# Direction-Optimizing Breadth-First Search

Scott Beamer Krste Asanović David Patterson Electrical Engineering and Computer Science Department University of California, Berkeley

Abstract—Breadth-First Search is an important kernel used by many graph-processing applications. In many of these emerging applications of BFS, such as analyzing social networks, the input graphs are low-diameter and scale-free. We present an efficient breadth-first search algorithm that is advantageous for low-diameter graphs. We adopt a hybrid approach, combining a conventional top-down algorithm along with a novel bottom-up algorithm. The bottom-up algorithm can dramatically reduce the number of edges examined, which in turn accelerates the search as a whole. On a multi-socket server, our hybrid approach demonstrates speedups of 3.3–7.8 on a range of standard synthetic graphs and speedups of 1.4–3.8 on graphs from real social networks compared to a strong baseline. We also show speedups of greater than 2.3 over the state-of-the-art multicore implementation when using the same hardware and input graphs.

#### I. Introduction

Graph algorithms are becoming increasingly important, with applications covering a wide range of scales. Warehouse-scale computers run graph algorithms that reason about vast amounts of data, with applications including analytics and recommendation systems [16] [19]. On mobile clients, graph algorithms are important components of recognition and machine-learning applications [18] [26].

Unfortunately, due to a lack of locality, graph applications are often memory-bound on shared-memory systems or communication-bound on clusters. In particular, Breadth-First Search (BFS), which is an important building block in many other graph algorithms, has low computational intensity, which exacerbates the lack of locality and results in low overall performance. To accelerate BFS, there has been significant prior work to change the algorithm and data structures, in some cases by adding additional computational work, to increase locality and boost overall performance [1] [9] [15] [25]. However, none of these previous schemes attempt to reduce the number of edges examined.

In this paper, we present a hybrid BFS algorithm that combines a conventional top-down approach with a novel bottom-up approach. By examining substantially fewer edges, the new algorithm obtains speedups of 3.3–7.8 on synthetic graphs and 1.4–3.8 on real social network graphs. In the top-down approach, nodes in the active frontier search for an unvisited child, while in our new bottom-up approach, unvisited nodes search for a parent in the active frontier. In general, the bottom-up approach will yield speedups when the active frontier is a substantial fraction of the total graph, as commonly occurs in small-world graphs such as social networks.

The bottom-up approach is not always advantageous, so we combine it with the conventional top-down approach, and use a simple heuristic to dynamically select the appropriate algorithm to use in each step of BFS. We show that our dynamic on-line heuristic achieves performance within 45% of the optimum achieved with an off-line oracle. Our hybrid implementation also provides a speedup of greater than 2.3 over the previous state-of-the-art for multicore when utilizing the same hardware and graphs [15]. An early version of this algorithm [5] running on a stock quad-socket Intel server was ranked 17<sup>th</sup> in the Graph500 November 2011 rankings [14], achieving the fastest single-node implementation and the highest per-core processing rate, and outperforming specialized architectures and clusters with more than 150 sockets.

#### II. GRAPH PROPERTIES

Graphs are a very powerful and general abstraction that allow a large number of problems to be represented and solved using the same algorithmic machinery. However, there is often substantial performance to be gained by optimizing algorithms for the types of graph present in a particular target workload.

We can characterize graphs using a few metrics, in particular their diameter and degree distribution. Graphs representing meshes used for physical simulations typically have very high diameters and degrees bounded by a small constant. As a result, they are amenable to graph partitioning when mapping to parallel systems, as they have mostly local communication and a relatively constant amount of work per vertex, which simplifies load-balancing.

In contrast, graphs taken from social networks tend to be both *small-world* and *scale-free*. A small-world graph's diameter grows only proportional to the logarithm of the number of nodes, resulting in a low effective diameter [22]. This effect is caused by a subset of edges connecting parts of the graph that would otherwise be distant. A scale-free graph's degree distribution follows a power law, resulting in a few, very high-degree nodes [4]. These properties complicate the parallelization of graph algorithms to efficiently analyze social networks. Graphs with the small-world property are often hard to partition because the low diameter and cross edges make it difficult to reduce the size of a cut. Meanwhile, scale-free graphs are challenging to load-balance because the amount of work per node is often proportional to the degree which can vary by several orders of magnitude.

#### function breadth-first-search(graph, source)

```
\begin{array}{l} \text{frontier} \leftarrow \{\text{source}\} \\ \text{next} \leftarrow \{\} \\ \text{parents} \leftarrow [\text{-1,-1,...-1}] \\ \textbf{while} \text{ frontier} \neq \{\} \textbf{ do} \\ \text{top-down-step(frontier, next, parents)} \\ \text{frontier} \leftarrow \text{next} \\ \text{next} \leftarrow \{\} \\ \textbf{end while} \\ \textbf{return} \text{ tree} \end{array}
```

Fig. 1. Conventional BFS Algorithm

## function top-down-step(frontier, next, parents)

```
\begin{array}{l} \textbf{for } v \in \text{frontier } \textbf{do} \\ \textbf{for } n \in \text{neighbors}[v] \ \textbf{do} \\ \textbf{if } parents[n] = -1 \ \textbf{then} \\ parents[n] \leftarrow v \\ next \leftarrow next \cup \{n\} \\ \textbf{end } \textbf{if} \\ \textbf{end } \textbf{for} \\ \textbf{end } \textbf{for} \\ \end{array}
```

Fig. 2. Single Step of Top-Down Approach

#### III. CONVENTIONAL TOP-DOWN BFS

Breadth-First Search (BFS) is an important building block of many graph algorithms. BFS can be used to test for connectivity or compute the single-source shortest paths of unweighted graphs. Starting from the source vertex, the frontier expands outwards during each step, visiting all of the vertices at the same depth before visiting any at the next depth (Figure 1). During a step of the conventional top-down approach (Figure 2), each vertex checks all of its neighbors to see if any of them are unvisited. Each previously unvisited neighbor is added to the frontier and marked as visited by setting its parent variable. This algorithm yields the BFS tree, which spans the connected component containing the source vertex. Other variants of BFS may record other attributes instead of the parent in the BFS tree, such as a simple boolean variable that marks whether it was visited, or an integer representing its depth in the tree.

