

# A Computational Study of External-Memory BFS Algorithms\*

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## Abstract

Breadth First Search (BFS) traversal is an archetype for many important graph problems. However, computing a BFS level decomposition for massive graphs was considered nonviable so far, because of the large number of I/Os it incurs. This paper presents the first experimental evaluation of recent external-memory BFS algorithms for general graphs. With our STXXL based implementations exploiting pipelining and disk-parallelism, we were able to compute the BFS level decomposition of a web-crawl based graph of around 130 million nodes and 1.4 billion edges in less than 4 hours using single disk and 2.3 hours using 4 disks. We demonstrate that some rather simple external-memory algorithms perform significantly better (*minutes* as compared to *hours*) than internal-memory BFS, even if more than half of the input resides internally.

## 1 Introduction

Solving real world optimization problems often boils down to traversing graphs in a structured way. Breadth First Search (BFS) is among the most fundamental such traversal strategies. It decomposes the input graph  $G = (V, E)$  of  $n$  nodes and  $m$  edges into at most  $n$  levels where level  $i$  comprises all nodes that can be reached from a designated source  $s$  via a path of  $i$  edges, but cannot be reached using less than  $i$  edges.

BFS is well-understood in the *von Neumann* model which assumes uniform memory access costs. Actual machines, however, increasingly deviate from this model. While waiting for a memory access, modern microprocessors can execute a number of register operations. Even worse, in an external-memory (EM) setting where the input graph is too big to fully fit in internal memory certain accesses have to be served by hard disks, thus incurring an I/O loss factor of up to seven orders of magnitude.

Typical real-world applications of EM BFS (and some of its generalizations like shortest paths or  $A^*$ )

include crawling and analyzing the WWW [29, 32], route planning using small navigation devices with flash memory cards [20], state space exploration [18], etc. In some of these cases the I/O-bottleneck is alleviated by heuristics, precomputations, accepting approximate solutions or profiting from special graph properties. In contrast, despite major efforts, I/O-efficient exact BFS for general sparse graphs was considered nonviable until recently: the MM\_BFS approach in [26] is the first to yield an  $o(n)$ -I/O bound for general undirected sparse graphs. However, we are not aware of any empirical study to evaluate the practical merits of either MM\_BFS or its alternatives (most prominently MR\_BFS [28]).

In this paper, we attempt at narrowing the gap between theory and practice by engineering and extensively testing implementations of MR\_BFS and MM\_BFS on a low-cost machine. The code will become publicly available to be used with the STXXL library [15, 16]. Even though we use some special features of STXXL, we conjecture that (modulo some constants) our results transfer to other libraries that efficiently implement scanning and sorting like, e.g., TPIE [4, 6].

In a nutshell, our findings for EM BFS on sparse graphs are as follows: Most importantly, our study demonstrates that I/O-efficient external-memory BFS is feasible at all: for example, we computed the BFS level decomposition of an external web-crawl based graph of around 130 million nodes and 1.4 billion edges in less than 4 hours using a single disk and 2.3 hours using 4 disks. It turned out that both MR\_BFS and MM\_BFS perform significantly better (*minutes* as compared to *hours*) than internal-memory BFS, even if less than half of the input resides externally. Furthermore, MR\_BFS exhibits some advantages over MM\_BFS on low-diameter graphs (saving a few *hours*) whereas MM\_BFS wins hands-down on most large-diameter graphs (a few *days* versus a few *months*). Altogether, in many situations, the moderate overhead of MM\_BFS on ‘easy’ instances seems to be an acceptable investment given its significant gains on really ‘tough’ inputs. We expect our results to prove helpful in judging the potential gains and limitations of more complicated (future) approaches, too.

The rest of the paper is organized as follows: In Section 2, we give a short description of the external memory model and the BFS algorithms considered in

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this study. In Section 3, we deal with various implementation issues - STXXL, data-structures, pipelining. We also describe our framework for generating graphs, BFS traversal routines and BFS verifier. In Section 4 we discuss our results on various graph classes.

## 2 Models and Algorithms

In connection with graph algorithms, the commonly accepted external-memory model by Aggarwal and Vitter [1] defines some parameters:  $M$  ( $< n + m$ ) is the number of vertices/edges that fit into internal memory, and  $B$  is the number of vertices/edges that fit into a disk block. In an I/O operation, one block of data is transferred between disk and internal memory. The measure of performance of an algorithm is the number of I/Os it performs. The number of I/Os needed to read  $N$  contiguous items from disk is  $\text{scan}(N) = \Theta(N/B)$ . The number of I/Os required to sort  $N$  items is  $\text{sort}(N) = \Theta((N/B) \log_{M/B}(N/B))$ . For all realistic values of  $N$ ,  $B$ , and  $M$ ,  $\text{scan}(N) < \text{sort}(N) \ll N$ . Advanced models include parallel disks [35] or hide the parameters  $M$  and  $B$  from the algorithms (cache-oblivious model [19]): due to their generality, cache-oblivious algorithms are efficient on each level of the memory hierarchy. A comprehensive list of results for the I/O-model and memory hierarchies have been obtained – for recent surveys see [14, 27, 34] and the references therein.

