LDBC Graphalytics: A Benchmark for Large-Scale Graph Analysis on Parallel and Distributed Platforms

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\(^\triangle\)UPC Barcelona \(^\odot\)Georgia Tech \(^\odot\)CWI Amsterdam

ABSTRACT

In this paper we introduce LDBC Graphalytics, a new industrial-grade benchmark for graph analysis platforms. It consists of six deterministic algorithms, standard datasets, synthetic dataset generators, and reference output, that enable the objective comparison of graph analysis platforms. Its test harness produces deep metrics that quantify multiple kinds of system scalability, such as horizontal/vertical and weak/strong, and of robustness, such as failures and performance variability. The benchmark comes with open-source software for generating data and monitoring performance. We describe and analyze six implementations of the benchmark (three from the community, three from the industry), providing insights into the strengths and weaknesses of the platforms. Key to our contribution, vendors perform the tuning and benchmarking of their platforms.

1. INTRODUCTION

Responding to increasingly larger and more diverse graphs, and the need to analyze them, both industry and academia are developing and tuning graph analysis software platforms. Already tens of such platforms exist, among them PowerGraph [17], GraphX [39], and PGX.D [23], but their performance is often difficult to compare. Moreover, the random, skewed, and correlated access patterns of graph analysis, caused by the complex interaction between input datasets and applications processing them, expose new bottlenecks on the hardware level, as hinted at by the large differences between Top500 and Graph500 rankings. Addressing the need for fair, comprehensive, standardized comparison of graph analysis platforms, in this work we propose the LDBC Graphalytics benchmark.

The Linked Data Benchmark Council (ldbcouncil.org, LDBC), is an industry council formed to establish standard benchmark specifications, practices and results for graph data management systems. Its goal is to inform IT professionals on the properties of the various solutions available on the market; to stimulate academic research in graph data storage, indexing, and analysis; and to accelerate the maturing process of the graph data management space as a whole. LDBC organizes a Technical User Community (TUC) that gathers benchmark input and feedback, and as such has investigated graph data management use cases across the fields of marketing, sales, telecommunication, production, publishing, law enforcement and bio-informatics. LDBC previously introduced the Social Network Benchmark [13] (SNB), which models a large social network but targets database systems (graph, SQL or SPARQL) that provide interactive updates and query answers. However, the LDBC scope goes beyond such database workloads: it also includes graph analysis frameworks that facilitate complex and holistic graph computations which may not be easily modeled as database queries, but rather as (iterative) graph algorithms, such as global metrics (e.g., diameter, triangle count) or clustering. Algorithmically analyzing large graphs is an important class of problems in “Big Data” processing, with applications such as the analysis of human behavior and preferences in social networks, root cause analysis in large-scale computer and telecommunication networks, and interactions between biological compounds and genetic structures.

In this paper, LDBC introduces Graphalytics, a benchmark for evaluating graph analysis platforms, that builds on the data generators from LDBC SNB and Graph500, making the following original contributions:

1. The first industrial-grade graph analysis benchmark specification. We carefully motivate the choice of algorithms in the benchmark, using the LDBC TUC and literature surveys to ensure good coverage of scenarios. Graphalytics consists of six core algorithms: breadth-first search, PageRank, weakly connected components, community detection using label propagation, local clustering coefficient, and single-source shortest paths. The workload includes real and synthetic datasets, which are classified into intuitive “T-shirt” sizes (e.g., XS, S, M, L, XL). The benchmarking process is made future-proof, through a renewal process.

2. A detailed process for running the benchmark. Our test harness characterizes performance and scalability with deep metrics (vertical vs. horizontal and strong vs. weak scaling), and also characterizes robustness by measuring SLA compliance, performance variability, and crash points.
3. A comprehensive tool-set developed using modern software engineering practices released as open-source benchmarking software, including a harness capable of supporting many types of target-systems, the scalable LDBC social-network generator Datagen, and the versatile Granula performance evaluation tool.

4. An extensive experimental evaluation of six state-of-the-art graph analysis systems: three community-driven (Giraph, GraphX, and PowerGraph) and three industry-driven (PGX.D, GraphMat, and OpenG). Benchmarking and tuning of the industry-driven systems in our evaluation has been performed by their respective vendors.

We describe the first three contributions, which combine the conceptual and technical specification of Graphalytics, in Section 2. The experimental evaluation is split among Section 3, which introduces the tested platforms and the benchmarking hardware, and Section 4, which presents and analyzes the real-world benchmarking results. We cover related work in Section 5, before concluding in Section 6.

2. Graphalytics

Graphalytics tests a graph analysis framework, consisting of a software platform and underlying hardware system. Graphalytics models holistic graph analysis workloads, such as computing global statistics and clustering, which run on the entire dataset on behalf of a single user.

2.1 Requirements

A benchmark is always the result of a number of design choices, responding to a set of requirements. In this section we discuss the main requirements addressed by LDBC Graphalytics:

(R1) Target platforms and systems: benchmarks must support any graph analysis platform operating on any hardware system. For platforms, we do not distinguish between programming model and support different models, including vertex-centric, gather-apply-scatter, and sparse matrix operations. For systems, we target the following environments: distributed systems, multi-core single-node systems, many-core GPU systems, hybrid CPU-GPU systems, and distributed hybrid systems. Without R1, a benchmark could not service the diverse industrial following of LDBC.

(R2) Diverse, representative benchmark elements: data model and workload selection must be representative and have good coverage of real-world practice. In particular, the workload selection must not only include datasets or algorithms because experts believe they cover known system bottlenecks (e.g., they can stress real-world systems), but also because they can be shown to be representative of the current and near-future practice. Without representativeness, a benchmark could bias work on platforms and systems towards goals that are simply not useful for improving current practice. Without coverage, a benchmark could push the LDBC community into pursuing cases that are currently interesting for the industry, but not address what could become impassable bottlenecks in the near-future.

(R3) Diverse, representative process: the set of experiments conducted by the benchmark automatically must be broad, covering the main bottlenecks of the target systems.

In particular, the target systems are known to raise various scalability issues, and also, because of deployment in real-world clusters, be prone to various kinds of failures, exhibit performance variability, and overall have various robustness problems. The process must also include possibility to validate the algorithm output, thus making sure the processing is done correctly. Without R3, a benchmark could test very few of the diverse capabilities of the target platforms and systems, and benchmarking results could not be trusted.

(R4) Include a renewal process: unlike many other benchmarks, benchmarks in this area must include a renewal process, that is, not only a mechanism to scale up or otherwise change the workload to keep up with increasingly more powerful systems (e.g., the scale parameters of Graph500), but also a process to automatically configure the mechanism, and a way to characterize the reasonable characteristics of the workload for an average platform running on an average system. Without R4, a benchmark could become less relevant for the systems of the future.

