

SUCCINCT TREE

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Summary: We have implemented Succinct Tree in 3 different methods. Succinct data structures require the amount of space that is close to the information-theoretic lower bound [A]. That is, there is very little "extra space". There are three methods for implementing succinct trees: balanced parentheses (BP), depth first unary degree sequence (DFUDS) and level-ordered unary degree sequence (LOUDS).

1. Introduction

Data structures are used to organize and store information in order to efficiently interact with the data. Most data structures are compared by the efficiency of the operations that can be performed. In order to complete these operations effectively, additional space may be used.

There continues to be a massive increase both in size and availability of large data sets that are processed by various applications. The underlying problem is: how can I compress the data but still query it quickly? What if there were a way to store the data in a compressed format, and be able to interact with it, without first having to decompress it?

In this project, we have implemented succinct data structure, that represents a given tree in succinct encoded form.

Succinct - Data - Structure: A data structure is called succinct if it uses an amount of space that uses nearly the theoretical lower wound of space with lower order additive terms, and still supports the desired operations.

Succinct representation of tree:

We have implemented three methods for succinct representation of tree:

- balanced parentheses (BP)
- \bullet depth first unary degree sequence (<code>DFUDS</code>)
- level-ordered unary degree sequence (LOUDS)

The pointer form of an n-node tree requires (n log n) bits. However, according to Information Theory only 2n (log n) bits are required to distinguish n nodes. A lot of research has focused on succinct representations which require just 2n + o(n) bits while still being able to implement sophisticated operations in constant time.

2. Succinct Representation of Trees

2.1. Balanced Parentheses (BP)

The balanced parentheses representation was first advocated by Jacobson in 1989 [1] and then expanded upon by Munro and Raman in 2001 [2]. The premise is that while the tree is traversed in depth-first preorder traver-

sal, an open parenthesis is written when a node is reached the first time, and a closing parenthesis is written when that node is reached again. This results in a sequence of 2n balanced parentheses, such that each node is represented by a pair of matching parentheses.

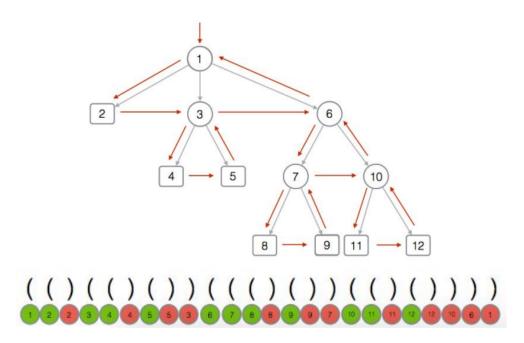


Figure 1: Representing tree as balanced parentheses.

Operations:

 \bullet parent bp(v): This function is used to find the parent of node whose opening parentheses is at position at v. It uses the fact that parent of node is the node corresponding to the opening parentheses that just encloses it.

Time complexity: o(number of nodes)

- \bullet first child bp(v): This function returns the index of parentheses corresponding to the first child of node whose parentheses is at v. It exploits the fact that if a vertex has child then all the parentheses corresponding the child lies between the opening and closing parentheses of the give node. Further, first child corresponds to the immediately next opening parentheses after opening parentheses of the node. Time complexity: o(number of nodes)
- \bullet last child bp(v): This function returns the index of parentheses corresponding to the last child of node whose parentheses is at v. If exist, the last child corresponds to the node whose closing parentheses is just immediately before the closing parentheses of the given node whose opening parentheses is at v. Time complexity: o(number of nodes)
- sibling(v): It returns the index of the parentheses corresponding to the sibling of node, if it exists. Sibling of a node is either the node corresponding to the parentheses(say m) just after the closing parentheses of the v, provided m is '('. Else it the node corresponding to the parentheses(say n) just before v, provided n is ')'.

Time complexity: o(number of nodes)

 \bullet subtree size bp(v) : The sub-tree size is calculated by finding the bounding parenthesis of v, adjusting it by v and then dividing by 2.

Time complexity: o(number of nodes)

- \bullet degree bp(v): This function returns the number of children of the node corresponding to v. Time complexity: O(number of children)
- depth bp(v): Depth is the length of the path from root to that node. This function returns the depth

of node corresponding to the parentheses at v. It can be calculated as rank of opening parentheses of v - rank of closing parentheses of v

Time complexity: o(number of nodes)

2.2. Depth-First Unary Degree Sequence (DFUDS)

The depth-first unary degree sequence was given in 1999 by Benoit, Demain, Munro, Raman, Raman and Rao. Here we follow depth first traversal for each node. At each node we first append i number of opening parentheses where i is the number of children and then 1 closing parenthesis. This way we use 2n parentheses to depict the entire tree. We use opening paranthesis to identify each node.

