

# Cache-Oblivious Dynamic Search Trees

by

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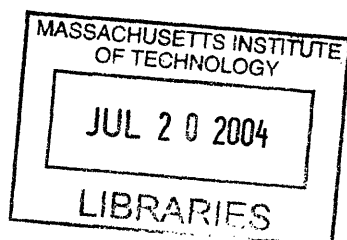
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## Abstract

I have implemented a cache-oblivious dynamic search tree as an alternative to the ubiquitous B-tree. I use a binary tree with a “van Emde Boas” layout whose leaves point to intervals in a “packed memory structure”. We refer to the data structure as a COB-Tree. The COB-Tree supports efficient lookup, as well as efficient amortized insertion and deletion. Efficient implementation of a B-tree requires understanding the cache-line size and page size and is optimized for a specific memory hierarchy. In contrast, the COB-Tree contains no machine-dependent variables, performs well on any memory hierarchy, and requires minimal user-level memory management. For random insertion of data, my data structure performs 7.5 times faster than the Berkeley DB and 3 times faster than a memory mapped implementation of B-trees. The packed memory array of the COB-Tree maintains data in sorted order, allows sequential reads at high speeds, and data insertions and deletions with few data writes on average. In addition, the data structure is easy to implement because I employed memory mapping rather than explicit file operations.

Thesis Supervisor: Dr. Bradley Kuszmaul  
Title: Research Scientist



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# Chapter 1

## Introduction

For decades, B-trees [2, 11] have been the predominant dictionary data structure. The B-tree is a  $k$ -ary tree where  $k$  is selected such that each node of the B-tree is the size of a block of disk. B-trees minimize the number of disk block transfers from disk to main memory needed to traverse down a path of the tree. Although efficient, B-trees require the programmer to experimentally tune the value of  $k$  on any machine used. Because machines are complex, finding the optimal  $k$  is a complex task that may even be impossible.

This thesis presents the implementation and testing of an alternative data structure that requires no such tuning, a “cache-oblivious” dynamic search tree. We refer to this data structure as a *COB-Tree*. The data structure is cache-oblivious because the implementation does not depend on any machine dependent variables such as disk block size or main memory size. Still, the COB-Tree has been proven to perform within a constant factor of the least number of memory transfers possible between *any* consecutive levels of memory.

The COB-Tree, presented by Bender, Demaine, and Farach-Colton [7], has two components: a static cache-oblivious binary tree and a loosely packed array. Data is kept in sorted order in the array. The leaves of the binary tree correspond to certain sections of the array. To insert elements, we search the tree to find the appropriate section and try to insert the element while moving as few other elements as possible. Infrequently, when sections of the array become too dense, we redistribute a larger portion of the array evenly.

The COB-Tree is tested on large test cases, comparing performance with the standard solution, B-trees, and analyzing asymptotic properties. One set of experiments involving disk access shows comparisons between the cache-oblivious search tree and B-trees. I compare performances using two insertion patterns and analyze results. When inserting elements with a random insertion pattern, the COB-Tree performs 3 times faster than a memory-mapped implementation of B-trees and 7 times faster than the Berkeley DB [23]. When inserting elements repeatedly into the same location, the COB-Tree performs only 3.5 times worse than both memory mapped B-trees and the Berkeley DB. Because we require data to remain in sorted order on disk, we expected much worse. Another set of experiments analyzes asymptotic runtime behavior of the cache-oblivious search tree. I compare experimental behavior of runtime, number of data moves, and data distribution costs with costs derived by theoretical analysis of the data structure on simple memory models.

I present details of the implementation of the COB-Tree, including arguments for why the implementation is efficient. The binary tree is an array in memory with a non-intuitive layout, the van Emde Boas layout. I implemented a function that converts indices from a simple layout, the breadth-first layout, to the layout of the cache-oblivious binary tree. Thus, the details of the cache-oblivious layout are abstracted from the programmer. The programmer, using the function, may program as though he is using the simple breadth-first layout. I present details on implementing each phase of the insertion algorithm, along with reasons for each non-trivial algorithmic decision. I show how the data is stored in memory using a single-level store. Finally, I present several solutions for the most work-intensive operation, redistributing many elements across the array.

The rest of this thesis is organized as follows. Chapter 2 describes, in detail, the cache-oblivious dynamic search tree, as developed over the last few years. We review the performance model used in previous literature to analyze cost and summarize theoretical arguments for why the tree performs well. Chapter 3 presents experimental results. Chapter 4 presents details of the implementation of the static cache-oblivious binary tree. Chapter 5 describes how to implement the loosely packed array.

# Chapter 2

## Description

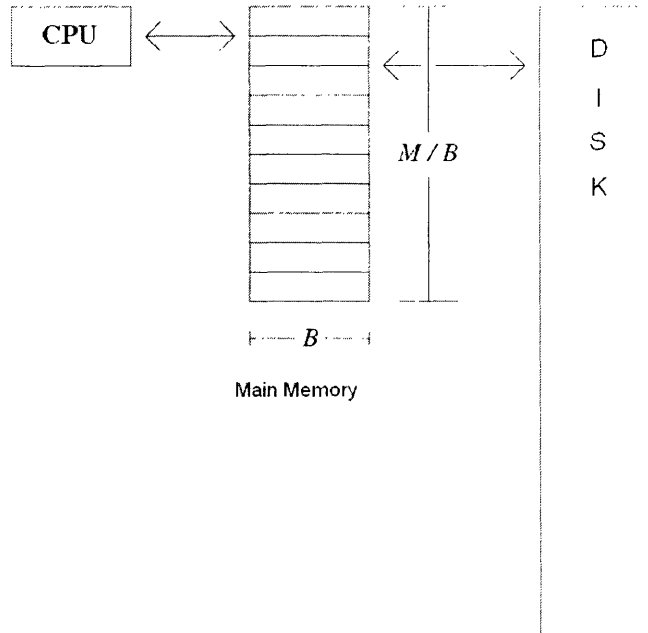
We focus on the problem of creating a data structure that supports efficient data scans, searches, insertions, and deletions. The traditional solution, B-trees, has limitations. B-trees perform sub-optimally on machines with complex memory hierarchies and employ machine dependent variables such as block size of disk. A simple and efficient alternative data structure, the COB-Tree, has been shown to perform asymptotically optimally on any memory hierarchy while not depending on any machine dependent variables. This chapter explains why, while Chapter 3 provides experimental results of the COB-Tree, and Chapters 4 and 5 present implementation details.

This chapter is organized as follows. Section 2.1 describes B-trees and their limitations. Section 2.2 presents the cache-oblivious model, which helps explain why COB-Trees theoretically perform well. Section 2.3 introduces the COB-Tree.

### 2.1 B-trees

B-trees are designed to perform optimally on the *Disk Access Machine (DAM) Model* [1], an idealized two-level memory model consisting of a CPU, main memory of size  $M$ , and disk, broken into *blocks* of size  $B$ . Unlike the *Random Access Memory* model, which assumes constant access time for all memory locations, the DAM model assumes *memory transfers*

of blocks from disk to main memory to be the most time consuming operation. Thus, cost is measured in memory transfers. Figure 2-1 shows an example of the two-level memory hierarchy.

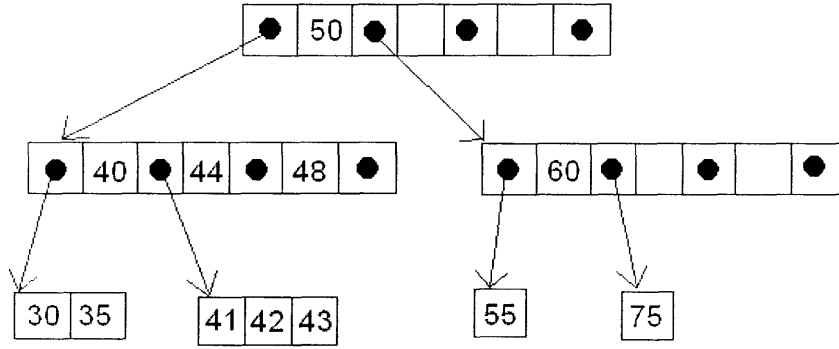


**Figure 2-1:** Two-Level Memory Hierarchy with a disk block size of  $B$  and memory size of  $M$ .

The B-tree works as follows. The B-tree nodes has degree proportional to the block size,  $B$ . Each node of size  $B$  holds as many pairs of keys and pointers as it may. The B-tree has uniform depth and is nearly balanced. Traversing an edge down the tree reduces the search space by a factor proportional to  $1/B$ . Thus, B-trees perform data searches in  $\Theta(1 + \log_B N)$  memory transfers. An information-theoretic argument proves this bound is asymptotically optimal. Figure 2-2 shows an example of a B-tree.

The biggest shortcoming of B-trees is that they are optimized for only an idealized two-level memory hierarchy where  $B$  is fixed. Real memory hierarchies have multiple levels and are more complex than the two-level memory hierarchy. Machines today have multiple levels of memory including registers, level-1 cache, level-2 cache, main memory, and disk. When





**Figure 2-2:** B-tree with capacity of 3 keys per node.

using a multilevel memory hierarchy, the programmer of a B-tree must decide which level of memory is the bottleneck and optimize accordingly. To program efficiently under a multilevel memory hierarchy requires the user to consider multiple block sizes  $B_1, B_2, \dots, B_n$ . To create a tree that is aware of these blocks sizes, aware of the number of memory levels, and performs optimally is very complex, at best.

Define the *effective block size* to be the value,  $B$ , B-trees should use as the size of a node to maximize performance. This value, if it even exists, may differ from the actual block size. Consider the effective block size of disk, the memory level for which most B-trees optimize. The disk consists of not only blocks, but tracks and cylinders as well. Disks engage in prefetching, that is, reading an entire track of data when only a disk block is requested. Tracks near the center of the platter are smaller than tracks near the end, thus the effective block size varies throughout the disk. The time to perform a seek to a track may vary by an order of magnitude, ranging from a track-to-track seek to a full seek. Remapped Bad Sectors change the effective block-transfer size and transfer cost. The Translation Lookaside Buffer complicates things. As a result of all of these complications, finding the optimal value,  $B$ , to optimize a B-tree for disk, is very difficult and may even be impossible.