**EM-BFS Algorithms covered in this study.** In this paper we compare the performance of three BFS algorithms on large inputs: the standard internal-memory approach (referred as IM\_BFS) [12], an algorithm by Munagala and Ranade (hereafter referred as MR\_BFS) [28], and an approach by Mehlhorn and Meyer (MM\_BFS) [26]. MR\_BFS and MM\_BFS are restricted to undirected graphs.

IM\_BFS visits the vertices in a one-by-one fashion; appropriate candidate nodes for the next vertex to be visited are kept in a FIFO queue  $Q$ . After a vertex  $v$  is extracted from  $Q$ , the adjacency list of  $v$  is examined in order to append unvisited nodes to  $Q$ . IM\_BFS in external memory has two drawbacks: (1) remembering visited nodes needs  $\Theta(m)$  I/Os in the worst case; (2) unstructured indexed access to adjacency lists may result in  $\Theta(n)$  I/Os. MR\_BFS neglects (2) but addresses (1) by I/O-efficiently comparing the BFS level under creation with the two most recently completed BFS-levels. The resulting worst-case I/O-bound is  $O(n + \text{sort}(n + m))$ .

MM\_BFS applies a preprocessing phase in which it groups the vertices of the input graph into disjoint clusters of small diameter and then runs an appropriately modified version of MR\_BFS. It exploits the fact that whenever the first node of a cluster is visited then the remaining nodes of this cluster will be reached soon

after. By spending only one random access (and possibly, some sequential access depending on cluster size) in order to load the whole cluster and then keeping the cluster data in some efficiently accessible data structure (pool) until it is all used up, on sparse graphs the total amount of I/O can be reduced by a factor of up to  $\sqrt{B}$ : edges may be scanned more often in the pool but unstructured I/O in order to fetch adjacency lists is reduced. The preprocessing of MM\_BFS can be done in several ways. We implemented a simple randomized variant which yields an overall expected I/O-bound of  $O(\sqrt{n \cdot (n + m) \cdot \log(n)/B} + \text{sort}(n + m))$ .

**EM-BFS Algorithms not (yet) covered in this study – and why.** An important goal of our study was to prove that I/O-efficient external-memory BFS is feasible at all. Therefore, we chose to implement those approaches with presumably small constant factors first. Also, due to the intrinsic time consumption for large-scale series of tests in external-memory, a certain self-restraint was unavoidable.

The next more complicated approaches are the counterparts of MR\_BFS for directed graphs [10]. While applying extra data structures to remember visited nodes, these algorithms share the weakness of MR\_BFS concerning unstructured access to adjacency lists, thus incurring  $\Omega(n)$  I/Os in the worst case. On non worst-case instances we expect performance similar to that of MR\_BFS. An equivalent to MM\_BFS featuring  $o(n)$  I/Os on general directed sparse graphs is still to be discovered (if it exists).

There is also a bunch of EM-BFS algorithms featuring  $O(\text{sort}(n))$  I/Os on special graphs classes like trees [10], grid graphs [5], planar graphs [25], outer-planar graphs [23], and graphs of bounded tree width [24]. One might be tempted to run the planar BFS approach on general graphs - accepting to loose the worst-case guarantee of small separators, thus causing additional I/O. Leaving aside probably large constants the approach of [3, 25] as it stands takes  $\Omega(n \cdot B)$  internal operations, which more or less translate in the time needed to perform  $\Omega(n)$  I/Os. On the other hand, MR\_BFS and MM\_BFS take  $O(m \cdot \log n)$  and  $O(m \cdot \log n + \min\{\log n, \sqrt{\frac{n \cdot B \cdot D \cdot \log n}{n + m}}\} \cdot \text{scan}(n + m))$  internal operations, respectively.

Another general technique to obtain I/O-efficient algorithms applies some kind of simulation (e.g., [11, 13, 21]) of suitable parallel approaches. As for general BFS this failed so far due to the lack of appropriate parallel BFS algorithms (as a rule of thumb they need to be fast and work-efficient at the same time).

Finally, there is a recent paper [8] transferring MM\_BFS into a cache-oblivious algorithm of similar theoretical performance. It would be interesting to see how the performance of this approach executed

on virtual memory compares to our implementation of MM\_BFS. It might even become interesting for internal-memory graph traversal [22]. The algorithm is somewhat involved but with more and more cache-oblivious basic routines like high-performance sorting becoming available (e.g., [9]), a prototypical implementation of cache-oblivious BFS is on our short-list for future work.

### 3 Implementation Design

**STXXL.** The key component of STXXL used by us is the stream sorter, which runs in two phases - **Runs Creator (RC) Phase**, in which the input vector/stream is divided into chunks of  $M$  elements and each chunk is sorted within itself, thereafter written to the disk space and **Runs Merger (M) Phase**, in which the first blocks of all the sorted chunks are brought to internal memory and merged there to produce the output stream which does not necessarily have to be stored on the disk.

The STXXL stream sorter (from version 0.75 onwards) does not need any I/O if  $n < B$ . Also, for this case, the internal work is proportional to  $n \log n$ , independent of  $B$ . Converting a vector into stream or initialization of runs creator or runs merger do not cause any I/O. Since for graphs with  $O(n)$  BFS levels, we don't want to spend one I/O per level and all these functions are called in each level, these features are crucial for our implementation.