(R5) Modern software engineering: benchmarks must include a modern software architecture and run a modern software-engineering process. They must make it possible to support R1, provide easy ways to add new platforms and systems to test, and allow practitioners to easily access the benchmark and compare their platforms and systems against those of others. Without R5, a benchmark could easily become unmaintainable or unusable.

2.2 Specification of Benchmark Elements

Addressing requirement R2, the key benchmarking elements in Graphalytics are the data model, the workload selection process, and the resulting algorithms and datasets.

2.2.1 Data Model

The Graphalytics benchmark uses a typical data model for graphs; a graph consists of a collection of vertices, each identified by a unique integer, and a collection of edges, each consisting of a pair of vertex identifiers. Graphalytics supports both directed and undirected graphs. Edges in directed graphs are identified by an ordered pair (i.e., the source and destination of the edge). Edges in undirected graphs consist of unordered pairs. Every edge must be unique and connect two distinct vertices. Optionally, vertices and edges have properties, such as timestamps, labels, or weights.

To accommodate requirement R2, Graphalytics does not impose any requirement on the semantics of graphs. That is, any dataset that can be represented as a graph can be used in the Graphalytics benchmark if it is representative of real-world graph-analysis workloads.

2.2.2 Two-Stage Workload Selection Process

To achieve both workload representativeness and workload coverage, we used a two-stage selection process to select the workload for Graphalytics. The first stage identifies classes of algorithms and datasets that are representative for real-world usage of graph analysis platforms. In the second stage, algorithms and datasets are selected from the most common classes such that the resulting selection is diverse, i.e., the algorithms cover a variety of computation and communication patterns, and the datasets cover a range of sizes and a variety of graph characteristics.
**Table 1: Results of surveys of graph algorithms.**

<table>
<thead>
<tr>
<th>Graph</th>
<th>Class (selected candidates)</th>
<th>#</th>
<th>%</th>
</tr>
</thead>
<tbody>
<tr>
<td>Unweighted</td>
<td>Statistics (PR, LCC)</td>
<td>24</td>
<td>17.0%</td>
</tr>
<tr>
<td></td>
<td>Traversal (BFS)</td>
<td>69</td>
<td>48.9%</td>
</tr>
<tr>
<td></td>
<td>Components (WCC, CDLP)</td>
<td>20</td>
<td>14.2%</td>
</tr>
<tr>
<td></td>
<td>Graph Evolution</td>
<td>6</td>
<td>4.2%</td>
</tr>
<tr>
<td></td>
<td>Other</td>
<td>22</td>
<td>15.6%</td>
</tr>
</tbody>
</table>

**Weighted**

<table>
<thead>
<tr>
<th></th>
<th></th>
<th>17</th>
<th>13.4%</th>
</tr>
</thead>
<tbody>
<tr>
<td>Distances/Paths (SSSP)</td>
<td></td>
<td>17</td>
<td>13.4%</td>
</tr>
<tr>
<td>Clustering</td>
<td></td>
<td>7</td>
<td>14%</td>
</tr>
<tr>
<td>Partitioning</td>
<td></td>
<td>5</td>
<td>10%</td>
</tr>
<tr>
<td>Routing</td>
<td></td>
<td>5</td>
<td>10%</td>
</tr>
<tr>
<td>Other</td>
<td></td>
<td>16</td>
<td>32%</td>
</tr>
</tbody>
</table>

**Table 2: Mapping of dataset scale ranges to labels (“T-shirt sizes”) in Graphalytics.**

<table>
<thead>
<tr>
<th>Scale</th>
<th>Label</th>
<th>( &lt; 7 )</th>
<th>([7,7.5))</th>
<th>([7.5,8))</th>
<th>([8,8.5))</th>
<th>([8.5,9))</th>
<th>([9,9.5))</th>
<th>( \geq 9.5 )</th>
</tr>
</thead>
<tbody>
<tr>
<td>XS</td>
<td>2XS</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>LS</td>
<td>S</td>
<td>M</td>
<td>L</td>
<td>XL</td>
<td>2XL</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

**2.2.3 Selected Algorithms**

Addressing R1, according to which Graphalytics should allow different platforms to compete, the definition of the algorithms of Graphalytics is abstract. For each algorithm, we define its processing task and provide a reference implementation and reference output. Correctness of a platform implementation is defined as output equivalence to the provided reference implementation.

To select algorithms which cover real-world workloads for graph analysis platform, we have conducted two comprehensive surveys of graph analysis articles published in ten representative conferences on databases, high-performance computing, and distributed systems (e.g., VLDB, SIGMOD, SC, PPoPP). The first survey (conducted for our previous paper [20]) focused only on unweighted graphs and resulted in 124 articles. The second survey (conducted for this paper) focused only on weighted graphs and resulted in 44 articles. Table 1 summarizes the results from these surveys. Because one article may contain multiple algorithms, the number of algorithms exceeds the number of articles. In general, we found that a large variety of graph analysis algorithms are used in practice. We have categorized these algorithms into several classes, based on their functionality, and quantified their presence in literature.

Based on the results of these surveys, with expert advice from LDBC TUC we have selected the following five core algorithm for unweighted graphs, and a single core algorithm for weighted graphs, which we consider to be representative for graph analysis in general:

**Breadth-first search (BFS):** For every vertex, determines the minimum number of hops required to reach the vertex from a given source vertex.

**PageRank (PR)** [32]: Measures the rank (“popularity”) of each vertex by propagating influence between vertices using edges.

**Weakly connected components (WCC):** Determines the weakly connected component each vertex belongs to.

**Community detection using label propagation (CDLP):** Finds “communities” in the graph, i.e., non-overlapping densely connected clusters that are weakly connected to each other. We select for community detection the label propagation algorithm [34], modified slightly to be both parallel and deterministic [24].

**Table 3: Real-world datasets used by Graphalytics.**

| ID       | Name               | \( |V| \) | \( |E| \) | Scale | Domain   |
|----------|--------------------|-------|-------|-------|---------|
| R1(2XS)  | wiki-talk [26]     | 2.30M | 6.92M | 6.9    | Knowledge |
| R2(2XS)  | kgs [26]           | 0.83M | 17.9M | 7.3    | Gaming   |
| R3(2XS)  | cit-patents [26]   | 3.77M | 16.5M | 7.3    | Knowledge |
| R4(S)    | dota-league [18]   | 0.01M | 0.08M | 7.7    | Gaming   |
| R5(XL)   | com-friendster [26]| 65.6M | 1.81B | 9.3    | Social   |
| R6(XL)   | twitter_uapi [9]   | 52.6M | 1.97B | 9.3    | Social   |

**2.2.4 Selected Datasets**

Graphalytics uses both graphs from real-world applications and synthetic graphs which are generated using data generators. Table 3 summarizes the six real-world graphs included in Graphalytics. By including real-world graphs from a variety of domains, Graphalytics covers users from different communities. Our two-stage selection process led to the inclusion of graphs from the knowledge, gaming, and social network domains. Within the selected domains, graphs were chosen to span a variety of sizes and densities.