Most of these effects can be mitigated by improving data locality. Thus, many database-management systems heuristically group logically close disk blocks physically near each other on disk [12, 16, 17, 19, 22]. Alternatively, many authors build B-trees that are optimized for

more detailed models of the memory hierarchy, typically for the DAM model extended to three levels, disk, memory, and cache. Such modified B-trees consist of cache-line B-trees at each node of a larger disk-block B-tree [3, 4, 7, 10]. Indeed, these 2-parameter B-trees have been shown to perform better than 1-parameter B-trees [10] by reducing cache misses.

These 2-parameter B-trees do not model certain disk properties such as prefetching, track size diversity, track locality, work load effects, disk cache. Even when generalizing 2-parameter B-trees to  $k$ -parameters B-trees, address such issues is difficult, because such memory effects are unpredictable and difficult to measure. Finally, it has been shown that the improvement given by increasing  $k$  is limited at least in theory [3].

## 2.2 Cache-Oblivious Model

Before presenting our alternative data structure, we present the *cache-oblivious model* [13, 20]. Like the DAM model, the cache-oblivious model is based on the *Ideal Cache Model* [14, 20]. In the Ideal Cache Model, the cache of size  $M$  is divided into  $M/B$  blocks of size  $B$ . The cache is a *fully associative* cache which can store an arbitrary block of memory into any “slot” of cache. The cost of transferring data from main memory to cache is the most expensive operation. Like the DAM model, a simple, and often accurate, performance measure of algorithms is to count the number of memory transfers of blocks from main memory to cache. Although the model assumes an unrealistic fully-associative cache, the model may be simulated with actual hardware with a constant factor of overhead [14]. Thus, results proved on this model are applicable to pairs of levels in the multilevel memory hierarchy.

Unlike the DAM model, the cache-oblivious model proves results on complex multilevel-memory hierarchies. Algorithms using the cache-oblivious model are not dependent on the block size  $B$  and main memory size  $M$ . If an algorithm performs optimally on the cache-oblivious model, the algorithm performs asymptotically optimally on any unknown multilevel memory hierarchy. From this we define two types of algorithms. *Optimal* algorithms execute

within a constant factor of the least possible number of memory transfers possible. *Optimal cache-aware algorithms* are optimal algorithms aware of the memory parameters of the machine the algorithm is run on,  $M$  and  $B$ . *Optimal cache-oblivious algorithms* perform optimally with no knowledge of  $M$  and  $B$ . We refer to these as cache-aware and cache-oblivious algorithms respectively.

## 2.3 Cache-Oblivious Dynamic Search Trees

The first dynamic cache-oblivious B-tree was proposed by Bender, Demaine and Farach-Colton [7]. Simplified alternatives include [6, 8, 9, 21]. Some of these simplifications have been evaluated experimentally for small data sets [8, 9, 18, 21], but there has been relatively little attempt to analyze the performance of cache-oblivious search structures on very large sets, where disk performance dominates.

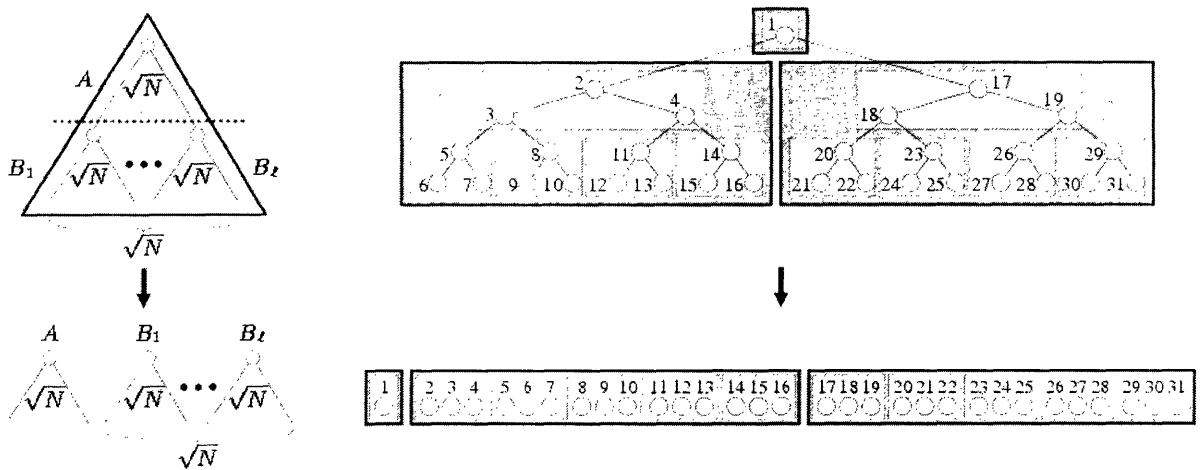
We implement two data structures, a static cache-oblivious binary tree [20] and a packed memory structure [7]. When combined, they form a COB-Tree.

### 2.3.1 Static Cache-Oblivious Binary Tree

The first data structure is a static binary tree with a cache-oblivious layout called a *van Emde Boas* layout. In memory, the tree is an array. The layout describes a mapping of nodes in the tree to positions in the array. Nodes are located in memory such that the tree may be traversed with  $O(1 + \log_B(N))$  memory transfers, which is asymptotically optimal [7].

**Definition of van Emde Boas Layout.** The tree is laid out in memory in a recursive fashion. Let  $h$  be the height of the binary tree. For simplicity, assume  $h$  is a power of 2. Let  $N$  be the number of nodes in the tree. We divide the tree into two sections. The first section is the “top half” containing a subtree, sharing the same root as the tree, of height  $h/2$  with  $\sqrt{N}$  nodes. The second section is the “bottom half” containing  $2^{h/2}$  subtrees of height  $h/2$ , each containing about  $\sqrt{N}$  nodes. This represents subtree  $A$  in Figure 2-3. The idea is to

first layout the top half recursively. Then layout the remaining  $2^{h/2}$  subtrees recursively in order. This represents subtrees  $B_1, B_2, \dots, B_t$  in Figure 2-3. In memory, the entire subtree  $A$  would be laid out first, followed by  $B_1, \dots, B_t$ . We assume the binary tree is full and balanced. If  $h$  is not a power of 2, the bottom half of the tree is chosen such that its height is a power of 2. Figure 2-3 shows the layout of a binary tree with height 5.



**Figure 2-3:** van Emde Boas example in general and of height 5. Figure taken from [7].

### 2.3.2 Packed Memory Structure

The idea behind a *packed memory structure* is to maintain elements of a loosely packed array in sorted order while keeping the average insertion cost low. The goal is to leave enough blank spaces in the array so that for most insertions, few elements need to be moved to accommodate the insertion. When portions of the array become imbalanced, we redistribute elements in a larger portion of the array. This happens infrequently. Bender, Demaine, and Farach-Colton show two important asymptotic bounds regarding this data structure [7]. Insertions and deletions are done with  $O(1 + \frac{\log^2 N}{B})$  memory transfers on average. To scan  $K$  consecutive elements, an additional  $O(1 + K/B)$  memory transfers are required.

A packed memory structure maintains  $N$  elements in order in an array of size  $cN$  for

some  $c > 1$ . The remaining fraction of the array,  $1 - 1/c$ , is blank. Let  $T$  be the size of the array. We specify  $T$  to be a power of 2 at all times. Divide  $T$  into equally sized sections of size  $s = \Theta(\log^2 T)$  such that  $s$  is a power of 2. This forces the number of sections to be a power of 2 as well. Let each section be represented by a leaf of a full and balanced binary tree of depth  $d$ .

Define the *portion* an internal node represents to be the union of all sections represented by the node's leaves. Thus, nodes whose two children are leaves represent two sections, whereas the root represents the entire array. The *capacity* of a node is the maximum number of elements the portion can hold. The *density* of a node is the number of elements maintained in its portion divided by its maximum capacity.

Each node has *upper and lower density threshold bounds* that are a function of its depth (distance from the root). Threshold bounds of a node are boundaries for acceptable densities of the portion the node represents. Their relevance becomes clear in the algorithm for insertion and deletion described below. Let the depth of the root node of the tree be 1 and the depth of all leaves be  $h$ . For each node at an arbitrary depth  $d$ , we have desired upper and lower bound thresholds,  $\tau_d$  and  $\rho_d$  respectively, for the density. We say a node,  $k$ , is within threshold if and only if  $\rho_d \leq \text{density}(k) \leq \tau_d$ . The densities of the root and leaves are  $\rho_1 = .4, \tau_1 = .5, \rho_h = .25$ , and  $\tau_h = 1$  respectively. We have tighter thresholds for nodes of smaller depth. Therefore, we preserve the property,

$$\rho_h < \rho_{h-1} < \dots < \rho_1 < \tau_1 < \tau_2 < \dots < \tau_{h-1} < \tau_h.$$

Thresholds for an internal node at depth  $i$  are selected such that the thresholds  $\tau_1, \tau_2, \dots, \tau_h$  form an arithmetic sequence. Thus,  $\tau_i = \tau_0 + i(\tau_h - \tau_0)/h$ . Similarly,  $\rho_i = \rho_0 - i(\rho_0 - \rho_h)/h$ .

**Insertion** The algorithm for insertion of an element  $i$  is as follows. Search the binary tree to find the appropriate section,  $j$ , where  $i$  belongs. If the section  $j$  is within threshold, insert by arbitrarily moving as few elements around as possible. Otherwise, find the least ancestor

node that is within threshold. *Rebalance* the elements in the portion, along with  $i$ , that is, evenly redistribute all elements in the ancestor's portion. Because this node has tighter threshold bounds than all of descendant nodes, the redistribution forces all descendent nodes to be within threshold. If no ancestor is within threshold, do one of two things. The density of the array is either too high or too low. If the density of the array is higher than the upper threshold of the root node, double the size of the array. If the density of the array is below the lower threshold of the root node, decrease the size of the array by half. In both cases, rebalance after resizing the array.