**Data Structures.** In the STXXL design framework, an adjacency array can be implemented using two vectors  $N$  and  $E$ .  $N$  contains the iterators to locations in  $E$ , marking the beginning of a new adjacency array. Each edge is stored twice in  $E$  - once in the adjacency array of each adjacent node. This representation allows scanning the edge vector in  $\text{scan}(2m)$  I/Os. To ensure that we do not spend one I/O per level, but rather incur at most  $\frac{2n}{B}$  I/Os in total, for writing the output, even the output itself — the BFS level decomposition — is stored in a similar format.

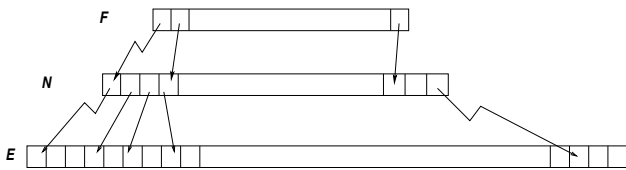


Figure 1: I/O-efficient data structure to represent a partitioned graph

For the case of MM\_BFS, we also need another vector to store iterators to different clusters in order to access an arbitrary cluster in  $O(1) + \frac{\text{cluster size}}{B}$  I/Os. Each cluster consists of an adjacency array containing

nodes and edges belonging to that cluster after the pre-processing phase. A better way (from the I/O efficiency perspective) to handle the graph-partitioning is to store the partitioned input graph as three vectors  $\mathcal{F}$ ,  $N$ , and  $E$  (as shown in Figure 1), containing the cluster iterators, nodes and adjacency arrays, respectively.  $N$  and  $E$  are kept sorted according to cluster indices.

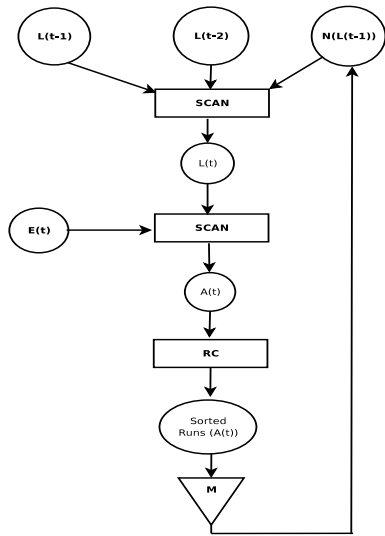
Although we reduced the amount of information kept with node and edge elements in this data-structure, our implementation is still generic: it can handle graphs with arbitrary number of nodes and the graph template is basic and can be used for other graph algorithms as well.

**Pipelining.** Pipelining is a well known principle used to design faster algorithms. The key idea behind pipelining is to connect a given sequence of algorithms with an interface so that the data can be passed-through from one algorithm to another without needing any external memory intermediate storage. This coalesced algorithm is still manageable because none of these algorithms needs to know about the inner structure of the other algorithms. Though pipelining saves some constant factors in the I/O complexity of an algorithm, it usually increases the computational cost. Therefore, the correct extent of pipelining needs to be carefully determined. For more details on the usage of pipelining as a tool to save I/Os, refer to [17].

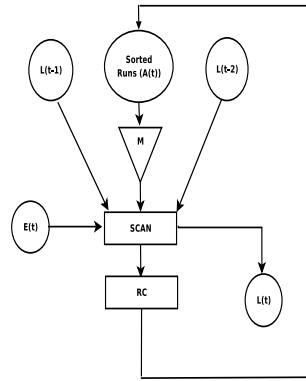
**MR\_BFS.** Figure 2 displays the flowchart of the pipelined and the non-pipelined versions of the main loop of our implementation of MR\_BFS. Let  $L(t)$  denote the set of nodes in BFS level  $t$  and  $N(S)$  denote the set of neighbors of nodes in  $S$ . The complexity of the pipelined MR\_BFS mainly lies in its scanner. The scanner reads the nodes in  $L(t-1)$ ,  $L(t-2)$  and the adjacency lists  $E(t)$  of nodes in  $L(t)$  from the disk and scans through the stream of  $N(L(t-1))$  and in just one pass outputs the nodes in the current level  $L(t)$  and the multi-set  $A(t)$  of neighbor nodes of  $L(t)$ . The stream  $A(t)$  is passed directly to runs creator and sorted runs are written on the disk. These are later merged and passed to the scanner as  $N(L(t))$  for the next iteration.

In this case, pipelining reduces the worst case number of I/Os from  $\sum_t (L(t-1) + L(t-2) + 2L(t) + E(t) + 2A(t) + \text{sorted runs of } A(t) + 2N(L(t-1))) = n + \text{scan}(4n + 14m)$  to  $\sum_t (L(t-1) + L(t-2) + L(t) + E(t) + \text{sorted runs of } A(t)) = n + \text{scan}(3n + 6m)$ .

