Parallel Breadth First Search for Scale-Free Social Network Graphs

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***Abstract*—Graphical representations are one of the most commonly used abstractions to model large networks. Social media networks like Facebook and Twitter contain massive scale free networks with billions of nodes. The scale free nature of the connections between nodes presents several challenges to the performance of traditional searching algorithms. In this paper, we present two implementations of Parallel Breadth First Search algorithms on scale free networks without the use of specialized hardware.**

I. Introduction

Graph abstractions are commonly used in large scale network analysis [5]. With the proliferation of "Big Data" applications, parallelized versions of common graph analysis algorithms have a very high demand.

Many graph analysis problems make use of Breadth First Search (BFS), as the efficiency of the algorithm scales linearly with the number of nodes and edges.

Parallelizing Breadth First Search algorithms offers an excellent opportunity to increase the efficiency of this common graph analysis technique. However, scale free networks such as those created by social media connections are challenging to parallelize efficiently as the high distribution of edges to a few nodes is difficult to balance in memory [1]. Our research is targeted at the implementation of three variations of Parallel Breadth First Search algorithms without the use of specialized hardware on scale free networks. The first algorithm, Parallel Breadth First Search with Partitioning (PPBFS), was originally designed to improve BFS performance in distributed memory architectures used by many modern supercomputers. Our implementation will specifically apply this solution to scale free networks, which was left as future work by the original authors. However, as we are not implementing specialized hardware, our research will be performed without the use of a distributed memory architecture.

The second algorithm we will implement, Multithreaded Breadth First Search (MTBFS), is designed to support fine-grained, low overhead synchronization in a massively multithreaded system [5]. This algorithm will be used for gauging the performance of PPBFS, as MTBFS was intended to support scale free graphs. Since the focus of MTBFS is on increased parallelism through multi threading, the differences between the Cray MTA-2 system MTBFS was designed for and a standard multi-core CPU should be less drastic than the shift from the distributed memory architecture used by PPBFS. Memory access by BFS and other graph algorithms is typically fine-grained and irregular. This leads to poor cache performance, especially in parallel versions of BFS as parallelization relies heavily on the cache performance. Some performance improvements can be made, but they can not be sufficiently generalized since cache performance depends largely on the structure of the graph [5].

We will compare both of these algorithms with the sequential version of BFS across two datasets. One dataset will be a sufficiently large import of a Facebook social network, and the other a generic scale free network generator GDBench [2].

II. Related Work

A Scalable Distributed Parallel Breadth First Search Algorithm on BlueGene/L[5]

This paper talks about searching networks that are too large to fit into memory of a single machine. To handle this, it divides the graph into partitions where each node processes a set of vertices assigned to it. If it finds a vertex that does not belong to it, the node that owns it is notified. One major drawback this approach seems to have is that it waits for all processes to reach the same level before moving on. While this is important to truly follow a breadth first pattern, it may not be necessary to limit it in this way.

Designing Multithreaded Algorithms for Breath First search and st-connectivity on the Cray MTA-2[1]

The Multithreaded BFS algorithm implemented by Bader and Madduri will serve as our baseline parallel BFS algorithm. Although initially implemented on a massively multithreaded shared memory system without a data cache, we believe the fine-grained parallelization of this algorithm will still perform efficiently in a single CPU multi-core system. We have chosen this algorithm to serve as baseline parallelized implementation because it was specifically designed for the traversal of large scale-free graphs similar to those used in our benchmarks.

Parallel Breadth First Search on Distributed Memory Systems[2]

The partitioning Parallel Breadth First Search algorithm for distributed memory systems implemented by Buluc and Madduri extends the research performed by Yoo on the Scalable Distributed Parallel Breadth First Search, and highlights inefficiencies in the supporting structure of the other research. This research provided prior work summaries and criticism on the two algorithms we will adopt, but implementing the improvements made here is outside the scope of our research.

Efficient Parallel Graph Exploration on Multi-Core CPU and GPU[3]

The breadth-first search algorithm presented in this paper was proposed by Sungpack Hong, Tayo Oguntebi, and Kunle Olukotun. Their algorithm uses a hybrid approach to dynamically determine the best search method to use. They combine a level synchronous approach with a partitioning approach. For their combined approach, they determine whether to use a queue based search or an array based search based on criteria observed from the graph.

III. Technical Approach

Our projects technical approach focuses on comparing the graph traversal time of the three versions of the BFS algorithm. The original implementations of our parallel BFS algorithms were intended for use on super-computing hardware. The reason we have implemented multiple parallel BFS algorithms is to allow for additional performance comparisons of the algorithms, since they are abstracted away from their specialized hardware. Additionally all of our parallel algorithms will have their performance compared with a traditional sequential implementation of BFS.