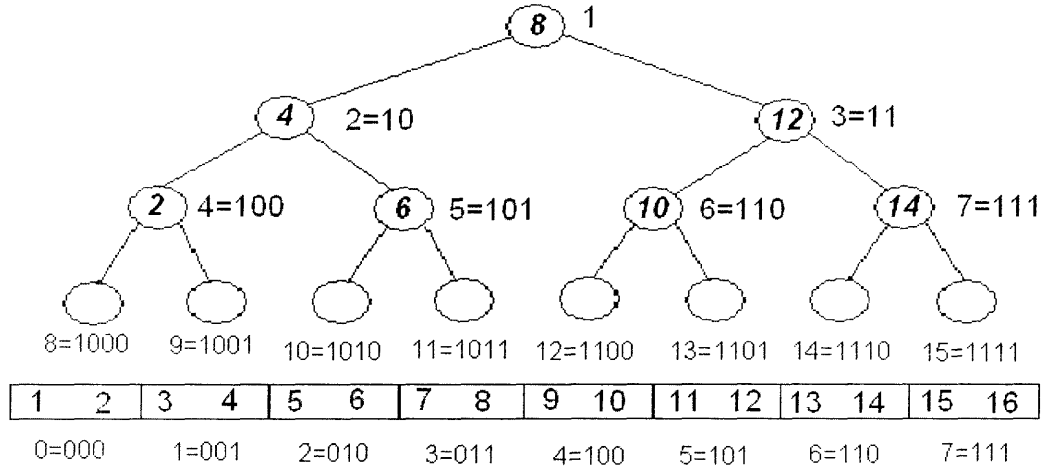
**Deletion** Upon deletion of an element  $i$ , there are two approaches. One is to find the element and perform the same threshold tests as for insertion. The other is to disregard thresholds and remove the element.

### 2.3.3 Combined Data Structure

The COB-Tree is similar to the data structure presented in [8]. We have a binary tree whose leaves correspond to sections (and not individual elements, as in [8]) of a packed memory structure. The binary tree has a van Emde Boas layout. Suppose the tree has  $N$  leaves and height  $h$ . The  $i$ th leaf of the tree, corresponds to the  $i$ th section of the tree. Each internal node of the tree represents a portion, the union of all sections represented by the node's leaves. The value held at each internal node is the highest value of the left child's portion.

### 2.3.4 Related Work.

The packed memory structure and its analysis closely follow [15]. The paper [15] considers the same problem of maintaining elements in order in an array of linear size, but in a different cost model and without the scanning requirement. The structure moves  $O(\log^2 N)$  amortized elements per insertion, but has no guarantee on the number of memory transfers for inserting, deleting, or scanning. This structure has been de-amortized in [24, 25, 26] and subsequently simplified by [5].



**Figure 2-4:** An example of the packed memory structure containing the values 1 through 16. The array contains 8 sections. The binary tree is labeled with a breadth-first layout along with the bit representations of the layout. The sections are labeled below the array. The numbers in bold italics in the nodes are values held by the node.

## Algorithms

**Data Query** Data query is simple. To search for a particular element  $i$ , first search the binary tree to find which appropriate section  $i$  belongs. To do so, we traverse a path of the tree. If  $i$  is less than or equal to the key at a node of the tree, go left. Otherwise, go right. Figure 2-4 shows an example of the packed memory structure. Once the section is found, perform a binary search within the section.

To search for a range of elements  $[a, b)$ , search for the element  $a$  in the data structure, and scan the array until an element greater than or equal to  $b$  is found. Because the data access is sequential once  $a$  is found, range queries should be very fast, operating at near  $1/c$  times the disk drive's maximum achievable bandwidth.

**Insertion/Deletion** Upon insertion or deletion, use the algorithm for inserting or deleting into a packed memory structure presented above. Because the binary tree is cache-oblivious, the section  $j$  is found using few memory transfers.





# Chapter 3

## Results

Chapter 2 introduced the COB-Tree and proof that the COB-Tree theoretically performs well without the same drawbacks as B-trees. In this chapter, we experimentally analyze the performance of the COB-Tree. Chapters 4 and 5 provide implementation details of the COB-Tree.

The experiments have two objectives: compare the performance speed of COB-Trees with B-trees on experiments involving external memory and to analyze the asymptotic behavior of COB-Trees. All experiments conducted have one of two insertion patterns:

1. Random Insertion: Each key is selected uniformly at random from the sample space of keys.
2. Insertion-at-Head: Each key inserted is less than all previously inserted keys. This insertion pattern forces each key to be inserted at the beginning of the array.

All keys and values are 64-bit integers. The range of keys and values are  $[0, 2^{64})$ . To simulate larger size keys, each key is padded with some number of bytes. All experiments start with 64-element arrays. Arrays are resized when needed.

The rest of the chapter is organized as follows. Section 3.1 provides results comparing COB-Trees and B-trees on experiments involving disk access. Section 3.2 shows graphs demonstrating the COB-Tree's asymptotic performance for large tests.

### 3.1 Comparing COB-Trees and B-Trees

Experiments were run on a Pentium II 400MHz Processor with 128MB of main memory. We inserted elements into the cache-oblivious search tree, a memory mapped implementation of B-trees, and the Berkeley DB [23]. Keys had 512 byte pads to simulate a 520 byte key. We used this small, old, machine so that we could experiments both on trees that fit in main memory and on trees that do not. In section 3.2 we use a faster machine to explore the asymptotic behavior of our data structure. We ran experiments inserting 40,000 elements and 160,000 elements using both insertion patterns presented above. Inserting 40,000 elements takes roughly 20-40 MB of space. Therefore, the insertion may be done entirely in memory. Inserting 160,000 elements takes 80-160 MB of space, therefore forcing data to be written to disk. Table 3.1 shows insertion times for 40,000 elements. Table 3.2 shows insertion times for 160,000 elements.

	COB-Tree	Memory Mapped B-Tree	Berkeley DB
Random Insertion	2.4s	5.03s	93.67s
Insertion-At-Head	5.2s	3.73s	10.85s

**Table 3.1:** Time needed to insert 40,000 520 byte keys into an empty database, in seconds.

	COB-Tree	Memory Mapped B-Tree	Berkeley DB
Random Insertion	202s	639s	1485s
Insertion-At-Head	140s	43.6s	42.91s

**Table 3.2:** Time needed to insert 160,000 520 byte keys into an empty database, in seconds.

Note that for a random insertion pattern, the COB-Tree outperforms both implementations of B-trees. This result is attributed to the fact that in the random insertion case, on average, many insertions occur between periods of heavy rebalancing. Thus, rebalancing occurs quite infrequently.

For the insertion-at-head pattern, the B-tree outperforms the COB-Tree by only a factor of roughly 3.5. We expected much worse. For the B-tree, this insertion pattern allows

for all relevant data elements to be stored in cache at all times. The COB-Tree requires inserting into the beginning of a sorted array every time. Given how often the COB-Tree must rebalance, performing only a factor of 3.5 worse than B-trees is quite remarkable.

## 3.2 Asymptotic Behavior of COB-Tree

We present graphs of the performance of the COB-Tree to show evidence of asymptotic runtime behavior. All experiments were run on a 1.4MHz AMD Opteron(tm) Processor with 16GB of main memory. Keys and values were each 8 bytes.

### 3.2.1 Random Insertion Pattern

We first focus on runtime. Figure 3-1 shows the average time for random insertion. The domains of the graph demonstrating large increases in the average runtime represent times large portions of the array are rebalanced frequently. This graph demonstrates that for random insertion patterns, periods of rebalancing are infrequent, yet costly. Figure 3-2 shows the average runtime for random insertion normalized by dividing by  $\lg^2 N$ . Once again, the sharp increases are attributed to heavy rebalancing, while the dips in the graph are attributed to long areas of time of infrequent rebalancing. The graph seems to be slightly increasing, not quite demonstrating  $\Theta(\log^2 N)$  amortized insertion cost the RAM model suggests.

We focus on number of moves. Define a *data move* to be moving of an element from one location to another within the array. Note this does not count the actual insertion of an element. Thus, it is possible to insert an element with no data moves. Figure 3-5 shows the average number of data moves per insertion is roughly constant.

We now focus on analyzing how often we rebalance and how large the rebalances are. We calculate the sum of all the portion sizes we rebalance. Call this sum the *rebalancing sum*. Figure 3-8 shows for a random insertion pattern, the average rebalancing sum to be constant. This implies that for case of random insertion, the cost of rebalancing, amortized over all of the insertions, is roughly constant.

### 3.2.2 Insertion-at-Head Pattern

We focus on runtime. Figure 3-3 shows the average time for inserting elements using the insertion-at-head pattern. We don't see the same dips and sharp increases as Figure 3-1 because rebalances occur much more frequently. Figure 3-4 shows the average runtime for the insertion-at-head pattern normalized by dividing by  $\lg^2 n$ .

We analyze the number of moves and average rebalancing sum together. Figure 3-6 shows the average number of data moves per insertion for the insertion-at-head pattern. For this pattern, Figure 3-7 shows the average number of data moves per insertion is roughly  $\Theta(\log^2 N)$ . For the insertion-at-head pattern, Figure 3-9 shows the average cost of rebalancing per element. Figure 3-10 shows the average cost of rebalancing is roughly  $\Theta(\log^2 N)$ . For all of these graphs, note a sharp change of behavior around  $n = 10^8$ . At  $n = 10^8$ , the section size doubles from 512 to 1024. The next time the section size is to double is roughly  $n = 6 \cdot 10^{11}$ . Given such large intervals between section size doubling, perhaps a smarter method of resizing the array is needed to achieve more consistent asymptotic behavior.