The behavior of BFS on social networks follows directly from their defining properties: small-world and scale-free. Because social networks are small-world graphs, they have a low effective diameter, reducing the number of steps required for a BFS and requiring that a large fraction of the vertices are visited during each step. The scale-free property requires some nodes to have much higher degrees than average, allowing the frontier growth rate to outpace the average degree. As a result of these two properties, the frontier ramps up and down exponentially during a BFS search of a social network. Even if a social network graph has hundreds of millions of vertices, the vast majority will be reached in the first few steps.

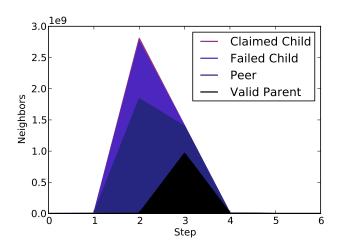


Fig. 3. Breakdown of edges in the frontier for a sample search on kron27 (Kronecker generated 128M vertices with 2B undirected edges) on the jkt system.

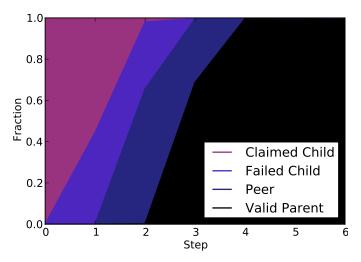


Fig. 4. Breakdown of edges in the frontier for a sample search on  ${\tt kron27}$  (Kronecker generated 128M vertices with 2B undirected edges) on the jkt system.

The majority of the computational work in BFS is checking edges of the frontier to see if the endpoint has been visited. The total number of edge checks in the conventional top-down algorithm is equal to the number of edges in the connected component containing the source vertex, as on each step every edge in the frontier is checked.

Figure 3 shows a breakdown of the result of each edge check for each step during a conventional parallel queue-based top-down BFS traversal on a Kronecker-generated synthetic graph, as are used for the Graph500 benchmark [14]. The middle steps (2 and 3) consume the vast majority of the runtime, which is unsurprising since the frontier is then at its largest size, requiring many more edges to be examined. During these steps, there are a great number of wasted attempts to become the parent of a neighbor. Failures occur when the neighbor has already been visited, and these can be broken down into

four different categories based on their depth relative to the candidate parent: valid parent, peer, and failed child. A valid parent is any neighbor at depth d-1 of a vertex at depth d. A peer is any neighbor at the same depth. A failed child is any neighbor at depth d+1 of a vertex at depth d, but at the time of examination it has already been claimed by another vertex at depth d. Successful checks result in a claimed child. Figure 3 shows most of the edge checks do fail and represent redundant work, since a vertex in a correct BFS tree only needs one parent.

Implementations of this same basic algorithm can vary in a number of performance-impacting ways, including: data structures, traversal order, parallel work allocation, partitioning, synchronization, or update procedure. The process of checking if neighbors have been visited can result in many costly random accesses. An effective optimization for shared-memory machines with large last-level caches is to use a bitmap to mark nodes that have already been visited [1]. The bitmap can often fit in the last-level cache, which prevents many of those random accesses from touching off-chip DRAM. These optimizations speed up the edge checks but do not reduce the number of checks required.

The theoretical minimum for the number of edges that need to be examined in the best case is the number of vertices in the BFS tree minus one, since that is how many edges are required to connect it. For the example in Figure 3, only 63,036,116 vertices are in the BFS tree, so at least 63,036,115 edges need to be considered, which is about  $\frac{1}{67}$  of all the edge examinations that would happen during a top-down traversal. This factor of 67 is substantially larger than the input degree of 16 for two reasons. First, the input degree is for undirected edges, but during a top-down search, both endpoints of each edge will check it, which doubles the number of examinations. Secondly, there are a large number of vertices of zero degree, which reduces the size of the main connected component and also further increases the effective degree of the vertices it contains. There is clearly substantial room for improvement by checking fewer edges, although in the worst case, every edge might still need to be checked.

Figure 4 zooms in on the edge check results of Figure 3 for the sample search. This progression of neighbor types is typical among the social networks examined. During the first few steps, the percentage of claimed children is high, as the vast majority of the graph is unexplored, enabling most edge checks to succeed. During the next few steps, the percentage of failed children rises, which is unsurprising since the frontier has grown larger, as multiple valid parents are fighting over children. As the frontier reaches its largest size, the percentage of peer edges dominates. Since the frontier is such a large fraction of the graph, many edges must connect vertices within the frontier. After the apex of frontier size, the percentage of valid parents rises since such a large fraction of edges were in the previous step's frontier.

