8.90765000000000000e+05
            max_nedge: 8.90765000000000000e+05
           mean_nedge: 8.90765000000000000e+05
         stddev nedge: 0.0000000000000000e+00
Kernel 2 TEPS
            min TEPS: 2.96467220212343894e+07
  firstquartile_TEPS: 3.03745608150749356e+07
         median_TEPS: 3.06689642658791840e+07
  thirdquartile_TEPS: 3.10911601670094803e+07
            max_TEPS: 3.18324191444078088e+07
 harmonic mean TEPS: 3.07239012190395705e+07
harmonic_stddev_TEPS: 1.69827840156330167e+04
Python 2.4.3 (#1, Nov 11 2010, 13:30:19)
Type "copyright", "credits" or "license" for more information.
IPython 0.8.4 -- An enhanced Interactive Python.
         -> Introduction and overview of IPython's features.
%quickref -> Quick reference.
         -> Python's own help system.
object? -> Details about 'object'. ?object also works, ?? prints
more.
In [5]: parents
Out[2]: Limiting print-out to first 100 elements
Elements stored on proc 0: {25802,1,1,19693,32686,624,21394,27838,
28156, -1, 14783, 8652, -1, -1, 26729, -1, 30564, 29617, -1, -1, 19294, -1,
6541,26552,23513,31138,2797,2567,30313,32502,-1,-1,8402,-1,32686,-1,
```

```
21311,30589,-1,16940,23819,-1,28403,32686,21906,25699,32645,30794,\\16244,28393,30280,8146,16578,23171,23683,31583,26762,29294,-1,32010,\\32686,27230,30280,-1,-1,28311,27501,-1,30589,30280,32736,-1,31834,\\12570,31009,14818,28403,30280,5489,31834,27838,-1,-1,31280,-1,-1,\\31009,9680,23819,21028,28403,16244,-1,-1,30279,30280,32100,32153,\\21906,32700,\}
```

Looking at the code in \$KDTINSTALL/examples/Graph500.py, we can see how the Graph500 specification is implemented with KDT methods.

The following code selects vertices, each to be used as the root of a BFS tree, as specified by the benchmark. KDT-specific functions are shown in **blue**; overloaded Python operators are not highlighted.

The code segment below repeatedly calls the KDT bfsTree method, which creates a BFS tree for the graph and given starting vertex, and then validates the BFS tree via the user-written k2Validate function. Each element of the parents vector points to its (unique) parent in the tree.

```
for start in starts:
    start = int(start)
    before = time.time()

# the actual BFS
    parents = G.bfsTree(start, sym=True)

itertime = time.time() - before
    nedges = len((parents[origI] != -1).find())

K2elapsed.append(itertime)
    K2edges.append(nedges)
```

```
K2TEPS.append(nedges/itertime)

i += 1
  verifyInitTime = time.time()
  verifyResult = "succeeded"
  if not k2Validate(G, start, parents):
       verifyResult = "FAILED"
  verifyTime = time.time() - verifyInitTime

if kdt.master():
       print "iteration %d: start=%d, BFS took %fs, verification took %fs and %s, TEPS=%s"%(i, start, (itertime), verifyTime, verifyResult, splitthousands(nedges/itertime))
```

The k2Validate user method calls the KDT isBfsTree method, which further illustrates the use of KDT with edge and vertex vectors. For instance, one section of isBfsTree code validates that each edge's endpoints are one level apart in the BFS tree. It achieves this by building a new DiGraph instance containing just the edges in the tree, validating in the process that no tree edge has the root as its destination. (Note that since isBfsTree is within the DiGraph class itself, which inherits the Graph class that contains the ParVec class, the code does not refer to the DiGraph module explicitly. Note further that this code can be viewed in its installed position (usually in /usr/lib64/python2.4/site-packages/kdt/DiGraph.py) and also in \$KDTINSTALL/kdt/DiGraph.py).

These very brief examples illustrate key points of KDT. First, the operations are graph operations, performed on graphs and (distributed edge- and vertex-) vectors. Second, to the extent practical, graph objects are accessible via standard Python methods such as subscripting, comparisons, and utility functions such as len.

# 3 Algorithms, Methods and Classes

#### **Version 0.1W**

The v0.1W release of KDT supports only directed graphs with single edges between any two vertices and hypergraphs with undirected edges incident upon multiple vertices, via the DiGraph and HyGraph classes respectively.

Collections of vertices and edges are represented as ParVec class instances. See section 3.3.9.6 for details about the class and method structure.

To save repetition, for the remainder of this section we assume that KDT has been imported as import kdt

and that the DiGraph or HyGraph instance being operated on has the variable name G.

#### 3.1 Execution Model

KDT supports parallel execution via the single-program-multiple-data (SPMD) model, meaning that every core runs its own copy of the Python/KDT program and communicates (within KDT) with the other cores as appropriate. The constraint this places on the KDT user is that the arguments passed to KDT functions must be the same on all cores. Most times this is trivial, as when an argument is the name of a variable. One place where this may require thought is in the use of random numbers, where having each core calculate its own random numbers locally may result in different sets of random numbers on each core. The ParVec.randPerm function is useful for ensuring random numbers are the same on all cores; the examples/Graph500.py file and DiGraph:\_approxBC method have examples of its use.

The algorithms and methods below by default execute across all vertices and edges in a graph, but can be constrained to execute only on vertices and/or edges whose attributes meet user-specified criteria; sometimes known as semantic graph processing. See section 3.3.9 for details.

# 3.2 Algorithms

The KDT DiGraph class includes the algorithms in this section.

# 3.2.1 Centrality

Centrality is the degree to which, by some measure, a vertex is *central* to a graph. There is a wide variety of measures used and means of calculating those measures and hence numerous centrality algorithms.

# **Syntax**

```
c = G.centrality('<algorithm>'[, <algorithm-specific keyword arguments>])
```

# **Description**

The centrality function takes as input a DiGraph object and an algorithm identifier (see below) and returns a ParVec (of length equal to the number of vertices in the graph) that contains, for each respective vertex of the graph, the vertex's centrality value. Optional algorithm-specific keyword arguments may also be specified as described by the algorithm-specific sections below.

### 3.2.1.1 Exact betweenness centrality algorithm

Betweenness centrality is the degree to which a vertex is *between* all other vertices in the graph, calculated as the fraction of shortest paths between two vertices that pass through the given vertex [Freeman 1977].

# **Syntax**

```
cl = G.centrality('exactBC'[, normalize=True])
```

# **Description**

The exactBC algorithm calculates exact betweenness centrality on the graph. The optional algorithm-specific normalize argument, which defaults to True, causes the function to normalize the betweenness values by dividing each value by (#vertices-1)\*(#vertices-2).

#### **Performance**

Exact betweenness centrality can be prohibitively expensive for large graphs, while approximate betweenness centrality is less computationally expensive.

#### Version 0.1

The performance of exact and approximate BC have not been fully characterized, but the following fragmentary results may be useful. The input graph was generated by twoDTorus, whose diameter grows as the diameter of the torus. All times are in seconds.

<pre>N = # graph vertices [twoDTorus(n)]</pre>	N = 4096 (n = 64)			N = 16384 (n = 128)			N = 65536 (n = 256)
#cores	1	4	16	1	4	16	16
exactBC	229	75	33	5649	1970	1163	35010
approxBC (sample=0.05)	155	68		260		330	2717

# 3.2.1.2 Approximate betweenness centrality algorithm

This algorithm [Bader] approximates betweenness centrality for each vertex by using a sample of vertices between which to calculate the fraction of shortest paths, rather than using all vertices as in exact betweenness centrality.

#### **Syntax**

```
cl = G.centrality('approxBC'[, normalize=True, sample=])
```

#### **Description**

The approxBC algorithm performs approximate betweenness centrality on the graph. The optional algorithm-specific normalize argument, which defaults to True, causes the function to normalize the betweenness values by scaling each value by the inverse of the sample factor (# total vertices / (#vertices actually calculated) and the inverse of (#vertices-1)\*(#vertices-2). The optional sample expert argument is a floating-point value between 0 and 1 that denotes the fraction of vertices whose connecting paths are calculated in the approximation; a value of 1.0 equates to exact betweenness centrality; the default value is 0.05.

#### **Performance**

See v0.1 timing results above in section 3.2.1.1. The numerical values of 'approxBC', sample=0.05 compared to 'exactBC' are as follows for a graph generated by twoDTorus(256), which has equal centrality for all vertices. Exact result: 0.001938. Approximate result: mean 0.001944, standard deviation 0.000784.

### 3.2.1 PageRank

The PageRank algorithm [Page] computes a Web page's importance based on the number and importance of links to it.

# **Syntax**

```
c = G.pageRank([epsilon = 0.1][, dampingFactor = 0.85])
```

# **Description**

The pageRank function takes as input a DiGraph object and optional arguments and returns a ParVec (of length equal to the number of vertices in the graph) that contains, for each respective vertex of the graph, the vertex's rank. The optional argument epsilon controls the stopping condition; iteration stops when the 1-norm of the difference in two successive result vectors is less than epsilon; the default is 0.1. The optional argument dampingFactor alters the results and speed of convergence, denoting the probability that the random surfer hops to an adjacent vertex (rather than hopping to a random vertex in the graph); the default value is 0.85.

The PageRank algorithm computes the percentage of time that a "random surfer" spends at each vertex in the graph. If the random surfer is at vertex V, she will take one of two actions:

- She will hop to another vertex to which vertex V has an outward edge (self loops are ignored).
- She will randomly hop to any vertex in the graph. This action is taken with probability (1 dampingFactor)

#### **3.2.2 Search**

Several graph algorithms search for connectedness within the graph, such as trees, connected components, independent sets, paths, etc.

#### 3.2.2.1 Breadth-first Search Tree

The bfsTree method creates a breadth-first search tree of a DiGraph instance from a starting vertex.

#### **Syntax**

```
parents = G.bfsTree(start[, sym=False])
```

#### Description

The bfsTree method takes as input a DiGraph instance and the index of the vertex from which to start the search. It returns a ParVec (of length equal to the number of vertices in the graph) that denotes, for each respective vertex of the graph, the vertex's parent in the BFS tree. The start vertex's parent is itself. A vertex that is unreachable from the start vertex has a parent of -1. The optional expert argument sym denotes whether the graph is a symmetric graph, which if True enables the operation to run faster.

#### **Performance**

The performance of the Graph500 benchmark is dominated by the performance of the bfsTree method. The fragmentary Graph500 performance available to date is described in section 1.4.

# 3.3 Methods

The KDT DiGraph and Vec classes implement the methods in this section.

# 3.3.1 Semantic graph overview: operations and filters

The descriptions of algorithms and methods so far have used graphs that are homogeneous in their vertices and edges. Real-world data often includes vertices and edges of different types, so-called *semantic* graphs. For instance, in a social network, people communicate via cell-phone calls, text and email messages, and face-to-face encounters. Each of these different modes of communication could have its own type of edge in a KDT graph, and the methods can be customized to consider only certain type(s) of vertices or edges during a calculation. Because Vec instances are used for vertex information, similar filter operations are usable on Vec instances as for DiGraph instances, with the same data and execution models and customizability.

# 3.3.1.1 Data model

To balance the needs for customizability and performance for semantic graphs, KDT supports two built-in object (structure) types, making it possible to use one object-type for edges and the other vertices (though there's nothing about the object-types that requires that). The details of the built-in objects are defined in kdt/ObjMethods.py. To read graph data in from files, you may need to customize this file to describe your data precisely.

### 3.3.1.2 Execution model

The execution model for semantic <code>DiGraph</code> and <code>Vec</code> instances is that they are the same as non-semantic <code>DiGraphs</code>, though they have some extra information about edges and vertices, respectively. To use this extra information, users create <code>filters</code> that describe which vertices/edges should be used in a particular operation. A filter is a function that looks at the contents of the vertex (edge) object and returns a True if the vertex (edge) is to be used and False otherwise. When a filter is added to a graph or vertex-vector it will effect the vertices/edges that are used for all operations until the filter is removed. Multiple filters can be added to a graph; they can be viewed as being run in series within each operation, so that only vertices/edges that pass (return True from) all filters will be used in the operation. With this execution model, the KDT team expects that materializing the filtered data (and consuming the potentially large amount of memory of the graph or vector copy) can usually be avoided.

#### 3.3.1.3 Elemental operations

KDT's two built-in elemental object types (Objl and Obj2) provide a few fields that can be used as needed for your particular graph data. Because of this flexibility, the KDT developers cannot know what it means to (e.g.) add two edge- or vertex-objects in your particular context. The built-in object types come with a sample implementation of the basic operations, defined in the file kdt/ObjMethods.py. The approach taken is that edge-/vertex-objects will often have a numerical value (that can be added, ORed, etc., in the normal way) and one or more attribute values (that would not make sense to add or OR in the normal way). The sample operations typically perform the corresponding scalar operation on the weight element of the edge-/vector-object and either ignore or just copy the other elements, but you can customize this behavior for your specific needs.

#### 3.3.2 Built-in methods for Vec vectors

#### Version 0.2

The Vec class represents vertex and directed-edge vectors. Its implementation for distributed memory includes a significant but not complete set of methods, which are mapped onto standard Python operators or methods consistent with clear understandability.

#### **Syntax**