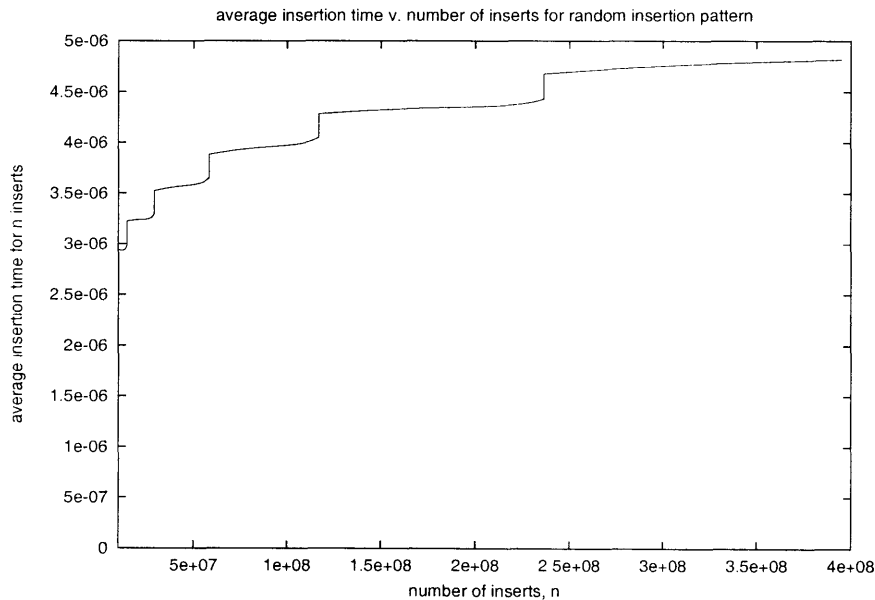
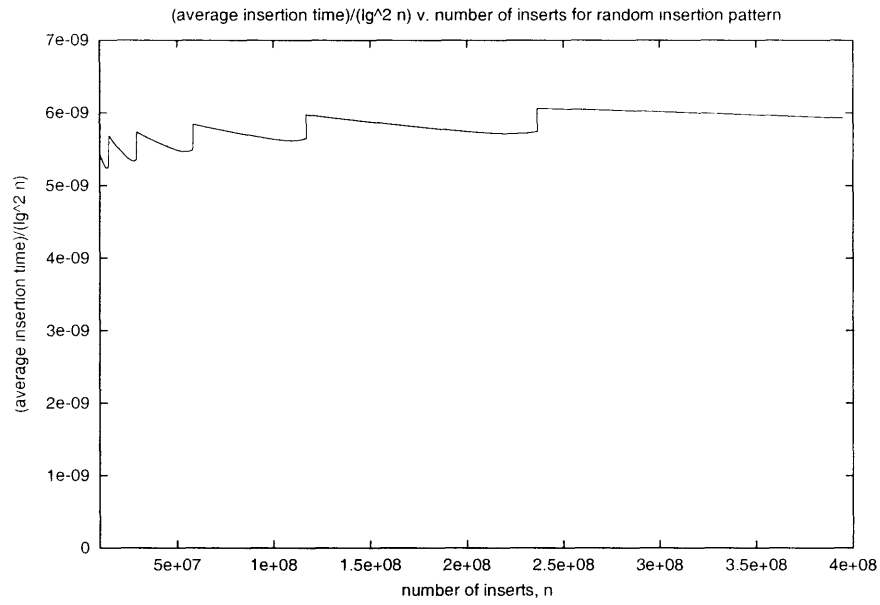
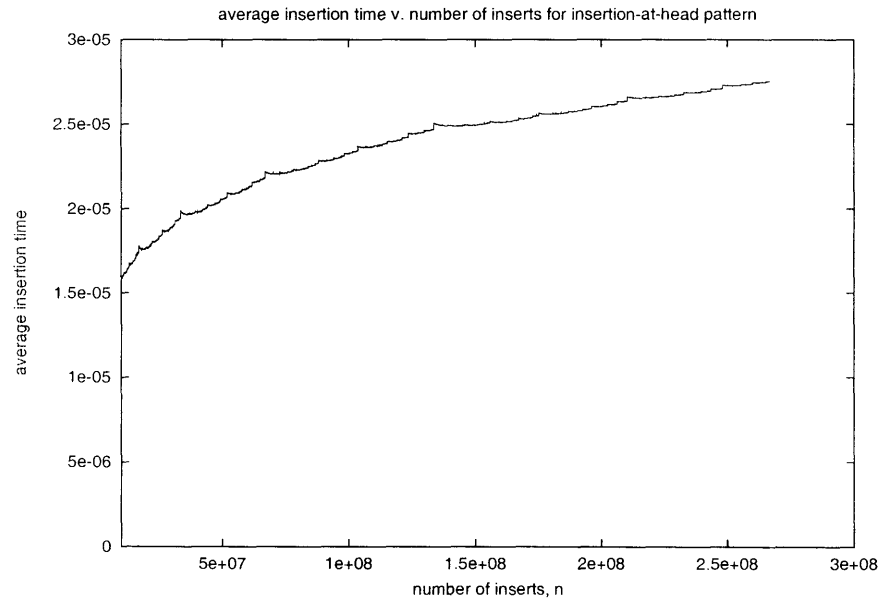


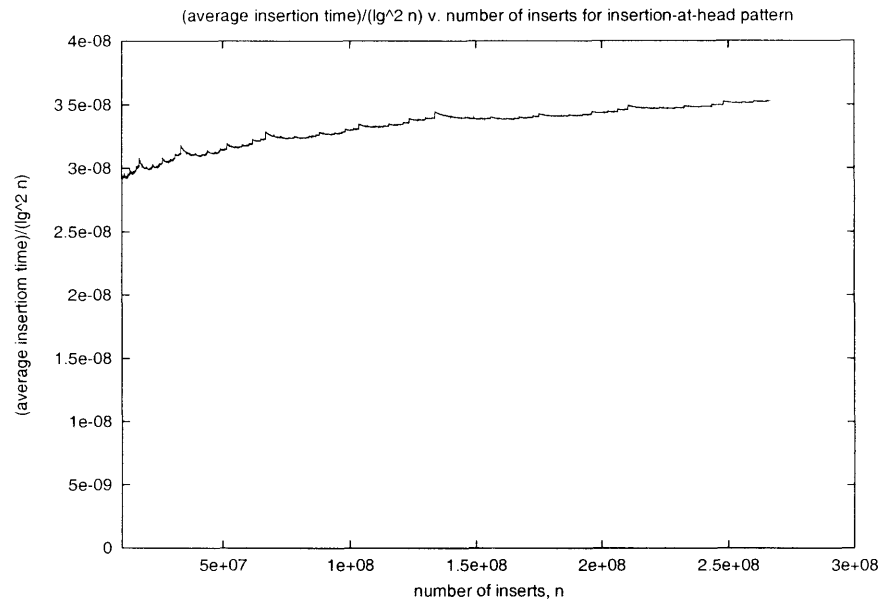
Figure 3-1: Average time for insertion with random insertion pattern.



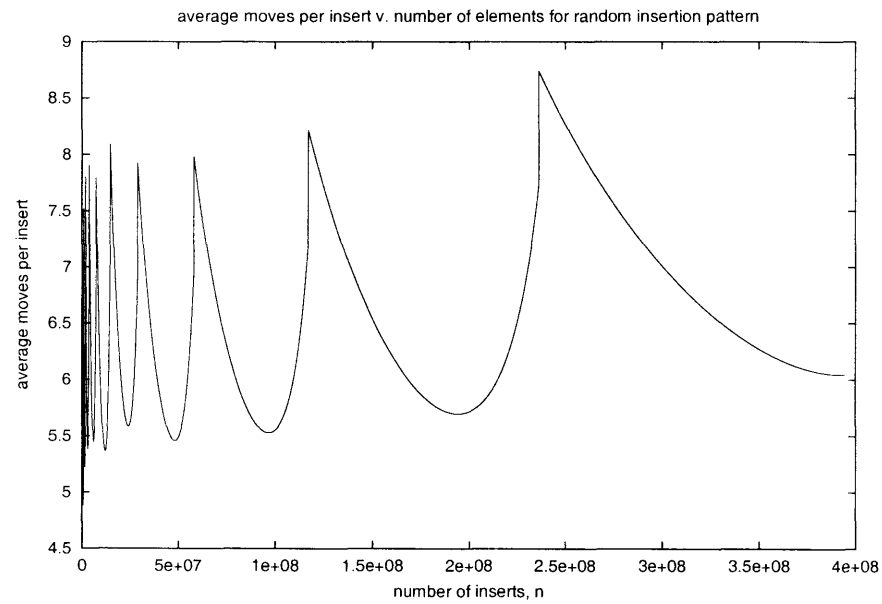
**Figure 3-2:** (Average time for insertion)/( $\lg^2(\text{number of elements})$ ) with random insertion pattern.



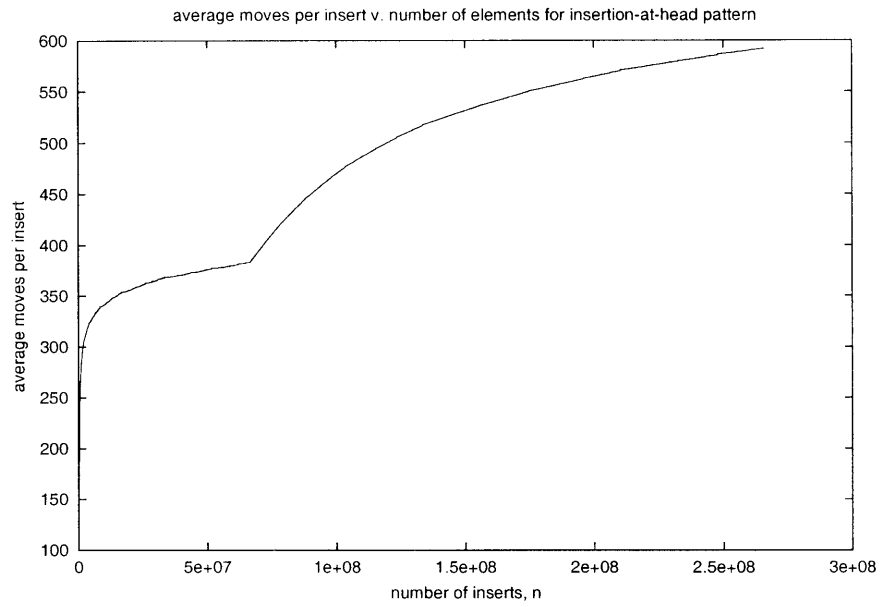
**Figure 3-3:** Average time for insertion with insertion-at-head pattern.



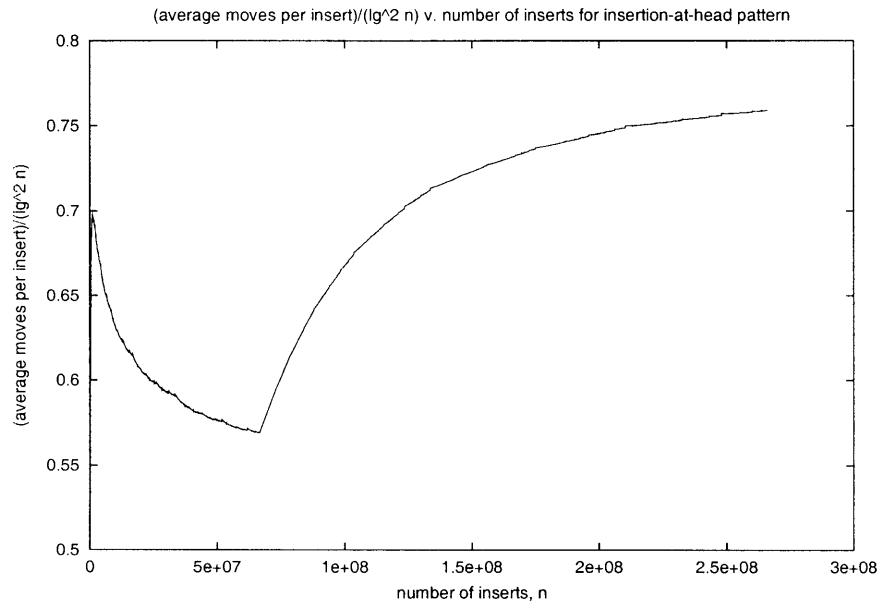
**Figure 3-4:** (Average time for insertion)/( $\lg^2(\text{number of elements})$ ) with insertion-at-head pattern.