The performance comparison of the algorithms will be determined by the amount of time each algorithm takes to fully traverse every node in two scale-free graphing benchmarks. We have implemented our solution using a third party application, NodeXL[4], to generate large scale-free graphs within our application. The NodeXL program is an open source application written in the C# language. In general, it is used as a network visualization tool which gives users the ability to view a graph's connectivity on a graphical interface. Additionally, there is a social network plugin for the NodeXL application which provides the capability of importing a social network graph from Facebook. We have utilized the class libraries for NodeXL and the associated social network plugin within our application to assist with the generation of the graphs.

Our first benchmark will be based on a graph of a sufficiently large social network imported through the use of the Facebook social network plugin for NodeXL. Our second benchmark is a based on an internet network dataset with over 124,000 nodes and 200,000 edges. Both of these datasets will be imported using NodeXL into a graph consumable by the algorithms. These benchmarks should both be sufficiently large to incur the cache performance challenges associated with many parallel BFS solutions.

We had intended to include the generic scale-free graph generated by the social network data generator GDBench FIX_CITATION[4]. The GDBench tool is capable of generating scale-free graphs with millions of nodes, and saving it in a common GraphML format, but the graphs generated by the tool were significantly disjoint. This made inclusion of the benchmark difficult as over half the nodes were unreachable.

IV. Technical Details

Our research is based on a distributed parallel approach to BFS, originally implemented on the IBM BlueGene/L supercomputer. We nearly attain performance increases using the same methods on a single CPU using multiple cores. Additionally, we will attempt to find optimizations to the algorithm such as allowing the processor to continue without waiting for send/receive messages before continuing the search for a given level/depth.

Parallel With Partitioning BFS (PPBFS)

The original algorithm used as a reference for our project was implemented as a distributed BFS on the BlueGene/L architecture. The algorithm takes advantage of the architecture's structure to develop efficient inter-processor communication, which is generally the bottleneck in distributed systems. Our implementation uses the techniques described in the paper to develop our parallel version of the algorithm, but with modifications that make more sense for working on a shared memory machine.

The primary difference between our implementation and the method used by the paper is how it transmits neighbor nodes to other threads. While a thread is executing the breadth first search, it may come across nodes that it does not own when searching adjacent nodes. The original paper solves this problem by having each thread create lists of neighboring nodes that belong to other processes and send them over a network to their respective processes and wait to receive lists from other processes. This sending and receiving of node lists forces the processes to need to wait for each other before descending into the next level.

Our implementation uses the same general methodology of partitioning the graph, but without the overhead of a network to perform on, we decided that creating queues for each thread, and enqueueing nodes as they were found made a lot more sense. This way each thread does not have to wait for all other threads to finish before searching the next depth level. The tradeoff is that the path found may not be the shortest possible path for to a node and the path found will depend on the scheduler.

The algorithm starts by partitioning the graph equally between the available threads. All of the vertices will be assigned an ID indicating which thread is responsible for searching it. One node will be marked as depth zero. This marks the starting node for the search, and is enqueued into the queue of its owner. Each thread takes the partitioned graph as input of size n/p. (Where 'n' is the number of vertices in the graph, and 'p' is the number of threads.) The algorithm works by simply dequeuing an vertex v_1, and enqueuing its neighbors $v'_1 \rightarrow v'_k$ into the local queues of the thread it belongs to. This process is repeated until all queues are empty. This modified version is fully illustrated as Algorithm 2.

Multi-threaded BFS (MTBFS)

The MTBFS implementation used for reference in this paper was implemented on a Cray MTA-2 multithreaded architecture (Reference paper 2). Similar to our approach, the algorithm was tested against scale-free graphs. When tested with a graph containing 400 million nodes, a 40 processor system showed

Algorithm 1 Distributed Breadth First expansion with 1D Partitioning[5]

1) Initialize $L_{v_s}(v) = \begin{cases} 0, & v = v_s, \text{where } v_s \text{ is a source} \\ \infty & \text{otherwise} \end{cases}$
2) **for** $l = 0$ *to* ∞ **do**
3) $F \leftarrow \{v \mid L_{v_s}(v) = l\}$, the set of local vertices with level l
4) **if** $F = \emptyset$ for all processors **then**
5) Terminate main loop
6) **end if**
7) $N \leftarrow$ {neighbors of vertices in F (not necessarily local)}
8) **for all** processors q **do**
9) $N_q \leftarrow$ {vertices in N owned by processor q}
10) **Send** N_q to processor q
11) **Receive** $\bar{N}_q$ from processor q
12) **end for**
13) $\bar{N} \leftarrow \bigcup_q \bar{N}_q$ (The $\bar{N}_q$ may overlap)
14) **for** $v \in \bar{N}$ and $L_{v_s}(v) = \infty$ **do**
15) $L_{v_s}(v) \leftarrow l + 1$
16) **end for**
17) **end for**

