

# A SPACE-EFFICIENT ALGORITHM FOR PAGING UNBALANCED BINARY TREES

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Abstract: This work presents a new approach for paging large unbalanced binary trees which frequently appear in computational biology. The proposed algorithm aims at reducing the number of pages accessed for searching, and at decreasing the amount of unused space in each page as well as reducing the total number of pages required to store a tree. The algorithm builds the best possible paging when it is possible and employs an efficient strategy based on bin packing for allocating trees that are not complete. The complexity of the algorithm is presented. Experimental results are reported and compared with other approaches, including balanced trees. The comparison shows that the proposed approach is the only one that presents an average number of page accesses for searching close to the optimal and, at the same time, the page filling percentage is also close to the optimal.

## 1 INTRODUCTION

Binary trees are data structures popular for allowing efficient data searching (Gonnet and Baeza-Yates, 1991). A binary tree can get very large, and in this case it may be impossible to keep the whole tree in primary memory. A similar situation occurs when the tree is stored remotely and is accessed through a network. Many practical applications involve large unbalanced trees, particularly computational biology, in which trees are employed for string processing (Cohen, 2004). Such trees are constructed from biological sequences, which cannot be messed with, i.e. these trees cannot be balanced, so B-trees, for instance, cannot be employed (Pedersen, 2000). In such cases, it is necessary to define an efficient strategy to transfer the tree either from secondary memory or from the remote computer to primary memory where the search is executed. Instead of transferring one data item at a time, data collections called *pages* are usually employed to improve the transfer latency.

As the time required to obtain a page is much larger than the time required to process that page once it is allocated in primary memory, the criteria for allocating data in pages are essential for the efficiency of executing search procedures. The smallest the number of pages transferred, the fastest the search procedure is executed.

This work presents a new algorithm for paging binary trees. The algorithm aims at reducing the number of pages accessed for searching, and at the same time decreasing the amount of unused space in each page as well as reducing the total number of pages required to store a tree. In this way, the algorithm avoids wasting space to store a paged tree. The algorithm obtains the best possible tree paging, when it is possible, i.e. when the tree is complete and information is not only stored at the leaves. Figure 1 shows an example of the ideal paging for a tree with 63 nodes grouped in pages of 7 nodes. Besides that, an efficient strategy for allocating non-complete trees based on bin packing (Garey and Johnson, 1979) is presented.

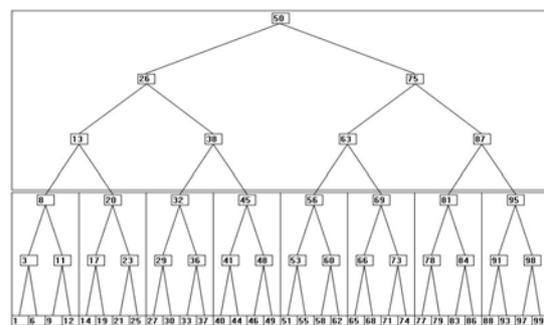


Figure 1: The ideal tree paging.

The algorithm starts at the root of the tree to be paged, allocating to the first page a subtree large

enough to completely fill the page. Next, every subtree that completely fills a page is allocated to a new page. The other subtrees that do not fill one page completely are collectively called the *fringe* of the tree. The algorithm allocates these subtrees in pages using bin packing.

Packing, in this case, consists of the allocation of a set of subtrees into a set of pages each with a previously known amount of available space. The number of pages required is minimized with this strategy. Furthermore, each subtree is guaranteed to be allocated in only one page, in order not to increase the number of pages accessed for searching.

An alternative data structure used to organized data in secondary memory is the B-tree (Gonnet and Baeza-Yates, 1991). We show through experimental results that our strategy is equivalent to B-trees in terms of the average number of pages accesses for searching. On the other hand our approach produces a page filling percentage by more than 30% in comparison with B-trees. Thus the total amount of space required to transfer a tree from a remote site is 30% better when our approach is used, in comparison with B-trees.

