# A Forest of Hashed Binary Search Trees with Reduced Internal Path Length and better Compatibility with the Concurrent Environment 

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#### Abstract

We propose to maintain a Binary Search Tree in the form of a forest in such a way that - (a) it provides faster node access and, (b) it becomes more compatible with the concurrent environment. Using a small array, the stated goals were achieved without applying any restructuring algorithm. Empirically, we have shown that the proposed method brings down the total internal pathlength of a Binary Search Tree quite considerably. The experiments were conducted by creating two different data structures using the same input - a conventional binary search tree, and a forest of hashed trees. Our empirical results suggest that the forest so produced has lesser internal path length and height in comparison to the conventional tree. A binary search tree is not a well-suited data structure for concurrent processing. The evidence also shows that maintaining a large tree in form of multiple smaller trees (forest) increases the degree of parallelism.


Keywords: Binary Search Tree Path Length, Parallel
Processing Binary Search Tree, Balanced Tree

## 1. INTRODUCTION

The Binary Search Tree (BST) is a widely used storage medium in the primary memory. A reasonably balanced $B S T$ provides fast data access and retrieval. Despite its wide popularity, it has few serious problems. First, its shape depends on the nature of the input. Second, it is a sensitive data structure for insertion and deletions -the tree shape could be destroyed, following a series of insertions and deletions. Third, it is a highly incompatible data structure for concurrent processing. Over the years these problems have been investigated extensively. For the tree shape, refer to [1]-[4], [13]-[18]; and, for the parallel processing of the tree, refer to [5]-[11]. Eppinger [12] investigated tree's behavior for a large number of insertions and deletions.

A tree could be maintained in better shape either dynamically or statically. In dynamic maintenance, following each operation tree is inspected, and readjusted, if the operation causes any structural damage. AVL Tree [1], Martin \& Ness [2], Red-Black Tree [3], and Splay trees [15] fall into this category. In Splay Trees, the frequently accessed node is pushed towards the root of the tree for future immediate access. Gonnet [12], and Gerasch [17] proposed algorithms to maintain the tree with reduced
internal path length. In static maintenance, periodically, the entire tree is taken as input, and some maintenance work is applied. Examples of static maintenance are: Day's algorithm [4], Chang \& Iyengar's algorithm [14], and, Stout \& Warren's algorithm [16]. Both dynamic and static solutions have their advantages and disadvantages. The dynamic maintenance of the tree is slow because of frequent inspections and rotations. Apart from that, every node in the tree has to store some additional information. Static algorithms, on the other hand, demand lots of extra space. In some cases, the algorithm consumes extra workspace that is twice the size of the input. For example, Chang \& Iyengar's algorithm [16] requires an additional array, equal to the size of input, as extra workspace.

As far as the parallel processing of the tree is concerned, substantial work has been done to develop concurrent algorithms, refer to [5]-[11]. In a binary search tree, the root is the only gateway for all the active processes making it difficult to achieve maximum parallelism. Imagine a situation where we have to update the root of the tree. A process performing this operation has to lock the root, making the entire tree unavailable for rest of the processes. No matter how efficient our concurrent algorithms are, other processes have to wait until the previous releases the lock. Ellis [9], [10] presented solutions for concurrent searches and insertions in the 2-3 and AVL trees. Kung and Lehman [11] investigated ordinary binary search trees, and proposed solutions so that the system could support any number of processes performing searches, insertions, deletions, and rotations. To ensure that the searches are never blocked they used special nodes and pointers.

Most of the presented solutions use some kind of locking scheme to allow multiple processes to act upon a single binary search tree simultaneously. The common goal of all proposed solutions was to increase the degree of concurrency by having a lesser portion of the tree locked, and thus exposing a major portion to the rest of the processes. Such algorithms can increase the degree of concurrency up to a certain extent. However, better results could be obtained, if the underlying data structure is modified to accommodate large number of processes. Substantial work has been done on algorithms, but hardly any attempt has been made to create a flexible data structure.

In this paper, we propose a forest of binary search trees to deal with the stated problems - tree balance, and its incompatibility with the concurrent environment. To examine the overall balance of the proposed forest, we have used internal path
length (IPL), and the height of the tree as measurement parameters. (Concurrency-related issues are discussed in the section 6.)