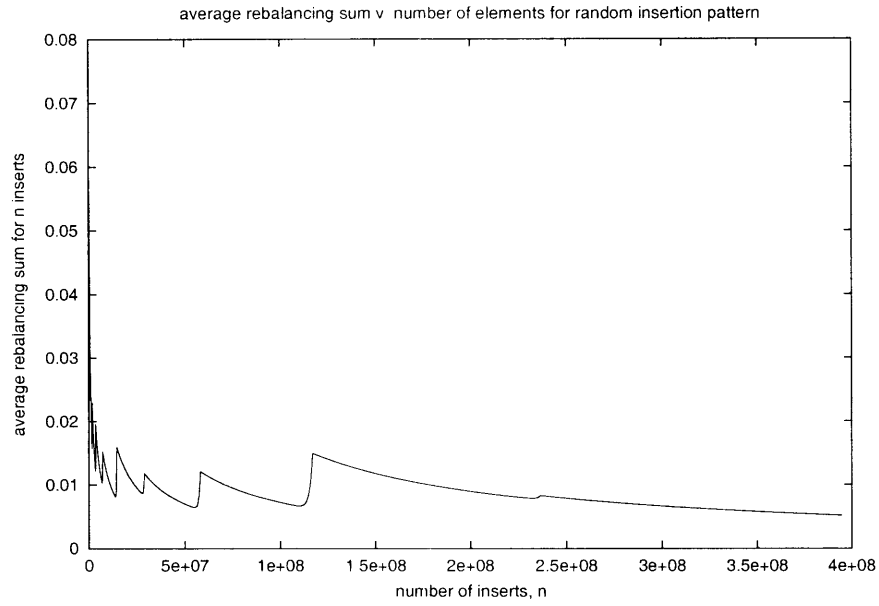
**Figure 3-5:** Average number of data moves with random insertion pattern.



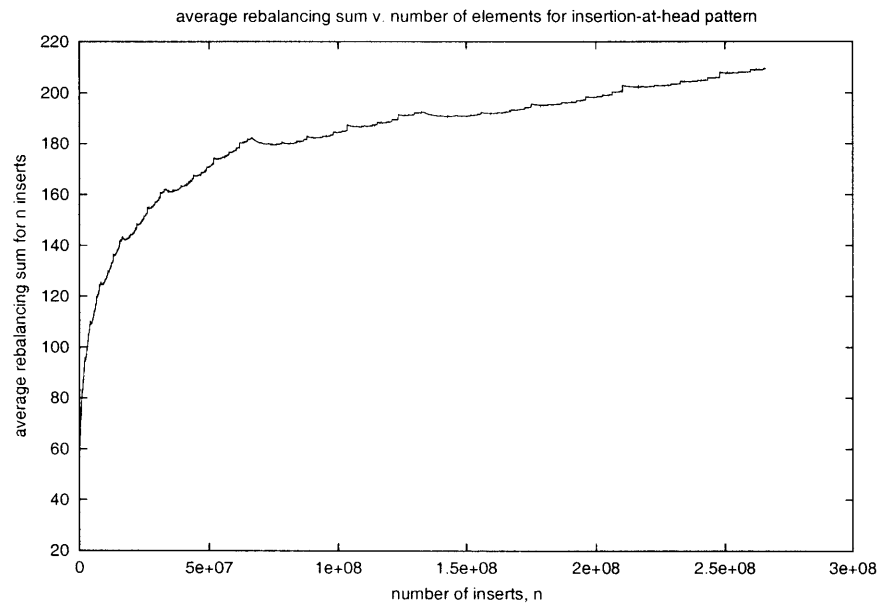
**Figure 3-6:** Average number of data moves with insertion-at-head pattern.



**Figure 3-7:** (Average number of data moves)/(lg<sup>2</sup>(number of elements)) with insertion-at-head pattern.

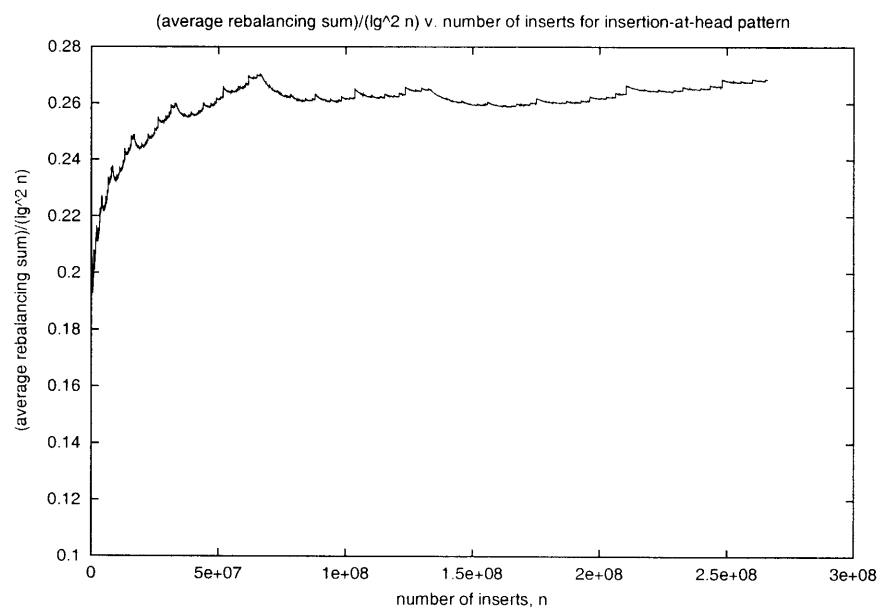


**Figure 3-8:** Average rebalancing sum with random insertion pattern.



**Figure 3-9:** Average rebalancing sum with insertion-at-head pattern.





**Figure 3-10:**  $(\text{Average rebalancing sum})/(\lg^2(\text{number of elements}))$  with insertion-at-head pattern.



## Chapter 4

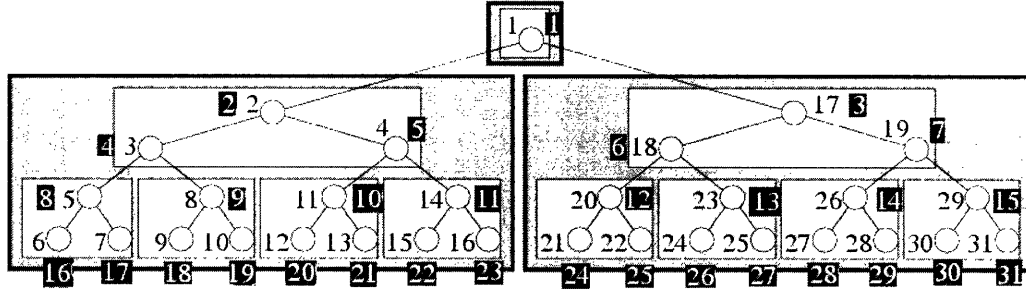
# Static Cache-Oblivious Binary Tree Implementation

Chapter 2 described the two data structures that form the COB-Tree, a static cache-oblivious binary tree with a van Emde Boas layout, and a packed memory structure. Chapter 3 provided experimental results. This chapter presents implementation details of a tree with a van Emde Boas layout. Chapter 5 presents implementation details of the packed memory structure.

The tree is represented in memory as an array. The value at location  $i$  of the array corresponds to some node of the tree. We need a way of computing the location of the left and right children of node  $i$ . One solution is to have the array store pointers, but pointers cost space. Instead, we wish to have an array such that the root of the tree is the first element of the array, and for a given node located at array location  $i$ , the locations of the node's two children are easily found. This chapter provides details.

Consider the breadth-first layout, a simple tree layout. In a breadth-first layout, a binary tree of  $N$  nodes is represented as an array. Each element of the array corresponds to a node. The values held in the array are values of nodes. The root node is located at the first position of the array. The locations of the children of node  $i$  are  $2i$  and  $2i + 1$ . Thus, the location of children may be implicitly calculated. Implicit calculations of children makes the

breadth-first layout simple to use and conserves space by not allocating pointers for children of nodes. Figure 4-1 shows an example of a binary tree with breadth-first indices, along with van Emde Boas indices.



**Figure 4-1:** van Emde Boas and Breadth First indices on tree of height 5. van Emde Boas indices are in black letters. Breadth first indices are in white letters. Figure taken from [7].

We create the same advantages for the van Emde Boas layout by creating an abstraction for the van Emde Boas layout using the breadth-first layout. We maintain an array of values such that the order of nodes is the same as the van Emde Boas layout. However, we provide a function so that the programmer deals with the array as though it has a breadth-first layout. The function converts breadth-first layout indices to van Emde Boas layout indices. My function, `normToBoas`, takes  $O(\log \log N)$  word operations. For example, given the tree in Figure 4-1, `normToBoas(12)` returns 20. Thus, a tree with a van Emde Boas layout will be an array in memory.

Having a function calculate indices from the breadth first layout to the van Emde Boas layout has several advantages. The function makes the creation of a tree with a van Emde Boas layout very simple. To create a tree with  $N$  nodes, one allocates an array of size  $N$ . The programmer may program with the ease of using a tree with a breadth-first layout, but gains the locality advantages of using a tree with a van Emde Boas layout. The actual layout of the tree is abstracted away from the user. As far as the programmer is concerned, the system provides a breadth-first tree that runs very efficiently.

To illustrate the ease and efficiency of such a function, consider C code below to perform

a binary search on a tree with a breadth-first layout. The variables, `depth` and `height`, are the depth of the current node and height of the tree, respectively. If the value is present, the program returns 1, otherwise, 0. To run the program on a tree of height 5, execute `search(tree,1,1,5,value)`.

```
int search(int* tree, int node, int depth, int height, int value){
    if (depth > height) return 0;
    else if (tree[node]==value)return 1;
    else if (tree[node]<value)
        return search(tree, 2*node, depth+1,height,value);
    else if (tree[node]>value)
        return search(tree, 2*node+1, depth+1,height,value);
}
```

Given an efficient `normToBoas` function, here is the code to do a search on a tree with a van Emde Boas layout.

```
int search(int* tree, int node, int depth, int height, int value){
    int boasNode;
    boasNode=normToBoas(node);
    if (depth > height) return 0;
    else if (tree[boasNode]==value)return 1;
    else if (tree[boasNode]<value)
        return search(tree, 2*node, depth+1,height,value);
    else if (tree[boasNode]>value)
        return search(tree, 2*node+1, depth+1,height,value);
}
```

Note that only two lines of code are added.