The rest of the paper is organized as follows. In section 2 we give preliminary definitions. In section 3 the algorithm is described, its specification is given as well as the complexity analysis. Section 4, contains experimental results. Section 5 points to related work, and section 6 concludes the paper.

## 2 PRELIMINARY DEFINITIONS

Binary trees are defined recursively as follows (Gonnet and Baeza-Yates, 1991):

- i) A binary tree  $T_0$  of zero nodes is a binary tree.
- ii) A binary tree  $T_n$  of  $n > 1$  nodes is a tuple  $(T_{left}, R, T_{right})$ , where  $R$  is a single node called the root of  $T_n$ .  $T_{left}$  and  $T_{right}$  are binary trees, respectively called left and right subtrees of the root. Considering that  $T_{left}$  has left nodes and  $T_{right}$  has right nodes, then  $left \geq 0$ ,  $right \geq 0$  and  $left + right = n - 1$ .

When it is not possible or desirable to keep the whole tree in main memory, the tree nodes are grouped in pages which are transferred to the main memory one at a time. Each page is formed by cells, each tree node is stored in a cell. As the time required to process a page is mainly the time required to transfer that page, the performance of tree manipulation algorithms is strictly related to the number of transferred pages.

Consider a binary tree with  $n$  nodes. Consider that a page stores a maximum of  $p$  nodes. The

allocation of nodes to pages must be done so that when a search algorithm or a tree traversal algorithm are executed, the number of accessed pages is as small as possible.

This work introduces a new algorithm for paging binary trees. The algorithm initially allocates subtrees which completely fill a page. The remaining subtrees are collectively called the fringe of the tree. The algorithm employs bin packing to allocate the fringe to as small a number of pages as possible, also keeping each subtree in only one page.

The bin packing problem is defined as follows. Given a constant  $C$  and a finite list of items  $L = p_1, p_2, \dots, p_n$ , where each item  $p_i$  is associated to a  $w(p_i)$  value satisfying  $0 < w(p_i) < C$ , find the smallest integer  $m$  such that  $L$  may be partitioned in  $m$  lists  $L_1, L_2, \dots, L_m$  where each list  $L_i$ , satisfies

$$w(L_i) = \sum_{p_j \in L_i} w(p_j) \leq C, i = 1, \dots, m.$$

In other words, the bin packing problem is expected to partition of a list of items into sublists in order to minimize the number of partitions considering the capacity of each sublist.

In this work bin packing is employed to allocate a set of subtrees to a set of pages. The algorithm determines both the sizes of the subtrees to be allocated, and the amount of space available in the pages. The subtrees, with sizes  $s_1, s_2, \dots, s_n$  must be allocated into  $C$  sized pages. By employing bin packing, the algorithm obtains an allocation that minimizes the number of pages required.

As an example, consider figure 2; subtree  $s_1$  is formed by nodes 3, 5, 7 and 12;  $s_2$  is formed by nodes 36, 38 and 41;  $s_3$  is formed by nodes 46, 49, 53 and 57;  $s_4$  is formed by node 73;  $s_5$  is formed by nodes 83, 85 and 87 and  $s_6$  is formed by nodes 93, 95, 97 and 98. These subtrees must be allocated in the smallest possible number of pages. So the sizes of subtree's are 4, 3, 4, 1, 3 and 4, respectively. Consider that the page size is 7.

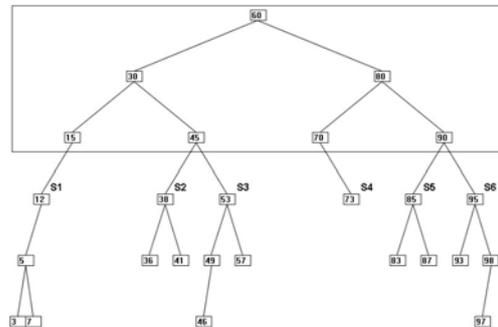


Figure 2: Bin packing: application example.