```
# v, v2, and v3 are Vec instances
# k is an integer scalar, m is a scalar (integer or floating-point)
v2 = v > k
                      # v2 is a Boolean vector, where each element
v2 = v >= k
                      # is the comparison of the corresponding
                      # element of v with the scalar k
v2 = v < k
v2 = v \le k
v2 = v == k
v2 = v != k
v2 = v > v3
                      # v2 is a Boolean vector, where each element
v2 = v >= v3
                      # is the comparison of the corresponding
                      # elements of v and v3
v2 = v < v3
v2 = v <= v3
v2 = v == v3
v2 = v != v3
                      # bitwise inversion
v2 = \sim v
                                 And
v3 = v \& v2
v3 = v \mid v2
                                Or
                      #
v3 = v ^ v2
                                 Xor
v3 = v + k
v3 = k + v
v3 = v + v2
v += k
v += v2
v3 = -v
```

```
v3 = v - k
v3 = k - v
v3 = v - v2
v -= k
v -= v2
v3 = v * k
v3 = k * v
v3 = v * v2
v *= v2
v3 = v / k
v3 = k / v
v3 = v / v2
v3 = v % k
v3 = v % v2
v[k] = m
m = v[k]
v[vec] = m
                      # vec an integer vector
v2 = v[vec]
del v[k]
del v[vec]
v.abs()
v2 = abs(v)
print v
v.printAll()
```

# **Description**

These methods have their standard definitions in Python with the following exceptions:

- Only a key set of indexing ("[]") modes are supported. The right-hand side modes all return a Vec instance with length equal to the number of requested elements, unless otherwise noted.
  - o Indexing on the right-hand side of an equation by a scalar, returning a scalar.
  - o Indexing on the right-hand side of an equation by a Boolean dense Vec instance.
  - o Indexing on the right-hand side of an equation by a non-Boolean dense Vec instance.
  - o Indexing on the right-hand side of an equation by a Boolean sparse Vec instance.
  - o Indexing on the right-hand side of an equation by a non-Boolean sparse Vec instance, returning a sparse Vec instance.

- Indexing on the left-hand side by a scalar index with a scalar value, changing a single value.
- o Indexing on the left-hand side by a Boolean dense Vec index with a scalar value, changing the indicated values of the base dense Vec instance to the value of the scalar.
- o Indexing on the left-hand side by a Boolean dense Vec index with a dense Vec value, changing the values of the base dense Vec instance corresponding to the True values of the dense Vec index to the corresponding values of the dense Vec value.
- o Indexing on the left-hand side by a non-Boolean sparse Vec index vector with a scalar value.
- o Indexing on the left-hand side by a Boolean sparse Vec index with a sparse Vec value, changing the indicated values of the base sparse Vec instance to the corresponding values from the value vector.
- The modulus ("%") function implements the behavior of Python's modulus function for negative numbers, where the sign of the result is the same as the sign of the divisor (e.g., "-1 % 8" yields "7"). This is different from the behavior of Python's math.fmod or C's fmod.
- The standard print statement will only print the first 100 (\_REPR\_MAX) elements of a Vec instance. Printing all the elements can be achieved via printAll.

# 3.3.3 Non-built-in methods for Vec vectors

The Vec class implements several other methods, some of which are common Python names and some which are not.

#### **Syntax**

```
# v and v2 are Vec instances, m a scalar
v = Vec(length=0, element=0.0, sparse=True)
v = kdt.Vec.ones(size, element=1.0, sparse=False)
v = kdt.Vec.zeros(size, element=0.0, sparse=False)
v = kdt.Vec.range([start,] stop, element=0, sparse=False)
v2 = v.copy(element=None) # deep copy
                           # convert a Vec to a sparse Vec
vSparse = v.sparse()
vDense = v.dense()
                           # convert a Vec to a dense Vec
boolScalar = v.isDense()
boolScalar = v.isSparse()
boolScalar = v.isObj()
boolScalar = v.isBool()
v.toBool()
v2 = v.abs()
```

```
v2 = v.ceil()
v2 = v.floor()
v2 = v.round()
v2 = v.sign()
boolResult = v.any() # boolResult is a Boolean scalar
boolResult = v.all()
boolResult = v.allCloseToInt()
v2 = v.logicalNot() # also accessible via not keyword
v3 = v.logicalAnd(v2) # element-wise logical And
v3 = v.logicalOr(v2)
v3 = v.logicalXor(v2)
k = v.len()
k = v.nn()
                     # number of nulls
k = v.nnn()
                    # number of non-nulls
                     # number of nonzero nonnulls
k = v.nnz()
v2 = v.find(pred=None)
v2 = v.findInds(pred=None)
m = v.max(initInf=None)
m = v.min(initNegInf=None)
m = v.sum()
m = v.mean()
m = v.std()
m = v.norm(ord=k) # k-norm, where k > 0
v.sort()
[sorted, perm] = v.sorted()
v2 = v.topK(k)
v2 = v.hist()
v.randPerm() # only for dense
v = kdt.load(fName, element=0.0, sparse=True)
v.save(fName)
print v
v.printAll()
                     # applies the Python function op to every nonnull
v.apply(op)
v.applyInd(op)
m = v.reduce(op, uniOp=None, init=None)
```

```
k = v.count(op, pred=None)
v.set(value)
# functions that work only for sparse Vec instances
v.spOnes
v.spRange()
```

#### **Description**

These methods have their standard Python or SciPy definitions, with the following exceptions and clarifications.

The Vec constructor creates a Vec instance of the length, element-type, and sparsity (density) specified. Note that length denotes the maximum number of positions in the vector; for a sparse vector, all of these positions are initially null. The ones and zeros methods creates a Vec instance of the length specified with all elements set to 1 and 0, respectively. The range method creates a Vec instance of the length specified with each element set to its index in the vector, with the optional start argument as in standard Python.

Assignment ("=") of a Vec to another variable name does not create a copy of the object, following Python usage for complex objects. The copy method can be used if a copy is needed. The element argument, if specified, can coerce a Vec instance whose element is an Objl or Objl instance to a Vec instance whose element is a 64-bit element. The sparse method converts a dense Vec to a sparse Vec, or returns the input if already sparse. Similarly, the dense method converts a sparse Vec to a dense Vec, or returns the input if already dense. The iDense and isSparse methods return True if the input vector is dense or sparse, respectively. The isObj method returns True if the elemental value of the Vec instance is either an Objl or Objl instance and False if it's the default 64-bit element. The isBool method returns True if all elements of the vector are True (1) or False (0). The toBool method converts each nonnull element of the vector in place to False (0) if its value is zero and True (1) otherwise.

The allCloseToInt method returns a Boolean scalar denoting whether all the elements of the vector are within machine precision of an integer value.

The logicalNot method returns the logical negation of each (nonnull) element of the input vector. The logicalAnd, logicalOr, and logicalXor methods return the element-wise logical And, Or, and Xor, respectively, of the two input vectors. By contrast, for input values 0x55 and 0xAA, the built-in bitwise & operator will return 0x00 (no corresonding bit is set in both values), while the logicalAnd method will return True (at least one bit is set in both values).

The nn and nnn methods return the number of nulls and nonnulls, respectively, with nnn behaving the same as SciPy's getnnz. The nnz method returns the number of nonnulls that are not equal to zero. The findInds method returns the indices of nonnull elements of the input Vec instance, the same as NumPy/SciPy's nonzero method. The distinct names were chosen to provide future flexibility for a sparse Vec class with a configurable null (zero) element.

The max and min methods allow the specification of what the initial value is for positive and negative infinity, respectively. The mean and std methods calculate the mean and standard deviation, respectively, of the (nonnull values of the) input vector. The norm method (modeled on the NumPy method) calculates the k-norm of the vector, where k is greater than zero.

The sort method sorts the elements of the input vector, in-place, in ascending order. The sorted method returns a tuple whose first element is a copy of the input vector, sorted in ascending order, and whose second element is the indices of the sorted elements in the original vector. If the input vector is sparse, the output vectors will be as well. The topk method returns the k largest values of the input vector, sorted in ascending order, in a Vec instance of length k.

The hist method returns the histogram of the input in a dense Vec instance of length equal to the maximum value in the vector rounded to an integer.

The randPerm method randomly permutes the elements of the ParVec instance.

The load and save methods load and save, respectively, a Vec instance from (to) a file of the given name. The file must be in the Coordinate Format of the Matrix Market Exchange Formats (see also Section 3.6.1). By default, the Vec instance will be created with an elemental 64-bit value; this can be overridden with the element argument specifying either an Objl or Obj2 instance. Also by default the Vec instance will be created sparse; this can be overridden with the sparse argument. [[FIX: how to specify Obj values per-line in the file?]]

The apply, applyInd, reduce, and count methods are highly configurable via their op (and uniOp) arguments, which are Python operations that define the action to be taken for each element of the Vec instance. See Section xxx [[FIX: insert cross-reference]] for more details. The apply method applies the op function to each (nonnull, in the case of a sparse vector) element of the Vec instance, changing its value in place. The applyInd method applies the op function to the index of each (nonnull, in the case of a sparse vector) element of the Vec instance, changing its value in place. The reduce method applies the uniOp function to each (nonnull, in the case of a sparse vector) element of the Vec instance and then applies the binary op function to that result and the current reduction value; the result of the binary function will be a single element (which can be a 64-bit element or an Obj1 or Obj2 instance). The count method applies the pred function to each (nonnull, in the case of a sparse vector) element of the Vec instance; if the Boolean return value is True, it will then apply the binary op function to that result and the current reduction value; the result of the binary function will be a single element (which can be a 64-bit element or an Obj1 or Obj2 instance).

The set method sets every (nonnull, in the case of a sparse vector) element of the Vec instance to value, which must be of the same type (64-bit or Obj1 or Obj2 instance) as the element of the Vec instance.

The spOnes method modifies the bound sparse Vec instance by setting all its nonnull elements to 1.0. The set method modifies the bound Vec instance by setting all its nonnull elements to the passed value. The spRange method modifies the bound sparse Vec instance by setting each of its nonnull elements to its index in the vector.

#### 3.3.4 Built-in methods for DiGraph objects

The DiGraph class represents directed graphs. Its implementation for distributed memory includes a small set of methods, some of which are mapped onto standard Python operators or methods consistent with clear understandability.

#### Version 0.1

DiGraph instances support a 64-bit floating-point or integer elemental datatype and a Boolean datatype, with an explicit conversion required to the Boolean form.

#### **Syntax**

```
# G1, G2 and G3 are DiGraphs
G3 = G1 + G2  # elemental addition
G3 += G2  # elemental addition in place
G3 = -G1  # elemental negation
G3 = G1 * G2  # elemental multiplication
G3 *= G2  # elemental multiplication in place
G3 = G1 / G2  # elemental division
G3 /= G2  # elemental division in place
```

# **Description**

These methods have their obvious definitions, with the caveat that the multiplication and division operators yield a null (zero) value in any element for which at least one of the graphs has a null entry. Because DiGraph are often large objects, in-place operators may often make sense to conserve the amount of memory consumed by temporary or transient objects.

The toBool method converts each nonnull element of a DiGraph instance to a Boolean True, thereby consuming less space and enabling faster operations.

The print method for DiGraph objects can be problematic, as they can often be extremely large (billions of elements), for which text display is rarely useful. For v0.1, for small graphs (defined as 100 edges or fewer), the print method will call the toParVec method and print those results. For larger graphs, you can accomplish this yourself by invoking the toParVec method manually and printing those results via ParVec.printAll.

#### 3.3.5 Non-built-in methods for DiGraph objects

In addition to built-in methods, a number of other utility methods are implemented for the DiGraph class.

## 3.3.5.1 Simple methods

#### **Syntax**