The rest of the chapter is organized as follows. Section 4.1 presents the algorithm to convert a node's breadth-first index to the corresponding van Emde Boas index, without

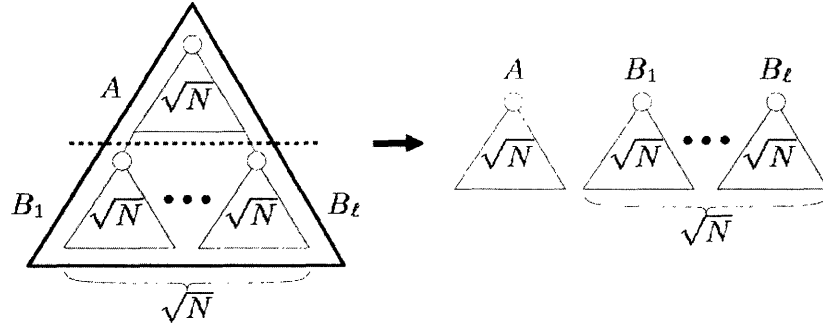
specifying how to evaluate certain intermediate variables. Sections 4.2-4.4 present the algorithms to evaluate these variables.

## 4.1 Algorithm to Convert Breadth-First Index to van Emde Boas Index

The recursive algorithm to convert a node with breadth-first index  $n$  to the corresponding van Emde Boas index is as follows. Given depth  $d$ , and height  $h$  of node  $n$ , we solve for  $\text{normToBoas}(n, d, h)$ . Let the breadth-first layout and the van Emde Boas layout be indexed at 1. For the base case, if  $h \leq 3$ , the van Emde Boas index is identical to the breadth-first index. Otherwise, let  $A, B_1, B_2, \dots, B_l$  be the subtrees recursively laid out in order, as shown in Figure 4-2. Let  $h_1$  be the height of  $A$ , and  $h_2$  be the height of  $B_1, \dots, B_l$ . If  $n \in A$  (which only happens if  $d \leq h_1$ ), recursively call  $\text{normToBoas}(n, d, h_1)$ . Otherwise, let  $n \in B_i$  for some  $i$ . Let  $n'$  be the breadth-first index of  $n$  in  $B_i$ . Let  $d'$  be the depth of  $n$  in  $B_i$ . Let  $x$  be the total number of nodes in  $A, B_1, \dots, B_{i-1}$ . The final result is  $x + \text{normToBoas}(n', d', h_2)$ . Given we can solve for the variables  $h_1, h_2, n', d', x$  in constant word computations, this algorithm does  $O(\log \log N)$  word computations, where  $N$  is the number of nodes in the tree. In the following subsections we present the methods to solve for  $h_1, h_2, n', d'$ , and,  $x$  using a constant number of word computations.

## 4.2 Evaluating Heights of Subtrees

We present the algorithm to solve for the height of subtree  $A$ ,  $h_1$ , and the height of remaining subtrees,  $h_2$ . We know  $h_1 + h_2 = h$  and  $h$ , so we focus our efforts on only evaluating  $h_2$ . Let the *hyperceil* of  $n$  be the smallest power of 2 greater than or equal to  $n$ . Let the *hyperfloor* of  $n$  be the largest power of 2 less than or equal to  $n$ . Note that if  $n$  is a power of 2, then the hyperfloor and hyperceil are equivalent. Otherwise, the hyperceil is twice the hyperfloor.  $h_2$  is the hyperceil of  $h/2$ . Therefore, the problem of solving  $h_1$  and  $h_2$  reduces to solving



**Figure 4-2:** General method of van Emde Boas layout. In memory, the subtrees  $A, B_1, B_2, \dots, B_t$  are recursively laid out in order. Figure taken from [7].

the hyperfloor of  $h/2$ .

The implementation for solving the hyperfloor of a number  $n$  is as follows. Let  $n$  be an unsigned integer. Consider the bit representation of  $n$  and of  $\text{hyperfloor}(n)$ . The hyperfloor has only one 1, the highest positioned 1 in the bit representation of  $n$ . For example, let  $n$  be 25. The bit representation of  $n$  would be  $00\dots011001$ . The hyperfloor, 16, would be  $00\dots010000$ . We have two variables,  $\text{val}=1$  and  $\text{temp}=n$ . We alternately right-shift  $\text{temp}$  one space (effectively dividing  $\text{temp}$  by 2) and left-shift  $\text{val}$  one space (multiplying  $\text{val}$  by 2) until  $\text{temp}=1$ . Upon termination,  $\text{val}$  holds the hyperfloor of  $n$ . An example of this execution for  $n = 25$  is in Table 4.1. The code for `hyperceil` is presented below. The variable

	initial values				final values
<code>temp</code>	$00\dots011001$	$00\dots001100$	$00\dots000110$	$00\dots000011$	$00\dots000001$
<code>val</code>	$00\dots000001$	$00\dots000010$	$00\dots000100$	$00\dots001000$	$00\dots010000$

**Table 4.1:** Values of `temp` and `val` in executing `hyperfloor(25)`.

`val` stores the hyperfloor. The last line returns the `hyperceil`, based on the value of `val`.

```
inline unsigned int hyperceil_int(unsigned int num) {
    unsigned int val = 1;
    unsigned int temp = num;
    while(temp != 1){
```

```

    temp = temp>>1;
    val = val<<1;
}
return (val == num) ? val : (val<<1);
}

```

### 4.3 Evaluating New Depth and Number of Preceding Nodes

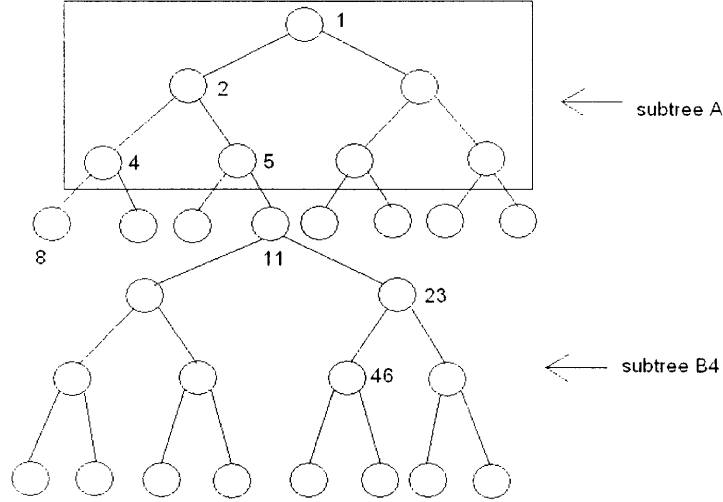
We show how to evaluate the new depth,  $d'$  of the recursive case and the number of nodes in preceding subtrees,  $x$ . Given  $n \in B_i$  for some  $i$ , and we have evaluated  $h_1$ , we know  $d' = d - h_1$ . Otherwise,  $d' = d$ . We now focus on solving  $x$ . Let  $a$  be the number of nodes in subtree  $A$  and  $b$  be the number of nodes any subtree  $B_1, \dots, B_l$ . We know  $x = a + (i - 1)b$ .

We focus on solving  $a$  and  $b$ . Let  $y$  be the breadth-first index of the root node of  $B_1$ . We know  $y = 2^{h_1}$  because of the nature of the breadth-first layout. We also know  $y$  is the first node to be placed in memory immediately after all nodes in subtree  $A$ . Thus,  $a = y - 1$ . Similarly, we know  $b = 2^{h_2} - 1$ . To solve for  $i$ , we take advantage of the fact we know the breadth first indices of the roots of  $B_1, \dots, B_l$ . We know the root of  $B_1$  is  $2^{h_1}$ , the root of  $B_2$  is  $2^{h_1} + 1$ , and so on. The root of  $B_i$  is  $2^{h_1} + i - 1$ . Thus, solving the breadth first index of the root of  $B_i$  solves the value of  $i$ . We know the difference in depth between node  $n$  and the root of  $B_i$  is  $d - h_1$ . We know the parent of an arbitrary node,  $j$ , is  $\lfloor j/2 \rfloor$ . We can evaluate the parent by right-shifting  $j$  by one bit. Thus, to evaluate the root of  $B_i$ , which is a distance  $d - (h_1 + 1)$  from  $n$ , we right-shift  $n$  by  $d - h_1 - 1 = d' - 1$  bits. This solves the root index of  $B_i$ . From this, we can evaluate  $i$ .

To illustrate, take a tree of height 7. Therefore,  $h_1 = 3$  and  $h_2 = 4$ . The recursive subtrees are  $A, B_1, B_2, \dots, B_8$ . Given  $h_1 = 3$ , we know the root of  $B_1$  is node 8 of the breadth-first layout. Suppose  $n = 46$  and  $d = 6$ . We know  $n$  is located in some subtree  $B_i$ . We see  $d' - 1 = d - h_1 - 1 = 2$ ,  $a = 2^3 = 8$ , and  $b = 2^4 = 16$ . Right-shifting  $46 = 000\dots0101110$



two bits gives us the root of  $B_i$  to be  $000 \dots 0001011 = 11$ . Therefore,  $x = 7 + (11 - 8) * 15$ . Figure 4-3 illustrates the tree.



**Figure 4-3:** A partial diagram of a tree with height 7 in breadth-first order. The seven nodes at the top three depths make up subtree  $A$  of the van Emde Boas layout. The subtree with root node 11 forms subtree  $B_4$  of the van Emde Boas layout.

## 4.4 Evaluating Breadth-First Index of Recursive Case

Given  $n \in B_i$  for some  $i$ , the breadth-first index of node  $n$  in subtree  $B_i$  is  $n'$ . We show how to evaluate  $n'$ . We know right-shifting  $n$  by  $d' - 1$  bits give us the root of  $B_i$ . We want right-shifting  $n'$  by  $d' - 1$  bits to give us 1. Thus, the last  $d' - 1$  bits of  $n$  and  $n'$  are identical. We wish to change the remaining bits to the integer value 1. For example, take Figure 4-3. Suppose  $h = 7$ ,  $n = 46$  and  $d' = 3$ . We wish to set  $n'$  to 6. Let integers be represented by  $k$  bits. We know the bit representation of 46 is  $000 \dots 0101110$ . We know the bit representation of 6 is  $0 \dots 0000110$ . We wish to keep the last two bits, 10, unchanged, but change the first  $k - 2$  bits from  $000 \dots 01011$  to  $000 \dots 00001$ .