**MM\_BFS.** Figure 3 shows the flow-chart of the pipelined version of both phases of MM\_BFS. The randomized graph partitioning phase begins with randomly selecting each node to be a master node with a probability  $\mu$ . The main scanner of this phase takes the sorted sequence of the nodes on the fringe of expanding clusters, stores the cluster index (by including the fringe nodes into their corresponding clusters) with

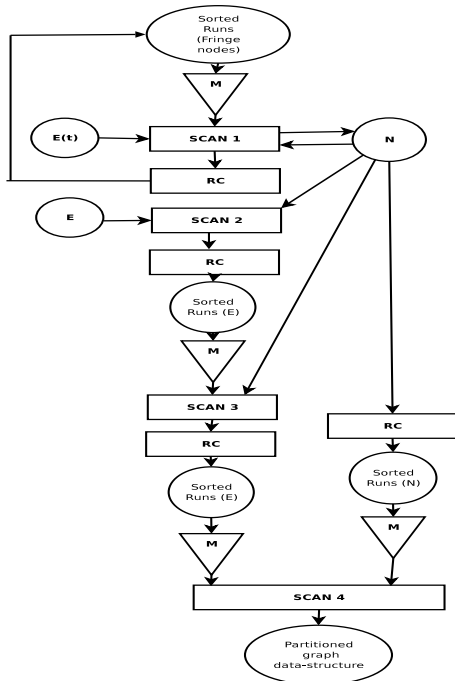


Flow-chart of MR\_BFS

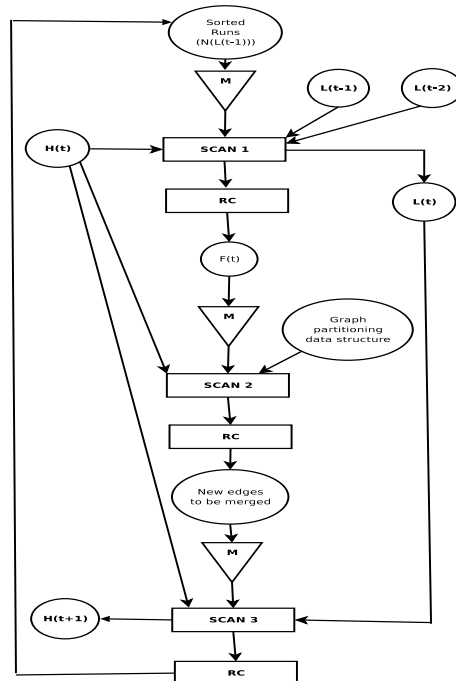


Flow-chart of pipelined version of MR\_BFS

Figure 2: Flow-chart of MR\_BFS implementation



Graph partitioning phase of MM\_BFS



BFS phase of MM\_BFS

Figure 3: Flow-chart of MM\_BFS implementation

these nodes and reads their adjacency lists to compute the new sequence of fringe nodes to be sent to the two-phase sorter. After the partitioning of nodes into clusters is complete, we store the cluster index of the incident nodes with each edge, sort  $N$  and  $E$  w.r.t. the cluster index and then adjust the cluster and the node iterators appropriately. Since the diameters of any cluster is less than  $\frac{\log n}{\mu}$  w.h.p., the total number of I/Os for this phase is bounded by  $\text{scan}(16m + 6n + \mu n + \frac{2 \cdot (m+n) \cdot \log n}{\mu})$  w.h.p.

In the pipelined BFS phase, the first scanner receives the sorted sequence  $N(L(t-1))$  of neighbor nodes of  $L(t-1)$  from the merger pipe, reads  $L(t-1)$ ,  $L(t-2)$  from the disk and the adjacency lists of nodes in  $L(t-1)$  from the hot pool  $\mathcal{H}$  and computes  $F(t)$  - the multi-set of cluster indices of nodes in  $L(t)$  - and in the process, also writes  $L(t)$  to disk. The second scanner takes the sorted  $F(t)$  and the hot pool  $\mathcal{H}$  to compute the multi-set of cluster indices that need to be merged into  $\mathcal{H}$ . The next scanner reads the sorted sequence of  $F(t)$ , eliminates duplicate cluster-indices, computes  $A(t)$  - the neighbor multi-set of  $L(t)$  - and updates  $\mathcal{H}$ :

$$\mathcal{H}_{new} = \mathcal{H}_{old} + \mathcal{H}1(\text{Merged clusters}) - \text{Adj}(L(t))$$

where  $\text{Adj}(S)$  represents the adjacency arrays of nodes in  $S$ . The total number of I/Os for this phase is bounded by  $\mu n + \text{scan}(14m + 4n + \frac{8m \log m}{\mu})$  w.h.p.

**Graph generators.** For the purpose of this study, we designed and implemented a pipelined version of an I/O efficient framework for generating large graphs. The key steps of this framework are edge-sequence creation, sorting the edge-sequence, duplicate removal and conversion into requisite adjacency array representation. For an I/O-efficient random permutation needed in the generation process of many graphs, we use [31]. More details for this framework are available in [2].

**BFS decomposition verifier.** As another side tool, we designed an I/O efficient verifier routine to determine whether or not a BFS level decomposition is correct for a given graph.

For an undirected graph, any BFS level labeling is correct if it satisfies the following criterion:

- BFS level 0 contains the source node only.
- Every node has a unique BFS level.
- $\forall (u, v) \in E, |bfs\_level(u) - bfs\_level(v)| \leq 1$ .
- $\forall u \in V$  in BFS level  $k$  ( $k \neq 0$ ),  $\exists edge(u, v)$  such that  $v$  is in BFS level  $k - 1$ .