The height of the tree is the length of the longest path from the root to the leaf. The height is an important parameter to study the worst-case behavior of the tree. For average case analysis, the internal path length of the tree (IPL) can be used. IPL is the average distance of every node in the tree from the root, and is defined as the sum of the depths of all the nodes in the tree. Thus, for a tree with just one node, the $I P L$ is equal to 0 , and for a tree with two nodes, the IPL is equal to 1 . Let $I_{n}$ be the $I P L$ of a tree with ' $n$ ' nodes, and $C_{n}$ be the average number of comparisons required for a successful search, then we have a relation: $I_{n}=n\left(C_{n}-1\right)$. From the given relation, it is clear that the IPL of the tree directly affects its performance. The lesser the height, the lesser will be the IPL, and hence, the faster would be the search. Knuth [19] has given a formula that relates the height of a ' $n$ ' node random binary search tree $H_{n}$, and $C_{n}$ as $C_{n}=2(1$ $+1 / n) H_{n}-3$.

## 2. CREATION OF THE FOREST

It is possible to maintain a random $B S T$ in form of a set of trees called 'forest'. The number of trees in the forest would depend on the application. For the purpose of simulation, we have used a forest of 11 trees. To hold multiple trees, we need multiple roots. We have used an array of pointers of size $k=11$. Each cell of the array acts as a tree root. Like a usual tree, when the tree is empty, each root (array cell) points to a null value (refer to Figure 1). When a key has to be inserted or deleted, it is first divided by ' $k$ ', and the remainder is calculated to find the array location. Assuming the array location to be the root of the tree, the desired operation is performed as usual. In other words, keys need to be hashed by the function: loc=key $\% k$. Where 'loc' is the array location to/from which the key has to be inserted/deleted. Using this technique, we get a set of $k$ possible trees, which resembles a forest, but acts as a single binary search tree. The only difference between a usual single $B S T$ and a forest is determining the tree location using hashing. In Figure 1, a forest of two trees is shown. The collective IPL of the forest = the sum of the IPLs of all the trees in the forest $=12+6=18$. Maximum height of the tree in the forest is


Figure 1: A Forest of Hashed Trees

## 3. METHODOLOGY

Using the same random input, two different structures were created - a conventional tree, and a forest. For every insertion, a random number was generated that was supplied to two separate algorithms: one that creates a conventional tree; and, the other that creates a forest. Duplicate keys were ignored. Once all the keys were inserted, the IPL of both of the structures was calculated. The IPL of the conventional tree was calculated as usual by calculating the sum of the depths of all the nodes in the tree. While, for the forest, the $I P L$ was calculated by adding all the IPLs of the individual trees in the forest, at the same time, the heights of all the trees in the forest were recorded. For example, in Fig.1, the total IPLs of the forest was $12+6=18$, the average height of the forest $=(3+2) / 2=2.5$, the height of the worst tree in the forest $=3$. For the division remainder
method of hashing, it has been shown that a prime number distributes the keys more uniformly - that is why the array size of 11 was our choice. Another reason was to maintain the array size to be roughly equal to $1 \%$ of the input size. We chose the tree size to be 1023 and 2047 so that the forest so produced could also be compared with a perfect, balanced tree, and not just with the random $B S T$. (For quick reference, please note that perfect balanced trees with 1023 and 2047 nodes would have heights 9 and 10, respectively; their IPLs would be 8194 and 18434, respectively.)

## 4. RESULTS

The following are parameters related to the forest that were used to compare the forest to a conventional and perfect balanced tree:
Average height of the forest $=$ Sum of the heights of all 11 trees in the forest $/ 11$.
Height of the worst tree in the forest $=$ A tree with the maximum height in the forest.
Collective path-length of the forest $=$ The sum of the pathlengths of all 11 trees in the forest

Table 1 is obtained as a result of several tests conducted with Borland C++ compiler 5.5 under windows, and GNU C++ compiler 3.4.3 (g++) under Linux. Table 1 (a) shows results for $n=1023$, and $k=11$. The average height of the forest was quite close to 9 - somewhat like a perfect, balanced tree with the same number of nodes. If we compare the average height of the forest to the height of the conventional tree, there is a huge reduction ( $39 \%$ to a whopping $52 \%$ ). The height of the worst tree in the forest was 17 , where the conventional tree went up to 24 . However, the worst tree in the forest would not have same number of nodes as the conventional tree; therefore, comparing their heights is not justifiable, but this will definitely give the worst-case behavior of both of the structures. As far as the $I P L$ is concerned, a reduction in the forest $I P L$ is more than considerable. Among the 10 sample runs, the worst IPL of the forest was recorded as 6623 - that is, far better than 11299 (the IPL of the corresponding BST), and much better than 8194 (the $I P L$ of the perfect, balanced tree with the same number of nodes). The reduction in IPL was recorded to be from $39 \%$ to $52 \%$. Though the $I P L$ is enough for an average case analysis, it would be interesting to apply the formula that relates $\operatorname{IPL}\left(I_{n}\right)$, and the average number of comparisons required for a successful search $C_{n}$. Putting $I_{n}=6623$, and $n=1023$, we get for the forest: $C_{n}=\left(I_{n} / n\right)+1=(6623 / 1023)+1=7.47$. For the corresponding conventional tree: $C_{n}=(11299 / 1023)+1=12$. For a perfect, balanced tree with the same number of nodes: $C_{n}=(8194 / 1023)+1=8$. Hence, in terms the average number of comparisons required for a successful search, it is quite clear that the forest needs a minimal number of comparisons. These results indicate that the behavior of the forest is far better than that of a random $B S T$. If we could ignore the extra cost of time consumed in hashing, on the average, the forest clearly seems to outperform even a perfect, balanced tree.