```
# G and G2 are DiGraphs
G = kdt.DiGraph(v1, v2, v3, n) # construct a DiGraph instance
G2 = G.copy()
                    # deep copy
k = G.nvert()
                   # number of vertices
                 # number of edges
k = G.nedge()
G.ones()
                   # set all non-null edge-weights to 1.0
G.toBool()
                    # set all non-null edge-weights to True
                    # set all non-null edge-weights to k
G.set(k)
                # see below
G.mulNot(G2)
G.removeSelfLoops() # remove all edges from a vertex to itself
```

# **Description**

These methods have their obvious definitions with the following elaborations.

Assignment ("=") of a DiGraph object to another variable name does not create a copy of the object, following Python usage for complex objects. The copy method can be used if a copy is needed.

The toBool method converts each nonnull element of a DiGraph instance to a Boolean True, thereby consuming less space and enabling faster operations.

The mulNot method takes the logical inverse of each element of the second DiGraph before doing the multiplication. Because it does this elementally, there is no extra memory consumed by the inverse values.

#### 3.3.5.2 DiGraph

The DiGraph method creates a directed graph from the edges passed to it.

#### **Syntax**

```
G = kdt.DiGraph(source, dest, weight, nVertOut[, nVertIn])
G = kdt.DiGraph()
```

#### **Description**

The DiGraph method creates a DiGraph instance. The required input parameters source and dest are ParVec objects created by the program or by generators such as genGraph500Edges. The required input parameter weight can be a ParVec or a scalar. The required input parameter nVertOut is an integer defining the number of vertices that have out edges in the graph. The optional input parameter nVertIn is an integer defining the number of vertices that have in edges in the graph; if all vertices potentially have both in and out edges, nVertIn may be omitted. The output argument is a DiGraph object. The values of any duplicate edges (same source and destination) are summed in the creation of the DiGraph object. The DiGraph method is almost the converse of the toParVec method, with the difference that, since the DiGraph method sums duplicate edges, the output of toParVec may have fewer edges, though the same set of vertices.

The alternate form of calling the DiGraph method with no argument creates a DiGraph instance with an empty underlying graph object. This is useful for certain constructors, like genGraph500Edges, which populate the underlying graph object themselves.

DiGraph implements the functionality of Kernel 1 of the Graph500 benchmark.

# **Example**

The code below creates a star graph with N vertices, with directed edges of weight 1 going only from vertex 0 to all vertices (including vertex 0).

```
source = kdt.ParVec.zeros(N)
dest = kdt.ParVec.range(N)
G = kdt.DiGraph(source, dest, 1, N)
```

toParVec

The toParVec method of the DiGraph class decomposes a DiGraph instance to its edges.

#### **Syntax**

3.3.5.3

```
[source, dest, weight] = G.toParVec()
```

# **Description**

The toParVec method of the DiGraph class decomposes a DiGraph instance to its edges, returning ParVec instances containing the source vertices, destination vertices, and edge-weights, respectively. Other than the bound DiGraph instance, there are no input parameters.

The toParVec method is almost the converse of the DiGraph method, with the difference that, since the DiGraph method sums duplicate edges, the output of toParVec may have fewer edges, though the same set of vertices.

# 3.3.5.4 reverseEdges

The reverseEdges method of the DiGraph class reverses the direction of each edge of a DiGraph instance.

#### **Syntax**

```
G.reverseEdges()
```

#### **Description**

The reverseEdges method of the DiGraph class reverses the direction of each edge of a DiGraph instance. Other than the bound DiGraph instance, there are no input parameters. The method works on the DiGraph instance in place, so overwrites its input contents.

#### 3.3.5.5 subgraph

The subgraph method of the DiGraph class creates a new graph from a selected set of vertices of an existing DiGraph object and those vertices' incident edges.

#### **Syntax**

```
G2 = G1.subgraph(vertrange[, vertrange2])
```

# **Description**

The subgraph method of the DiGraph class creates a new DiGraph instance by selecting a set of vertices of an existing DiGraph instance and all the edges incident to those vertices. The required input argument vertrange specifies a range (i.e., a consecutive set of vertex indices) of vertices (and out-edges incident to them) to be used for the new DiGraph instance. The optional input argument vertrange2 specifies another range of vertices (and in-edges incident to them) to be used as the invertices for the new DiGraph instance. If vertrange2 is not specified, vertrange designates both out- and in-vertices.

#### **Example**

The code below creates a new DiGraph from the first half of the vertices in G and edges whose source and destination are both one of those vertices.

```
G2 = G.subgraph(kdt.DiGraph.range(G.nvert()/2))
```

The code below creates a new DiGraph with out-vertices of the first half of the vertices in G, invertices equal to the second half of the vertices in G, and edges whose source is in the first set and destination is in the second.

```
nvG = G.nvert()
G3 = G.subgraph(kdt.DiGraph.range(nvG/2), kdt.DiGraph.range(nvG/2,
nvG))
```

# 3.3.5.6 degree / sum / max / min

The degree, sum, max, and min methods of the DiGraph class calculate, respectively, the degree (count), sum, maximum, and minimum edge-weight of the edges of each vertex of a DiGraph instance.

#### **Syntax**

```
inoutdegs = G.degree([dir=kdt.InOut])
inoutsums = G.sum([dir=kdt.InOut])
inoutmaxs = G.max([dir=kdt.InOut])
inoutmins = G.min([dir=kdt.InOut])
```

#### **Description**

The sum, max, and min methods of the DiGraph class calculate, respectively, the sum, maximum, and minimum edge-weights of the edges of each vertex of a DiGraph instance, returning a ParVec object. The optional dir argument specifies whether the operation is performed on InOut (default), In, or Out edges.

#### **Example**

The code below calculates the sum of the edge-weights of the out-edges of the vertices of a DiGraph.

The scale method of the DiGraph class multiplies the edge weights of each vertex of a DiGraph instance by the corresponding element of a vector of scale factors.

#### **Syntax**

```
G.scale(scaleV[, dir=kdt.Out])
```

#### **Description**

The scale method of the DiGraph class multiplies the edge weights of each vertex of a DiGraph instance by the corresponding element of a SpParVec vector of scale factors. The optional dir argument specifies whether the operation is performed on Out (default) or In edges.

#### Example

The code below normalizes the out-edge weights of each vertex of a DiGraph instance such that the sum of the edge-weights of each vertex is 1.0.

```
scalefac = kdt.ParVec.ones(G.nvert()) / G.sum()
G.scale(scalefac)
```

# 3.3.5.8 normalizeEdgeWeights

The normalizeEdgeWeights method of the DiGraph class normalizes the out-edge weights of each vertex V of a DiGraph instance such that each out-edge's weight is 1 / out-degree(V).

#### **Syntax**

```
G.normalizeEdgeWeights()
```

#### **Description**

The normalizeEdgeWeights method of the DiGraph class normalizes the out-edge weights of each vertex V of a DiGraph instance such that each out-edge's weight is 1 / out-degree(V). Besides the DiGraph instance there are no other arguments.

#### 3.3.6 Advanced methods for DiGraph objects

The DiGraph class implements several advanced methods.

# 3.3.6.1 neighbor

The neighbor method of the DiGraph class calculates the neighbors of a set of starting vertices in a DiGraph instance.

#### **Syntax**

```
neighbors = G.neighbor(start[, nhop=1, sym=False])
```

#### **Description**

The neighbor method of the DiGraph class calculates, for a ParVec of input starting vertices, which vertices are neighbors; *i.e.*, connected by out-edges. The optional nhop argument determines how many hops from the starting vertices are used to calculate the neighbors; the default is 1. The optional expert argument sym denotes whether the graph is a symmetric graph, which if True enables the operation to run faster. The return value is a ParVec with the indices of neighboring vertices.

#### **Example**

If start is a Boolean ParVec of starting vertices, the call below will return all vertices connected via out-bound edges within one hop.

```
neighbors = G.neighbor(start)
```

#### 3.3.6.2 pathsHop

The pathsHop method of the DiGraph class calculates the vertices that can be reached from a set of starting vertices in a DiGraph instance, also returning which of the start vertices has an edge to each new vertex. pathsHop is equivalent to one step in a breadth-first search.

#### **Syntax**

```
[fromV, toV] = G.pathsHop(start[, sym=False])
```

#### **Description**

For a ParVec of input starting vertices, the pathsHop method calculates which vertices can be reached across a single edge in the DiGraph instance and, for each reachable vertex, which starting vertex has an edge to it. The source vertices and the new vertices are returned in ParVec and Boolean ParVec instances, respectively. The set of reachable vertices may include starting vertices. In the case of more than one start vertex having an edge to a reachable vertex, the highest-numbered vertex is selected. pathsHop can be used repeatedly to implement a breadth-first search. The optional expert argument sym denotes whether the graph is a symmetric graph, which if True enables the operation to run faster.

# **Example**

If start is a Boolean ParVec of starting vertices, the code below finds all reachable vertices with the call, and then selects just the newly found vertices from the set of reachable vertices.

```
[fromV, toV] = G.pathsHop(start)
newV = toV & ~start
```

# 3.3.6.3 Breadth-first Search Tree Validation

The isBfsTree method verifies that a breadth-first search tree of a DiGraph instance is valid.

# **Syntax**

```
[valid, levels] = G.isBfsTree(start, parents)
```

#### **Description**

The isBfsTree method takes as input a DiGraph instance, the index of the vertex from which the BFS-tree construction started, and the parent of each vertex (as returned by bfsTree). It returns a 2-tuple whose first element is 1 if the BFS tree is valid and the negative of the number of the test below that failed if the tree is invalid. The second element of the tuple is a ParVec instance (of length equal to the number of vertices in the graph) that denotes, for each respective vertex of the graph, the vertex's level in the BFS tree. Just as for bfsTree, the start vertex is its own parent, and a vertex that is unreachable from the start vertex has a parent of -1.

The validity tests include:

- [1] The tree does not contain cycles, every vertex with a parent is in the tree, and the root is not the destination of any tree edge.
- [2] Tree edges are between vertices whose levels differ by exactly 1.

# 3.3.7 Support for HyGraph objects

The HyGraph class represents undirected hypergraphs, *i.e.*, graphs where edges can be incident upon more than one or two vertices. The HyGraph class is included to explore hypergraph usefulness as a container (converted into DiGraph instances before calling computational methods) and as the direct basis for operations as well. The HyGraph class' implementation for distributed memory includes a small set of methods.

#### 3.3.7.1 Methods

#### **Syntax**

```
# G is a HyGraph, diG is a DiGraph,
# v is a ParVec, k is a scalar integer,
# s is a scalar integer or floating-point value
G = kdt.HyGraph() # constructor from incidence vectors
G.load(filename)  # load incidence vectors to create HyGraph
G2 = G.toDiGraph() # convert to a DiGraph
ret = G.toParVec()  # deconstruct into its incidence vectors
                  # number of vertices
k = G.nvert()
              # number of edges
k = G.nedge()
                  # set all non-null edge-weights to s
G.set(s)
G.toBool()
                  # set all non-null edge-weights to True
G.invertEdgesVertices # invert edges to vertices and vice versa
G.bfsTree(start) # create a breadth-first-search tree
G.isBfsTree()
                  # verify a breadth-first-search tree
```

#### **Description**

These methods have their obvious definitions with the following elaborations.

The load method is described in section 3.6.1.

The toBool method converts each nonnull element of a HyGraph instance to a Boolean True, thereby consuming less space and enabling faster operations.

#### 3.3.7.2 HyGraph

The HyGraph method creates a hypergraph from the incidence vectors passed to it.

#### **Syntax**