An optimal paging solution results in a high filling page percentage, allocating subtrees  $s_1$  and  $s_2$  to page 1, subtrees  $s_3$  and  $s_5$  to page 2 and subtrees  $s_4$  and  $s_6$  to page 3, as illustrated on figure 3.

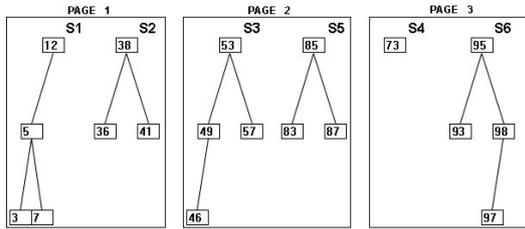


Figure 3: Paging by using bin packing.

The bin packing problem is a *NP-hard* combinatorial optimization problem (Garey and Johnson, 1979). Among the feasible alternatives for bin packing implementation, several approximation algorithms based on heuristics have been proposed (Garey and Johnson, 1979). In those cases, there is no guarantee that the optimal solution will be obtained, but the execution time is polynomial. The implementation used to obtain the experimental results found in section 5 of this work is based on a greedy approximation algorithm, that searches the best allocation for each current page.

### 3 PRELIMINARY DEFINITIONS

This section presents the proposed algorithm for paging binary trees. The algorithm can be applied when the information set to be treated is static, the access frequencies are not known and the storage is remote or secondary.

#### 3.1 Overview of the Algorithm

The proposed algorithm initially tries to group nodes in pages such that the nodes in one page are as close to each other as possible in the original tree. The algorithm reaches the ideal paging when the binary tree is complete and the number of nodes is a multiple of the page size. Furthermore, it establishes an efficient policy to page non-complete trees.

Before describing the algorithm, it is necessary to give the definition of a *patriarch*. A patriarch is the root of a subtree that is the first node of that subtree to be allocated in a new page. As many descendants of the patriarch as possible will be allocated in that page. Consider that a page stores up to  $x$  levels or generations of a tree, where  $x$  is a

positive integer, and the patriarch is at the first level. The algorithm stores in one page  $2^x - 1$  nodes.

The algorithm starts allocating the root of the tree as the patriarch of the first page. The descendants of the patriarch in the next  $x - 1$  levels are then allocated to the same page. At this point, if unused space was left in this page, it is filled with subtrees of the subsequent levels. If the page is completely filled, a new patriarch is chosen for a new page, and the process is repeated. If the subtrees of the subsequent levels do not completely fill the unused space, they belong to the fringe of the tree. The algorithm later uses bin packing to allocate all fringe subtrees.

#### 3.2 Algorithm Specification

The proposed algorithm uses two data structures called *SQ* (*Stack-Queue*) and *FL* (*Fringe List*). *FL* is a linear list (Gonnet and Baeza-Yates, 1991) that keeps the roots of the subtrees that belong to fringe.

*SQ* is a linear list  $SQ = (a_1, a_2, \dots, a_n)$  in which insertions and removals are possible at one end, called either rear or top, while at the other end, called front, only removals can be executed. The operations defined for *SQ* are the following: *create(SQ)*, initializes *SQ* as an empty data structure; *enqueue(x, SQ)*, inserts an element  $x$  at the rear of *SQ*, returning the resulting structure; *dequeue(SQ)*, removes the element at the front end of *SQ*, returning the element and the resulting structure; *pop(SQ)*, removes an element from the top of *SQ*, returning the element and the resulting structure; *empty(SQ)*, returns true when *SQ* is empty and false otherwise.

The algorithm is now described in terms of these data structures. Initially, the root of the tree is enqueued in *SQ*. The patriarch of a new page is dequeued from *SQ*. The patriarch descendants in the next  $x - 1$  levels are then allocated to this page.

If the page is completely filled, every element of the subsequent level is enqueued in *SQ* if it is the root of a subtree that has size greater than or equal to the page size; otherwise it is inserted in *FL*.

When a page is not completely filled, i.e. there is available space, every element of the subsequent level that is the root of a subtree that has size greater than or equal to the available space in the current page is enqueued in *SQ*; otherwise it is inserted in *FL*.