## function bottom-up-step(frontier, next, parents)

```
\begin{array}{l} \textbf{for} \ v \in \ vertices \ \textbf{do} \\ \textbf{if} \ parents[v] = -1 \ \textbf{then} \\ \textbf{for} \ n \in neighbors[v] \ \textbf{do} \\ \textbf{if} \ n \in frontier \ \textbf{then} \\ parents[v] \leftarrow n \\ next \leftarrow next \cup \{v\} \\ \textbf{break} \\ \textbf{end} \ \textbf{if} \\ \textbf{end} \ \textbf{for} \\ \end{array}
```

Fig. 5. Single Step of Bottom-Up Approach

#### IV. BOTTOM-UP BFS

When the frontier is large, there exists an opportunity to perform the BFS traversal more efficiently by searching in the reverse direction, that is, going bottom-up. Note that in Step 3 there are many valid parents, but the other ends of these edges mostly resulted in failed children during Step 2. We can exploit this phenomenon to reduce the total number of edges examined. Instead of each vertex in the frontier attempting to become the parent of all of its neighbors, each unvisited vertex attempts to find any parent among its neighbors. A neighbor can be a parent if the neighbor is a member of the frontier, which can be determined efficiently if the frontier is represented by a bitmap. The advantage of this approach is that once a vertex has found a parent, it does not need to check the rest of its neighbors. Figure 4 demonstrates that for some steps, a substantial fraction of neighbors are valid parents, so the probability of not needing to check every edge is high. Figure 5 shows the algorithm for a single step of this approach.

The bottom-up approach also removes the need for some atomic operations in a parallel implementation. In the top-down approach, there could be multiple parallel writers to the same child, so atomic operations are needed to ensure mutual exclusion. With the bottom-up approach, only the child writes to itself, removing any contention. This advantage, along with the potential reduction of edges checked, comes at the price of serializing the work for any one vertex, but there is still massive parallelism between the work for different vertices. The bottom-up approach is advantageous when a large fraction of the vertices are in the frontier, but will result in more work if the frontier is small. Hence, an efficient BFS implementation must combine both the top-down and bottom-up approaches.

If the graph is undirected, performing the bottom-up approach requires no modification to the graph data structures as both directions are already represented. If the graph is directed, the bottom-up step will require the inverse graph, which could nearly double the graph's memory footprint.

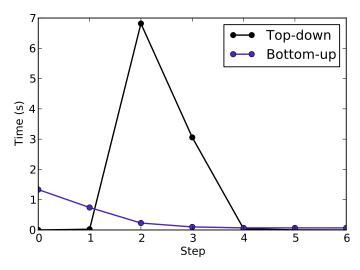


Fig. 6. Sample search on kron27 (Kronecker 128M vertices with 2B undirected edges) on the jkt system.

#### V. HYBRID ALGORITHM

The pairing of the top-down approach with the bottom-up approach is complementary, since when the frontier is its largest, the bottom-up approach will be at its best whereas the top-down approach will be at its worst, and vice versa. The runtime for either the top-down approach or the bottom-up approach is roughly proportional to the number of edges examined. The top-down approach will examine every edge in the frontier while the bottom-up approach could examine every edge attached to an unvisited vertex, but hopefully fewer. Figure 6 illustrates this behavior by showing the time per step for each approach using the same example search as in Section III. As the size of the frontier ramps up, the time per step of the top-down approach rises correspondingly, but the time per step for the bottom-up approach drops.

Our hybrid algorithm uses the top-down approach for steps when the frontier is small and the bottom-up approach for steps when the frontier is large. We begin each search with the top-down approach, and continue until the frontier becomes too large when we switch to the bottom-up approach. Although it is difficult to tell from Figure 6, it is usually worthwhile to switch back to the top-down approach for the final steps. During some searches, there can be a long tail, and edges that are not in the connected-component continue to consume runtime using the bottom-up approach.

The step when transitioning from the top-down approach to the bottom-up approach provides an enormous opportunity, since all of the edge checks that would happen during the top-down step (all of the edges in the frontier) are skipped. For this reason, the optimum point to switch is typically when the number of edges in the frontier is at its largest. This is fortuitous, since the first bottom-up step will probably benefit from a high percentage of valid parents, as Figure 4 shows.

To control the hybrid approach, we use a heuristic based on: the number of edges to check from the frontier  $(m_f)$ , the number of vertices in the frontier  $(n_f)$ , and the number of

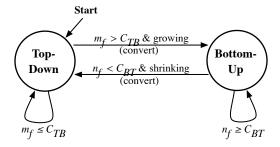


Fig. 7. Control algorithm for hybrid algorithm. (convert) indicates the frontier must be converted from a queue to a bitmap or vice versa between the steps.

edges to check from unexplored vertices  $(m_u)$ . These metrics are efficient to compute, since they only require: summing the degrees of all vertices in the frontier, counting the number of vertices added to the frontier, or counting how many edges have been checked. Note that with an undirected graph,  $m_f$  or  $m_u$  might be counting some edges twice since each edge will be explored from both ends. Figure 7 shows the overall control heuristic, which employs two thresholds,  $C_{TB}$  and  $C_{BT}$ , together with  $m_f$  and  $n_f$ .

