Big data for all!

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Abstract

The modern internet generates petabytes of data per day. Processing vast amounts of data is an increasingly common task both for scientists and modestly experienced programmers. Often this data is naturally represented as a graph, such as social media networks, webpage links or city networks, and requires clusters of machines to process. Concurrent trends in data centre architecture suggest that the rack is the new server, and shared memory is now a feasible interface between collocated rack servers. These trends made us wonder: *How simple can fast graph processing be on a rack of servers?*. We investigated the tradeoffs of the conventional Pregel *"think like a vertex"* programming model, and found its performance unacceptable. In contrast, we explored the merits of a *"Think like a subgraph"* model, which respects graph locality in common graphs, provides a more holistic programming interface, and runs faster!

1 Introduction

Big data processing is complicated. Scientists and experienced programmers alike struggle with managing and configuring clusters of machines to processes large amounts of data. Frameworks like Hadoop and Pregel have significantly eased the difficulty of big data processing, but they remain intimidating for the layman. We noticed a trend in scientists and researchers, finding that they wanted to *"Just write python code that ran on a bunch of computers"*. For the benefit of science we investigated how to make this dream a reality.

Running all Python code on clusters of machines is impractical would lead to sluggish un-optimal code, due to immense network overhead. Instead, we concentrated our efforts on a common but difficult big data processing task: graph processing. Many frameworks exist for distributed graph processing [18, 6, 15, 17, 23, 9], and many general frameworks exist which are used for graph processing [22, 24, 13, 20]. These frameworks vary in their complexity, but none are *"accessible"* for programmers that lack relevant experience.

With the exception of [15], the aforementioned systems suffer a common pitfall to accessibility; they expose the complexity of a distributed message passing system to the user. Extensive work has been done to hide this complexity in the abstraction of distributed shared memory (DSM) [14, 21, 19, 10, 12]. The benefits of DSM have been ignored in recent years due to its flaws, most notably fate sharing and sub optimal performance. Dismissing DSM may have been a shortsighted mistake. Ultra dense memory and the approach of terabit-level bandwidth within a rack give modern racks the appearance of a single machine, and has lead towards disaggregated architectures [1, 2, 3, 5, 11]. Such futuristic systems lend themselves naturally to DSM which motivates our proposal for a corresponding computation framework.

The largest disadvantage of DSM is performance. Programmers can write terribly performant programs by failing to reason about the location of memory, leading to memory thrashing. In computation where a high degree of consistency between shared resources is needed, DSM is the wrong tool for the job. In contrast, when large amounts of computation can be performed between memory synchronizations, DSM provides a simple and efficient programming model. Graph processing suffers from a lack of locality. In computations such as PageRank, a single iteration may perform edge updates which require the synchronization of all machines in a cluster. This problem can be largely avoided in practice by carefully pre-processing graphs into equally-sized partitions, where the minimum number of graph edges exist across machines. The cost of pre-processing a graph can be large; in some cases, the complexity of finding a good partition is greater than solving the initial problem! Here we demonstrate that the cost of graph partitioning is worth it for the benefits that DSM provides.

In this paper we attack the problem of developing a simple and efficient graph processing interface for DSM. Specifically we make the following contributions.

- A simple graph processing API
- A graph partitioning scheme optimal for DSM

- An evaluation of processing performance between partitioned DSM processing, and Pregel-style graph processing

The remainder of this paper is organized as follows. In Section 2 we over view related work. In Section 3 we describe our graph processing API. Section 4 we describe our approach to graph partitioning.In Section 5 we evaluate our framework against a Pregel style *think like a vertex model*. Section 6 describes our experiences with our system, and Section 7 concludes the paper.

2 Related work

Distributed shared memory has been previously explored in several domains

3 API

4 Partitioning

Our partitioning algorithm aims to provide an optimal environment for the parallel processing of graphs. We leverage the METIS [16] library, an MPI implementation for parallel partitioning of large graphs, to determine an optimal partitioning for the system. Metis is built in and runs in C, and the corresponding Python binding, PyMetis [4] was used to determine the optimal graph partitioning it is read into memory.

Naturally, partitioning a graph will cause edges in the graph to cross partitions. Applying techniques seen in [8] and [7], we use a "mirroring" technique for storing edges between partitions. For each edge that crosses partitions, only vertices on two separate partitions will be affected. The partition which contains the destination vertex will be considered the master of the given edge. All edges which connect to the master vertex from a different partition will create a "mirror" vertex. These edges are stored in shared memory, such that they can be written to and read from different machines.

5 Evaluation

6 Discussion

7 Conclusion

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