To change the first  $k - 2$  bits to 0000...0001 may be done easily. In the bit representation of  $n$ , we know the  $d$ th bit is a 1 and all higher order bits are 0. Thus, to evaluate  $n'$ , we need to right-shift bits up to position  $d$  to position  $d'$ . For example, for  $n = 46$  and  $d = 6$ , we need to take all bits from 6 and above and shift them to location  $d' = 3$ . This would, in effect, eliminate bits in  $[3, 6)$ . A simple function, shown below, achieves this goal. This evaluates  $n'$ .

```
inline size_t fixBits(size_t x, unsigned int i, unsigned int j) {
    unsigned long y = (x>>j)<<i;
    unsigned long z = x&((1<<i)-1);
    return y+z;
}
```

# Chapter 5

## Packed Memory Structure Implementation

Chapters 2 and 3 presented COB-Trees and their performance results. Chapter 4 presented implementation details of one component of the COB-Tree, the static cache-oblivious binary tree. This chapter presents implementation details of the packed memory structure.

The chapter is organized as follows. Section 5.1 lists several decisions important to making the implementation efficient. Section 5.2 presents detailed descriptions of the implementation of each phase of the insertion algorithm, with the exception of rebalancing. Section 5.3 introduces several possible rebalancing algorithms.

### 5.1 Data Representation

We present our data representation decisions for the packed memory structure. Below is a list of decisions made to make the packed memory structure efficient.

1. The packed memory structure and binary tree are separately stored as arrays. We employ memory mapping to store the two arrays consecutively in memory. This allows the data to be used as a single level store.

2. The size of the array  $T$ , is a power of 2. The array is divided into sections of size  $S$  such that  $S$  is the power of 2 arithmetically closest to  $\lg^2 T$ . Note that  $(\sqrt{2}/2) \lg^2 T \leq S \leq \sqrt{2} \lg^2 T$ , thus  $S = \Theta(\lg^2 T)$ . As a result, the number of sections,  $N$ , is a power of 2 as well.  $S$ ,  $T$ , and  $h$  are all global variables.
3. The relationship between a leaf of the binary tree and its corresponding section is implicit. Given the breadth-first index of the leaf, we simply need to subtract  $2^{h-1}$  from the index to evaluate the corresponding section.
4. The binary tree of height  $h$  is stored as an array. Note that no information needs to be held at the leaf of a tree. We do not need to access a key because no searching remains. We do not need to store corresponding sections because they may be evaluated implicitly. Thus, there is no reason to allocate space for the leaves of the array. Therefore, although we implicitly use a tree of height  $h$  with a breadth-first layout, we represent the tree in memory as a tree of height  $h - 1$  with a van Emde Boas layout.

## 5.2 Insertion

We present implementation details for each part of the insertion algorithm. Suppose we wish to insert  $i$  and have found the appropriate section  $j$ . The insertion algorithm has several parts. First, the precise location within  $j$  must be found. Second, threshold must be tested. Then, there are three possible cases. The first case is that the section,  $j$ , is within threshold, and the element is inserted. The second case is  $j$  is not within threshold, but some portion containing  $j$  is. The third case is no ancestor is within threshold and the array needs to be resized. We present issues and solutions for each phase and case.

**Finding the Proper Location of  $i$  within the Section.** The goal is to find the proper location for  $i$  as quickly as possible. A poor, yet simple, solution is to perform a linear scan of the section. This strategy works well with small databases, but not with large ones. Although small in comparison to the entire packed memory structure, for large databases, sections

become large enough so that performing a linear scan noticeably slows down performance. Instead, perform a binary search within the section to find the proper location of  $i$ . Although tricky to program because the section has empty spaces, a binary search within section  $j$  is efficient.

**Testing a Section to be within Threshold.** When inserting, we do not explicitly test if a section is within threshold, because testing threshold requires scanning the entire section. Just as scanning to find the appropriate location of an element  $i$  is too costly, so is scanning to find the density. Instead, an attempt is made to insert  $i$ . If an empty space is found, the element is inserted. Otherwise, the density is 1 and the section is not within threshold. We proceed to find a portion to rebalance. Thus, we require the upper bound threshold for a section to be 1.

**Insertion within a Section.** Given the section,  $j$ , is within threshold, we insert  $i$  as efficiently as possible. We shift as few elements as possible to insert  $i$ . Via a binary search, we find between what two values the element  $i$  should be inserted. If empty spaces between the values exist,  $i$  is inserted somewhere between the two values. Otherwise, we scan left until we find an empty location, shift all elements up to that location left one space, and insert  $i$ . If no empty space exists, we attempt the same thing in the right direction. One of these procedures must work because the section is within threshold.

When inserting into a section, we never need to update the binary tree. When traversing the tree, we traverse down a left branch if and only if the key we wish to insert is strictly less than the key located at the node. Therefore, we never insert a key into a section that is larger than all existing keys in the section. Thus, for this case, the tree never needs to be updated. If we delete the largest element in a section, we choose not to update the tree. Correctness is not affected and we believe the effect on performance is negligible.

**Inserting an Element when Rebalancing is Required.** We wish to insert the key and rebalance the elements within some portion of the array. Rebalancing and resizing the array

has several issues, along with several possible solutions, all of which are presented in the next section. We focus on updating the search tree. Suppose we have found and rebalanced the proper portion that will contain  $i$ . The subtree rooted at the node in the tree representing the rebalanced portion is no longer valid. We must update the subtree representing this portion. This can be done recursively. If the node is a leaf, return the maximum value of the represented section. Every internal node solves for two values from its children, the largest key in each child's portion. Thus, the node returns the value received by the right child, the largest key in the node's own portion, and stores the value returned by the left child, the largest key in the left child's portion. The code is presented below.

```
long long int updateTree(node* tree, size_t index, pair* array){
    long long int left, right;
    if(isLeaf(index)) {
        return //maximum value of section represented;
    }
    else {
        left = updateTree(tree, 2*index, array);
        right = updateTree(tree, 2*index+1, array);
        tree[boasloc].val = left;
        return right;
    }
}
```

### 5.3 Rebalancing

The problem is as follows. Suppose we have an element,  $i$ , we wish to insert. Suppose the section where  $i$  is to be inserted, section  $j$ , is completely full. The problem is finding the smallest portion containing  $j$  that is within threshold and rebalancing that portion as efficiently as possible. We do not address rebalancing and inserting the element concurrently.

The portion is first rebalanced, and then the element is inserted into the portion. Resizing the array is trivial within any of the algorithms.

We present several possible solutions and analyze their respective costs in terms of data reads and data writes. Of the algorithms presented, only the read-efficient algorithm presented in Section 5.2.2 is implemented. For all of these algorithms, suppose  $n$  is the number of elements contained in the portion that is rebalanced.

### 5.3.1 Simple Rebalancing Algorithm

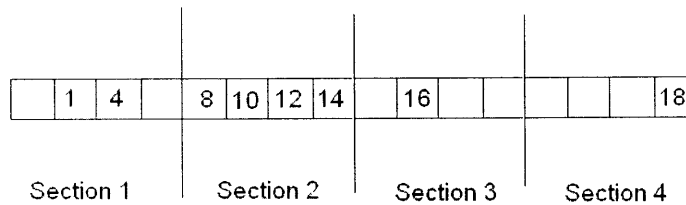
We first propose a simple algorithm. Given sections of size  $s$ , we first scan adjacent portions of size  $s, 2s, 4s, \dots, 2^{k-1}s$ , testing densities, until we find a portion of size  $2^k s$  within threshold. Given the portion, we move every element as far left as possible, maintaining order. So, if  $n$  elements exist in the portion, after the move, the  $n$  elements take up the first  $n$  spaces in the portion. We call this a *crunch* to the left. Lastly, starting from the rightmost element, move every element to its proper location.

We provide an example. Suppose we wish to insert the key 11 into the array in Figure 5-1. The array has four sections. Each section has a capacity of four elements. We see the second section is completely full, so the first section of size  $s = 4$  is scanned. Suppose the combined density of sections 1 and 2 is still too high. Sections 3 and 4 of size  $2s = 8$  are scanned. The combined density of the four sections is within threshold. Thus, we crunch to the left. Figure 5-2 shows the resulting state of the array after the crunch. After redistribution, the key 11 may be inserted into the array. Figure 5-3 shows the resulting state after the redistribution.

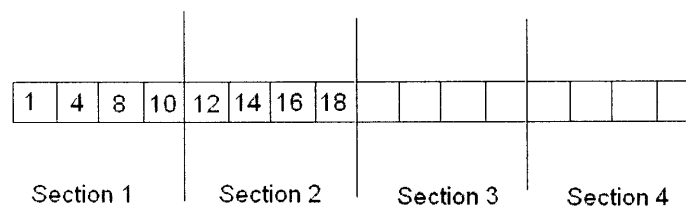
This algorithm performs three passes over the data and  $2n$  data writes. In the following subsections, we propose improvements on this algorithm.

### 5.3.2 Read-Efficient Algorithm

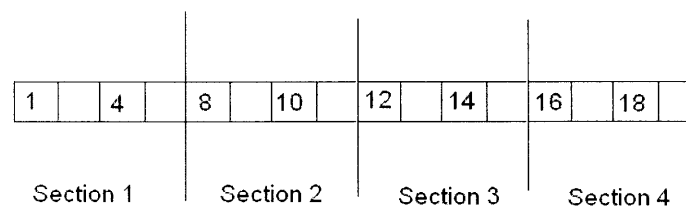
We present an algorithm that performs fewer passes over the data, therefore performing fewer data reads. The idea is to combine two steps of the previous algorithm into one. In



**Figure 5-1:** The initial state of the array before rebalancing begins. The array has four sections, each with a capacity of four elements.



**Figure 5-2:** The state of the array after all elements have been crunched to the left.



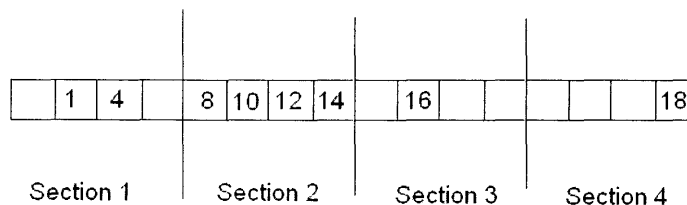
**Figure 5-3:** The final state of the array after elements have been redistributed from the crunched array.