Since the top-down approach will always check  $m_f$  edges, and the bottom-up approach will check at most  $m_u$ , if  $m_u$  is ever less than  $m_f$ , there is a guarantee that switching to the bottom-up approach at that step will check less edges. Since  $m_u$  is an overly pessimistic upper-bound on the number of edges the bottom-up step will check, we use a tuning parameter  $\alpha$ . This results in the condition for switching from top-down to bottom-up being:

$$m_f > \frac{m_u}{\alpha} = C_{TB}$$

Switching back to the top-down approach at the end should occur when the frontier is small and there is no longer benefit to the bottom-up approach. In addition to the overhead of checking edges outside the main connected component, the bottom-up approach becomes less efficient at the end because it scans through all vertices to find unvisited ones. The heuristic to switch back attempts to detect when the frontier is too small, and we use another tuning parameter  $\beta$  to get:

$$n_f < \frac{n}{\beta} = C_{BT}$$

When switching between approaches, the representation of the frontier must be changed from the frontier queue used in the conventional top-down approach to the frontier bitmap used for the bottom-up approach. Different data structures are used since the frontiers are of radically different sizes, and the conversion costs are far less than the penalty of using the wrong data structure. The bottom-up approach uses a frontier bitmap to allow a constant-time test for whether a particular node is in the frontier. The time to convert the frontier is typically small, since it should be done when the frontier is small.

Abbreviation	Graph	# Vertices	# Edges	Input Degree	Directed	References
kron25	Kronecker	33,554,432	536,870,912	16.0	N	[14][17]
erdos25	Erdős–Réyni (Uniform Random)	33,554,432	268,435,456	8.0	N	[2][13]
rmat25	RMAT	33,554,432	268,435,456	8.0	Y	[2][10]
faceA	Facebook Trace A	3,097,165	28,377,481	9.2	N	[24]
faceB	Facebook Trace B	2,937,612	24,236,701	8.3	N	[24]
hollywood	Hollywood Movie Actor Network	1,139,906	57,515,616	50.5	N	[6], [7], [8][12]
ljournal	LiveJournal Social Network	5,363,261	79,023,142	14.7	N	[6], [7], [8][12]
wikipedia	Wikipedia Links	5,716,808	130,160,392	22.8	Y	[23]
twitter	Twitter User Follow Links	61,578,415	1,468,365,182	23.8	Y	[16]

TABLE I GRAPHS USED FOR EVALUATION

Name	gaines	jkt	box
Architecture	Nehalem-EP	Sandy Bridge-EP	Westmere-EX
Intel Model	X5550	E5-2680	E7-4860
Release	Q1'09	Q1'12	Q2'11
Clock rate	2.67GHz	2.7GHz	2.26GHz
Cores/socket	4	8	10
Threads/socket	8	16	20
LLC/socket	8MB	20MB	24MB
# Sockets	2	2	4
DRAM Size	12GB	128GB	128GB

TABLE II SYSTEM SPECIFICATIONS

## VI. EVALUATION

#### A. Methodology

We evaluate the performance of our hybrid algorithm using the graphs in Table I. Our suite of test graphs includes both synthetic graphs as well as real social networks. The synthetic graph generators as well as their parameters are selected to match the Graph500 Competition [14] (Kronecker (A,B,C) = (0.57,0.19,0.19)), and prior work [1][15] (Uniform Random and RMAT (A,B,C) = (0.45,0.25,0.15)). The real social networks are taken from a variety of web crawls [6], [7], [8][12][16][23][24].

We report results from three multi-socket server systems (gaines, jkt, box) as shown in Table II. The jkt system is used for most of the evaluation, as it is representative of the next generation of compute nodes for clusters. The box system is used to evaluate the parallel scalability of our approach up to 80 threads. The gaines system is used to perform a direct comparison with prior work [15] utilizing the same hardware and input graphs.

Our algorithm implementations use C++ and OpenMP, and store the graphs in Compressed Sparse Row (CSR) format after removing duplicates and self-loops. For each plotted data point, we perform BFS 64 times from pseudo-randomly selected non-zero degree vertices and average the results. The time for each search includes the time to allocate and initialize search-related data structures (including parents).

To benchmark performance, we calculate the search rate in Millions of Edges Traversed per Second (MTEPS) by taking the ratio of the number of edges in the input graph to the runtime. This metric is artificial in the sense that either the top-down approach or the hybrid approach might not check every edge, but this standardized performance metric is similar in spirit to measuring MFLOPS for optimized matrix multiplication in terms of the classic  $O(N^3)$  algorithm.

## B. Tuning $\alpha$ and $\beta$

First we determine values of  $\alpha$  and  $\beta$  to use for *hybridheuristic*, our hybrid implementation of BFS described above. We first tune  $\alpha$  before tuning  $\beta$ , as it has the greatest impact. Sweeping  $\alpha$  across a wide range demonstrates that once  $\alpha$  is sufficiently large (>12), BFS performance for many graphs is relatively insensitive to its value (Figure 8). This is because the frontier grows so rapidly that small changes in the transition threshold do not change the step at which the switch occurs. When trying to pick the best value for the suite, we select  $\alpha=14$  since it maximizes the average and minimum. Note that even if a less than optimal  $\alpha$  is selected, the *hybridheuristic* algorithm will still execute within 15–20% of its peak performance on most graphs.

Tuning  $\beta$  is less important than tuning  $\alpha$ . We select  $\beta=24$ , as this works well for the majority of the graphs (Figure 9). The value of  $\beta$  has a smaller impact on overall performance because the majority of the runtime is taken by the middle steps when the frontier is at its largest, even when those steps are accelerated by the bottom-up approach. We explore a much larger range for  $\beta$  since it needs to change by orders of magnitude in order to have an impact on which step the heuristic will switch.