Algorithm 2 Modified Breadth First search with 1D Partitoning

```
subgraph<Vertex>[numVerts/numThreads]
subgraph[k].level = 0
localQueue[numThreads]
bfs(subGraph, threadID)
    finishMask |= 1 << threadID
    for Vertex v in subgraph
        v.id = threadID
        if (v.level = 0)
            localQueue[v.id] = threadID
    while ( finishMask != currentMask )
        while ( localQueue[threadID].size > 0 )
            srcVert = localQueue[threadID].dequeue()
            for Edge e in srcVert.edges()
                localQueue[e.dest.id].enqueue(e.dest)
                currentMask = 0
        currentMask |= 1 << threadID
```

the multi-threaded algorithm to have a system speedup of about 30 time over the sequential implementation.

The algorithm takes in a graph with source node s, and returns a shortest-path array d such that d[v] contains the length of the shortest path from source node s to destination node v, where v is a node in the graph. The algorithm uses the standard approach for implementing a sequential BFS where you start by adding the source node to a queue, then continue looping until the queue is empty. Each iteration dequeues the first node, finds it's neighboring nodes, and adds all unvisited neighbors to the queue. The main difference is that looping through the queue and through the neighboring nodes is done in parallel. In detail, the algorithm works as shown in Algorithm 3. ADD_CITATION

This algorithm takes full advantage of the Cray MTA-2 system's architecture by using its fine-grained parallelism and zero-overhead synchronization while looping through queue and looping over each node's set of neighbors. This multi-threaded BFS technique along with the hardware architecture

Algorithm 3 Level-synchronized Parallel BFS[1]

1) *for all_v* $\in$ *V in parallel do*
2) $d[v] \leftarrow -1$;
3) $d[s] \leftarrow 0$;
4) $Q \leftarrow \phi$;
5) *Enqueue* $s \leftarrow Q$;
6) *while* $Q \neq \phi$ *do*
7) *for all* $u \in Q$ *in parallel do*
8) *Delete* $u \leftarrow Q$;
9) *for each* v *adjacent to* u *in parallel do*
10) *if* $d[v] = -1$ *then*
11) $d[v] \leftarrow d[u] + 1$;
12) *Enqueue* $v \leftarrow Q$;

design offer a considerable speedup advantage over any sequential implementation of BFS on graphs of similar size.

MTBS has been implemented using nested AsParallel commands on methods that perform the functions highlighted above. Our implementation is significantly slower than the version implemented on the Cray MTA-2 as the nested parallelization limits the performance of the algorithm. The parallelization of the edges is responsible for most of the performance decrease as the workload of collecting adjacent edges is a relatively small.

Level Synchronous BFS (LSBFS)

The LSBFS algorithm uses level synchronization to keep the processing of nodes in order. A level property with an initial value of infinity is applied to each of the vertices in a graph. This property represents the distance a given vertex is away from root node. When the LSBFS starts, it sets the level of the root node to zero (0) and adds the root node to the list of visited nodes. Then for each iteration of the breadth first search, all nodes at the current level are searched in parallel. For each vertex in the current level, all of the vertex's neighbors are obtained and visited. For every visited vertex, the level property is set to the current level plus one (level + 1). After all the neighbors have been visited, the current level counter is incremented and all the neighboring nodes we just visited are the next set of nodes to continue searching on. All threads will execute this task in parallel.

The bottleneck in this algorithm comes from the fact that a finished thread must wait until all other threads finish their portion of the search at the current level. This is because all threads operate on a single level at a time. A general observation is that the number of nodes increases at higher levels of the graph. The algorithm benefits from this property as the parallel threads have more nodes to execute on as the level increases. However, another drawback of this method is that the algorithm does not take full advantage of breadth-first search properties. It would benefit from a partitioning scheme that allowed for multiple threads to operate on subgraphs of connected verticies. According to the parallel exploration paper[3], the algorithm still performs well in real-world systems despite it's synchronization overhead.