Obviously, as the size of input increases, we will need a larger array to get similar results. What is the optimal ratio of ' $n$ ' and ' $k$ '? This could be an interesting question. In our case, presented results in table 1(a) are for 1:93 (input size is 93 times of array size) and, 1 (b) is for $1: 186$. Let's consider a simpler question; how large should an array be, if we wanted to reduce the average height of the forest by $50 \%$ ? Using random input: we know that the tree height is $O(l g(n))$. We have $k$ trees, assuming that our hash function distributes keys uniformly among $k$ trees, we would have each tree with a height of approximately $O(\lg (n / k))$, i.e., we have the relationship :$\lg (n) / \lg (n / k)=2$
$\Rightarrow \lg (n)=2 \lg (n / k) \Rightarrow n=(n / k)^{2} \Rightarrow k=\sqrt{n}$

Theoretically, that means for $n=1023$, to reduce the average height of the forest to half, we need an array of size $\sqrt{1023}=32$. However, we have seen that the empirical results do not confirm this, and are more encouraging. With an array of a size of $k=11$, and $n=1023$, several runs have shown that, in most of the cases, the reduction in average height of the forest was between $39 \%$ to $52 \%$. The total IPL of the forest fell to $40 \%$ in comparison to that of the conventional tree. In most of the cases, the total IPL of the forest beat even a corresponding perfect, balanced tree.

In Table 1(b), we have provided the results of when input size was doubled, i.e., now $n=2047$, but array-size is kept unchanged. This would give us some idea of how the performance of a forest is affected, when we increase the size of the input. Comparing the average height of the forest with the height of the conventional tree, it is evident that the reduction in height was still $26 \%$ to $48 \%$. The worst IPL of the forest was 16216 - that is, far better than 24133 (the IPL of the corresponding BST), and still better than 18434 (the IPL of the perfect, balanced tree with 2047 nodes). Indeed, there was some deterioration in the performance, but this confirms that the forest still works better when input size is increased to 186 times of the array size.

## 5. TIME AND SPACE REQUIREMENTS OF THE FOREST

We will compare the time and space requirements of the proposed forest with those of the usual BST. On the average, a random BST with $n$ nodes requires $\lg (n)$ time to perform most of the operations. However, in the case of a forest, hashing is required to jump to the correct tree, i.e. additional $O$ (1) is required for hashing. Following this, the entire process is the same as that of a usual $B S T$ operation. Hence, it is straightforward that, on the average, the forest with $n$ nodes and $k$ trees takes $O(1)+\lg (n / k)$ time. The worst-case behavior of a $B S T$ is $O(n)$ - which remains the same as for the forest. This happens when all the keys are in sorted order, and hashed to one single tree in the forest, letting the tree grow up to $n$. A natural question is: what is the additional space requirement of the forest? When compared to a normal $B S T$, the forest does not require additional space. Whatever space is used in form of the array is actually used to store the roots of the trees in the forest; hence, the array is not an overhead. From the given results, it is evident that the proposed forest is capable of providing an AVL tree-like performance at the cost of $O$ (1) time. An $A V L$ tree does have additional overheads, such as: time consumed in examining the tree balance and restructuring. Apart from that, each node in the $A V L$ tree requires two additional bits to store balance information. This is all that is required, in terms of time and space. As a result, the forest behaves as if it was a balanced tree without using any rebalancing algorithm.

The biggest advantage of this technique is that the entire forest acts as if it was a single $B S T$. That means we do not need to change any existing $B S T$-related algorithm in order to apply with the forest; insertion, deletion and updating algorithms remain the same. The tree traversal requires the roots of the individual tree to be passed to the traversal algorithm. In fact, all the trees in the forest can be traversed in parallel.