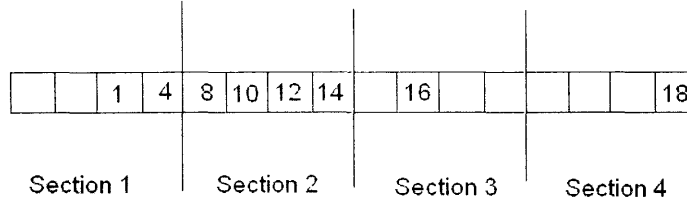
the previous algorithm, portions of size  $s, 2s, \dots, 2^{k-1}s$ , are scanned to test densities. Once the appropriate portion of size  $2^k s$  is found, that portion is crunched. Doing these separately is inefficient because any area scanned will eventually be crunched as well. Also, if we crunch an area, in the process, we learn how many elements exist. Thus, when crunching, we can learn the density as well.

The algorithm is as follows. The idea is instead of only scanning to find a portion within threshold, crunch the portions as they are being scanned around the section where  $i$  is to be inserted. Do so until the portion to be rebalanced is found. This step takes one pass over the data and performs at most  $n$  data writes. The portion to be rebalanced is already crunched. Next, redistribute the elements across the portion. This step takes one pass over the data and performs at most  $n$  data writes as well. Overall, we save one pass over the data from the previous algorithm.

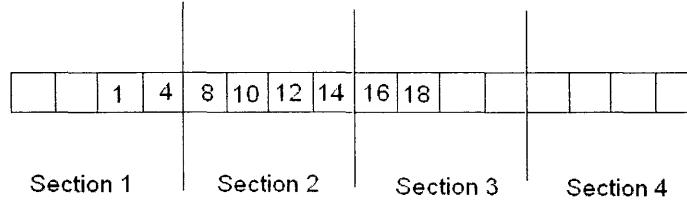
We demonstrate the algorithm using the same example as in the previous section. Once again, we wish to insert the key 11 into the array shown in Figure 5-4. Figure 5-5 shows the state of the array after section 1 has been crunched towards the middle of the array. Figure 5-6 shows the state of the array after sections 3 and 4 have been crunched towards the middle of the array. After crunching is complete, redistribution takes place and 11 is inserted.



**Figure 5-4:** The initial state of the array before rebalancing begins. The array has four sections, each with a capacity of four elements.



**Figure 5-5:** The state of the array after section 1 is crunched towards Section 2. Sections 3 and 4 remain to be crunched.



**Figure 5-6:** The state of the array after Sections 3 and 4 are crunched towards section 2.

### 5.3.3 Write-Efficient Algorithm

We present an algorithm that performs fewer data writes than the simple algorithm. We still scan portions of size  $s, 2s, \dots, 2^{k-1}s$  to find the appropriate portion to be scanned, but we do not redistribute the elements by first crunching then redistributing. Instead, we redistribute in a manner so that at most only  $n$  total data writes take place.

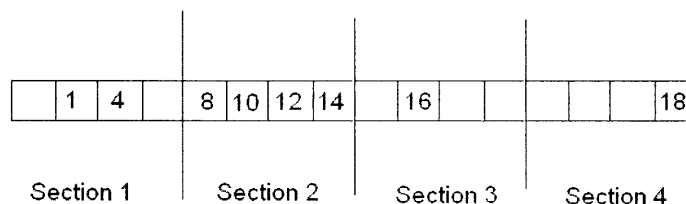
First, we define types of element. Take an arbitrary element of the portion at location  $k$ . Suppose the location of this element in the redistribution will be some location other than  $k$ . If the new location of this element will be greater than  $k$ , call the element a *right moving element*, because the element will be moved to the right of its previous location. Similarly, if the new location will be less than  $k$ , call it a *left moving element*.

We show safe ways of relocating two elements based on which type of elements so that no data may be overwritten. Suppose we have two elements, in locations  $a$  and  $b$ , such that  $a < b$  in the array. Suppose the new locations for  $a$  and  $b$  in the redistribution are  $a'$  and  $b'$ . If they are both right moving elements, move  $b$  to  $b'$  such that  $b' > b$ , then move  $a$  to  $a'$  such that  $b' > a'$ . We move the elements in this order because if  $a' = b$ , then no data

is lost. Similarly, if  $a$  and  $b$  are both left moving elements, first move  $a$  and then  $b$ . If  $a$  is a left moving element and  $b$  is a right moving element, it does not matter which order the elements are moved, because  $a$  and  $b$  are moved farther away from each other. If  $a$  is a right moving element and  $b$  is a left moving element, it does not matter which order the elements are moved, because  $a < a' < b' < b$ .

The redistribution algorithm is as follows. First, starting from the beginning of the array, scan right and move all left moving elements to their new proper locations. Then, starting from the end of the array, scan left and move all right moving elements to their new proper locations. New locations may be evaluated given  $n$ , the capacity, and the order of an element in the portion. The order may be calculated dynamically in the scan. The relative order which any two elements are moved follow the restrictions described in the paragraph above. Because each element of the array is moved at most once, this performs at most only  $n$  data writes. The algorithm still performs three passes over the data.

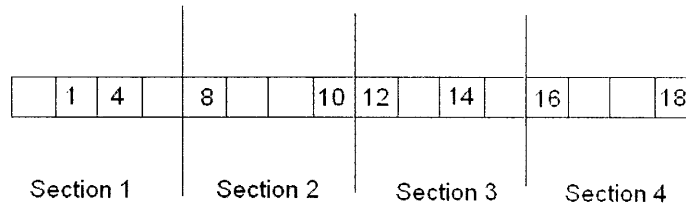
We demonstrate the algorithm using the same example as in the previous sections of inserting the key 11 into the array in Figure 5-7. Figure 5-8 shows the state of the array after relocating right moving elements. Only left moving elements 1, 10, and 18 remain to be moved. Figure 5-9 shows the state of the array after relocating these elements.



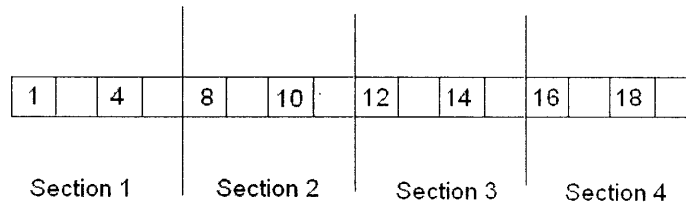
**Figure 5-7:** The initial state of the array before rebalancing begins. The array has four sections, each with a capacity of four elements.

### 5.3.4 Partially Correct Efficient Algorithm

Another possible solution is to rebalance only partially. That is, redistribute most, not all, elements to their proper locations. Given we plan to rebalance a portion of size  $2^k s$ , we



**Figure 5-8:** The state of the array after all right moving elements have been relocated.



**Figure 5-9:** The final state of the array after left moving elements 1, 10, and 18 have been moved as well.

know half of the portion has a much lower density the other half. Otherwise, we would have rebalanced only half of the portion. A good solution may be to only move as many elements as possible in the direction of the low density half. Suppose the low density half is located prior to the high density half. Perhaps only scanning once to move all left moving elements while leaving the right moving elements alone may be good enough. If the low density half is located after the high density half, move all right moving elements. In the example above, 5-8 would be the *final* state of the array.

This algorithm takes two passes over the data and does at most  $n$  data writes. Time restrictions did not allow for implementation tests or theoretical analysis to examine the effect of such a rebalancing algorithm.

# Chapter 6

## Conclusion

I have implemented and tested a cache-oblivious dynamic search tree as an alternative to B-trees. The data is stored in a sorted array on disk, allowing for very fast range queries. The COB-Tree runs 3 times faster than B-trees on random insertion and only 3.5 times slower for repeated insertion into one place, which is the worst-case for the COB-tree.

I have shown how to implement the COB-Tree efficiently. To implement the static cache-oblivious tree layout, the van Emde Boas layout, I describe a function that converts the breadth-first index of a tree node to the corresponding van Emde Boas index, allowing a programmer to use the tree with minimal difficulty. I outline the implementation of the packed memory structure. I provide several possible algorithms for rebalancing a portion of a packed memory structure.

### 6.1 Future Work

We outline several suggestions for future work.

**Deamortize Insertion Cost.** Although the COB-Tree performs the majority of insertions very fast, users may not accept *any* long insertion times. Specifically, spending long periods of time to heavily rebalance the packed memory structure, no matter how infrequently, may

be intolerable to users. Thus, one area of research is to deamortize the cost of insertion. That is, find a way to reduce the variance of the time to insert elements.

**Determine the Best Rebalancing Algorithm.** We have yet to determine which rebalancing algorithm presented in Section 5.3 is best. Sections 5.3.2 and 5.3.3 describe two correct rebalancing algorithms. Only the read-efficient algorithm of Section 5.3.2 has been implemented. Currently, we do not know which of the two algorithms performs faster. In addition, Section 5.3.4 describes a possible rebalancing algorithm that is theoretically faster than all others presented, but only partially correct. Whether this partially correct algorithm performs well in theory or practice is unknown.

**Determining a Smarter Rebalancing Algorithm.** In cases where elements seem to not be inserted in a random order, perhaps an uneven rebalancing algorithm is needed. Suppose a user repeatedly inserts elements into the same section,  $j$ . Perhaps the program should recognize the section  $j$  is the repeated target for insertion. Upon rebalancing, section  $j$  should have more empty spaces than other sections. Such a rebalancing algorithm improves performance when repeatedly inserting elements into one place, a common insertion pattern.

**Variable Size Keys.** The COB-Tree currently supports only fixed-sized keys. An efficient implementation of a COB-Tree that supports variable-sized keys has yet to be created.

**Concurrency** An efficient implementation of a COB-Tree that supports concurrent data query, insertion and deletion has yet to be implemented.

## 6.2 Future Vision

Massive-data is at the center of many important applications. Speeding up operations on that data impacts file systems and databases in all computer systems. COB-Trees go further in helping to eliminate much of the complexity that programmers currently face when trying

to code for memory effects. If further studies show COB-Trees truly outperform B-trees with significantly less complexity, the results will have an effect both on industrial practice and on the way computer science is taught.

## **6.3 Software Download**

To download the software, go to <http://supertech.csail.mit.edu>.





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