The real-world graphs in Graphalytics are complemented by two synthetic dataset generators, to enable performance comparison between different graph scales. The synthetic dataset generators are selected to cover two commonly used graphs: power-law graphs generated by Graph500, and social network graphs generated using LDBC Datagen (see Section 2.5.1). The graphs generated for the experiments are listed in Table 4.

To facilitate performance comparisons across datasets, we define the scale of a graph in Graphalytics as a function of the number of vertices \((|V|)\) and the number of edges \((|E|)\) in a graph: \( s(V,E) = \log_{10}(|V|+|E|) \), rounded to one decimal place. To give its users an intuition of what the scale of a graph means in practice, Graphalytics groups dataset scales into classes. We group scales in classes spanning 0.5 scale units, e.g., graphs in scale from 7.0 to 7.5 belong to the same class. The classes are labelled according to the familiar system of “T-shirt sizes”: small (S), medium (M), and large (L), with extra (X) prepended to indicate smaller and larger classes to make extremes such as 2XS and 3XL possible.

The reference point is class L, which is intuitively defined by the Graphalytics team to be the largest class such that
the BFS algorithm completes within an hour on any graph from that class in the Graphalytics benchmark using a state-of-the-art graph analysis platform on a single common-off-the-shelf machine. The resulting classes used by Graphalytics are summarized in Table 2.

2.3 Process

Addressing R3, the goal of the Graphalytics benchmark is to objectively compare different graph analysis platforms, facilitating the process of finding their strengths and weaknesses, and understanding how the performance of a platform is affected by aspects such as dataset, algorithm, and environment. To achieve this, the benchmark consists of a number of different experiments. In this section, we introduce these experiments, which we detail and conduct in Section 4.

The baseline experiments measure how well a platform performs for different workloads on a single machine. The core metric for measuring the performance of platforms is run-time. Graphalytics breaks down the total run-time into several components:

- **Upload time**: Time required to preprocess and convert the graph into a suitable format for a platform.
- **Makespan**: Time required to execute an algorithm, from the start of a job until termination.
- **Processing time** ($T_{\text{proc}}$): Time required to execute an actual algorithm as reported by the Graphalytics performance monitoring tool, Granula (Section 2.5.2). This does not include platform-specific overhead, such as allocating resources, loading the graph from the file system, or graph partitioning.

In our experiments we focus on $T_{\text{proc}}$ as a primary indication of the performance of a platform. We complement this metric with two user-level throughput metrics:

- **Edges per second (EPS)**: Number of edges in a graph divided by $T_{\text{proc}}$ in seconds. EPS is used in other benchmarks, such as Graph500.
- **Edges and vertices per second (EVPS)**: Number of edges plus number of vertices (i.e., $10^{10}$ scale, see Section 2.2.4), divided by $T_{\text{proc}}$ in seconds. EVPS is closely related to the scale of a graph, as defined by Graphalytics.

To investigate how well a platform performs when scaling the amount of available resources, the size of the input, or both, Graphalytics includes scalability experiments. We distinguish between two orthogonal types of scalability: strong vs. weak scalability, and horizontal vs. vertical scalability. The first category determines whether the size of the dataset is increased when increasing the amount of resources. For **strong scaling**, the dataset is kept constant, whereas for **weak scaling**, the dataset is scaled. The second category determine how the amount of resources is increased. For **horizontal scaling**, resources are added as additional computing machines, whereas for **vertical scaling** the added resources are cores within a single machine. Graphalytics expresses scalability using a single metric:

- **Speedup**: The ratio between $T_{\text{proc}}$ for scaled and baseline resources. We define the baseline for each platform and workload as the minimum amount of resources needed by the platform to successfully complete the workload.

Finally, Graphalytics assesses the **robustness** of graph analysis platforms using two metrics:

- **Stress-test limit**: The scale and label of the smallest dataset defined by Graphalytics that the system cannot process.
- **Performance variability**: The coefficient of variation (CV) of the processing time, i.e., the ratio between the standard deviation and the mean of the repeatedly measured performance. The main advantage of this metric is its independence of the scale of the results.

For all experiments, Graphalytics defines a service-level agreement (SLA): generate the output for a given algorithm and dataset with a makespan of up to 1 hour. A job breaks this SLA, and thus does not complete successfully, if its makespan exceeds 1 hour or if it crashes (e.g., due to insufficient resources).

2.4 Renewal Process

Addressing requirement R4, include in Graphalytics a renewal process which leads to a new version of the benchmark every two years. This renewal process updates the workload of the benchmark to keep it relevant for increasingly powerful systems and developments in the graph analysis community. This results in a benchmark which is future-proof. Renewing the benchmark means renewing the algorithms as well as the datasets. For every new version of Graphalytics, we follow the same two-stage workload selection process as presented in Section 2.2.2.

The algorithms of Graphalytics have been selected based on their representativeness. However, over time, graph algorithms might lose or gain popularity in the community. For example, community detection is an active field of graph research nowadays, even though our label propagation algorithm [34] was only introduced less than a decade ago. To ensure that algorithms stay relevant, for every version of the benchmark, we will select a new set of core algorithms using the same process as presented in Section 2.2.3. We will perform a new comprehensive survey on graph analysis in practice to determine new algorithm classes and select new algorithms from these classes using expert advice from LDBC TUC. If a new algorithm is found to be relevant which was not part of the set of core algorithms, it will be added. If an older core algorithm is found to be no longer relevant, it is marked as obsolete and will be removed from the specification in the next version.

The datasets of Graphalytics have been selected based on their variety in size, domain, and characteristics. Using the same process as described for algorithms, the Graphalytics team will introduce additional real-world datasets and synthetic dataset generators as they become relevant to the community. In addition, with every new version of the specification the notion of a "large" graph is reevaluated. In particular, class L is redefined as the largest class of graphs such that at a state-of-the-art platform can complete the BFS algorithm within one hour on all graphs in class L using a single common-off-the-shelf machine. The selection of platforms used to determine class L is limited to platforms implementing Graphalytics that are available to the Graphalytics team when the new specification is formalized.

2.5 Design of the Graphalytics Architecture

The Graphalytics architecture, depicted in Figure 1, consists of a number of components, including the system under test and the testing system.
As input for the benchmark, the Graphalytics team provides a benchmark description (1). This description includes definitions of the algorithms, the datasets, and the algorithm parameters for each graph (e.g., the root for BFS or number of iterations for PR). In addition, the system customer, developer, or operator can configure the benchmark (2). The benchmark user may select a subset of the Graphalytics workload to run, or they may tune components of the system under test for a particular execution of the benchmark.