```
G = kdt.HyGraph(edgeNumV, incidentVertexV, weightV, nVert[, nEdge])
G = kdt.HyGraph()
```

#### **Description**

The HyGraph method creates a HyGraph instance. The required input parameters edgeNumV and incidentVertexV are ParVec objects. The required input parameter weightV can be a ParVec or a scalar. Each element of edgeNumV denotes an edge number; each corresponding element of incidentVertexV denotes the number of a vertex to which the (undirected) edge is incident. A single edge number can occur an arbitrary number of times in the first argument; all the vertices denoted for the same edge collectively define the hyperedge. The ParVec instances edgeNumV and incidentVertexV (and weightV, if a ParVec) must be of the same length. The required input parameter nVert is an integer defining the number of vertices in the graph (not all vertices must have incident edges). The optional input parameter nEdge is an integer defining the number of edges in the graph. The output argument is a HyGraph object. If multiple entries refer to the same edge and vertex number, their weights are summed in creating the HyGraph object.

The alternate form of calling the HyGraph method with no argument creates a HyGraph instance with an empty underlying graph object. This may be useful for certain constructors that populate the underlying graph object themselves.

#### 3.3.7.3 toDiGraph

The toDiGraph method converts a HyGraph instance to its analogous DiGraph instance.

#### **Syntax**

```
G2 = G.toDiGraph()
```

#### **Description**

The toDiGraph method of the HyGraph class converts a HyGraph instance to its analogous DiGraph, where each hyperedge is replaced by a pair of directed edges between each pair of vertices in the hyperedge. Other than the bound HyGraph instance, there are no input parameters.

#### 3.3.7.4 toParVec

The toParVec method of the HyGraph class decomposes a HyGraph instance to three vectors representing the individual linkes of its hyperedges.

#### **Syntax**

```
[edgeNumV, incidentVertexV, weightV] = G.toParVec()
```

#### **Description**

The toParVec method of the HyGraph class decomposes a HyGraph instance to three vectors respectively denoting the number of the edge to which the incident link belongs, the vertex to which the link is incident, and the weight of the incident link. Other than the bound HyGraph instance, there are no input parameters.

The toParVec method is almost the converse of the HyGraph method, with the difference that, since

the HyGraph method sums the weights of duplicate incident links, the output of toParVec may have fewer elements, though the same set of vertices and edges.

#### 3.3.7.5 invertEdgesVertices

The invertEdgesVertices method of the HyGraph class inverts the sense of each edge and vertex of a HyGraph instance.

#### **Syntax**

```
G.invertEdgesVertices()
```

### **Description**

The invertEdgesVertices method of the HyGraph class inverts the sense of each edge and vertex of a HyGraph instance. An n-vertex and m-edge HyGraph instance becomes an m-vertex and n-edge instance. Other than the bound HyGraph instance, there are no input parameters. The method works on the HyGraph instance in place, so overwrites its input contents.

# 3.3.7.6 degree

The degree method calculates the degree of each vertex of a hypergraph.

#### **Syntax**

```
deg = G.degree()
```

#### **Description**

The degree method calculates the degree (count of incident edges) of each vertex of a hypergraph, returning a ParVec object.

# 3.3.7.7 Breadth-first Search Tree

The bfsTree method creates a breadth-first-search tree of a HyGraph instance from a starting vertex.

#### **Syntax**

```
parents = G.bfsTree(start)
```

#### **Description**

The bfsTree method takes as input a HyGraph instance and the index of the vertex from which to start the search. It returns a ParVec (of length equal to the number of vertices in the graph) that denotes, for each respective vertex of the graph, the vertex's parent in the BFS tree. The start vertex is its own parent. A vertex that is unreachable from the start vertex has a parent of -1.

Note the definition of tree used by bfsTree. In a given round of the bfsTree algorithm, a previously-discovered vertex may have a hyperedge incident to both another previously-discovered

vertex and an undiscovered vertex. The algorithm ignores the potential cycle created by including the entire hyperedge in the tree and denotes the original vertex as the parent (the parent-child link can be viewed as a simple edge).

#### 3.3.7.8 Breadth-first Search Tree Validation

The isBfsTree method verifies that a breadth-first search tree of a HyGraph instance is valid.

#### **Syntax**

```
[valid, levels] = G.isBfsTree(start, parents)
```

#### **Description**

The isBfsTree method takes as input a HyGraph instance, the index of the vertex from which the BFS-tree construction started, and a ParVec instance where each element denotes the parent of each vertex (as returned by bfsTree). It returns a 2-tuple whose first element is 1 if the BFS tree is valid and the negative of the number of the test below that failed if the tree is invalid. If the tree is valid, the second element of the tuple is a ParVec instance (of length equal to the number of vertices in the graph) that denotes, for each respective vertex of the graph, the vertex's level in the BFS tree. Just as for bfsTree, the start vertex is its own parent, and a vertex that is unreachable from the start vertex has a parent of -1.

The validity tests include:

- [1] The tree does not contain cycles and every vertex with a parent is in the tree
- [2] Tree edges are between vertices whose levels differ by exactly 1.

#### 3.3.8 Miscellaneous methods

The following methods exist in the KDT package but not in any class.

- version: returns the KDT version number as a string.
- revision: returns the SVN revision number (of the KDT release contents) as a string.
- sendFeedback: as described in section 1.6, sends feedback to the KDT development team.
- master: as described in section 3.6.4, can be used to restrict output to the master process.

#### 3.3.9 Semantic graph details

#### 3.3.9.1 Semantic graphs via DiGraph and Vec constructors

Semantic graph data will typically be coming from the real world, and so will be loaded from files. The DiGraph.load and Vec.load methods depend on the load function in ObjMethods.py to understand the layout of the file data. Semantic graphs can also be built from a vector of edge-object elements by using the element argument to the DiGraph constructor.

The DiGraph.copy (Vec.copy) methods accept an element argument that can be used to convert a DiGraph (Vec) instance with Objl or Obj2 elements to a DiGraph (Vec) instance with the non-semantic 64-bit element.

3.3.9.2 addEFilter

Syntax

G.addEFilter(filter)

### **Description**

The addEFilter method attaches the passed edge-filter to the given DiGraph instance. The filter must be a PyObject object that accepts as input the elemental object of the DiGraph instance (Obj1 or Obj2) and returns a Boolean value. The filter will be invoked internally to all subsequent methods on the DiGraph instance until the filter is deleted via delEFilter or the DiGraph instance is deleted or goes out of scope. Multiple filters may be attached; they can be viewed as executing in series (for each edge) until all have returned True or until a filter returns False. Only edges for which the filter(s) returns a Boolean True (and which are not incident to vertices which do not pass a filter) will be used for the remainder of the method. This method executes in-place and does not provide a return value.

3.3.9.3 delEFilter

**Syntax** 

G.delEFilter([filter])

#### **Description**

The delEFilter method deletes edge-filter(s) from the given DiGraph instance. If the optional argument is passed, it must be a PyObject object previously passed to addEFilter on the same DiGraph instance. If the optional argument is not passed, all filters are deleted from the DiGraph instance. This method executes in-place and does not provide a return value.

3.3.9.4 addVFilter

Syntax

v.addVFilter(filter)

# **Description**

The addVFilter method attaches the passed vertex-filter to the given Vec instance (which may be the vertex attributes of a DiGraph instance (see section 3.3.9.6)). The filter must be a PyObject object that accepts as input the elemental object of the Vec instance (Obj1 or Obj2) and returns a Boolean value. The filter will be invoked internally to all subsequent methods on the Vec instance (and any DiGraph instance to which the Vec instance is attached as a vertex filter) until the filter is deleted

via delVFilter or the Vec instance is deleted or goes out of scope. Multiple filters may be attached; they can be viewed as executing in series (for each vertex) until all have returned True or until a filter returns False. Only vertices for which the filter(s) returns a Boolean True will be used for the remainder of the method; vertices for which the filter(s) returns a Boolean False will have their incident edges viewed as not having passed the filter as well and thereby ignored for the remainder of the method. This method executes in-place and does not provide a return value.

#### 3.3.9.5 delVFilter

#### **Syntax**

```
v.delVFilter([filter])
```

# **Description**

The delVFilter method deletes vertex-filter(s) from the given DiGraph instance (which may be the vertex attributes of a DiGraph instance (see section 3.3.9.6)). If the optional argument is passed, it must be a PyObject object previously passed to addVFilter on the same Vec instance. If the optional argument is not passed, all filters are deleted from the Vec instance. This method executes inplace and does not provide a return value.

# 3.3.9.6 Adding or modifying vertex attributes to a DiGraph instance

Attributes of the vertices of a DiGraph instance can be specified by setting the vAttr attribute of the DiGraph instance to a Vec instance whose element is the desired object-type and data.

## **Syntax**

```
G.vAttr = v
G.vAttr.addVFilter(filter)
G.vAttr.delVFilter([filter])
```

#### **Description**

The vAttr attribute will be queried for vertex filters whenever operations are performed on the DiGraph instance. Only vertices for which the filter(s) returns a Boolean True will be used for the remainder of the method; vertices for which the filter(s) returns a Boolean False will have their incident edges viewed as not having passed the filter as well and thereby ignored for the remainder of the method.

A Vec instance that is the vAttr attribute of a DiGraph instance may still be modified by adding or removing filters, as shown by the syntax above.

# 3.3.9.7 Semantic-graph example

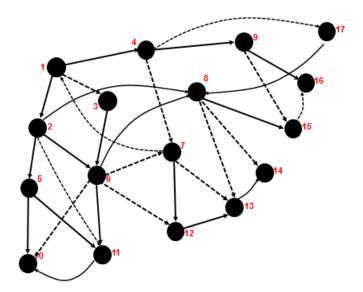


Figure 1. Sample semantic graph with two types of edges

A simple synthetic example of the use of filters is given below. The code is from the test/TestDiGraph.py file of the KDT distribution. It builds a semantic DiGraph instance of the graph in Figure 1 with Objl edge-elements split between categories 1 and 2 and calculates the degree of each vertex looking first at just category 1 and then category 2 edges.

```
def test_degree_Out_Obj1_filteredTwoWays(self):
   # filters that test if category is equal to 1 or 2, respectively
   def category_eq_1(x):
         if isinstance(x, (Obj1, Obj2)):
                return x.category==1
         else:
                raise NotImplementedError
   def category_eq_2(x):
         if isinstance(x, (Obj1, Obj2)):
                return x.category==2
         else:
                raise NotImplementedError
   nvert = 18
   nedge = 31
   7, 7, 8,12, 8,13, 8, 9, 9,15, 4]
   j = [1, 2, 3, 4, 5, 6, 6, 7, 7, 8, 8, 8, 9, 10, 10, 10, 11, 11, 11, 12,
12,13,13,13,14,14,15,15,16,16,17]
   1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1]
```

```
1, 2, 2, 1, 2, 1, 1, 2, 1, 2, 2]
   self.assertEqual(len(i), nedge)
   element = Obj1()
   G = self.initializeGraph(nvert, nedge, i, j, (w, c),
           element=element)
   G.addEFilter(category_eq_1)
   degOutCat1 = G.degree(DiGraph.Out)
   G.delEFilter(category_eq_1)
   expectedOutDegCat1 = [ 0, 2, 3, 1, 1, 2, 2, 1, 1, 1, 0, 1, 1, 1,
0, 0, 0, 1]
   self.assertEqual(len(degOutCat1), nvert)
   for ind in range(G.nvert()):
           self.assertEqual(expectedOutDegCat1[ind], degOutCat1[ind])
   G.addEFilter(category_eq_2)
   degOutCat2 = G.degree(DiGraph.Out)
   G.delEFilter(category_eq_2)
   expectedOutDegCat2 = [ 0, 1, 1, 0, 2, 0, 3, 2, 2, 1, 0, 0, 0,
0, 1, 0, 0]
   self.assertEqual(len(degOutCat2), nvert)
   for ind in range(G.nvert()):
           self.assertEqual(expectedOutDegCat2[ind], degOutCat2[ind])
```

# 3.3.9.8 Performance

Calling Python functions to filter and calculate element-wise is substantially slower than calling the built-in C++ functions used for non-semantic graphs. [[FIX: any data?]]

# 3.4 Package and class structure

KDT is structured as shown in Table 1, with some methods omitted for brevity. The complete list of functions can be seen by executing (within IPython)