The algorithm proceeds as follows. The last element enqueued in *SQ* is popped and stored in the page. Its sons are enqueued in *SQ*. If there is still available space in the current page, and *SQ* is not

empty, again the last element enqueued in  $SQ$  is popped and stored in the page, and its sons are enqueued in  $SQ$ . This process is repeated until the current page is completely filled or  $SQ$  is empty.

When the current page is completely filled, that is, with no available space, the algorithm starts filling a new page. The patriarch of this new page is dequeued from  $SQ$ . The process above is repeated in order to fill the page.

When  $SQ$  is empty, the algorithm starts to allocate the fringe subtrees. The algorithm considers both the sizes of these subtrees, the available space in the last page, as well as the page size. The algorithm employs bin packing to determine the smallest number of pages that allocates those subtrees.

The proposed algorithm is specified in high level pseudocode in figure 4.

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```

Let the page size be 2**x-1

BEGIN
  create(SQ);
  enqueue(the tree root, SQ);
REPEAT
  create a new page;
  patriarch <- dequeue(SQ);
  allocate to current page the patriarch and
  its descendants of the next x-1 levels;
  IF (the page is completely filled)
  THEN
    FOR-ALL nodes at the subsequent level
      IF (node is root of subtree with size >=
page size)
        THEN enqueue(node, SQ);
        ELSE insert node in FL;
    ELSE
  /* there is space available in current page
  */
    FOR-ALL nodes at the subsequent level
      IF (node is root of subtree with size >=
available space)
        THEN enqueue(node, SQ);
        ELSE insert node in FL;
    WHILE (there is available space in current
page)and (not empty(SQ)) DO
      node <- pop(SQ);
      allocate node to current page;
      FOR-ALL sons of the allocated node
        enqueue(son, SQ);
    END-WHILE;
UNTIL empty(SQ);
IF (FL is not empty)
  THEN
    apply bin packing for paging the fringe
subtrees;

```

Figure 4: The proposed algorithm.

### 3.3 Algorithm Complexity

The complexity analysis is performed considering the algorithm divided in two phases. The first phase allocates the whole tree in pages, except the fringe. The second phase allocates the fringe using bin packing.

Let  $f$  be a worst-case complexity function, such that  $f(n)$  is the largest number of node accesses that the algorithm requires when the total number of nodes is  $n$ .

The complexity of the algorithm's first phase is linear, actually  $4n$ . Such linearity can be confirmed as follows. To compute the number of descendants of each node  $n$  accesses are required. To store all nodes of the tree in pages,  $2n$  accesses are required:  $n$  accesses to allocate the nodes themselves and  $n$  accesses to record the page address at the node's parent. Finally, at most  $n$  accesses are required in order to insert and remove nodes from the data structures employed,  $SQ$  and  $FL$ ; less than  $n$  nodes are ever inserted in one of these data structures. A node that is inserted in one of them, is not inserted in the other. Once a node is removed from the structure, it is not inserted again.

The algorithm's second phase depends mainly on the bin packing algorithm employed. Consider  $g$  the complexity of such algorithm, where  $g(n)$  describes the number of node accesses required to allocate the fringe subtrees. A number of practical approximation algorithms with quadratic complexity function are reported in the literature (Garey and Johnson, 1979). In this case, the algorithm complexity is quadratic.