The workload of Graphalytics is executed on a specific graph analysis platform (3), as provided by the user. This platform is deployed on user-provided infrastructure, e.g., on machines in a self-owned cluster or on virtual machines leased from IaaS clouds. The graph analysis platform and the infrastructure it runs on form the system under test (4). The graph analysis platform may optionally include policies that the right one is clearly better defined than the one in the paper. Both graphs exhibit a community structure (shown in different colors, detected by the Louvain method), but we see that the right one is clearly better defined than the one in the left, a consequence of the larger average clustering coefficient.

2.5.1 LDBC Datagen: Graph Generation

Graphalytics relies not only on real but also on synthetically generated graphs. Synthetic graph generators provide a means of testing data configurations not always available in the form of real datasets (e.g., due to privacy concerns). Thus, Graphalytics adopts the LDBC Social Network Benchmark Data generator (Datagen) [13], a scalable, synthetic social network generator, whose output preserves many realistic graph features: correlated data (i.e., persons with similar characteristics are more likely to be connected), skewed degree distribution (it generates a Facebook-like friendship distribution), non-uniform activity volume, etc. However, the static nature of Datagen did not allow for the generation of graphs with different degree distributions or structural characteristics. Thus, as envisioned previously [8], we have extended this work Datagen to generate graphs with these characteristics. Moreover, we have optimized the critical execution path of Datagen, to improve its performance and scalability. We summarize this two-fold contribution as follows:

Tunable Clustering Coefficient: Besides supporting different degree distributions [8], we now also allow changing the friendship generation algorithm. With the goal of generating realistic yet diverse graphs, we have implemented an edge generator which allows tuning the average clustering coefficient of the resulting friendship graph. The method relies on constructing a graph with a core-periphery community structure. Such communities are ubiquitous in social networks and their presence is strongly related to other real-world graph properties, such as a small diameter and a large connected component. Figure 2 shows two small graphs generated with Datagen, with two different target average clustering coefficients of 0.05 (left) and 0.3 (right). Both graphs exhibit a community structure (shown in different colors, detected by the Louvain method), but we see that the right one is clearly better defined than the one in the left, a consequence of the larger average clustering coefficient.

1Available at github.com/ldbc/ldbc_snbor datagen
Datagen generates friendships between persons falling in the same block. For details on how blocks are constructed please refer to LDBC SNB [13]. Given a block of persons and their expected degree, the new goal set for this work is to build communities including these persons, while maintaining the correlated nature of the produced graph (consecutive persons in a block must have a larger probability to connect) and achieving a given average clustering coefficient.

Optimization of execution flow: Figure 3 shows the old versus the new execution flow implemented for the Person-Person graph generation in Datagen. In the old flow, the output produced by step $i$ (Persons and all edges generated in steps from 0 to $i$) is read by $i + 1$, which sorts it by the corresponding correlation dimension and produces new edges. Thus, the cost of running a step grows as more edges are produced in previous steps, because more data needs to be sorted. This is exemplified in the figure with the lengths of the steps. This design guarantees that no duplicate edges are generated. In the new flow, each edge generation step is independent of the rest, and its output is written into a different file. Later, all files are merged to remove the duplicates. This approach is more efficient because the cost of executing a step remains constant, as does the amount of I/O required for the sorting. As we will see in Section 4.8, the performance improvements are significant.

2.5.2 Granula: Fine-grained Evaluation

Performance evaluation is a critical part in developing a graph analysis platform, as it helps developers gain a better understanding of the platform’s performance. However, the comprehensive evaluation of graph-analysis platforms still faces many challenges: using a coarse-grained “black-box” approach does not provide sufficient insight into the platform performance; using a fine-grained approach is too time-consuming and requires in-depth knowledge of the platform architecture; and finally it is difficult for users to apply the results of empirical performance studies for their specific use cases.

To extend Graphalytics with fine-grained performance evaluation, we developed Granula [31], a performance evaluation framework consisting of three main modules: the modeler, the archiver, and the visualizer.

Modeler: Fine-grained evaluation of graph-analysis platforms requires domain-specific expertise and can be time-consuming. The Granula modeler allows experts to explicitly define once their evaluation method for a graph analysis platform, such that the evaluation process can be fully automated. This includes defining phases in the execution of a job (e.g., graph loading), and recursively defining phases as a collection of smaller, lower-level phases (e.g., graph loading includes reading and partitioning), up to the required level of granularity. The performance model may also include other information, such as the number of vertices processed in a phase.

Archiver: The Granula archiver uses the performance model of a graph analysis platform to collect and archive detailed performance information for a job running on the platform. Such information is either gathered from log files produced by the platform, or derived using rules defined in the performance model. The archiver produces a performance archive which encapsulates the comprehensive set of performance information captured for each job. The archive is complete (i.e., all observed and derived results are included), descriptive (i.e., all results are described to non-experts) and examinable (i.e., all results are derived from a traceable source).

Visualizer: While a performance archive is sufficiently informative, it is not the most natural way of examining performance results. The Granula visualizer presents the performance archive in a human-readable manner and allows efficient navigation through the performance results at varying levels of granularity using an interactive Web interface. Results presented using the Granula visualizer can be easily communicated and shared among performance analysts with different levels of expertise.

In this work, we use Granula to build comprehensive performance models for several graph processing platforms, e.g., Apache Giraph (see detailed analysis in our technical report [24]). And for each platform, we have developed a basic performance model which allow us to define, capture, and report fine-grained performance breakdown metrics, e.g., processing time (See Section 2.3).

3. EXPERIMENTAL SETUP

A major contribution of this work is the evaluation and comparison of graph analysis platforms whose development is industry-driven, and of other, community-driven, platforms. In this section, we present the setup of our experiments.

3.1 Selected Platforms

We evaluate and compare in this work six different graph analysis platforms, three community-driven (C) and three industry-driven (I), see Table 5. These platforms are based on six different programming models, spanning an important design space for real-world graph analysis.

The platforms can be categorized into two classes: distributed (D) and non-distributed (S) platforms. Distributed platforms use when analyzing graphs multiple machines connected using a network, whereas non-distributed platforms can only use a single machine. Distributed systems suffer from a performance penalty because of network communication, but can scale to handle graphs that do not fit into the memory of a single machine. Non-distributed systems cannot scale as well because of the limited amount of resources of a single machine.

Table 5: Selected graph analysis platforms.