## 6. PARALLEL PROCESSING OF THE FOREST

Another benefit is compatibility with the concurrent environment. The forest of trees is far more compatible with the concurrent environment than a single BST. Here, we have a forest with multiple trees, and each tree in the forest is independent, and hence, can be operated independently. No proof is required to demonstrate that a forest allows more numbers of processes to act upon the different trees simultaneously. Without using any locking schemes, ' $k$ ' processes can act simultaneously on ' $k$ ' different trees. From the structure of the forest, it is reasonable to conclude that maintaining the tree this way increases the degree of parallelism by ' $k$ ' times. The choice of ' $k$ ' depends on the required degree of parallelism. For huge data sets and massively parallel systems, we would like to have a greater number of trees in the forest, requiring larger arrays. Although, to reduce the IPL, we haven't used any tree-restructuring technique here, still the individual tree in the forest can be restructured for even better results. As stated earlier, in global restructuring, the entire tree is taken as input and restructured. This process may take a lot of time particularly if the tree size is large. But in case of a forest, only a
part of the forest (single tree) would be needed for restructuring at a time. In fact, tree restructuring can be done simultaneously with the other operations. Furthermore, a binary search tree is a sensitive data structure, with regard to insertions and deletions. A series of insertions and deletions can harm the balance structure of the tree. Eppinger [10] has shown that performing a large number of insertions and asymmetric deletions increases the IPL of a tree to up to $\theta\left(n g^{3}(n)\right)$, resulting in the tree being no longer random. In the case of a forest, insertions and deletions will get distributed over the different trees, increasing the immunity of the structure to such operations.

## 7. CONCLUSION

We have shown that, without losing structural information, a tree could be converted into a forest with reduced internal path length and better compatibility with the concurrent environment. This has been achieved without using any restructuring algorithm. At the cost of hashing, without compromising for space, small modifications in the data structure results in faster node access, and an increased degree of parallelism.

Table: A comparison between the Conventional Tree and the Forest of Trees

| Sample <br> run | Conventional Tree |  |  | Forest |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Height | Path- <br> length | Average Height <br> of the Forest | Height of <br> the Worst <br> Tree | Collective Path- <br> length of the <br> Forest | \% Reduction in <br> Height | \% Reduction in <br> the Path-length |
| 1 | 18 | 11018 | 10.8 | 12 | 6193 | 40 | 44 |
| 2 | 20 | 10782 | 12.0 | 14 | 6594 | 40 | 39 |
| 3 | 21 | 11299 | 11.6 | 13 | 6623 | 44 | 41 |
| 4 | 24 | 13728 | 12.0 | 17 | 6542 | 50 | 52 |
| 5 | 21 | 11840 | 11.1 | 14 | 6081 | 47 | 48 |
| 6 | 22 | 11158 | 12.2 | 15 | 6567 | 44 | 41 |
| 7 | 20 | 11185 | 11.4 | 16 | 6467 | 43 | 42 |
| 8 | 20 | 10814 | 12.2 | 17 | 6533 | 39 | 40 |
| 9 | 22 | 11359 | 12.0 | 16 | 6481 | 45 | 43 |
| 10 | 24 | 12174 | 11.4 | 13 | 6414 | 52 | 47 |

a): Number of Nodes: 1023

| Sample <br> run | Conventional Tree |  |  | Forest |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Height | Path- <br> length | Average Height <br> of the Forest | Height of <br> the Worst <br> Tree | Collective Path- <br> length of the <br> Forest | \% Reduction in <br> Height | \% Reduction in <br> the Path-length |
| 1 | 21 | 24918 | 15.0 | 17 | 15865 | 28 | 36 |
| 2 | 21 | 24133 | 15.5 | 17 | 16216 | 26 | 32 |
| 3 | 25 | 24179 | 14.7 | 19 | 15603 | 41 | 35 |
| 4 | 26 | 26814 | 14.9 | 17 | 16110 | 42 | 40 |
| 5 | 22 | 24728 | 14.1 | 18 | 15936 | 36 | 35 |
| 6 | 27 | 28267 | 14.0 | 17 | 15207 | 48 | 46 |
| 7 | 24 | 26528 | 14.5 | 17 | 15489 | 39 | 42 |
| 8 | 24 | 25742 | 13.9 | 18 | 15705 | 42 | 39 |
| 9 | 25 | 24478 | 14.7 | 18 | 15508 | 41 | 37 |
| 10 | 23 | 24219 | 14.5 | 18 | 15418 | 37 | 36 |

(b): Number of Nodes: 2047

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