All these conditions can be checked in  $O(\text{sort}(n + m))$  I/Os.

## 4 Experiments

In this section, we consider the total running time and the I/O wait time - the total time spent by an implementation waiting for an I/O to complete, and not I/O time - the total time spent by an implementation on I/Os. This distinction is necessary as STXXL maximizes the overlap of I/O with computation.

**Configuration.** We have implemented the algorithms in C++ using the g++ 3.3.2 compiler (optimization level -O3) on the GNU/Linux distribution *Debian* with a 2.4 kernel and the external memory library STXXL version 0.77. Our experimental platform has two 2.0 GHz Intel Xeon processors (we use only one), one GByte of RAM, 512KB cache and 250 GByte Seagate Barracude hard-disks [33]. These hard-disks have 8 MB buffer cache. The average seek time for read and write is 8.0 and 9.0 msec, respectively, while the sustained data transfer rate for outer zone (maximum) is 65 MByte/s. This means that  $2^{28}$  random read and write I/Os will take 596.5 and 671.1 hours, respectively.

**Graph classes considered.** *Random graph:* On a  $n$  node graph, we randomly select  $m$  edges with replacement (i.e.,  $m$  times selecting a source and target node such that source  $\neq$  target) and remove the duplicate edges to obtain random graphs.

*B-level random graph:* This graph consists of  $B$  levels, each (except the level 0 containing only the source node) having  $\frac{n}{B}$  nodes. The edges are randomly distributed between consecutive levels, such that these  $B$  levels approximate the BFS levels. The initial layout of the nodes on the disk is such that the  $\frac{n}{B}$  nodes in the same level are all in different blocks. This graph causes MR\_BFS to incur its worst case of  $\Omega(n)$  I/Os.

*B-level spider web graph:* This graph class is a specialization of web graph defined in [37]. Again, it consists of  $B$  levels, each having  $\frac{n}{B}$  nodes. All nodes in a level are connected in a cyclic fashion and a node has an edge to its corresponding node in the level before and after. The initial layout of the nodes on the disk is random.

*Grid graph:* It consists of a  $\lfloor n \rfloor \times \lfloor n \rfloor$  grid.

*MM BFS worst graph:* This graph [7] causes the (randomized pre-processing variant) MM\_BFS to incur its worst case of  $\Theta(n \cdot \sqrt{\frac{\log n}{B}} + \text{sort}(n))$  I/Os.

*Line graphs:* A line graph consists of  $n$  nodes and  $n - 1$  edges such that there exists two nodes  $u$  and  $v$ , with the path from  $u$  to  $v$  consisting of all the  $n - 1$  edges. We took three different initial layouts - simple, in which all blocks consists of  $B$  consecutively lined nodes; B-interleaved in which the consecutive nodes are all in different but consecutive blocks; random in which the arrangement of nodes on disk is given by a random permutation.

**Fine-tuning Parameters.** Most practitioners of

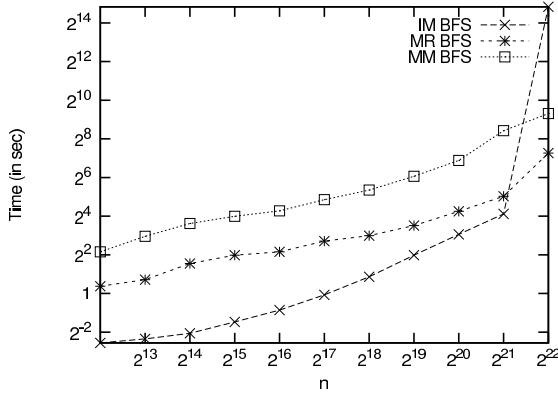


Figure 4: Variation of running time of IM\_BFS, MR\_BFS and MM\_BFS (in logarithmic scale) with graph size (also in logarithmic scale)

external memory algorithms know that the block size  $B$  is a parameter that needs to be finely tuned for optimal performance. This is all the more relevant in the STXXL design framework, as the STXXL vector is organized as a collection of blocks (of size  $B$ ) residing on the external storage media (parallel disks). Access to the external blocks is organized through the fully associative *cache* which consists of  $Pg\_Nr$  in-memory pages where a page is a collection of  $Blk\_Nr$  consecutive blocks. The internal memory consumption of a vector is  $B \cdot Blk\_Nr \cdot Pg\_Nr$ . Another important parameter to be fine-tuned is  $M\_use$  - the internal memory reserved for a runs creator (or equivalently, runs merger). While tuning these parameters, a key constraint is that the internal memory allocated for all the vectors, runs creator and runs merger active simultaneously, at any time, should be less than the internal memory available for the user. Typically, half of the internal memory is kept for OS requirements. For our implementations, we chose  $B = 512$  KB (after experimenting between 1 KB and 8 MB),  $Pg\_Nr = 4$ ,  $Blk\_Nr =$  number of parallel disks in use, allocation strategy = randomized striping and LRU page replacement strategy.