<table>
<thead>
<tr>
<th>Type</th>
<th>Name</th>
<th>Vendor</th>
<th>Lang.</th>
<th>Model</th>
<th>Vers.</th>
</tr>
</thead>
<tbody>
<tr>
<td>C, D</td>
<td>Giraph [28]</td>
<td>Apache</td>
<td>Java</td>
<td>Pregel</td>
<td>1.1.0</td>
</tr>
<tr>
<td>C, D</td>
<td>GraphX [39]</td>
<td>Apache</td>
<td>Scala</td>
<td>Spark</td>
<td>1.6.0</td>
</tr>
<tr>
<td>C, D</td>
<td>PowerGraph [17]</td>
<td>CMU</td>
<td>C++</td>
<td>GAS</td>
<td>2.2</td>
</tr>
<tr>
<td>I, S/D</td>
<td>GraphMat [34]</td>
<td>Intel</td>
<td>C++</td>
<td>SpMV</td>
<td>Feb ’16</td>
</tr>
<tr>
<td>I, S</td>
<td>OpenG [30]</td>
<td>Intel</td>
<td>C++</td>
<td>Native code</td>
<td>Feb ’16</td>
</tr>
<tr>
<td>I, D</td>
<td>PGX.D [23]</td>
<td>Oracle</td>
<td>C++</td>
<td>Push-pull</td>
<td>Feb ’16</td>
</tr>
</tbody>
</table>

2Available at github.com/tudelft-atlarge/granula
Table 6: Experiments used for benchmarks.

<table>
<thead>
<tr>
<th>Category</th>
<th>Sec.</th>
<th>Experiment</th>
<th>Algorithms</th>
<th>Datasets</th>
<th>#nodes</th>
<th>#threads</th>
<th>Metric</th>
</tr>
</thead>
<tbody>
<tr>
<td>Baseline</td>
<td>4.1</td>
<td>Dataset variety</td>
<td>BFS, PR</td>
<td>All, up to L</td>
<td>1</td>
<td>-</td>
<td>(T_{\text{proc}}, E(V))PS</td>
</tr>
<tr>
<td></td>
<td>4.2</td>
<td>Algorithm variety</td>
<td>All</td>
<td>R4(S), D300(L)</td>
<td>1</td>
<td>-</td>
<td>(T_{\text{proc}})</td>
</tr>
<tr>
<td>Scalability</td>
<td>4.3</td>
<td>Vertical</td>
<td>BFS, PR</td>
<td>D300(L)</td>
<td>1</td>
<td>1-32</td>
<td>(T_{\text{proc}}, S)</td>
</tr>
<tr>
<td></td>
<td>4.4</td>
<td>Strong/Horizontal</td>
<td>BFS, PR</td>
<td>D1000(XL)</td>
<td>1-16</td>
<td>-</td>
<td>(T_{\text{proc}}, S)</td>
</tr>
<tr>
<td></td>
<td>4.5</td>
<td>Weak/Horizontal</td>
<td>BFS, PR</td>
<td>G22(S)-26(XL)</td>
<td>1-16</td>
<td>-</td>
<td>(T_{\text{proc}}, S)</td>
</tr>
<tr>
<td>Robustness</td>
<td>4.6</td>
<td>Stress test</td>
<td>BFS</td>
<td>All</td>
<td>1</td>
<td>-</td>
<td>SLA</td>
</tr>
<tr>
<td></td>
<td>4.7</td>
<td>Variability</td>
<td>BFS</td>
<td>D300(M), D1000(L)</td>
<td>1, 16</td>
<td>-</td>
<td>CV</td>
</tr>
<tr>
<td>Self-Test</td>
<td>4.8</td>
<td>Data Generation</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>(T_{\text{gen}})</td>
</tr>
</tbody>
</table>

Apache Giraph [28] uses an iterative vertex-centric programming model similarly to Google’s Pregel. Giraph is open source and built on top of Apache Hadoop’s MapReduce.

Apache GraphX [39] is an extension of Apache Spark, a general platform for big data processing. GraphX extends Spark with graphs based on Spark’s Resilient Distributed Datasets (RDDs).

PowerGraph [17], developed by Carnegie Mellon University, is designed for real-world graphs which have a skewed power-law degree distribution. PowerGraph uses a programming model known as Gather-Apply-Scatter (GAS).

GraphMat [37, 4], developed by Intel, maps Pregel-like vertex programs to high-performance sparse matrix operations, a well-developed area of HPC. GraphMat supports two different backends which need to be selected manually: a single-machine shared-memory backend [37] and a distributed MPI-based backend [4].

OpenG [30] consists of handwritten implementations for many graph algorithms. OpenG is used by GraphBIG, a benchmarking effort initiated by Georgia Tech and inspired by IBM System G.

PGX.D [23], developed by Oracle, enables vertices to “pull” (read) data from neighbors, as opposed to conventional graph analysis systems which only allow vertices to “push” (write) data. Additionally, PGX.D uses a fast cooperative context-switching mechanism and focuses on low-overhead, bandwidth-efficient network communication.

3.2 Environment

Experiments have been performed on the DAS-5 [1] (Distributed ASCII Supercomputer), consisting of 6 clusters with over 200 dual 8-core compute nodes. DAS-5 is funded by a number of organizations and universities from the Netherlands and is actively used as a tool for computer science research in the Netherlands. We use individual clusters for the experiments and we test all platforms on the same hardware. The hardware specifications of the machines in the clusters are listed in Table 7.

4. EXPERIMENTAL RESULTS

Graphalytics conducts automatically the complex set of experiments summarized in Table 6. The experiments are divided into four categories: baseline, scalability, and robustness (all introduced in Section 2.3); and self-test. Each category consists of a number of experiments, for which Table 6 lists the parameters used for the benchmarks (algorithm, dataset, number of machines, and number of threads) and the metrics used to quantify the results.

4.1 Dataset Variety

For this experiment, Graphalytics reports the processing time of all platforms executing BFS and PageRank on a variety of datasets using a single node. Key findings:

- GraphMat and PGX.D significantly outperform their competitors in most cases.
- PowerGraph and OpenG are roughly an order of magnitude slower than the fastest platforms.
- Giraph and GraphX are consistently two orders of magnitude slower than the fastest platforms.
- Across datasets, all platforms show significant variability in performance normalized by input size.

The workload consists of two selected algorithms and all datasets up to class L. We present the processing time (\(T_{\text{proc}}\)) in Figure 4, and the processed edges per second (EPS) and processed edges plus vertices per second (EVPS) in Figure 5. The vertical axis in both figures lists datasets, ordered by scale (results for missing datasets are available in our technical report [24]).

Figure 4 depicts the processing time of BFS and PageRank for all platforms on a variety of datasets. For both algorithms, GraphMat and PGX.D are consistently fast. Giraph and GraphX are the slowest platforms and both are two orders of magnitude slower than GraphMat and PGX.D for most datasets. Finally, OpenG and PowerGraph are generally slower than both PGX.D and GraphMat, but still significantly faster than Giraph and GraphX. A notable exception is OpenG’s performance for BFS on dataset R2(XS).