```
import kdt
dir(kdt)
dir(kdt.DiGraph)
dir(kdt.DeVec)
dir(kdt.SpVec)
```

Entity	Name	Elements	Comments
Module	KDT	version, revision, sendFeedback, master	Methods unrelated to any specific data structure
Class	DeVec		Distributed parallel dense vector
		DeVec, [], =, +, -, += , -=, <>, >, findInds, abs, any, all, nnn, addVFilter,	Methods
Class	SpVec		Distributed parallel sparse vector

		SpVec, [], =, +, -, += , -=, <>, >, findInds, abs, any, all, nnn, addVFilter,	Methods
Class	Vec		Base class for DeVec, SpVec
Class	DiGraph		Directed graph
		DiGraph, genGraph500Edges, load, fullyConnected	Constructors
		centrality, cluster	Algorithms
		bfsTree, isBfsTree, neighbors, pathsHop, toParVec, reverseEdges, subgraph	Graph primitives
		load	1/0
		[], nvert, nedge, degree, +, *, ones, set	General-purpose routines
		addEFilter, delEFilter	Utility routines
		In, Out	Constants
	HyGraph		Hypergraph
		HyGraph, load	Constructors
		bfsTree, isBfsTree, toParVec, toDiGraph, invertEdgesVertices, toBool	Graph primitives
		nvert, nedge, degree, set	General-purpose routines

Table 1. KDT library hierarchy

# 3.5 Graph Generators

The DiGraph class includes the graph generators in this section. With v0.1, edges may be created as an ParVec object directly with a subsequent call to the DiGraph method, by the load or UFget method, by a constructor such as fullyConnected, or by using an application-specific method like genGraph500Edges.

# 3.5.1 genGraph500Edges

The genGraph500Edges function creates a graph following the specifications for the V1.1 Graph500 benchmark's input graph [Graph500]. The edges are inserted into the DiGraph object passed and represent an RMAT graph with specific values provided by the benchmark.

#### **Syntax**

time = G.genGraph500Edges(scale)

#### **Description**

The genGraph500Graph method creates an input graph as defined by the Graph500 benchmark. The required input parameter scale (logarithm base 2 of the number of desired vertices) defines the number of vertices. The edges are directed, though each edge has a twin going in the other direction because the specification requires the graph to be symmetric. Some vertices may have no edges incident to them. The time returned from genGraph500Edges includes the execution time of converting the edge vector to a graph but does not include the time to create the edge vector, which is exactly the time measured by Kernel 1 of the Graph500 benchmark.

#### Example

The following code will creates a DiGraph instance and inserts edges into it that match the Graph500 specification, of size scale.

```
G = kdt.DiGraph()
time = G.genGraph500Edges(scale)
```

#### 3.5.2 twoDTorus

The twoDTorus method creates a DiGraph object reflecting the connectivity pattern of a 2D torus.

#### **Syntax**

```
G = kdt.twoDTorus(nnode)
```

#### **Description**

The twoDTorus method creates a DiGraph with the connectivity pattern of a 2D torus; *i.e.*, connections to its north, west, south, and east neighbors, where the neighbor may be wrapped around to the other side of the torus. The required input parameter nnodes defines the number of nodes along one dimension of the torus; the torus and the graph representing it will have nnodes\*\*2 vertices. The newly created DiGraph object is the return value from the method. A 2D torus has an easily analyzed betweenness centrality value that can be useful for simple tests; specifically, each vertex of the DiGraph created by twoDTorus(nnodes) has an identical betweenness centrality value of 0.5 \* 2\*\*(3\*scale\*0.5) - 2\*\*scale + 1 for an even nnodes.

#### **Example**

The following code creates a DiGraph instance G with nnodes \* \* 2 vertices and inserts edges from every vertex to every other vertex, including itself.

```
G = kdt.DiGraph.twoDTorus(nnodes)
```

#### 3.5.3 fullyConnected

The fullyConnected method creates a DiGraph object in which all vertices are directly connected to all other vertices. The edges are inserted into a newly created DiGraph object, which is the return value from the method.

#### **Syntax**

```
G = kdt.fullyConnected(nvert)
```

#### **Description**

The fullyConnected method creates a DiGraph. with all vertices having a directed edge to all vertices, including itself. The required input parameter nvert defines the number of vertices. The newly created DiGraph object is the return value from the method.

#### Example

The following code creates a DiGraph instance G with nvert vertices and inserts edges from every vertex to every other vertex, including itself.

```
G = kdt.DiGraph.fullyConnected(nvert)
```

## 3.6 I/O

The DiGraph class includes the I/O-related methods in this section.

#### 3.6.1 load: Reading a graph from a file

A Matrix Market file can be loaded directly into a DiGraph or HyGraph object with the standard load I/O operation.

#### **Syntax**

```
G = kdt.DiGraph.load(fname)
G = kdt.HyGraph.load(fname)
```

#### **Description**

The load method loads a file in the Coordinate Format of the Matrix Market Exchange Formats [MatrixMarket] directly into a DiGraph or HyGraph object.

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The source-vertex / destination-vertex (for DiGraph) or edge-number / vertex-number (for HyGraph) information in the first two columns of the Matrix Market Exchange format is 1-based in the file, and is converted to 0-based in the load. With v0.1, the error-checking of a source-vertex or destination-vertex or edge-number or vertex-number being out of bounds is not robust and can result in KDT aborting with Segmentation Faults or malloc errors.

#### **Example**

The following code will load the contents of the file mymatrix.mtx into a DiGraph instance named G.

```
G = kdt.DiGraph.load('mymatrix.mtx')
```

## 3.6.2 save: Writing a graph into a file

A DiGraph object can be saved directly into a Matrix Market file with the standard save I/O operation.

#### **Syntax**

```
G.save(fname)
```

#### **Description**

The save method save a a DiGraph object directly into a file in the Coordinate Format of the Matrix Market Exchange Formats [MatrixMarket].

#### Example

The following code will save the contents of a DiGraph instance named G into the file mymatrix.mtx.

```
G.save('mymatrix.mtx')
```

## 3.6.3 UFget: Fetching a file from the University of Florida Sparse Matrix Library and loading it as a DiGraph

The UFget method downloads a named Matrix Market file from the University of Florida Sparse Matrix Library (link) and loads it into a DiGraph instance.

#### **Syntax**

```
G.UFget(fname)
```

#### **Description**

The UFget method downloads a file from the UF Sparse Matrix Library, untars the contents, and loads the contained file (in the Coordinate Format of the Matrix Market Exchange Formats) directly into a DiGraph instance.

Note: See Errata #3 in section 5.5.

#### **Example**

The following code will load the contents of the file Andrianov/ex3stal.tar.gz from the UF Sparse Matrix Library into a DiGraph instance named G.

```
G = kdt.DiGraph.UFget('Andrianov/ex3sta1')
```

#### 3.6.4 master: Avoiding redundant print output

When running in parallel, each process will execute the Python code, including any print statements. For user output, this can lead to numerous copies of the output, often intermingled so as to make the output unintelligible. The master method is useful for avoiding this situation.

Note: Printing of DiGraph, ParVec, or SpParVec instances will, in general, require participation of all the active processes, and thus cannot be done solely on the master, and thus should not be done inside a master check. Doing so typically results in the program hanging.

#### **Syntax**

```
bool = kdt.master()
```

#### Description

The master method returns a Boolean result that is True in the master process of the underlying infrastructure and False in all other processes.

#### Example

The following code, taken from the Graph500 script in KDT, restricts printing output just to the master process.

```
if kdt.master():
    print 'Graph500 benchmark run for scale = %2i' % scale
    print 'Kernel 1 time = %8.4f seconds' % K1elapsed
    print "\nKernel 2 BFS execution times"
    printstats(K2elapsed, "time", False)
```

## 3.7 Advanced Usage

KDT v0.1 supports only graphs with a single edge between any two vertices, which doesn't address nearly all graph types. While fully general graphs will have to wait for future releases, v0.1 does support (via built-in Python mechanisms) multiple graphs of different types of data. For instance, one potential KDT use case is to analyze data from protein-protein, DNA-protein, and other interactions. Data about a particular organism (e.g., Shewanella) could be collected in a single DiGraph instance which has other DiGraph instances as members, as portrayed by the following code.

```
shew = kdt.DiGraph()
shew.prot2prot = kdt.DiGraph.load('shewanella/protein_protein.mtx')
shew.DNA2prot = kdt.DiGraph.load('shewanella/DNA_protein.mtx')
d2p_bc = shew.DNA2prot.centrality('approxBC')
p2p_bc = shew.prot2prot.centrality('approxBC')
d2p_deg = shew.DNA2prot.degree()
p2p_deg = shew.prot2prot.degree()
```

In this example, each graph is still operated on independently, but where graph operations make scientific or mathematical sense, DiGraph graphs can be joined together.

#### 4 Performance

KDT is intended to enable subject-matter experts who are not graph experts to solve large problems quickly, including both the development and execution phases of a program, and thus performance is a vital aspect of KDT. Its performance has not yet been fully characterized. Performance of specific methods, where available, is provided in their descriptions.

Today we have no means to execute KDT interactively in parallel, where "interactive" means having the full range of IPython's abilities (*e.g.*, breakpoints, stepping, variable inspection) at the user's disposal. Thus we consider interactive and parallel performance separately.

#### 4.1 Interactive

Since interactive execution means serial execution and memory capacities of a single node for v0.1, users will likely limit the problem sizes they develop with interactively to get quick response times.

The Graph500 example in KDT (see examples/Graph500.py) is configurable for problem size and number of iterations. For scale=15 (equating to the number of graph vertices being 2\*\*15 or 32K), the Graph500 kernel2 itself runs for a single starting vertex in about 0.3 seconds on a modern x86 core. The entire processing of the graph, including validation of the resulting BFS tree, takes about 3 seconds per iteration (starting vertex).

Betweenness centrality (e.g., the class CentralityTests in file test/TestDiGraph.py) in KDT is also configurable for problem size and sample rate (for approximate BC). Using the twoDTorus graph generator with the number of torus nodes equal to 32 (equating to the number of graph vertices being 1024), exact BC runs in about 8 seconds on a modern x86 core, and for this scale approximate BC is only slightly faster. Note that a 2D torus is a poor performance case for BC, as one factor in BC's computational complexity is the diameter of the graph, which for a 2D torus is the square root of the number of graph vertices, whereas typical RMAT graphs have a diameter more like the log of the number of graph vertices.

#### 4.2 Parallel

The parallel testing of KDT has occurred on three systems:

- Neumann, a 32-core cluster at UCSB
- Triton, a medium-sized 256-node cluster at the San Diego Supercomputer Center (configuration)
- Hopper II, a large 6392-node, 24 cores-per-node Cray XE6 system at the National Energy Research Supercomputer Center (<u>configuration</u>)

The performance characterization of KDT at large scale is still in process. The following performance has already been demonstrated.

Much of the early performance work with KDT has focused on the Graph500 benchmark. In Table 1, initial Graph500 performance numbers from the Triton cluster at the San Diego Supercomputer Center are provided. The units are millions of traversed-edges per second (TEPS); the first number is for each data element being a 64-bit double-precision value; the second is for each data element being a Boolean value; the performance for Boolean values is often a factor of 5-10 faster than 64-bit double-precision values. On 256 cores, the KDT implementation of Graph500 performs at over 1 GTEPS.

Scale	20	23
Number of cores		
16	- /200.5	25.9 / 171.4
64	- / 518.5	
256	- / 485.1	- /1,133.2

Table 1. Graph500 performance

Betweenness centrality is a much more compute-intensive applet than the Graph500, so the problem sizes run so far are much smaller. On Neumann a 2D torus with 128 torus nodes (equating to 16K graph vertices) runs exact BC in 1163 seconds and approximate BC (sample rate of 0.05) in 884 seconds on 16 cores.

#### 5 Practicalities

## 5.1 Downloading

To get the latest released package, go to <a href="http://kdt.sourceforge.net/download.html">http://kdt.sourceforge.net/download.html</a> and download it.

#### 5.2 Linux

#### **5.2.1** System Requirements

We recommend Python 2.4 or newer. KDT has been tested primarily with OpenMPI 1.4 and IPython 0.8.4.

#### 5.2.2 Installation and Build

The information in this section is repeated verbatim from the README.txt file in the base directory of the KDT package. In case of any discrepancies between this text and README.txt, README.txt should prevail.

KDT is distributed as a Python distutils package. It requires the MPI compiler to be specified in the \$CC and \$CXX environment variables. For example, in bash:

```
export CC=mpicxx export CXX=mpicxx
```

The build and installation is performed by the standard distutils setup command:

```
$ python setup.py build
$ sudo python setup.py install
```

If the build step fails fails see the "Choosing a compiler" section below.

KDT makes use of some features from C++ TR1. If your compiler does not support TR1 then the free Boost C++ library (http://www.boost.org/) supplies the required headers as well. Make sure boost/ is in your include path. If it is not, you can append the include path with the -I switch to the setup.py script. For example, if you installed Boost in /home/username/include/boost:

```
$ python setup.py build -I/home/username/include
```

The MPI library must be compiled with -fPIC. Python modules must be compiled with -fPIC, and that includes all libraries that get linked into the module, which includes the MPI library in the case of KDT. If your MPI was not compiled with -fPIC then the link step will fail. If this happens, contact your system administrator.

#### 5.2.2.1 Choosing a Compiler

KDT consists of pure Python classes and a C++ extension. This C++ extension, called pyCombBLAS, is MPI code and must be compiled as such. However, it is also a Python module, so must be compiled with a compiler that is compatible with your Python installation. By default, distutils (i.e. setup.py) will use the same compiler with which Python was compiled. That was probably not an MPI compiler, so the proper compiler must be specified to distutils via the CC and CXX environment variables.

Note that your system may give you a choice between GNU and Portland Group (PGI) compilers. These are not fully binary compatible with each other, so you must use the same one that your Python was compiled with. Note that mpicxx is often just a wrapper around GNU or PGI compilers.

#### 5.2.2.2 System Libraries

If you chose to use a non-default MPI compiler then be aware that the runtime may link to the default MPI libraries anyway. This is probably not what you want. The result may be something like this:

```
ImportError: ./kdt/_pyCombBLAS.so: undefined symbol: _ZN3MPI3Win4FreeEv
```

The solution is to set your LD\_LIBRARY environment variable such that the MPI library you compiled with appears before the incorrect defaults.

#### 5.2.2.3 Building without Distutils

If the setup.py script fails for you, you can build the C++ extension manually. The kdt/pyCombBLAS directory contains a Makefile (named Makefile-dist; rename as necessary) which you can tune for your system. The important things to change are:

- COMPILER compiler to use.
- INCADD include your Python build directory as well as your Boost installation if you need it.
- OPT any flags you wish to change.

Once pyCombBLAS is built, you may manually install the kdt/directory in your systemwide Python site-packages directory, or simply set your PYTHONPATH environment variable to point to the parent directory of kdt/(i.e., the base of the distribution).

#### **5.2.3** Tests and Examples

The KDT package includes test/ and example/ directories at the base level of the distribution, and in particular the example/Graph500.py script runs the Graph500 benchmark, which may be a good way to exercise KDT after installation. Because these files embody the use of KDT functionality, not the core functionality itself, they are not installed in the standard site-packages directory with the core functionality. A user who downloads KDT for her own use may just leave them in the directory where the package was unzipped/untarred, or may put them in a system-wide directory for others' use as well. For the rest of this section the test/ and example/ directories are assumed to reside in a directory pointed to by the KDTINSTALL environment variable.

The test/ directory contains three class-specific files: TestParVec.py, TestSpParVec.py, and TestDiGraph.py, which test the classes with the corresponding names. The current failures are noted in section 5.5 Errata.

The examples/ directory contains the Graph500.py script, which implements the Graph500 benchmark specification, including all validation steps.

#### **5.2.4** Execution with Python or IPython

This section provides details about how to execute KDT scripts via IPython and Python.

#### 5.2.4.1 *IPython*

KDT has been used interactively only with IPython, namely version 0.8.4, on a single core. The command-line and IPython commands to invoke it follow (note the same information appears in section 2).

#### In [1]: import Graph500

```
Activating auto-logging. Current session state plus future input saved.
```

```
#This output is from the startup of the IPython logstart

Mode : over # mechanism used for the sendFeedback function

Output logging : False # from Section 1.6

Raw input log : False # ""

Timestamping : False # ""

State : active # ""
```

```
Generating a Graph500 RMAT graph with 2^15 vertices...

Duplicates removed (or summed): 78590 and self-loops removed: 0

Generation took 0.371470s.

iteration 1: start=9171, BFS took 0.030166s, verification took
2.803589s and succeeded, TEPS=29,545,933

iteration 2: start=12434, BFS took 0.030009s, verification took
2.762023s and succeeded, TEPS=29,700,392

iteration 3: start=20292, BFS took 0.029619s, verification took
2.818182s and succeeded, TEPS=30,091,517

iteration 4: start=29823, BFS took 0.029394s, verification took
2.728882s and succeeded, TEPS=30,321,679
```

```
[...]
iteration 63: start=14394, BFS took 0.030498s, verification took
2.766191s and succeeded, TEPS=29,224,184
iteration 64: start=7008, BFS took 0.031285s, verification took
2.771903s and succeeded, TEPS=28,489,226
Graph500 benchmark run for scale = 15
Kernel 1 time = 0.3715 seconds
Kernel 2 BFS execution times
            min time: 2.93109416961669922e-02
  firstquartile_time: 3.01442742347717285e-02
         median_time: 3.08154821395874023e-02
  thirdquartile_time: 3.10997962951660156e-02
           max_time: 3.19378376007080078e-02
           mean_time: 3.06311249732971191e-02
         stddev time: 6.26722467712359749e-04
Kernel 2 number of edges traversed
            min_nedge: 8.91280000000000000e+05
  firstquartile nedge: 8.91280000000000000e+05
         median nedge: 8.91280000000000000e+05
  thirdquartile_nedge: 8.9128000000000000e+05
            max_nedge: 8.91280000000000000e+05
           mean_nedge: 8.91280000000000000e+05
         stddev nedge: 0.0000000000000000e+00
Kernel 2 TEPS
            min TEPS: 2.79067108782669082e+07
  firstquartile TEPS: 2.86587086147099845e+07
         median_TEPS: 2.89231245181400143e+07
  thirdquartile TEPS: 2.95671458547492772e+07
            max_TEPS: 3.04077572545734048e+07
 harmonic_mean_TEPS: 2.90972009933353513e+07
harmonic_stddev_TEPS: 1.99059058009882137e+04
In [2]:
```

## 5.2.4.2 Python, not via a job manager

KDT has been used in parallel only with the Python language processor, namely Python 2.4.3. The MPI used was OpenMPI 1.4. The Linux version was 2.6.18.

Note: The current Combinatorial BLAS, on which KDT is based, only supports core counts that are perfect squares, meaning that the argument to the mpirun -n command must be a perfect square.

For parallel execution with Python and MPI, you can use the following command-line.

```
[sam@neumann test]$ mpirun -n 4 python $KDTINSTALL/examples/Graph500.py Generating a Graph500 RMAT graph with 2^15 vertices...

Duplicates removed (or summed): 78883 and self-loops removed: 0 Generation took 0.080711s.
```

```
iteration 1: start=12627, BFS took 0.011984s, verification took
0.619967s and succeeded, TEPS=74,322,165
iteration 2: start=20502, BFS took 0.011391s, verification took
0.601116s and succeeded, TEPS=78,192,512
iteration 3: start=8839, BFS took 0.011701s, verification took
0.616723s and succeeded, TEPS=76,119,720
iteration 4: start=1999, BFS took 0.011484s, verification took
0.612136s and succeeded, TEPS=77,559,400
iteration 5: start=71, BFS took 0.011390s, verification took 0.619576s
and succeeded, TEPS=78,199,059
     [...]
iteration 62: start=32066, BFS took 0.011214s, verification took
0.605297s and succeeded, TEPS=79,427,725
iteration 63: start=27903, BFS took 0.011867s, verification took
0.608496s and succeeded, TEPS=75,055,323
iteration 64: start=16201, BFS took 0.011442s, verification took
0.609218s and succeeded, TEPS=77,843,838
Graph500 benchmark run for scale = 15
Kernel 1 time = 0.0807 seconds
Kernel 2 BFS execution times
            min_time: 1.09930038452148438e-02
  firstquartile time: 1.14405155181884766e-02
         median_time: 1.16879940032958984e-02
  thirdquartile_time: 1.18589401245117188e-02
            max time: 1.21469497680664062e-02
           mean_time: 1.16281397640705109e-02
         stddev_time: 2.91774617905905386e-04
Kernel 2 number of edges traversed
            min nedge: 8.90685000000000000e+05
  firstquartile nedge: 8.90685000000000000e+05
         median_nedge: 8.90685000000000000e+05
  thirdguartile nedge: 8.9068500000000000e+05
            max nedge: 8.90685000000000000e+05
           mean_nedge: 8.90685000000000000e+05
         stddev_nedge: 0.0000000000000000e+00
Kernel 2 TEPS
            min_TEPS: 7.33258156991442293e+07
  firstquartile TEPS: 7.51066276284680367e+07
         median_TEPS: 7.62052171539593935e+07
  thirdquartile_TEPS: 7.78535725269949883e+07
            max TEPS: 8.10228953379023224e+07
 harmonic_mean_TEPS: 7.65973765427299738e+07
```

```
harmonic_stddev_TEPS: 6.64094336990017182e+04
[sam@neumann test]$
```

#### 5.2.4.3 Python, via a job manager

Many large HPC systems require programs using a large number of cores to be submitted via a job management system. The script below works on SDSC's Triton cluster, using the TORQUE (formerly PBS) job management system.

```
$ qsub testscriptRun.sh
where testscriptRun.sh looks like
#!/bin/bash
#PBS -q batch
#PBS -N pyCombBLAS
#PBS -1 nodes=72:ppn=8
#PBS -l walltime=1:20:00
#PBS -o out.testscript23-576-b
#PBS -e err.testscript23-576-b
#PBS -V
#PBS -M alugowski@gmail.com
#PBS -m abe
#PBS -A alugowski-ucsb
export LD_LIBRARY_PATH=/opt/openmpi/gnu/mx/lib:$LD_LIBRARY_PATH
cd ~/trunk/kdt/pyCombBLAS
/opt/openmpi/gnu/mx/bin/mpirun -v -machinefile $PBS_NODEFILE -np 576
python ./testscript.py -1 /home/alugowski-ucsb/matrices/rmat23.mtx
```

#### 5.3 Windows

#### **5.3.1** System Requirements

KDT has been tested with Windows HPC Server 2008 R2, Python 2.7.1, and MPICH2 1.3.2p1.

#### 5.3.2 Installation and Build

The information in this section is repeated verbatim from the README\_win.txt file in the base directory of the KDT package. In case of any discrepancies between this text and README\_win.txt, README\_win.txt should prevail.

KDT is distributed as a Python Distutils package. The commands to build and install it via the typical "python.exe setup\_win.py build" approach are encapsulated in the PowerShell script install\_prereqs\_KDT.ps1.

KDT can be installed with the following steps:

- 1) The install procedure assumes that Python 2.7 and MPICH2 1.3.2p1 are not already installed. If they are, you may wish to ignore the steps related to them and edit install\_prereqs\_KDT.ps1 not to operate on those files.
- 2) Download the Python 2.7 MSI installer from <a href="here">here</a> and the MPICH2 1.3.2p1 MSI installer from here.
- 3) Download the KDT\_0.1W.zip file onto the head node of the HPC Server cluster into the same directory as the Python and MPICH2 MSI installers.
- 4) Unzip the KDT file, either with a utility or with the following PowerShell commands

```
PS > $shell_app=new-object -com shell.application
PS > $zip_file=$shell_app.namespace((Get-Location).Path+"\kdt_0.1W.zip")
PS > $zip_dest=$shell_app.namespace((Get-Location).Path+"\kdt_0.1W")
PS > $zipdest.Copyhere($zip_file.items())
```

5) Use clusrun to install (Python 2.7 and MPICH2 and) KDT on the head node and all the cluster nodes. *I.e.*, when executing on the head node,