## C. Comparing Algorithms

We first compare our own implementations of different algorithms on a range of graphs. As a baseline, we use two top-down parallel queue implementations. The *top-down* implementation uses a typical parallel queue, much like the omp-csr reference code [14]. We also created an optimized version, *top-down-check*, which adds a bitmap to mark completed nodes. Performance is very similar to the baseline reported in [15] when running on the same model hardware (gaines machine) with the same synthetic graphs from the same generator [2]. To determine the maximum benefit of our hybrid approach and to evaluate the effectiveness of our online heuristic, we also develop *hybrid-oracle*, that repeatedly runs the algorithm trying all possible switch points and reports the best performing to act as an oracle-controlled hybrid.

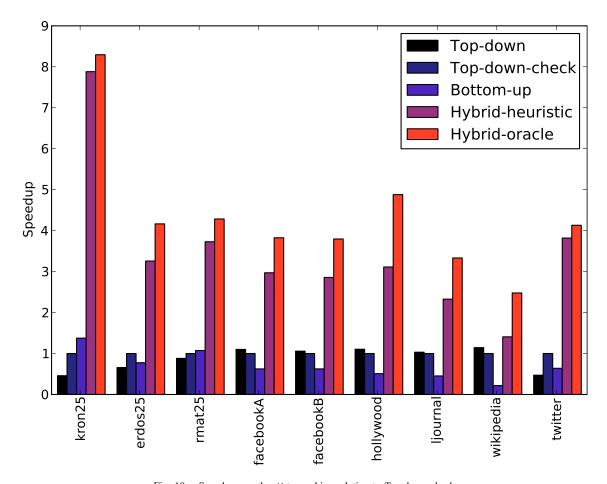


Fig. 10. Speedups on the jkt machine relative to Top-down-check.

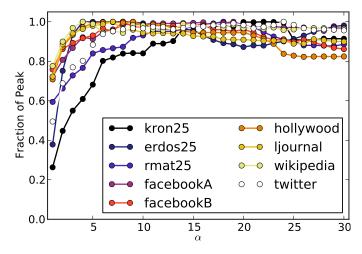


Fig. 8. Performance of *hybrid-heuristic* on each graph relative to its best on that graph for the range of  $\alpha$  examined.

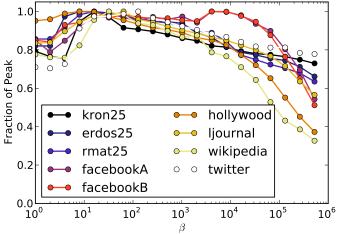


Fig. 9. Performance of *hybrid-heuristic* on each graph relative to its best on that graph for the range of  $\beta$  examined.

We use the jkt machine to compare both hybrid implementations (hybrid-heuristic & hybrid-oracle) against both baselines (top-down & top-down-check) in Figure 10. The hybrid provides large speedups across all of the graphs, with an average speedup of 3.5 and a speedup no lower than 1.4. The on-line heuristic often obtains performance within 10% of

the oracle. Also notice that the speedups are far greater than the impact of a mistuned  $\alpha$  or  $\beta$  in the heuristic.

## D. Explaining the Performance Improvement

The speedup of the hybrid approach demonstrated in Figure 10 is due to the reduction in edge examinations (Fig-

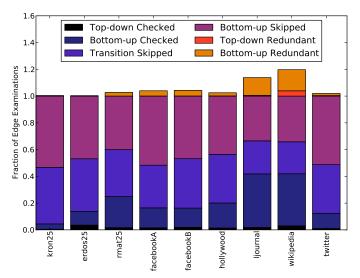


Fig. 11. Breakdown of edge examinations.

ure 11). In a classical top-down BFS, every edge in the connected component containing the starting vertex will be checked, and for an undirected graph each edge will be checked from both ends. The bottom-up approach skips checking some edges in two ways. First, it passes the responsibility of setting the parents variable from the parent to the child, who can stop early once a parent is found. Secondly, it skips all of the edges in the frontier on the step that transitions from top-down to bottom-up. As shown by Figure 11, the edge examinations skipped are roughly split between those skipped in the transition and those skipped by the first bottom-up step. Since the bottom-up approach is used when the frontier is large, the top-down approach in the hybrid implementation processes only a small fraction of the edges. Every edge examination after the first bottom-up step is redundant, since the first bottom-up step will have attempted to examine every edge attached to an unvisited vertex. Since the size of the frontier decreases so rapidly, this redundancy is typically negligible. Unsurprisingly, the graphs with the least speedup (ljournal and wikipedia), skip the fewest edges and check the most redundant edges. They have a substantially higher effective diameter, so there is a longer tail after the apex of the frontier size.

Examining where the time is spend during an entire search reveals the majority of it is spent in bottom-up implementation (Figure 12). Since the bottom-up approach skips so many examinations, the effective search rate for the bottom-up steps is much higher (order of magnitude) than the top-down approach. Conversion and  $m_f$  calculation take a non-negligible fraction of the runtime, but this overhead is worthwhile due to the extreme speedup provided by the bottom-up approach. The nearly constant degree of erdos25 results in a steady growth of the frontier, so more steps will be run top-down with a larger frontier, resulting in more work at the conversion point, more degrees to sum, and more checks run in top-down mode.