## 4 EXPERIMENTAL RESULTS

This section presents experimental results. The implementation used to obtain the experimental results is based on a greedy approximation algorithm, that searches the best allocation for each current page. The metrics used to measure the algorithm performance are described. Results are compared to those of other approaches, including sequential allocation, breadth-first allocation, depth-first allocation, theoretical optimal paging of balanced trees and B-trees.

Experiments were performed with random sequences of keys. The trees had from 10 to 2000 nodes, in intervals of 10 nodes. The experiments were divided according to the page size, considering the values 3, 7 and 15. For each page size 100 experiments were performed.

### 4.1 Evaluation Metrics

To evaluate the proposed algorithm’s performance, two metrics were defined: the amount of unused space left in the pages, and the number of pages accessed when searching is executed.

### 4.2 Evaluating the Number of Pages Accessed for Searching

The first experiment reported refers to the total number of pages accessed for searching all nodes. Table 1 shows the comparison.

Table 1: Average number of accessed pages in different strategies.

Strategies	Page Size = 3	Page Size = 7	Page Size = 15
Sequential	1495,99	3313,30	7410,69
Breadth-First	1681,04	3981,23	9309,85
Depth-First	1198,61	2500,61	5471,75
Proposed Algorithm	969,69	1726,17	3365,44
B-trees	841,04	1644,62	2914,78
Theoretical Optimum	771,45	1383,10	2760,65

Figure 5 presents the average number of accessed pages considering page size equal to 15. As shown in the figure, results were compared to those produced by the usage of other approaches.

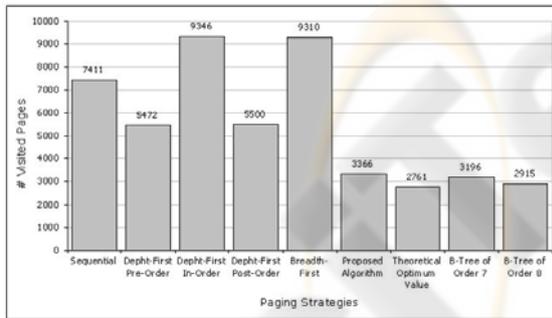


Figure 5: Average number of accessed pages; page size = 15.

Considering the presented results, the proposed algorithm is always better than sequential, breadth-first and depth-first allocation and worse than the theoretical optimal value, being however much closer to the optimal results than to others approaches.

In comparison with B-trees, the number of pages accessed for searching is similar. However, as presented in the next subsection, the proposed

approach is much more efficient than B-trees in terms of space efficiency.

### 4.3 Space-Efficiency Analysis

Another experimental result refers to the unused space left in the pages. Table 2 shows the average page filling percentage obtained in different strategies.

In the performed experiments it was observed that B-trees present a page filling percentage of 67.52% for randomized trees. On the other hand, the proposed algorithm presents an average page filling percentage of 98.62%, near to the optimal obtained with sequential paging.

Table 2: Page filling percentage in different strategies.

Strategies	Page Size = 3	Page Size = 7	Page Size = 15
Sequential, Breadth-First, Depth-First	98.82%	98.42%	98.69%
Proposed Algorithm	98.77%	98.42%	98.68%
B-trees	67.15%	67.30%	67.75%

Figure 6 shows the total amount of unused space produced by the different approaches. Considering B-trees the total bandwidth required to transfer a complete tree is proportionally larger than that required to transfer a tree paged with the proposed algorithm.

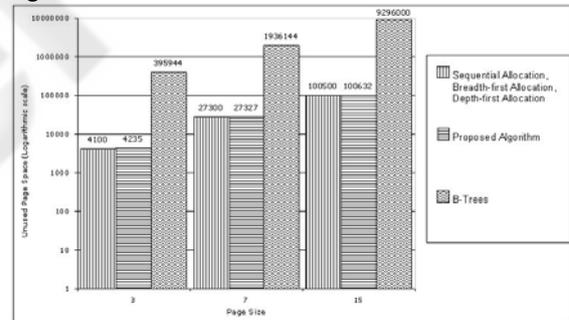


Figure 6: Amount of unused space measured experimentally.

## 5 RELATED WORK

Many algorithms and data structures have been proposed for treating massive data stored in external memory (Frakes and Baeza-Yates, 1992; Baeza-Yates and Ribeiro-Neto, 1999; Vitter, 2001), including the allocation of trees. A technique for allocating a binary tree partially paged using

external balancing is presented in (Henrich, SIX and Widmayer, 1990). Algorithms for maintaining a suffix tree structure in secondary storage are presented in (Clark and Munro, 1996). A clustering algorithm generating optimal worst-case external path length mapping of tree structures is described in (Diwan, Rane, Seshadri and Sudarshan, 1996). An efficient dynamic programming algorithm to pack trees is presented in (Gil and Itai, 1999). Another approach to pack trees in hierarchical memory using approximate algorithms is proposed in (Bender, Demaine and Farach-Colton, 2002).