Another important parameter to be optimized for MM\_BFS is  $\mu$  - the probability of choosing a node to be a master node. For worst case optimality, we choose  $\mu = \sqrt{\frac{(m+n) \cdot \log(n)}{n \cdot B \cdot D}}$ . On the other hand, the number of I/Os in the BFS phase for graphs with low (high) diameter can be quite less (high) and therefore, the optimal  $\mu$  value for these graphs is much higher (lower). However, in order to exploit this fact, we need to assume some a priori knowledge of the graph structure. We consider both the cases - one in which we choose our  $\mu$  value independent of the graph-structure (common  $\mu$ ) and one in which we assume a priori knowledge of it

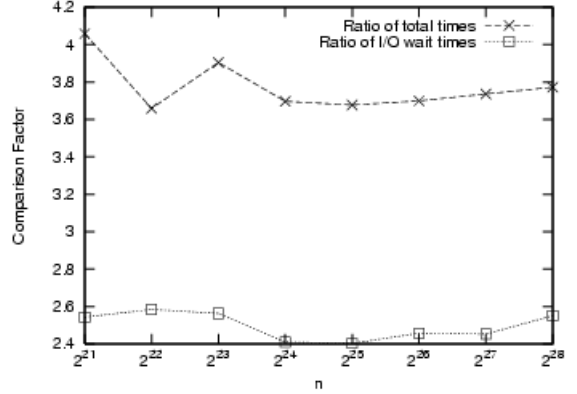


Figure 5: Variation of comparison factor (MM\_BFS/MR\_BFS) of running time and I/O wait time with random-graph size

(graph-structure dependent  $\mu$ ).

**IM\_BFS loses fast.** Figure 4 shows the total running time of IM\_BFS, MR\_BFS, and MM\_BFS on random graphs of varying sizes (keeping  $m = 4n$ ). An important point to note here is that even when half of the graph fits in internal memory, the performance of IM\_BFS is much worse than that of external BFS algorithms. For this case ( $2^{22}$  nodes and  $2^{24}$  edges), the I/O wait time of IM\_BFS (8.09 hours) dominates the total running time (8.11 hours), thereby explaining the worse behavior of IM\_BFS. On the other hand, MR\_BFS and MM\_BFS have much less I/O wait time (1.55 and 4.93 minutes) and consequently, the total running time (2.57 and 10.6 minutes) is also small. This clearly establishes the need for efficient implementations of external memory BFS algorithms.

Note that some of the results in Table 1- 9 have been interpolated using the symmetry in the graph structure.

**Single disk - common  $\mu$ .** Table 1 and 2 show the I/O wait time and running time for different graphs in the single disk common  $\mu$  case. Note that for these large graphs, even the efficient implementations of external memory algorithms are I/O dominant.

Let's first consider the case of random graphs. The total time for BFS traversal on random graphs is much less than that for most other graph classes. This is explained by the fact that MR\_BFS incurs  $O(\text{sort}(n+m))$  I/Os per level, and the number of BFS levels in random graphs (typically 10-15 for the graph sizes we studied) are very few. Infact, it's known [30] that a random graph  $G(n, c/n)$  has an expected diameter  $O(\log n)$ . On the other hand, since  $\mu$  for MM\_BFS is  $\sim \sqrt{\frac{\log n}{B}}$  for the asymptotic worst case I/O complexity, the edges remain in hot pool for quite long and MM\_BFS

Graph class	n	m	MR_BFS		MM_BFS	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	2.4	3.4	7.3	9.6
Grid	$2^{28}$	$\sim 2^{29}$	4733.0	4736.0	52.1	53.9
<i>B</i> -level random	$2^{28}$	$2^{30}$	3989.8	3994.8	42.2	49.7
<i>B</i> -level spider web	$2^{28}$	$\sim 2^{29}$	3364.2	3366.5	36.5	39.8
MM Worst	$2^{25}$	$\sim 2^{25}$	25.2	25.4	19.5	32.4
Simple line	$2^{28}$	$2^{28}$	0.6	10.2	84.8	275.9
Rand line	$2^{28}$	$2^{28}$	4156.2	4167.7	100.2	283.3
B-interleaved line	$2^{28}$	$2^{28}$	4210.3	4222.6	97.5	280.8

Table 1: Single Disk Common  $\mu$  - I/O wait time and running time (in hours) of MR\_BFS and MM\_BFS

Graph class	n	m	MM_BFS Phase 1		MM_BFS Phase 2	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	3.7	5.1	3.6	4.5
Grid	$2^{28}$	$\sim 2^{29}$	6.6	7.3	45.5	46.6
<i>B</i> -level random	$2^{28}$	$2^{30}$	3.6	5.1	38.6	44.6
<i>B</i> -level Spider Web	$2^{28}$	$\sim 2^{29}$	6.6	7.3	29.9	32.5
MM Worst	$2^{25}$	$\sim 2^{25}$	6.5	6.7	13.0	25.7
Simple line	$2^{28}$	$2^{28}$	84.3	85.1	0.5	190.8
Rand line	$2^{28}$	$2^{28}$	79.4	80.6	20.8	202.7
B-interleaved line	$2^{28}$	$2^{28}$	79.3	80.4	18.1	200.4