Figure 13 plots the speedups for the suite compared to

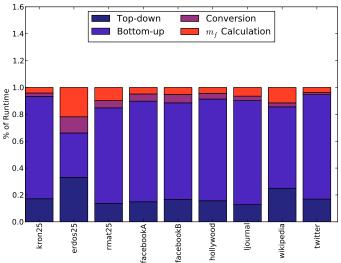


Fig. 12. Breakdown of time spent per search.

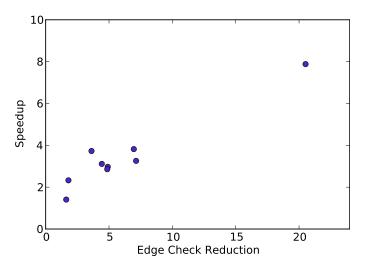


Fig. 13. Speedups on the jkt system for *Hybrid-heuristic* compared to the reduction in edges checked.

the reduction in number of edges checked. The speedup is reasonably correlated with the reduction in edges checked, with the slope of a best-fit line at approximately 0.3. The slope is less than 1.0 due to a variety of overheads. While the bottom-up approach skips edges, it reduces the spatial locality of the remaining memory accesses, which impacts memory performance in modern processors that bring data from memory in units of cache blocks rather than individual words. Conversion and  $m_f$  calculation add other overhead. Using top-down-check as a baseline rather than top-down, which further reduces the relative advantage since edge examinations become less expensive in the baseline as they will often hit the bitmap in the last-level cache.

## E. Scalability

We evaluate the scalability of our *Hybrid-heuristic* implementation on the box system in a fashion similar to prior work [1]. Our approach gets parallel speedup with each additional

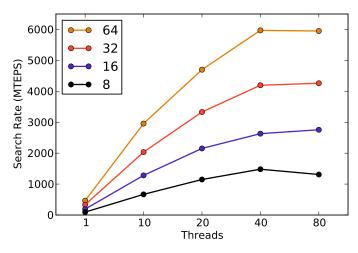


Fig. 14. Parallel scaling of *Hybrid-heuristic* on the box system for an RMAT graph with 16M vertices and varied degree.

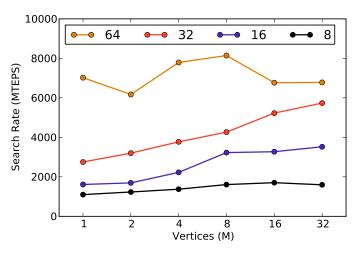


Fig. 15. Vertex scaling of *Hybrid-heuristic* on the box system for an RMAT graph with varied degree.

core, but does not gain much from hyperthreading on the box system (Figure 14). Our approach does benefit from larger graphs (more vertices), as it gives it more computation to drown out the overheads (Figure 15). Increasing the degree is particularly beneficial for our approach because it does not significantly increase the runtime but it does increase the number of input edges which produces a higher search rate.

#### F. Comparison Against Other Implementations

We also compare our hybrid implementation against the previous state-of-the-art for multicore by Hong et al. [15]. Note the implementation by Hong et al [15] is faster than that of Agarwal et al. [1] even while using slower hardware. We use the exact same model hardware and the same graph generator source code (SNAP [2]) to provide a point-for-point comparison as shown in Figures 16 and 17. Our implementation is  $2.4 \times$  faster on uniform random (erdos25) and  $2.3 \times$  faster on scale-free (rmat25). Our implementation experiences a slowdown after 8 threads due to the switch to hyperthreads in our tightly synchronized approach, but using 16 hyperthreads

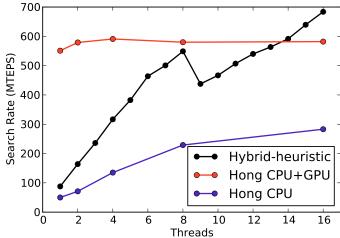


Fig. 16. Search rates on the gaines system on erdos25 (Uniform Random with 32M vertices and 256M edges). Other lines from Hong et al. [15]

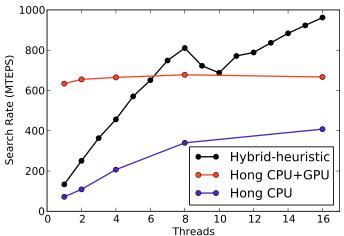


Fig. 17. Search rates on the gaines system on rmat25 (RMAT with 32M vertices and 256M edges). Other lines from Hong et al. [15]

is still noticeably faster than just 8 hyperthreads on the 8 available cores. Furthermore, our CPU implementation using 16 hyperthreads is even faster than the GPU-CPU hybrid implementation presented by Hong et al. [15], despite the GPU having greater memory bandwidth and supporting more outstanding memory accesses. Note also that not all of the graphs in our study above would fit in the reduced memory capacity of the GPU, and that our implementation records a full parent pointer rather than just the BFS depth [15].

We compare our results to that of Merrill et al. [20]. which is the highest published performance for shared memory (Table III). Our *Hybrid-heuristic* outperforms it by a significant margin due to the high input degrees of the graphs used [20]. The GTEP rates from the prior work have been halved since they count GTEPs based on directed edges instead of undirected edges.

System	kron_g500-logn20	random.2Mv.128Me	rmat.2Mv.128Me
Single-GPU	1.25	1.20	1.30
Quad-GPU	3.10	3.70	4.15
gaines	7.76	5.72	5.64
jkt	11.23	8.06	6.90
box	8.89	6.77	7.14

TABLE III

Hybrid-heuristic On MULTICORE SYSTEMS IN THIS STUDY COMPARED TO GPU RESULTS FROM MERRILL ET AL. [20] (IN GTEPS).