The BFS on this graph covers approximately 10% of the vertices in the graph, so OpenG’s queue-based BFS implementation results in a large performance gain over platforms that process all vertices using an iterative algorithm.

To better understand the sensitivity of the tested platforms to the datasets, we present normalized processing times for the BFS algorithm in Figure 5. The left and right subfigures depict EPS and EVPS, respectively. Ideally, a platform’s performance should be directly related to the size of the graph, thus the normalized performance should be close to constant. As evident from the figure, all platforms show signs of dataset sensitivity, as EPS and EVPS vary between datasets.

Table 7: Hardware specifications.

<table>
<thead>
<tr>
<th>Component</th>
<th>Name</th>
</tr>
</thead>
<tbody>
<tr>
<td>CPU</td>
<td>2 × Intel Xeon E5-2630 @ 2.40 GHz</td>
</tr>
<tr>
<td>Cores</td>
<td>16 (32 threads with Hyper-Threading)</td>
</tr>
<tr>
<td>Memory</td>
<td>64 GiB</td>
</tr>
<tr>
<td>Disk</td>
<td>2 × 4 TB</td>
</tr>
<tr>
<td>Network</td>
<td>1 Gbit/s Ethernet, FDR InfiniBand</td>
</tr>
</tbody>
</table>

Table 7: Hardware specifications.
Besides $T_{\text{proc}}$, it is also interesting to look at the makespan (i.e., time spent on the complete job for one algorithm). This includes platform-specific overhead such as resource allocation and graph loading. Table 8 lists the makespan, $T_{\text{proc}}$, and their ratio. The percentages show that the overhead varies widely for the different platforms and is between 66% and 99.8% of the makespan. However, we note that the platforms have not been tuned to minimize this overhead and in many cases it could be significantly reduced by optimizing the configuration.

### 4.2 Algorithm Variety

The second set of baseline experiments focuses on the algorithm variety in the Graphalytics benchmark, and on how the performance gap between platforms varies between workloads. Note that all algorithms are implemented for all platforms, with the exception of LCC for PGX.D. Key findings:

- Relative performance between platforms is similar for BFS, WCC, PR, and SSSP.
- LCC is significantly more demanding than the other algorithms, only OpenG and PowerGraph complete it without breaking the SLA.
- PGX.D’s performance on WCC degrades on a graph with a large number of components.
- OpenG performs best on CDLP, whereas GraphX is unable to complete CDLP. Other platforms perform similarly on CDLP.

*Figure 6* depicts $T_{\text{proc}}$ for the core algorithms in Graphalytics on two graphs with edge weights: R4(S), the largest real-world graph in Graphalytics with edge-weights; and D300(L). BFS, WCC, PR, and SSSP all show similar results. PGX.D and GraphMat are the fastest platforms. GraphMat can run either the S or D backend, but does not select so autonomously; SSSP is not supported in S, so we use D only for this algorithm. Giraph, GraphX and PowerGraph are much slower, with GraphX showing the worst performance, especially on D300(L). OpenG’s performance is close to that of PGX.D and GraphMat on WCC, and up to an order of magnitude worse for BFS, PR, and SSSP. The processing times for CDLP are much closer for all platforms. OpenG performs best on CDLP, and GraphX fails to complete even on R4(S). LCC fails for all platforms on the graphs used in this experiment, except for OpenG and PowerGraph. The complexity of the LCC algorithm depends on the degrees of vertices, so failures are expected on dense graphs, such as R4(S), and on large graphs, such as D300(L).

**Table 8:** $T_{\text{proc}}$ and makespan for BFS on DG300(L).

<table>
<thead>
<tr>
<th>Algorithm</th>
<th>GraphX</th>
<th>P’Graph</th>
<th>G’Mat(S)</th>
<th>OpenG</th>
<th>PGX.D</th>
</tr>
</thead>
<tbody>
<tr>
<td>Time (s)</td>
<td>205.3 s</td>
<td>22.3 s</td>
<td>22.8 s</td>
<td>4.8 s</td>
<td>205.3 s</td>
</tr>
<tr>
<td>Makespan</td>
<td>276.6 s</td>
<td>101.5 s</td>
<td>2.1 s</td>
<td>0.3 s</td>
<td>0.5 s</td>
</tr>
<tr>
<td>$T_{\text{proc}}$</td>
<td>22.3 s</td>
<td>21.7 s</td>
<td>0.3 s</td>
<td>0.5 s</td>
<td>205.3 s</td>
</tr>
<tr>
<td>Ratio</td>
<td>8.1%</td>
<td>34.0%</td>
<td>9.9%</td>
<td>1.4%</td>
<td>33.3%</td>
</tr>
</tbody>
</table>

**Figure 4:** Dataset variety: $T_{\text{proc}}$ for BFS and PR.

**Figure 5:** Dataset variety: EPS and EVPS for BFS.

**Figure 6:** Algorithm variety: $T_{\text{proc}}$. 
4.3 Vertical Scalability

To analyze the effect of adding additional resources in a single machine, we use Graphalytics to run the BFS and PageRank algorithms on D300(L) with 1 up to 32 threads on a single machine. Key findings:
- All platforms benefit from using additional cores, but only PGX.D and GraphMat approach optimal efficiency.
- Most platforms experience minor or no performance gains from Hyper-Threading.

Figure 7 depicts the processing time for this experiment. The majority of tested platforms show increasing performance as threads are added up to 16, the number of cores. Adding additional threads up to 32, the number of threads with hardware support through Hyper-Threading, does not appear to improve the performance of GraphX, GraphMat, or OpenG. Giraph and PGX.D benefit slightly from 32 threads. The maximum speedup obtained by each platform is summarized in Table 9. Overall, PGX.D scales best with a maximum speedup of 15.0. This means that PGX.D has the same performance for PageRank as OpenG when using 32 threads, even though PGX.D is significantly slower than OpenG for a single thread.

4.4 Strong Horizontal Scalability

We use Graphalytics to run BFS and PR for all distributed platforms on D1000(XL) while increasing the number of machines from 1 to 16 in powers of 2 to measure strong scalability. Key findings:
- PGX.D and GraphMat show a reasonable speedup.
- Giraph’s performance degrades significantly when switching from 1 machine to 2 machines, but improves significantly with additional resources.
- PowerGraph and GraphX scale poorly; GraphX shows no performance increase past 4 machines.