Table 2: Single Disk Common  $\mu$  - I/O wait time and running time (in hours) of different phases of MM\_BFS

Graph class	n	m	MR_BFS		MM_BFS	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	0.9	1.3	1.9	4.4
<i>B</i> -level Random	$2^{28}$	$2^{30}$	2094.3	2105.1	18.3	26.0
<i>B</i> -level Spider Web	$2^{28}$	$\sim 2^{29}$	1492.0	1497.9	13.5	17.1
MM Worst	$2^{25}$	$\sim 2^{25}$	13.4	13.7	6.5	10.5
Rand line	$2^{28}$	$2^{28}$	4702.8	4730.5	56.4	239.9
B-interleaved line	$2^{28}$	$2^{28}$	1243.0	1258.7	56.4	239.9

Table 3: Multi disk common  $\mu$  - I/O wait time and running time (in hours) of MR\_BFS and MM\_BFS

Graph class	n	m	MM_BFS Phase 1		MM_BFS Phase 2	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	0.9	2.5	1.0	1.9
<i>B</i> -level random	$2^{28}$	$2^{30}$	0.9	2.5	17.4	23.5
<i>B</i> -level Spider Web	$2^{28}$	$\sim 2^{29}$	2.4	3.2	11.1	13.9
MM Worst	$2^{25}$	$\sim 2^{25}$	5.3	5.6	1.2	4.9
Rand line	$2^{28}$	$2^{28}$	49.0	50.5	7.5	189.4
B-interleaved line	$2^{28}$	$2^{28}$	48.4	49.0	6.1	28.1

Table 4: Multi disk common  $\mu$  - I/O wait time and running time (in hours) of the two phases of MM\_BFS

can still take  $O(n \cdot \sqrt{\frac{\log n}{B}})$  I/Os. Consequently, as shown in Figure 5, MR\_BFS not only outperforms MM\_BFS in terms of total running time by a factor of  $\sim 3.8$ , but also in terms of I/O wait time by a factor of  $\sim 2.5$ .

As noted in Section 2, MR\_BFS takes  $O(n + \text{sort}(n + m))$  I/Os while MM\_BFS takes  $O(\text{sort}(n +$

$m) + \sqrt{n \cdot \text{scan}(n + m) \cdot \log(n)})$  I/Os w.h.p.. So, if the two implementations are I/O bound and  $\frac{m \cdot \log(n)}{n} < B$ , MM\_BFS provides better running-time guarantees w.h.p.. In other words, we should be able to find some sparse graphs for which MM\_BFS performs significantly better than MR\_BFS. Our search for such a graph led

Graph class	n	m	MR_BFS		MM_BFS	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	2.4	3.4	5.5	7.9
<i>B</i> -level Random	$2^{28}$	$2^{30}$	3989.8	3994.8	10.0	16.6
<i>B</i> -level Spider Web	$2^{28}$	$\sim 2^{29}$	3364.2	3366.5	25.1	29.3

Table 5: Single Disk, Graph structure dependent  $\mu$  - I/O wait time and running time (in hours) of MR\_BFS and MM\_BFS

Graph class	n	m	MM_BFS Phase 1		MM_BFS Phase 2	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	3.3	4.9	2.2	3.0
<i>B</i> -level Random	$2^{28}$	$2^{30}$	4.0	5.5	6.0	11.1
<i>B</i> -level Spider Web	$2^{28}$	$\sim 2^{29}$	12.9	13.7	12.2	15.6

Table 6: Single Disk, Graph structure dependent  $\mu$  - I/O wait time and running time (in hours) of the two phases of MM\_BFS

Graph class	n	m	MR_BFS		MM_BFS	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	0.9	1.3	1.5	3.9
<i>B</i> -level Random	$2^{28}$	$2^{30}$	2094.3	2105.1	3.1	9.7
<i>B</i> -level Spider Web	$2^{28}$	$\sim 2^{29}$	1492.0	1497.9	9.7	13.9

Table 7: Multi disk graph structure dependent  $\mu$  - I/O wait time and running time (in hours) of MR\_BFS and MM\_BFS

Graph class	n	m	MM_BFS Phase 1		MM_BFS Phase 2	
			I/O wait time	Total time	I/O wait time	Total time
Random	$2^{28}$	$2^{30}$	0.7	2.3	0.8	1.6
<i>B</i> -level random	$2^{28}$	$2^{30}$	1.1	2.6	2.0	7.1
<i>B</i> -level Spider Web	$2^{28}$	$\sim 2^{29}$	4.3	5.1	5.4	8.8

Table 8: Multi disk graph structure dependent  $\mu$  - I/O wait time and running time (in hours) of the two phases of MM\_BFS

	MR_BFS		MM_BFS - common $\mu$		MM_BFS - Graph dep $\mu$	
	I/O wait time	Total time	I/O wait time	Total time	I/O wait time	Total time
Single disk	3.7	4.0	7.4	9.4	6.3	8.4
Multiple disk	2.0	2.3	2.7	4.8	2.3	4.5

Table 9: I/O wait time and running time (in hours) of the two algorithms on web graph

us to *B*-level random graphs and *B*-level spider web graphs — high diameter sparse graphs, where MR\_BFS takes  $\Theta(n)$  I/Os. For these graphs, MR\_BFS takes 90-95 times more I/O wait time than MM\_BFS and as a result, while MM\_BFS takes a total running time of 49.7 and 39.8 *hours* respectively, MR\_BFS takes 166.5 and 140.3 *days*. A  $\lceil\sqrt{n}\rceil \times \lfloor\sqrt{n}\rfloor$  grid is yet another instance of a high diameter  $\lceil\sqrt{n}\rceil + \lfloor\sqrt{n}\rfloor$  sparse graph. Once again, MM\_BFS outperforms MR\_BFS by a factor of 90.8 in I/O wait time and 87.9 in total running time.