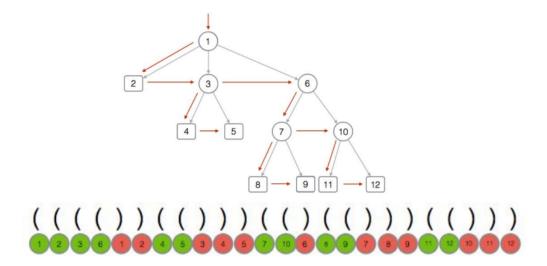


Figure 2: Representing tree as DFUDS.

Operations:

• parentdfuds(v) : This function is used to find the parent of node whose opening parentheses is at position at v. It uses the fact that parent of node is the node corresponding to the first closing parentheses after the node.

Time complexity: O(number of nodes)

 \bullet first child dfuds(v): This function returns the index of parentheses corresponding to the first child of node whose parentheses is at v.

Time complexity: O(number of nodes)

 \bullet last child dfuds(v): This function returns the index of parentheses corresponding to the last child of node whose parentheses is at v. If exist, the last child corresponds to the node whose opening parenthesis is just immediately before the closing parenthesis of the given node whose opening parenthesis is at v.

Time complexity: O(number of nodes)

- \bullet degree dfuds(v): This function returns the number of children of the node corresponding to v. Time complexity: O(number of nodes)
- \bullet leaf rank dfuds(v): It returns the number of leaves before the given node which is same as the number of occurrences of pattern "))" before the closing parenthesis of the node. Time complexity: O(number of nodes)
- \bullet leaf select dfuds(i): It returns the index of the ith leaf node which is same as the ith occurrence of the pattern "))".

Time complexity: O(number of nodes

2.3. Level-Ordered Unary Degree Sequence (LOUDS)

In this program we focus on the practical performance the fundamental Level-Order Unary Degree Sequence (LOUDS) representation [Jacobson, Proc. 30th FOCS, 549–554, 1989]. The premise is that the tree is traversed in level order. At each node 111..(d times)0 is appended to the string where d is the number of children of the node. To traverse the tree in the level order we heave used Breadth-First Search. At start, bit 10 is appended to the empty string to accommodate for the 1st node which is traversed from an imaginary super node. Then further traversal is carried using BFS. This results in a sequence of 2n+1 bits (n 1s and n+1 0s). A node, n, is identified by the position where nth 1 is in the string. We have used ones based numbering to represent the string. The numbering procedure is explained below. Ones-based numbering: Numbering the i th node in level-order by the position of the i th 1 bit and at the end of the corresponding level we append 0 to the string. This gives a node a number from 1, ..., 2n + 1. Number of 1 bits =n and number of 0 bits =n+1.

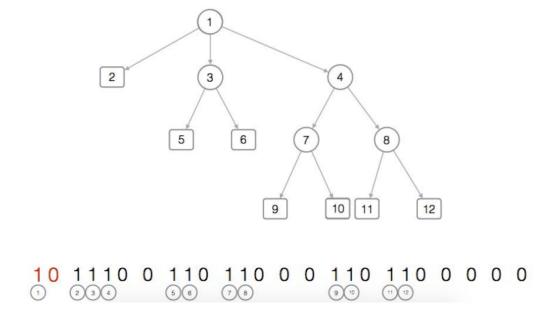


Figure 3: Representing tree as LOUDS.

Operations:

 \bullet parent lds(x): Algorithm = select1(rank0(S,x)). rank0(S,x) finds the number of "clumps" and then rank1() goes to the position of that clump. This gives index of the parent of the corresponding to the current node accessed through bit string stored.

Time complexity: O(n)

- \bullet first child $\mathrm{lds}(x)$: rank1(S,x) finds which node x is (in level order) and then select0 () goes that many 0's deep in the string, which is the 0 right before the child. The offset of 1 ensures that the first child is returned. If this returns 0, then return -1 else return the result. Time complexity: O(n)
- last child ds(x): rank1 (S,x) + 1 goes to the node following x, then select0 () visits the first child of that next node. The offset of -1 ensure we go back in the string 1, which will be the last child of x. If this returns 0, then return -1 else return the result. Time complexity: O(n)
- rightsibling(x): urn -1 else return the result. Right sibling(x) = if S[x+1] == 0 then -1 else x+1. Siblings are represented as 1's next to each other in the bit string so if a 0 follows x, then it is the last child. Otherwise, the right sibling is in the following index.