The processing times for this experiment are depicted in Figure 8. Ideally, $T_{proc}$ halves when the amount of resources (i.e., the number of machines) is doubled given a constant workload. Giraph suffers a large performance hit when switching from 1 machine to a distributed setup with 2 machines. For PR, this results in an SLA failure on 2 machines, even though it succeeds on 1 machine. At least 4–8 machines are required for Giraph to improve in performance over the single-machine setup. The speedup obtained by Giraph is 3.3 on BFS and 5.3 on PR. GraphX also scales poorly for the given workload. It requires 2 machines to complete BFS, and 4 machines to complete PR. GraphX achieves a speedup of 2.3 using 8 times as many resources on BFS, and a speedup of 1.2 with 4 times as many resources on PR. PowerGraph is able to process the D1000(XL) graph on any number of nodes, but only scales reasonably for BFS and poorly for PR. Its speedup is 6.9 for BFS and 1.8 for PR, compared to the ideal of 16. PGX.D fails to complete either algorithm on a single machine. For BFS, PGX.D scales poorly past 4 nodes, where it already achieves sub-second processing times. PGX.D scales better on PR, up to a speedup of 3.8 using 8 times as many resources as the baseline. GraphMat shows a clear outlier for PR on a single machine, most likely because of swapping.

4.5 Weak Horizontal Scalability

To measure weak scalability, Graphalytics runs BFS and PR for all distributed platforms on Graph500 G22(S) through G26(XL) while increasing the number of machines from 1 to 16 in powers of 2. The amount of work per machine is approximately constant, as each graph in the series generated using Graph500 is twice as large as the previous graph. As the workload per machine is constant, $T_{proc}$ is ideally constant. Key findings:
- None of the tested platforms achieve optimal weak scalability.
- Giraph’s performance degrades significantly on 2 machines, but scales well from 4 to 16 machines.
- GraphMat and PowerGraph scale reasonably, whereas GraphX scales poorly.
- PGX.D fails in multiple configurations due to memory limitations.

In Figure 9, GraphX and PowerGraph show increasing
processing times as the number of machines increases, peaking at a maximum slowdown (i.e., inverse of speedup) of 15.2 and 8.2, respectively. Similar to the strong scalability experiments, Giraph’s performance is worst with two machines and a maximum slowdown of 13.3 on PR. Performance improves slightly as more machines are added, for a slowdown of 6.4 with 16 machines on PR. PGX.D shows best performance and even fails to complete many experiments due to memory limitations. PGX.D’s performance in the distributed experiments could be improved by using a different graph partitioning scheme.

### 4.6 Stress Test

To test the maximum processing capacity of each platform, we use Graphalytics to run the BFS algorithm on all datasets, and report the scale of the smallest dataset that breaks the SLA (Section 6) on a single machine. Key findings:

- GraphX and PGX.D fail to process the largest class L graph on a single machine.
- Most platforms fail on a Graph500 graph, but succeed on a Datagen graph of comparable scale. This indicates sensitivity to graph characteristics other than graph size.
- PowerGraph and OpenG can process the largest graphs on a single machine, up to scale 9.0.

Table 10 lists the smallest graph, by scale, for which each platform fails to complete BFS. The results show that both GraphX and PGX.D are unable to complete the BFS algorithm on Graph500 scale 25, a class L graph. PGX.D is specifically optimized for machines with large amount of cores and memory, and thus exceeds the memory capacity of our machines. PGX.D can be tuned to be more memory-efficient, but does not do so autonomously. Like GraphX and PGX.D, Giraph and GraphMat fail on a Graph500 graph. Both platforms successfully process D1000 with scale 9.0, but fail on G26 of the same scale. This suggests that characteristics of the graphs affect the performance of graph analysis platforms, an issue not revealed by the Graph500 benchmark. Finally, PowerGraph and OpenG fail to complete BFS on the Friendster graph, a scale 9.3 graph and among the largest graphs currently used by Graphalytics.

### 4.7 Variability

To test the variability in performance of each platform, Graphalytics runs BFS 10 times on D300(L) with 1 machine for all platforms, and on D1000(XL) with 16 machines for the distributed platforms.

- All platforms have a CV of at most 10%, i.e., their standard deviation is at most 10% of the mean $T_{proc}$.
- GraphMat and PGX.D show higher than average variability in both S and D configurations. However, due to their much smaller mean, the absolute variability is small.

The mean and CV for $T_{proc}$ are reported in Table 11. In both S and D configurations, PowerGraph shows the least variability in performance. GraphX has similarly low variability, but due to its significantly longer mean processing time it can deviate by tens of seconds between runs. Conversely, GraphMat and PGX.D show much larger variability between runs, but in absolute values their deviation is limited to tens of milliseconds.

### 4.8 Data Generation

We also evaluate the performance and scalability of Datagen with the new execution flow presented in Section 2.5.1, by running several experiments using different cluster configurations and scale factors. We compare this version of Datagen (v0.2.6) to the latest version not including the performance optimizations discussed in this paper (v0.2.1). We use the previous generation of DAS, DAS-4 (2 Intel Xeon E5620, 24 GiB RAM, spinning disks, 1 Gbit/s Ethernet), to perform the experiments, which leads to conservative results.

We configure Datagen to run on top of a Hadoop 2.4.1 cluster. For each given cluster configuration, one machine is kept as master and does just management tasks, while the rest are workers. The number of mappers is controlled by Hadoop, and depends on the size of the input files of each Hadoop job. The number of reducers is set to 6 per worker (1 per core); e.g., when using 16 machines, we have 1 master and 15 workers, executing 90 reducers in total. Finally, scale factors reflect the approximate number of generated edges in millions.

Figure 10 (l) shows the execution time of Datagen v0.2.6 and v0.2.1 in a 16-machine configuration for five different scale factors: 30, 100, 300, 1000 and 3000. Datagen v0.2.6 is able to improve 1.16x, 1.33, 1.83, 2.15 and 2.9x the execution time with respect to version v0.2.1, respectively. Not only the new version is faster but also the speedup shows a clear increasing trend with the scale factor. Thus, we have not only improved the actual execution time, but also the scalability. Overall, Datagen v0.2.6 takes just 44 minutes to generate a billion edge graph using 16 machines, which represents a significant improvement over the 95 minutes required by v0.2.1.

Figure 10 (r) shows the execution time of Datagen v0.2.6 for different cluster sizes and scale factors (i.e., #edges). On the one hand, we see that Datagen scales very well. In-
creasing the scale factor size by a factor of 10 (from 1000 to 10000), increases the execution time by 10.6. As a result, we are able to generate a 10 billion graph in less than 8 hours, using hardware from 2010. The worse scalability observed for smaller scale factors is due to the overhead incurred by Hadoop when spawning the jobs, which becomes more negligible the larger the scale factor is. On the other hand, we also observe very good horizontal scalability. The speedup obtained from going from 4 to 16 machines, for scale factors 30, 100, 300 and 1000 is 1.1, 1.4, 2.0 and 3.0, respectively. This means we can add more hardware to generate larger datasets faster. To sum up, we see that with Datagen, we can generate large and complex graphs with reasonable amounts of time using small sized clusters with commodity hardware.