Apart from diameter, another important consideration affecting the relative performance of the two algo-

rithms is the initial graph layout on the disk. The pre-processing phase of MM\_BFS neutralizes the impact of an adverse layout. So, while we observe that the I/O wait time of MR\_BFS (0.6 hours) is much less than 84.8 hours of MM\_BFS (dominated by the 84.3 hours in the pre-processing phase) on a simple line graph, the I/O wait time of MR\_BFS (167.6 and 177.7 days) is much more than that of MM\_BFS (4.2 and 4.1 days) on a random and *B*-interleaved layouts. Thus, pre-processing makes MM\_BFS provide better worst case guarantees (saving *months*) at the cost of losing out on simple layouts (losing *days*).



Except for the line graphs, the pre-processing phase dominates the run-time of MM\_BFS for low diameter graphs while the BFS phase dominates it for large diameter graphs. For the line graphs, the hot pool is rarely bigger than  $B$  and therefore, always stays in internal memory.

**Multi disk - common  $\mu$ .** We ran the same experiments with vectors striped over 4 disks. Although the usage of multiple disks allows us to handle larger volumes of data, herein we restrict ourselves to smaller sizes for better comparison with the single disk case. Table 3 and 4 show the running time of the two algorithms on different graphs. For most graphs, the usage of parallel I/O channels alleviates the I/O problem further and computation starts becoming the bottleneck, in particular for MM\_BFS, which seems to gain more from the parallel I/O channels. However, with some new features of STXXL like a SMP multi-processor version of sorting routines, we hope to bring down the total running time fairly close to the I/O wait time. Besides, the computation speed increases at a much faster rate than the external memory bandwidth, thereby reducing the computation time relative to the I/O wait time.

While MR\_BFS on random line hardly seems to have any benefit from the multiplicity of disks, it is almost four times better with four disks on B-interleaved line. This is because a random access to a block brings the neighboring blocks on other disks automatically to the internal memory and therefore, the access to the adjacency lists of next three nodes comes without any external I/O.

**Single disk - graph structure dependent  $\mu$ .** Table 5 and 6 show the I/O wait time and running time for the two algorithms in the single disk case, where  $\mu$  could be optimized based on the graph structure. Since for random graphs, there are very few BFS levels, it is better to choose a very high  $\mu$  value ( $\mu = 0.7$ ) so that a cluster stays in the hot pool for a short duration and therefore, an edge is considered in the hot pool for only one or two levels. With this  $\mu$ , the I/O wait time of the BFS phase of MM\_BFS is less than MR\_BFS. So, in the case when we need to compute BFS from different source nodes knowing the graph-structure, the time for graph-partitioning can be amortized on multiple BFS runs, thereby making MM\_BFS the preferable algorithm, even for random graphs. Since for  $B$ -level random and spider web graph, the hot pool almost always stays in the internal memory, the larger diameter of the cluster is no longer a problem. Consequently, we choose small  $\mu$  value ( $\mu = 0.00025 \simeq \frac{1}{B}$  and  $\mu = 0.001$ ) in order to have fewer clusters and therefore, less time in loading them. In general, the  $\mu$  value should be chosen so as to balance the I/O time of the two phases of MM\_BFS.

**Multi disk - graph structure dependent  $\mu$ .** Next, we experimented with running MM BFS on multiple-disk with graph-structure dependent  $\mu$  to measure the extent to which we can alleviate the I/O bottleneck of BFS. Table 7 and 8 show the results for this setup.

**Web graph - A real world graph.** As an instance of a real world graph, we consider an actual crawl of the world wide web [36], where an edge represents a hyperlink between two sites. This graph has around 130 million nodes and 1.4 billion edges. For this graph, the total input and output volume of the BFS algorithm (with the data-structure described earlier) is around 23.5GB and 1.5GB, respectively. The bulk of the nodes are contained in the core of this web graph spread across 10-12 BFS levels (similar to random graphs). The remaining nodes are spread out over thousands of levels with 2-3 nodes per level (which behaves more like a line graph). However, the I/O wait time as well as the total running time for BFS traversal is dominated by the core of this graph and hence, the results are similar to the ones for random graphs. As Table 9 shows, both the algorithms can compute the BFS decomposition of this graph in a matter of few *hours*.

## 5 Conclusion

Empirical evidence suggests that MR\_BFS performs better on small-diameter random graphs. However, the better asymptotic worst-case I/O complexity of MM\_BFS helps it to outperform MR\_BFS for large diameter sparse graphs, where MR\_BFS incurs close to its worst case of  $\Omega(n)$  I/Os.

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