```
PS > clusrun PowerShell -ExecutionPolicy bypass
   -file \\<headnode-name>\<path>\kdt_0.1W\install_prereqs_KDT.ps1
```

It may be worth executing clusrun on a single node to ensure it's working properly before executing it on all the nodes.

For those administrators wishing more control, the build and installation may be performed (on each node) by the standard Distutils setup commands executed in the correct directory, *i.e.*,

```
$ C:\python27\python.exe setup_win.py build
$ C:\python27\python.exe setup_win.py install
```

The second command above must be performed with Administrator privileges.

KDT must be built with the same compiler used to build the MPICH2 library, which is currently the Visual Studio 2008 64-bit compiler. The default build process handles this implicitly. If the build step fails see the "Choosing a compiler" section below.

### 5.3.2.1 Choosing a Compiler

KDT consists of pure Python classes and a C++ extension. This C++ extension, called pyCombBLAS, is MPI code and must be compiled as such. However, it is also a Python module, so must be compiled with a compiler that is compatible with your Python installation. By default, distutils (i.e. setup\_win.py) will use the same compiler with which Python and MPICH2 were compiled, but if that doesen't happen properly problems can result.

#### 5.3.3 Tests and Examples

The KDT package includes test/ and example/ directories at the base level of the distribution, and in particular the example/Graph500.py script runs the Graph500 benchmark, which may be a good

way to exercise KDT after installation. Because these files embody the use of KDT functionality, not the core functionality itself, they are not installed in the standard site-packages directory with the core functionality. A user who downloads KDT for her own use may just leave them in the directory where the package was unzipped/untarred, or may put them in a system-wide directory for others' use as well. For the rest of this section the test/ and example/ directories are assumed to reside in a directory pointed to by the KDTINSTALL environment variable.

The test/ directory contains three class-specific files: TestParVec.py, TestSpParVec.py, and TestDiGraph.py, which test the classes with the corresponding names. The current failures are noted in section 5.5 Errata.

The examples/ directory contains the Graph500.py script, which implements the Graph500 benchmark specification, including all validation steps.

### 5.3.4 Execution with Python or IPython

This section provides details about how to execute KDT scripts via IPython and Python.

#### 5.3.4.1 IPython

KDT has been used extensively with IPython on Linux, but not on Windows. Much of the following section is probably relevant to the execution of IPython on Windows, but has not been exercised, so the rest of this section is stippled.

KDT has been used interactively only with IPython, namely version 0.8.4, on a single core. The command-line and IPython commands to invoke it follow (note the same information appears in section 2).

```
[sam@neumann ~]$ PYTHONPATH="$PYTHONPATH:$KDTINSTALL/examples"
[sam@neumann ~]$ ipython
Python 2.4.3 (#1, Nov 11 2010, 13:30:19)
Type "copyright", "credits" or "license" for more information.
IPython 0.8.4 -- An enhanced Interactive Python.
    -> Introduction and overview of IPython's features.
%quickref -> Quick reference.
help -> Python's own help system.
object? -> Details about 'object'. ?object also works, ?? prints
more.
In [1]: import Graph500
Activating auto-logging. Current session state plus future input
saved.
Filename : .KDT_log #This output is from the startup of the IPython logstart
Mode
               : over # mechanism used for the sendFeedback function
Output logging : False # from Section 1.6
Raw input log : False # ""
Timestamping : False # ""
State : active # ""
```

```
Generating a Graph500 RMAT graph with 2^15 vertices...
Duplicates removed (or summed): 78590 and self-loops removed: 0
Generation took 0.371470s.
iteration 1: start=9171, BFS took 0.030166s, verification took
2.803589s and succeeded, TEPS=29,545,933
iteration 2: start=12434, BFS took 0.030009s, verification took
2.762023s and succeeded, TEPS=29,700,392
iteration 3: start=20292, BFS took 0.029619s, verification took
2.818182s and succeeded, TEPS=30,091,517
iteration 4: start=29823, BFS took 0.029394s, verification took
2.728882s and succeeded, TEPS=30,321,679
iteration 63: start=14394, BFS took 0.030498s, verification took
2.766191s and succeeded, TEPS=29,224,184
iteration 64: start=7008, BFS took 0.031285s, verification took
2.771903s and succeeded, TEPS=28,489,226
Graph500 benchmark run for scale = 15
Kernel 1 time = 0.3715 seconds
Kernel 2 BFS execution times
           min_time: 2.93109416961669922e-02
  firstquartile_time: 3.01442742347717285e-02
        median_time: 3.08154821395874023e-02
  thirdquartile time: 3.10997962951660156e-02
           max_time: 3.19378376007080078e-02
          mean_time: 3.06311249732971191e-02
         stddev time: 6.26722467712359749e-04
Kernel 2 number of edges traversed
          min nedge: 8.91280000000000000e+05
  firstquartile_nedge: 8.9128000000000000e+05
        median_nedge: 8.91280000000000000e+05
  thirdquartile_nedge: 8.91280000000000000e+05
           max_nedge: 8.91280000000000000e+05
          mean nedge: 8.9128000000000000e+05
         stddev_nedge: 0.0000000000000000e+00
Kernel 2 TEPS
           min_TEPS: 2.79067108782669082e+07
  firstquartile_TEPS: 2.86587086147099845e+07
        median_TEPS: 2.89231245181400143e+07
  thirdquartile_TEPS: 2.95671458547492772e+07
           max_TEPS: 3.04077572545734048e+07
 harmonic_mean_TEPS: 2.90972009933353513e+07
harmonic_stddev_TEPS: 1.99059058009882137e+04
In [2]:
```

## 5.3.4.2 Python, not via a job manager

KDT has been used in parallel only with the Python language processor, namely Python 2.7.1.

Note: The current Combinatorial BLAS, on which KDT is based, only supports core counts that are perfect squares, meaning that the argument to the mpirun -n command must be a perfect square.

For parallel execution with Python and MPI, you will typically want to create a machinefile in the format described in the MPICH2 User Guide (<u>link</u> on p. 8), such as in hosts.txt, namely

```
# comment line
hostA:n  # assign n ranks to hostA
hostB:m  # assign m ranks to hostB
```

With such a machinefile in hand, you can use the following command-line:

```
[sam@neumann test]$ "c:\program files\mpich2\bin\mpiexec.exe" -n 4
-machinefile hosts.txt c:\python27\python.exe
$KDTINSTALL\examples\Graph500.py
Generating a Graph500 RMAT graph with 2^15 vertices...
Duplicates removed (or summed): 78883 and self-loops removed: 0
Generation took 0.080711s.
iteration 1: start=12627, BFS took 0.011984s, verification took
0.619967s and succeeded, TEPS=74,322,165
iteration 2: start=20502, BFS took 0.011391s, verification took
0.601116s and succeeded, TEPS=78,192,512
iteration 3: start=8839, BFS took 0.011701s, verification took
0.616723s and succeeded, TEPS=76,119,720
iteration 4: start=1999, BFS took 0.011484s, verification took
0.612136s and succeeded, TEPS=77,559,400
iteration 5: start=71, BFS took 0.011390s, verification took 0.619576s
and succeeded, TEPS=78,199,059
     [...]
iteration 62: start=32066, BFS took 0.011214s, verification took
0.605297s and succeeded, TEPS=79,427,725
iteration 63: start=27903, BFS took 0.011867s, verification took
0.608496s and succeeded, TEPS=75,055,323
iteration 64: start=16201, BFS took 0.011442s, verification took
0.609218s and succeeded, TEPS=77,843,838
Graph500 benchmark run for scale = 15
Kernel 1 time = 0.0807 seconds
Kernel 2 BFS execution times
            min_time: 1.09930038452148438e-02
  firstquartile_time: 1.14405155181884766e-02
         median time: 1.16879940032958984e-02
  thirdquartile time: 1.18589401245117188e-02
            max time: 1.21469497680664062e-02
           mean time: 1.16281397640705109e-02
         stddev_time: 2.91774617905905386e-04
```

```
Kernel 2 number of edges traversed
           min_nedge: 8.90685000000000000e+05
  firstquartile nedge: 8.90685000000000000e+05
         median_nedge: 8.90685000000000000e+05
  thirdguartile nedge: 8.9068500000000000e+05
            max nedge: 8.90685000000000000e+05
           mean nedge: 8.90685000000000000e+05
         stddev_nedge: 0.0000000000000000e+00
Kernel 2 TEPS
            min_TEPS: 7.33258156991442293e+07
  firstquartile TEPS: 7.51066276284680367e+07
         median_TEPS: 7.62052171539593935e+07
  thirdquartile_TEPS: 7.78535725269949883e+07
            max TEPS: 8.10228953379023224e+07
 harmonic_mean_TEPS: 7.65973765427299738e+07
harmonic_stddev_TEPS: 6.64094336990017182e+04
[sam@neumann test]$
```

## 5.4 Debugging

The KDT developers have developed all their Python code interactively on a single core with IPython and then executed it in parallel with Python. When parallel execution encounters bugs that don't appear serially, the primary debugging tool has been printing, either via Python's print statement or via the ParVec and SpParVec printAll methods.

## 5.5 Errata

The following bugs or anomalies are known to exist in v0.1:

- 1. The definition of the distinction between nonnull and nonzero elements in DiGraph and SpParVec instances is not completely coherent. The most glaring instance of this is SpParVec instances that represent indices, in which zero is a valid nonnull element. Because there is not a sparse Find method exposed from the Combinatorial BLAS, KDT converts a SpParVec instance to a ParVec instance before calling the dense Find from the Combinatorial BLAS. This conversion obliterates the distinction between a nonnull element whose value is zero and a null element.
- 2. The ParVec and SpParVec print methods will sometimes cause the message ADIOI\_UFS\_OPEN (line 83): \*\*io Too many open filesMPI\_FILE\_CLOSE (line 51): \*\*iobadfh Problem reading binary input file to be printed on stderr and stdout. This appears to effect only the output itself and not the correct functioning of the rest of the program.

3. The DiGraph UFget method does not act properly on files that have the Matrix Market keyword symmetric in their first (comment) line, so the resulting graphs only have half as many edges as expected. The Pajek/dictionary28 file illustrates this problem. Such a file can still be loaded properly by the following code:

```
G = kdt.UFget('Pajek/dictionary28')
G2 = G.copy()
G2.reverseEdges()
G = G+G2
```

- 4. In addition, a handful of unit tests (see files TestParVec.py, TestSpParVec.py, and TestDiGraph.py in directory test) fail:
  - a. Two tests expose that KDT does not always collapse multiple edges between the same source and destination to a single edge; TestDiGraph:test\_load and TestDiGraph:test\_DiGraph\_duplicates.
  - b. test\_UFget\_simple exposes the issue noted in item #3 above
  - c. TesteSpParVec:test\_indexing\_LHS\_SpParVec\_booleanParVec\_scal ar exposes an issue that indexing of a SpParVec by a Boolean ParVec key and a scalar value does not always produce the right numerical result.
- 5. On Windows, KDT programs that complete successfully will typically appear to fail with messages like the following:

```
job aborted:
rank: node: exit code[: error message]
0: CN01.clus.private.local: 0: process 0 exited without calling finalize
1: CN01.clus.private.local: 0: process 1 exited without calling finalize
2: CN01.clus.private.local: 0: process 2 exited without calling finalize
3: CN01.clus.private.local: 0: process 3 exited without calling finalize
4: CN02.clus.private.local: 0: process 4 exited without calling finalize
5: CN02.clus.private.local: 0: process 5 exited without calling finalize
6: CN02.clus.private.local: 0: process 6 exited without calling finalize
7: CN02.clus.private.local: 0: process 7 exited without calling finalize
8: CN03.clus.private.local: 0: process 8 exited without calling finalize
9: CN03.clus.private.local: 0: process 9 exited without calling finalize
10: CN03.clus.private.local: 0: process 10 exited without calling finalize
11: CN03.clus.private.local: 0: process 11 exited without calling finalize
12: CN04.clus.private.local: 0: process 12 exited without calling finalize
13: CN04.clus.private.local: 0: process 13 exited without calling finalize
14: CN04.clus.private.local: 0: process 14 exited without calling finalize
15: CN04.clus.private.local: 0: process 15 exited without calling finalize
```

6. On Windows, killing the execution of mpiexec via <ctl>C sometimes does not cleanly terminate the execution of all ranks on all nodes, and sometimes leaves Python-language files (e.g., "xyz.py") locked.

# Appendix A. Implementing Graph Algorithms with the Combinatorial BLAS

The Combinatorial BLAS [Buluc] is described by its authors as:

We describe the Parallel Combinatorial BLAS, which consists of a small but powerful set of linear algebra primitives specifically targeting graph and data mining applications. We provide an extendible library interface and some guiding principles for future development. The library is evaluated using two important graph algorithms, in terms of both performance and ease-ofuse. The scalability and raw performance of the example applications, using the combinatorial BLAS, are unprecedented on distributed memory clusters.

It provides C++ interfaces that are appropriate for many HPC users, but no interface from the high-level productivity languages such as Python and the M language of MATLAB™. KDT provides such an interface via Python, but the current KDT exposes just a small fraction of the power of the Combinatorial BLAS. This section describes the implementation of graph algorithms with the Combinatorial BLAS' primitives, and describes a few of the Combinatorial BLAS primitives whose full power is not yet exposed. The Combinatorial BLAS' basic structure is a sparse matrix, which KDT uses with From vertices as rows and To vertices as columns. The Combinatorial BLAS' C++ interfaces are wrapped into Python as the pyCombBLAS module, with classes pyDenseParVec, pySpParVec, and pySpParMat that correspond directly with KDT's ParVec, SpParVec, and DiGraph classes.

This information is provided so that KDT users can understand the generality of what could be implemented eventually, and so that heroic KDT users can implement their own extensions to KDT by using the pyCombBLAS directly.

## A.1 Implementing Breadth-first Search with the Combinatorial BLAS

One of the core algorithms implemented in KDT v0.1 is creating a breadth-first search tree from a graph and a root vertex. The code for this is found in DiGraph.bfsTree.

The abstract BFS algorithm starts with a "fringe" or "frontier" of vertices that has been first reached in the prior step of the algorithm, and in each step calculates all as-yet-unreached vertices that are reachable from the fringe and makes those vertices the new fringe. In sparse-matrix terms, the graph is the dual of the adjacency matrix, with destination vertices corresponding to rows and source vertices corresponding to columns as shown below. A standard sparse-matrix/vector multiplication multiplies corresponding elements of a row of the matrix and the vector and then adds those products. The operation needed for BFS is similar but different. We still want to identify positions where both the matrix and the vector have nonnulls, but instead of a multiplication the result just needs to be the position of the nonnull in the row/column (denoting which fringe vertex is the parent of the new vertex in the BFS tree), and instead of addition of multiple nonnulls in the row/column we just need to select one of them (*i.e.*, if a new vertex is reachable from multiple vertices in the fringe, in general it doesn't matter which one of those vertices is denoted as the parent in the BFS tree). The Combinatorial BLAS and KDT draw on the rich theory of linear algebra on semirings [Gilbert, p. 28-31] to implement this operation. Using the same computational structure and communication pattern but replacing the

multiplication and addition of standard matrix-vector multiplication with selection (of the column of the matrix element) and maximum, BFS achieves the needed computation.

Let's consider the example shown in, with a graph and its dual adjacency matrix, which is transposed to work properly with matrix-vector multiplication.

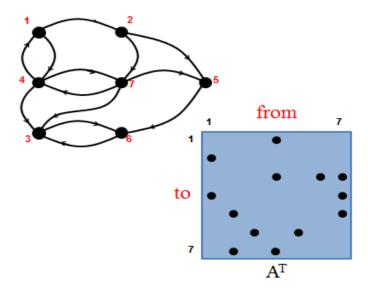


Figure 2. Illustration of bfsTree algorithm (initial state)

Let's make vertex 1 the root of the BFS tree and take the first step of the algorithm in Figure 3. The fringe vector has a nonnull only in position 1. Since the "multiplication" of row 2 of the matrix with the fringe vector results in a nonzero in position 1, element 2 of the result matrix is set nonzero, with its value equal to the value of the nonzero in the vector "product". More precisely, the algorithm selects any position with nonzeros in both in the row and the fringe vector, with the result being the value of the fringe vector element (which is previously set to its position in the element). Applying this algorithm, vertices 2 and 4 are calculated as the next level of the BFS tree (with vertex 1 as each's parent), and the red edges are added to the BFS tree. (The algorithm truncates from the new fringe any vertices that have already been visited, as with the self-loop from vertex 1 to itself.)

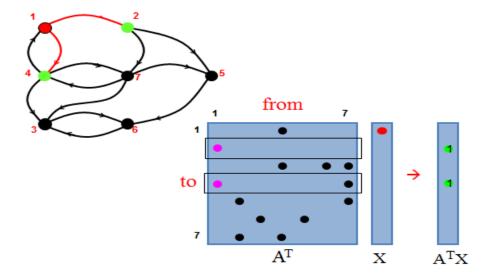


Figure 3. Illustration of bfsTree algorithm (after first iteration)

For the next step, the result column vector of the previous step is used as the fringe vector for the current step, with each nonnull element set to its position in the fringe. As illustrated in Figure 4, the same algorithm is applied. The calculation of the edge to vertex 7 illustrates the maximum step (in place of the addition in standard matrix-vector multiplication). Vertices 2 and 4 both have edges to vertex 7. The algorithm chooses the maximum-numbered vertex (vertex 4, denoted by the dark circle around the pink dot in the matrix) as the parent. Vertices 3, 5, and 7 are in the new fringe with parents of 4, 2, and 4, respectively. The resulting green edges are added to the BFS tree.

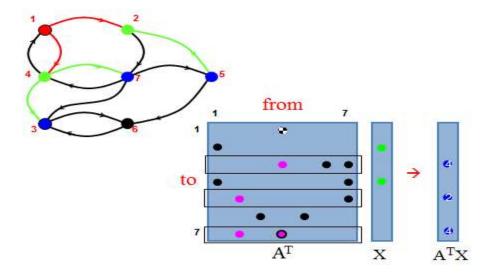


Figure 4. Illustration of bfsTree algorithm (after second iteration)

The next step of the algorithm for this graph is illustrated in Figure 5. The only as-yet-unvisited vertex is vertex 6, which is reachable from either vertex 3 or 5. The tie-breaker (maximum vertex number) determines that vertex 6's parent is vertex 5, and the blue edge is added to the BFS tree.

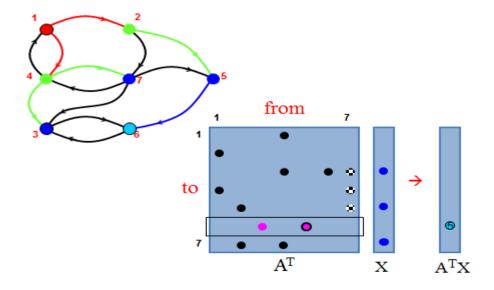


Figure 5. Illustration of bfsTree algorithm (after third iteration)

The last step of the algorithm uses the result vector of the third step, with only vertex 6 in the fringe, and detects no unreached vertices, at which point the algorithm terminates. The parent vector result is [0, 1, 1, 4, 1, 2, 3, 4]; note that the root vertex is its own parent and that in KDT vertices are numbered from 0, even though vertex 0 has no edges in this example.

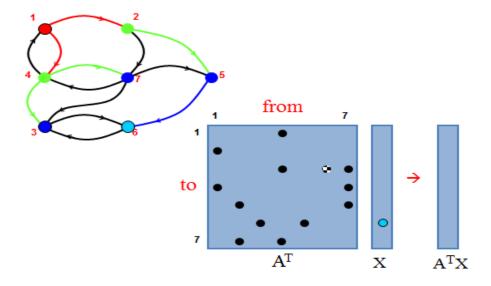


Figure 6. Illustration of bfsTree algorithm (after fourth iteration)

## A.2 Powerful Operations in the Combinatorial BLAS

The Combinatorial BLAS has several operations which have powerful generality, of which sparse-matrix/vector multiplication on semirings, illustrated in the previous section, is one. The full complement of pyCombBLAS operations is specified in the file pyCombBLAS.i in the kdt/pyCombBLAS directory.

#### A.2.1 Sparse-matrix/vector multiplication on semirings

As illustrated in the previous section, replacing the element-wise multiplication and summation reduction of the standard sparse-matrix/vector multiplication operation with other semiring operations can perform powerful operations on the graph. The standard operation is available in pyCombBLAS as SpMV\_PlusTimes. The operation used for the BFS tree calculation is available as SpMV\_SelMax. Eventually this operation will be generalized in the style of the Reduce and Apply operations described below.

## A.2.2 Applying an elemental operator

pyCombBLAS has a variety of elemental operations built-in (see the file pyCombBLAS.i for a complete list), both unary (e.g., abs) and binary (e.g., greater, or fmod with a constant). These operations can be applied to each element of a pySpParMat object by the pySpParMat.Apply method.

#### A.2.3 Applying an elemental operator with a column-specific value

pyCombBLAS can also apply these elemental operations with a second input that is column-specific. The pySpParMat.ColWiseApply method takes a second argument (a pySpParVec instance), which provides the second element for a binary operation. For instance, the DiGraph.scale method, which multiplies each out-/in-edge of a DiGraph by the corresponding element of a SpParVec, is implemented for the in-edge case as

self.spm.ColWiseApply(other.spv, pcb.multiplies())

## A.2.4 Reducing the out-edges (rows) of a DiGraph (pySpParMat) with configurable operators

pyCombBLAS has a pySpParMat.Reduce method that is configurable with the same unary and binary operations. The Reduce method, in addition to its bound pySpParMat instance, accepts a dimension (columns or rows), a binary (reduction) function, and a unary function that's applied elementally before the binary function. For example, the row-wise sum of the absolute values of the nonnull elements could be implemented by

```
ret = self.spm.Reduce(pcb.pySpParMat.Row(),pcb.plus(), pcb.abs())
```

## Appendix B. Glossary

The following terms are used with their given meanings throughout this document.

**Graph:** A collection of **vertices** and **edges** connecting the vertices.

**Edge vector:** A vector of tuples, with each tuple containing the indices of the vertices upon which an edge is incident and the weight of the edge. Represented by classes specific to the type of graph, *e.g.* DiEdgeV.

**Vertex vector**: A vector of integers, each being the index of a vertex being referred to. Represented by the DiEdgeV class generic to all types of graphs.

**DiGraph:** A collection of vertices and directed simple edges connecting pairs of vertices.

HyGraph: A collection of vertices and undirected edges, where any edge may connect any subset of the vertices.

## **Appendix C.** References

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## Appendix D. Revision History

R0.01	December 16, 2010	Initial draft
R0.02	December 20, 2010	Revisions including comments from Aydin Buluc and Drew Waranis
R0.03	January 22, 2011	Bring current with code base (almost), documenting neighbors() and pathHop() as well as several other methods.
R0.04- 05	February 12, 2011	Bring current with code base, documenting SpParVec class, many new ParVec and DiGraph methods, adding start-up and Implementing-graphs-on-CombBLAS information
R0.06	February 13, 2011	Finish Implementing-graphs-on-CombBLAS, add UFget, stipple cluster(), add DiGraph.save
R0.07	February 21, 2011	Add pageRank, toBool. Remove cluster. Fix several minor additions/deletions/errors.
R0.08	February 21, 2011	Add normalizeEdgeWeights, change names to spOnes/sprang, note that only serial work can be done inside a master() if-check.
R0.09	February 28, 2011	Clean up out-/in-edge directions and ParVec/SpParVec indexing modes.
R0.10	March 16, 2011	Add high-level note about SPMD execution; add note about UFget oddities; change SpParVec:getitem to reflect CombBLAS changes; add sort/sorted/topK for ParVec/SpParVec; add mean/std for SpParVec. Include last directory clean-up before v0.1 release and errata.
R0.11	April 13, 2011	Several typos

R0.12	August 2, 2011	Update to v0.1W content, namely the HyGraph class and Windows installation instructions.
R0.13	October 11, 2011	Update to with semantic graph content