## VII. RELATED WORK

Buluç and Madduri [9] provide a thorough taxonomy of related work in parallel breadth-first searches. In this section we focus on the work most relevant to this study, principally parallel shared-memory implementations.

Bader and Madduri [3] demonstrate large parallel speedups for BFS on an MTA-2. Parallelism is extracted both at the level of each vertex as well as at each edge check. The fine-grained synchronization mechanisms of the MTA are used to efficiently exploit this parallelism. Since the MTA does not use caches and instead uses many hardware threads to hide its latency to its shared main memory, the implementation does not need to optimize for locality.

In contrast, Agarwal et al. [1] optimize for locality to push the limits of a quad-socket system built from conventional microprocessors, and show speedups over previous results from clusters and custom supercomputers, including the MTA-2 [3]. Careful programming is used to reduce off-socket traffic (memory and inter-socket) as much as possible. Threads are pinned to sockets and utilize local data structures such that all the work that can be done within a socket never leaves the socket. Bitmaps are used to track completed vertices to avoid reading DRAM. Whenever an edge check must go to another socket, it utilizes a custom efficient inter-socket messaging queue.

Hong et al. [15] improve upon Agarwal et al. with hybrid algorithms that utilize multiple CPU implementations and a GPU implementation. As in our work, the algorithms switch at step boundaries and make decisions based on on-line heuristics. The CPU implementation is accelerated by the read-array approach, which combines the frontier with the output array. The output provided is the depth of each node rather than its parent, and by bounding the depth to be less than 256, the output array can be compacted by storing it as bytes. Instead of appending vertices to a frontier, their output is set to depth+1. On each step, all vertices are scanned searching for the current depth, and then visits are performed from any vertex found on the frontier. By using the output array to act as the frontier, duplicate entries are squashed, and spatial locality is increased due to the sequential array accesses to the graph. The GPU implementation outperforms the CPU implementation for a sufficiently large frontier, and heuristics select which implementation to use. In Section ??, we compare our new scheme directly against the performance in [15] utilizing the same synthetic data on the same model CPU hardware, but without the GPU. Note our scheme retains full parent information, not just search depth.

Merrill et al. [20] improve the performance of BFS on GPUs through the use of prefix sum to achieve high utilization of all threads. The prefix sum is used to compute the offsets from the frontier expansion, which reduces the contention for atomic updates to the frontier queue. This allows the graph exploration to have little control divergence between threads. They also leverage various filtering techniques enabled by bitmaps to reduce the number of edges processed. This approach is beneficial for all diameters of graphs, and they present results indicating they are the fastest published for shared memory systems, especially with their quad-GPU parallelization. Our approach is advantageous for low diameter graphs, and as shown in Section VI-F, our approach outperforms their quad-GPU implementation for all sizes of system used in our evaluation.

An early implementation of this algorithm on the mirasol system reached 17<sup>th</sup> place in the November 2011 rankings of the Graph500 competition [5][14]. This earlier implementation used a cruder heuristic and included an alternate top-down step that integrated the conversion for bottom-up. It achieved the highest single-node performance and the highest per-core processing rate. Using just a single quad-socket system similar to boxboro, the hybrid BFS algorithm outperformed clusters of >150 sockets and specialized architectures such as the Cray XMT2 [21] and the Convey HC-1<sup>ex</sup> [11]. Its processing rate per-core was over 30× the top-ranked cluster system, highlighting the performance penalty a cluster experiences when crossing the network. This implies that scaling across a cluster should be used to solve larger problems (weak scaling) rather than solving fixed-size problems faster (strong scaling), indicating large shared-memory systems are still useful for irregular difficult-to-scale problems like BFS.

Yoo et al. [25] present a 2D partitioning, which greatly reduces the communication costs for a cluster-scale implementation, in contrast to our work and the other related work above, which use 1D partitionings. With a 1D partitioning, each thread is allocated a subset of the vertices, but is responsible for all of their edges. With the 2D partitioning, each thread is allocated a subset of the edges for a subset of vertices. For massively parallel machines, this can not only reduce communication costs, but can also help with load balancing. Buluç and Madduri [9] improve upon this by reducing communication times for the 2D approach by using a hybrid of MPI for inter-node parallelism and OpenMP for intra-node parallelism. The success of these 2D approaches highlight the importance of reducing communication costs to improve BFS performance, and a bottom-up approach would further reduce the communication volume by reducing the number of edges checked.

# VIII. DISCUSSION

The results of this work demonstrate the performance improvement potential of integrating the bottom-up approach into BFS, but it will not always be advantageous. Fortunately, the

same top-down implementation that is used for the steps when the frontier is small can be used for entire searches when the bottom-up approach is not advantageous. The heuristic will allow this to happen automatically, as searches that do not generate a massive frontier will not trigger the switch. In this sense, the bottom-up approach can be seen as a powerful way to accelerate a BFS implementation when it is searching low-diameter graphs.

High diameter graphs will not benefit from the bottom-up approach, but they are much easier to partition, and thus easier to parallelize than low diameter graphs. This work presents an algorithmic innovation to accelerate the processing of difficult to parallelize BFS.