5. RELATED WORK

Table 12 summarizes and compares Graphalytics with previous studies and benchmarks for graph analysis systems. R1–R5 are the requirements formulated in Section 2.1. As Table 12 indicates, there is no alternative to Graphalytics in covering requirements R1–R4. We also could not find evidence of requirement R5 being covered by other systems than LDBC. While there have been a few related benchmark proposals (marked “B”), these either do not focus on graph analysis, or are much narrower in scope (e.g., only BFS for Graph500). There have been comparable studies (marked “S”) but these have not attempted to define—let alone maintain—a benchmark, its specification, software, testing tools and practices, or results. Graphalytics is not only industry-backed but also has industrial strength, through its detailed execution process, its metrics that characterize robustness in addition to scalability, and a renewal process that promises longevity. Graphalytics is being proposed to SPEC as well, and BigBench [16, 33] explicitly refers to Graphalytics as its option for future benchmarking of graph analysis platforms.

Previous studies typically tested the open-source platforms Giraph [28], GraphX [39], and PowerGraph [17], but our contribution here is that vendors (Oracle, Intel, IBM) in our evaluation have themselves tuned and tested their implementations for PGX.D [23], GraphMat [37] and OpenG [30]. We are aware that the database community has started to realize that with some enhancements, RDBMS technology could also be a contender in this area [14, 25], and we hope that such systems will soon get tested with Graphalytics.

Graphalytics complements the many existing efforts focusing on graph databases, such as LinkedBench [5], XGDBench [10], and LDBC Social Network Benchmark [13]; efforts focusing on RDF graph processing, such as LUBM [19], the Berlin SPARQL Benchmark [7], SP² Benchmark [36], and WatDiv [2] (targeting also graph databases); and community efforts such as the TPC benchmarks. Whereas all these prior efforts are interactive database query benchmarks, Graphalytics focuses on algorithmic graph analysis and on different platforms which are not necessarily database systems, whose distributed and highly parallel aspects lead to different design trade-offs.

6. CONCLUSION

Responding to an increasing use of large-scale graphs, industry and academia have proposed a variety of distributed and highly-parallel graph analysis platforms. To compare these platforms, but also to tune them and to enable future designs, the Linked Data Benchmark Council (LDBC) has been tasked by its industrial constituency to develop an offline (batch) graph analysis workload—the LDBC Graphalytics benchmark, which is the focus of this work. We believe Graphalytics brings both conceptual and technical contributions. We present implementations on the main community-driven graph analysis platforms, as well as three vendor-tuned platforms.

The Graphalytics workload was designed through a two-stage selection incorporating the concept of choke-point (expertise-driven) design, and a data-driven selection of relevant algorithms and input datasets. The specification of Graphalytics is innovative: its metrics go beyond the traditional performance metrics, and in particular enable deep studies of two key features of distributed and highly-parallel systems, scalability and robustness. Graphalytics is the first graph benchmark to cover stress-testing and performance variability. The benchmarking process is managed by an advanced harness, which includes flexible and scalable tools for data collection, analysis, and sharing, and for distributed generation of synthetic yet realistic graph datasets. In particular, the data generator tool Datagen is the first to generate graphs with a pre-specified clustering coefficient for benchmarking. Graphalytics also specifies a novel process for renewing its core parameters, to withstand the test of time while still being understandable for non-experts.

We present here the open-source implementation of the harness3, which is able to conduct over ten different experiments, collect in-depth data that can be further used for tuning, and then extract the relevant benchmarking metrics. We also provide reference implementations of the drivers and algorithms for six target systems, which differ widely in distribution and programming model. Three of the systems originate from industry: PGX.D from Oracle, GraphMat from Intel, and Open G from IBM. We hope and believe Graphalytics to be interesting for academics, IT practitioners, industry engineers, and system designers.

7. REFERENCES


3Available at github.com/tudelft-atlarge/graphalytics
Table 12: Summary of related work. (Acronyms: Reference type: S, study, B, benchmark. Target system, structure: D, distributed system; P, parallel system; MC, single-node multi-core system; GPU, using GPUs. Input: 0, no parameters; S, parameters define scale; E, parameters define edge properties; +, parameters define other graph properties, e.g., clustering coefficient. Datasets/Algorithms: Rnd, reason for selection not explained; Exp, selection guided by expertise; 1-stage, data-driven selection; 2-stage, 2-stage data- and expertise-driven process. Scalability tests: W, weak, S, strong, V, vertical, H, horizontal.)

<table>
<thead>
<tr>
<th>Name [Publication]</th>
<th>Target System (R1)</th>
<th>Design (R2)</th>
<th>Tests (R3)</th>
<th>(R4)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Structure</td>
<td>Programming</td>
<td>Input</td>
<td>Datasets</td>
</tr>
<tr>
<td>B CloudSuite [10], only graph elements</td>
<td>D/MC</td>
<td>PowerGraph</td>
<td>S</td>
<td>Rnd</td>
</tr>
<tr>
<td>S Montresor et al. [12]</td>
<td>D/MC</td>
<td>3 classes</td>
<td>0</td>
<td>Rnd</td>
</tr>
<tr>
<td>B Graph5000 [9]</td>
<td>P</td>
<td>S</td>
<td>Exp</td>
<td>Exp</td>
</tr>
<tr>
<td>B GreenGraph500</td>
<td>P/MC/GPU</td>
<td>—</td>
<td>S</td>
<td>Exp</td>
</tr>
<tr>
<td>B WGB [8]</td>
<td>D</td>
<td>—</td>
<td>SE</td>
<td>Exp</td>
</tr>
<tr>
<td>S Own prior work [20, 21, 8]</td>
<td>D/MC/GPU</td>
<td>10 classes</td>
<td>S</td>
<td>Exp</td>
</tr>
<tr>
<td>B BigDataBench [29, 38], only graph elements</td>
<td>D/MC</td>
<td>Hadoop</td>
<td>S</td>
<td>Rnd</td>
</tr>
<tr>
<td>S Satish et al. [35]</td>
<td>D/MC</td>
<td>6 classes</td>
<td>S</td>
<td>Exp</td>
</tr>
<tr>
<td>S Yi et al. [25]</td>
<td>D</td>
<td>4 classes</td>
<td>S</td>
<td>Exp</td>
</tr>
<tr>
<td>B GraphBG [30]</td>
<td>P/MC/GPU</td>
<td>System G</td>
<td>S</td>
<td>Exp</td>
</tr>
<tr>
<td>S Chekinaova et al. [11]</td>
<td>MC</td>
<td>Galois</td>
<td>0</td>
<td>Rnd</td>
</tr>
</tbody>
</table>

H LDBC Graphalytics (this work) | D/MC/GPU | 10+ classes | SE+ | 2-stage | 2-stage | Process | W/S/V/H | Yes | Yes |


[38] N. Sundaram et al. Graphmat: High performance graph analytics made productive. PVLDB, 8(11), 2015.