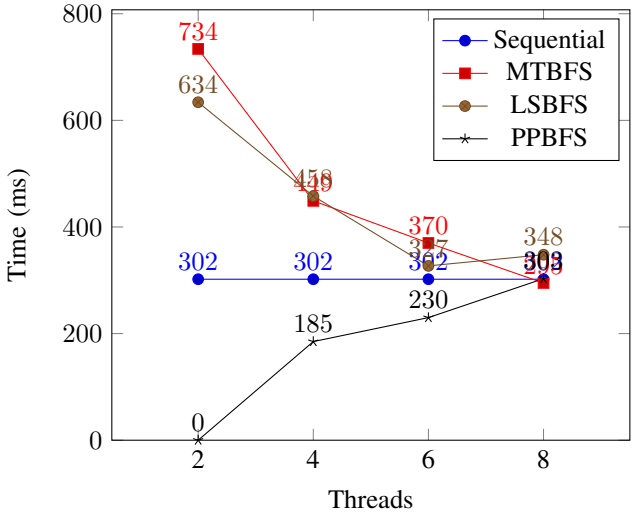
V. Experimental Results

Our experiments were implemented on an Intel i7 system with 4 cores. Our first benchmark included an import of a FaceBook social network with 7737 nodes and 12,694 edges. The second benchmark is a scale free dataset based on internet architecture with 124,651 nodes and 207,214 edges. The experiments were run multiple times over 2, 4, 6, and 8 threads.

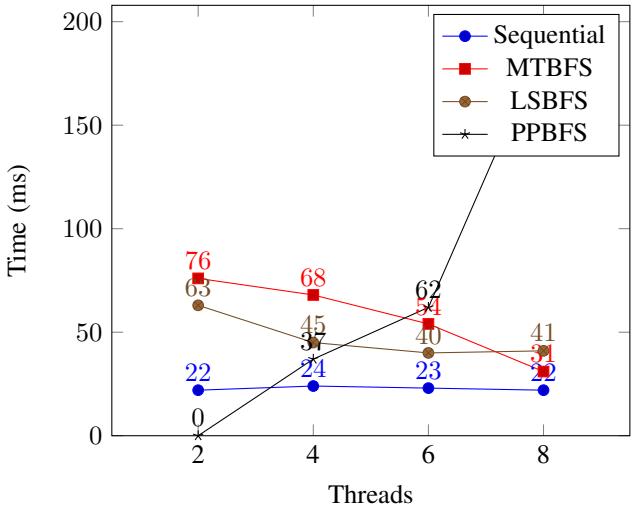
Our experimental results show that the PPBFS algorithm has a 62% improvement over the performance of the sequential algorithm when it is executed on the larger scale free graph with four threads.

The MTBFS and LSBFS algorithms do not perform as well as the sequential BFS algorithm, but the performance gap narrows as the number of threads increases.

Scale-Free Graph



Facebook



VI. Conclusions

Our experimental results show that PPBFS algorithm can improve the performance of scale free graph traversals, under certain conditions. The algorithm consistently has the best performance when it uses four threads, and if the dataset is large enough to encourage parallel analysis, PPBFS may be an appropriate choice. The MTBFS and LSBFS algorithms have generally lower performance than their sequential counterpart, but this was partially expected as it has been reported that scale free graphs don't benefit much from these simpler forms of parallelization.

VII. Challenges

Many benchmark datasets contain graphs with disjoint sets of vertices. Disjoint graphs with BFS won't allow the algorithm to visit all the nodes. The GDBench tool that we had intended to use for the production of substantially large datasets would produce disjoint graphs with roughly half the nodes in each set. This made the datasets produced nearly unusable as only half the graph could be traversed by the algorithms.

These datasets may have been usuable, but the NodeXL architecture we implemented becomes sensitive about the amount of memory used as the nodes increase above 100,000.

VIII. Future Work

There are a few minor optimizations we would like to focus on for the future works of this project. First, we would like to compare the differences in performance between restricting the algorithm to visiting nodes on a level synchronous bases, versus running a level asynchronous version. The asynchronous version would have the possibility of visiting nodes multiple times, but it would eliminate the overhead of threads needing to synchronize their execution after each level. Another small optimization we would like to implement is attempting to use array based queues. The size of the array for the queue could be easily determined by observing the number of nodes rationed to each partition of the graph. This would give us the benefit of spatial locality over the current list based queue that is used.

We would also like to use larger graphs to test our algorithm. We believe the algorithm would benefit from larger graphs since the parallelism could take full advantage of the larger range of nodes. As we observed from our test results, sometimes it's beneficial to use a sequential approach over a parallel approach, and sometimes it's beneficial to use a partitioned approach over a full graph search. With this in mind, we would like to optimize our implementation by taking a hybrid approach to determine which algorithm would be best to use based on the graph provided and the capabilities of the system which is being used to run the search.

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