#### IX. CONCLUSION

Performing a BFS in the bottom-up direction can substantially reduce the number of edges traversed compared with the traditional top-down approach. A child needs to find only one parent instead of each parent attempting to claim all possible children. This bottom-up technique is advantageous when the search is on a large connected component of low-effective diameter, as then the frontier will include a substantial fraction of the total vertices. The conventional top-down approach works well for the beginning and end of such a search, where the frontier is a small fraction of the total vertices. We have developed a hybrid approach to effectively combine these two algorithms, using a simple heuristic to guide when to switch. We believe this hybrid scheme yields the current best-known BFS performance for single shared-memory nodes.

# REFERENCES

- [1] V Agarwal, F Petrini, D Pasetto, and David A Bader. Scalable graph exploration on multicore processors. *Supercomputing*, 2010.
- [2] David Bader and Kamesh Madduri. SNAP: Small-world network analysis and partitioning: an open-source parallel graph framework for the exploration of large-scale networks. *International Parallel and Distributed Processing Symposium*, 2008.
- [3] David A. Bader and Kamesh Madduri. Designing multithreaded algorithms for breadth-first search and st-connectivity on the Cray MTA-2. International Conference on Parallel Processing, 2006.
- [4] Albert-László Barabási and Réka Albert. Emergence of scaling in random networks. Science, 286:509–512, Oct 1999.
- [5] Scott Beamer, Krste Asanović, and David A. Patterson. Searching for a parent instead of fighting over children: A fast breadth-first search implementation for graph500. Technical Report UCB/EECS-2011-117, EECS Department, University of California, Berkeley, 2011.
- [6] Paolo Boldi, Bruno Codenotti, Massimo Santini, and Sebastiano Vigna. Ubicrawler: A scalable fully distributed web crawler. Software: Practice & Experience, 34(8):711–726, 2004.
- [7] Paolo Boldi, Marco Rosa, Massimo Santini, and Sebastiano Vigna. Layered label propagation: A multiresolution coordinate-free ordering for compressing social networks. In *Proceedings of the 20th international conference on World Wide Web*. ACM Press, 2011.
- [8] Paolo Boldi and Sebastiano Vigna. The WebGraph framework I: Compression techniques. In Proc. of the Thirteenth International World Wide Web Conference (WWW 2004), pages 595–601, Manhattan, USA, 2004. ACM Press.
- [9] Aydın Buluç and Kamesh Madduri. Parallel breadth-first search on distributed memory systems. Supercomputing, 2011.
- [10] Deepayan Chakrabarti, Yiping Zhan, and Christos Faloutsos. R-MAT: A recursive model for graph mining. SIAM Data Mining, 2004.
- [11] Convey HC-1 family. http://www.conveycomputer.com/Resources/Convey \_HC1\_Family.pdf.

- [12] Timothy Davis and Yifan Hu. The University of Florida sparse matrix collection. ACM Transactions on Mathematical Software (to appear), http://www.cise.ufl.edu/research/sparse/matrices/.
- [13] Paul Erdős and Alfréd Réyni. On random graphs. i. Publicationes Mathematicae, 6:290–297, 1959.
- [14] Graph500 benchmark. http://http://www.graph500.org/.
- [15] Sungpack Hong, Tayo Oguntebi, and Kunle Olukotun. Efficient parallel graph exploration on multi-core CPU and GPU. Parallel Architectures and Compilation Techniques, 2011.
- [16] Haewoon Kwak, Changhyun Lee, Hosung Park, and Sue Moon. What is Twitter, a social network or a news media? *International World Wide Web Conference*, 2010.
- [17] Jurij Leskovec, Deepayan Chakrabarti, Jon Kleinberg, and Christos Faloutsos. Realistic, mathematically tractable graph generation and evolution, using Kronecker multiplication. European Conference on Principles and Practice of Knowledge Discovery in Databases, 2005.
- [18] Y Low, J Gonzalez, A Kyrola, D Bickson, C Guestrin, and J.M Hellerstein. GraphLab: A new framework for parallel machine learning. Uncertainty in Artificial Intelligence, 2010.
- [19] Grzegorz Malewicz, Matthew Austern, Aart Bik, James Dehnert, Ilan Horn, Naty Leiser, and Grzegorz Czajkowski. Pregel: A system for large-scale graph processing. SIGMOD, Jun 2010.
- [20] Duane Merrill, Michael Garland, and Andrew S. Grimshaw. Scalable GPU graph traversal. *Principles and Practice of Parallel Programming*, 2012.
- [21] David Mizell and Kristyn Maschhoff. Early experiences with large-scale cray xmt systems. Supercomputing, 2009.
- [22] D.J Watts and S.H Strogatz. Collective dynamics of 'small-world' networks. *Nature*, 393:440–442, June 1998.
- [23] Wikipedia page-to-page link database 2009. http://haselgrove.id.au/wikipedia.htm.
- [24] Christo Wilson, Bryce Boe, Alessandra Sala, Krishna P. N. Puttaswamy, and Ben Y. Zhao. User interactions in social networks and their implications. *EuroSys*, 2009.
- [25] Andy Yoo, Edmond Chow, Keith Henderson, William McLendon, Bruce Hendrickson, and Ümit Çatalyürek. A scalable distributed parallel breadth-first search algorithm on BlueGene/L. Supercomputing, 2005.
- [26] K You, J Chong, Y Yi, E Gonina, C Hughes, Y.K Chen, W Sung, and K Keutzer. Scalable hmm based inference engine in large vocabulary continuous speech recognition. *IEEE Signal Processing Magazine*, 2010.