Time complexity : O(n)

 \bullet degree lds(v): The degree of the node is equal to the number of children it has. It is calculated using the formula degree(x) = lastchild(x) firstchild(x) + 1 Time complexity: O(n)

2.4. Visual Comparisons of Trees

Below is a visual comparison of the BP, DFUDS and LOUDS sequences for the following tree:

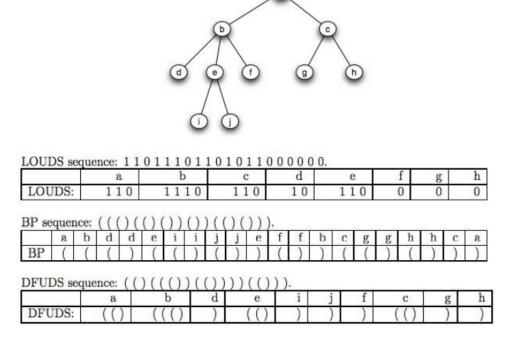


Figure 4: Traversal Comparison among 3 Representation.

Succinct Tree Comparison

LOUDS can implement a reduced set of navigation operations using just 5 percent extra space. In fact, it is the fastest choice in several operations.

DFUDS and LOUDS provide the best representations for navigating to a child quickly.

3. Algorithms

3.1. Rank and Select

Rank and Select algorithms have been implemented using Z-algorithm.

Algorithm 1 Rank (char* pattern, int size, int i)

```
1: integer p \leftarrow 0 and c \leftarrow 0
 2: integer l \leftarrow size + 1 + i
 3: allocate memory for Z array \leftarrow malloc(l sizeof(int))
 4: allocate memory for new string \leftarrow malloc((l + 1) sizeof(char))
 5: create new string \leftarrow pattern + "A" + parentheses string
 6: a \leftarrow 0 and b \leftarrow 0
 7: for j = 1tol do
       if j > b then
          a \leftarrow j \text{ and } b \leftarrow j
9:
10:
          while newstring[ba] == newstring[b]b < l do
            increment b
11:
          end while
12:
          Z \text{ array}[j] \leftarrow b \text{ a}
13:
         increment b
14:
       end if
15:
16: else
       k getsja;
17:
18:
       if Zarray[k] < bi + 1 then
          Z \text{ array}[j] \leftarrow Z \text{ array}[k];
19:
       end if
20:
21: else
22:
       a \leftarrow j;
       while newstring[ba] == newstring[b]b < l do
23:
         increment b
24:
       end while
25:
26:
       Z \text{ array}[j] \leftarrow b \text{ a}
       increment b
27:
28: end for
29: for p = 1tol - 1 do
       if Zarray[p] == size then
          increment c
31:
       end if
32:
       free(Z-array)
33:
34:
       free(new-string)
35:
       return c
36: end for
```

```
Algorithm 2 Select (char* pattern, int size, int i)
    integer p \leftarrow 0
 2: integer l \leftarrow size + 1 + i
    allocate memory for Z array \leftarrow malloc(l sizeof(int))
 4: allocate memory for new string \leftarrow malloc((l + 1) sizeof(char))
    create new string \leftarrow pattern + "A" + parentheses string
 6: store value of node and its corresponding '(' parenthesis in bp[count]
    a \leftarrow 0andb \leftarrow 0
 8: for j = 1 toldo do
       if j > b then
10:
         a \leftarrow jandb \leftarrow j
         while newstring[ba] == newstring[b]ANDb < l do
            increment b
12:
         end while
         Z \text{ array}[j] \leftarrow b \text{ a}
14:
         increment b
       end if
16:
    else
18:
       k \leftarrow j \ a
       if Zarray[k] < bi + 1 then
         Z \text{ array}[j] \leftarrow Z \text{ array}[k]
20:
       end if
22: else
       while newstring[ba] == newstring[b]ANDb < l do
24:
         increment b
26:
       end while
       Z \text{ array}[j] \leftarrow b \text{ a}
       increment b
28:
    end for
30: for p = 1tol1 do
       if Zarray[p] == size then
32:
         increment c
       end if
       if c == i then
34:
         free(Z-array)
         free(new-string)
36:
         return p-size
       end if
38:
       free(Z-array)
40:
       free(new-string)
```

return -1

42: end for

3.2. Balanced Parentheses (BP)

Algorithm 3 balanced - parenthesis

```
    start from root of the tree(r)
    store value of node and its corresponding '(' parenthesis in bp[count]
    increment count
    for foreachchildofnode(v) do
    balanced parentheses(v)
    end for
    store value of node and its corresponding ')' parenthesis in bp[count]
    increment count
```

Algorithm 4 find close bp (i)

```
    store value of node at bp[i] in some variable temp(say)
    for for j = i + 1to2numberofnodes do
    if valueofnodeatbp[j]isequaltotemp then
    return j
    end if
    end for
```