## 6 CONCLUSION

This work described an efficient algorithm for paging unbalanced binary trees. The algorithm can be particularly applied for computational biology, in which large trees constructed from biological sequence, that cannot be balanced, are frequently found. The algorithm obtains the best possible allocation of nodes to pages when it is possible and proposes an efficient policy for filling pages of non-complete trees, based on the application of bin packing to the fringe of the tree. The complexity of the algorithm is given, which depends on the packing algorithm's complexity. The algorithm was implemented and experimental results were presented.

Considering the average number of accessed pages per search, the algorithm produces a page allocation up to 55% better than sequential allocation, up to 64% better than breadth-first and depth-first allocation, and results that are very close to those obtained with B-trees. On the other hand, considering the amount of unused space per page, and the total number of pages required, the algorithm presents an average page filling percentage of 98.62%. The comparison shows that the proposed approach is the only one that presents an average number of page accesses for searching close to the optimal and, at the same time, the page filling percentage is also close to the optimal.

Future work includes investigating experimentally the behavior of the algorithm considering other approximation algorithms for packing the fringe, and comparing those results to those obtained with variations of B-trees. Another topic left as future work is the evaluation of the behavior of the algorithm considering dynamic data, with frequent insertions and removals of nodes, as well as the impact of concurrent data access. The

evaluation of the algorithm considering real data instead of random data is also left as future work.

## REFERENCES

- Gonnet, G. H. ; Baeza-Yates, R. Handbook of Algorithms and Data Structures: in Pascal and C. Addison-Wesley, 1991, 424 p.
- Cohen, J. Bioinformatics - An Introduction for Computer Scientists. ACM Computing Surveys, v. 36, n. 2, p. 122-158, 2004.
- Pedersen, C. N. S. Algorithms in Computational Biology. PhD Dissertation, University of Aarhus, Denmark, 2000, 210 p.
- Garey, M. R. ; Johnson, D. S. Computers and Intractability: A Guide to the Theory of NP-Completeness. W. H. Freeman and Company, 1979, 338 p.
- Frakes, W. B. ; Baeza-Yates, R. Information Retrieval Data Structures and Algorithms. Prentice Hall, 1992, 464 p.
- Baeza-Yates, R. ; Ribeiro-Neto, B. Modern Information Retrieval. Addison-Wesley, 1999, 513 p.
- Vitter, J. S. External Memory Algorithms and Data Structures: Dealing with Massive Data. ACM Computing Surveys, v. 33, n. 2, p. 209-271, 2001.
- Henrich, A. ; SIX, H.W. ; Widmayer, P. Paging Binary Trees with External Balancing. Proceedings of the 15th International Workshop on Graph-theoretic Concepts in Computer Science, p. 260-276, Netherlands, 1990.
- Clark, D. R. ; Munro, J. I. Efficient Suffix Trees on Secondary Storage. Proceedings of the 7th Annual ACM-SIAM Symposium on Discrete Algorithms, p. 383-391, Atlanta, 1996.
- Diwan, A. A. ; Rane, S. ; Seshadri, S. ; Sudarshan, S. Clustering Techniques for Minimizing External Path Length. Proceedings of the 22nd VLDB Conference, p. 342-353, India, 1996.
- Gil, J. ; Itai, A. How to Pack Trees. Journal of Algorithms, v. 32, n. 2, p. 108-132, 1999.
- Bender, M. A. ; Demaine, E. D. ; Farach-Colton, M. Efficient Tree Layout in a Multilevel Memory Hierarchy. Proceedings of the 10th Annual European Symposium on Algorithms, p. 165-173, Italy, 2002.