Algorithm 5 find open bp (i)

```
    store value of node at bp[i] in some variable temp(say)
    for forj_ilto0 do
    if valueofnodeatbp[j]isequaltotemp then
    return j
    end if
    end for
```

Algorithm 6 enclose bp (i)

```
1: count \leftarrow 0
2: for forj i1to0 do
      if parenthese satbp[j] is( then
3:
         count \leftarrow count + 1
4:
      end if
5:
      if parenthese satbp[j]is) then
6:
        count \leftarrow count \ 1
7:
      end if
8:
9: end for
10: if count = 1 then
      return j
12: end if
```

3.3. Depth-first Unary Degree Sequence (DFUDS)

```
Algorithm 7 depth - first

1: start from root of the tree(r)
2: for foreachchildofnode(v) do
3: store value of node and its corresponding '(' parenthesis in dfuds[count]
4: increment count
5: end for
6: store value of node and its corresponding ')' parenthesis in dfuds[count]
7: increment count
8: for foreachchildofnode(v) do
9: depth first(v)
10: end for
```

Algorithm 8 find close dfuds (i)

```
    store value of node at dfuds[i] in some variable temp(say)
    for for j = i + 1to2numberofnodes do
    if value of node at dfuds[j] is equal to temp then
    return j
    end if
    end for
```

Algorithm 9 find open dfuds (i)

```
1: store value of node at dfuds[i] in some variable temp(say)
2: for forj_ilto0 do
3: if value of node at dfuds[j] is equal to temp then
4: return j
5: end if
6: end for
```

Algorithm 10 enclose dfuds (i)

```
1: count ← 0
2: for forj i1to0 do
3: if parentheses at dfuds[j] is ( then
4: count ← count + 1
5: end if
6: end for
```

3.4. Level-Order Unary Degree Sequence (LOUDS)

Algorithm 11 LOUDS (traversal)

```
    start from root of the tree(r)
    store value of node and its corresponding '10' bits in lds[count]
```

3: increment count

4: call ldstraversal(root)

5: increment count

Algorithm 12 ldstraversal (root)

- 1: for foreachchildofnode(v) do
- 2: store the value of the node
- 3: add '1' bit to the string
- 4: enqueue(the current child for BFS traversal)
- 5: increment count by 1
- 6: end for
- 7: add bit'0' to the string
- 8: increment count by 1
- 9: **if** queue is empty **then**
- 10: return
- 11: **else**
- 12: call ldstraversal(dequeue)
- 13: **end if**

Algorithm 13 parent (y)

- 1: y = select1(rank0(x))
- 2: if y<0 then
- 3: return -1
- 4: **else**
- 5: return y
- 6: end if

Algorithm 14 firstchild (y)

- 1: y = select0(rank1(x))
- 2: if ds[y] = 0 then
- 3: return -1
- 4: else
- 5: return y
- 6: end if

Algorithm 15 lastchild (y)

- 1: y = select0(rank1(x)+1)-2
- 2: if ds[y] = 0 then
- 3: return -1
- 4: **else**
- 5: return y
- 6: end if

Algorithm 16 rightsibling (y)

- 1: if ds[y+1] = 0 then
- 2: return -1
- 3: **else**
- 4: return y+1
- 5: end if

4. Applications

There exist a range of real-world applications that can utilize this space-efficient storage method, most centered around information retrieval:

Search Engines : - Search engines can index billions of web pages and respond to queries about those pages in real-time. Therefore, it is crucial to decrease the space used by their index while still allowing efficient queries.

Mobile applications : - Mobile applications have a limited amount of storage available on the device and efficient storage allows for added functionality.

DNA representation : - Medical databases are massive and contain sequences that contain patterns and need to be queried quickly. One can successfully represent DNA in a succinct manner.

Streaming environments : - In a streaming environment, the next frame(s) need to be accessed quickly and efficiently. The smaller the size of the content, the faster it can be processed, and at a smaller cost.

5. Some further useful suggestions

Succinct representation of data structures like trees and graph had been a field of great interest. Many scientists and researchers had published considerable amount of work on same.

Few of them are, Guy Jacobson, J. I. Munro and V. Raman, using their algorithm Rank and select function can be found in nearly constant time, i.e O(1).

Thus using them, other operations which are dependent on them can be found efficiently both in terms of time and space.

6. Conclusions

Through this project, we implemented a space efficient data structure to store data and performed navigational operations on the given static tree.

The number of n-node binary trees is known to be the nth Catalan number [C]. Therefore, an encoding of binary trees with n nodes requires at least $\log(Cn)$ 2n $\log n + O(1)$ number of bits. Further we used three different succinct representation of a given tree to do so. We also analysed how we saved the space in representation of tree in succinct form.

7. Bibliography and citations

7.1. References:

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