COT 5407: Introduction to Algorithms

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Animations

- https://visualgo.net/
- http://www.cs.jhu.edu/~goodrich/dsa/trees/
- https://www.youtube.com/watch?v=Y-5ZodPvhmM
- http://www.algoanim.ide.sk/
## More Dynamic Operations

<table>
<thead>
<tr>
<th></th>
<th>Search</th>
<th>Insert</th>
<th>Delete</th>
<th>Comments</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Unsorted Arrays</strong></td>
<td>$O(N)$</td>
<td>$O(1)$</td>
<td>$O(N)$</td>
<td></td>
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<tr>
<td><strong>Sorted Arrays</strong></td>
<td>$O(\log N)$</td>
<td>$O(N)$</td>
<td>$O(N)$</td>
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<tr>
<td><strong>Unsorted Linked Lists</strong></td>
<td>$O(N)$</td>
<td>$O(1)$</td>
<td>$O(N)$</td>
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<tr>
<td><strong>Sorted Linked Lists</strong></td>
<td>$O(N)$</td>
<td>$O(N)$</td>
<td>$O(N)$</td>
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<tr>
<td><strong>Binary Search Trees</strong></td>
<td>$O(H)$</td>
<td>$O(H)$</td>
<td>$O(H)$</td>
<td>$H = O(N)$</td>
</tr>
<tr>
<td><strong>Balanced BSTs</strong></td>
<td>$O(\log N)$</td>
<td>$O(\log N)$</td>
<td>$O(\log N)$</td>
<td>As $H = O(\log N)$</td>
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</tbody>
</table>

<table>
<thead>
<tr>
<th></th>
<th>Se/In/De</th>
<th>Rank</th>
<th>Select</th>
<th>Comments</th>
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<tr>
<td><strong>Balanced BSTs</strong></td>
<td>$O(\log N)$</td>
<td>$O(N)$</td>
<td>$O(N)$</td>
<td></td>
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<tr>
<td><strong>Augmented BBSTs</strong></td>
<td>$O(\log N)$</td>
<td>$O(\log N)$</td>
<td>$O(\log N)$</td>
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</tbody>
</table>
RB-Tree Augmentation

- Augment x with \(\text{Size}(x)\), where
  - \(\text{Size}(x) = \text{size of subtree rooted at } x\)
  - \(\text{Size}(\text{NIL}) = 0\)
Augmented Data Structures

- Why is it needed?
  - Because basic data structures not enough for all operations
  - Storing extra information helps execute special operations more efficiently.

- Can any data structure be augmented?
  - Yes. Any data structure can be augmented.

- Can a data structure be augmented with any additional information?
  - Theoretically, yes.

- How to choose which additional information to store.
  - Only if we can maintain the additional information efficiently under all operations. That means, with additional information, we need to perform old and new operations efficiently maintain the additional information efficiently.
How to augment data structures

1. choose an underlying data structure
2. determine additional information to be maintained in the underlying data structure,
3. develop new operations,
4. verify that the additional information can be maintained for the modifying operations on the underlying data structure.
Augmenting RB-Trees

Theorem 14.1, page 309

Let $f$ be a field that augments a red-black tree $T$ with $n$ nodes, and $f(x)$ can be computed using only the information in nodes $x$, $\text{left}[x]$, and $\text{right}[x]$, including $f[\text{left}[x]]$ and $f[\text{right}[x]]$. Then, we can maintain $f(x)$ during insertion and deletion without asymptotically affecting the $O(\log n)$ performance of these operations.

For example,

\[
\text{size}[x] = \text{size}[\text{left}[x]] + \text{size}[\text{right}[x]] + 1
\]

\[
\text{rank}[x] = ?
\]
Augmenting information for RB-Trees

- Parent
- Height
- Any associative function on all previous values or all succeeding values.
- Next
- Previous
Room Scheduling Problem

- Given a set of requests to use a room
  - \([0, 6], [1, 4], [2, 13], [3, 5], [3, 8], [5, 7], [5, 9], [6, 10], [8, 11], [8, 12], [12, 14]\)
- Schedule largest number of above requests in the room
- Different approaches
  - Try by hand, exhaustive search, improve an initial solution, iterative methods, divide and conquer, greedy methods, etc.

**Simple Greedy Selection**

- Sort by start time and pick in “greedy” fashion
- Does not work. WHY?
  - \([0, 6], [6, 10]\) is the solution you will end up with.

**Other greedy strategies**

- Sort by length of interval
- Does not work. WHY?
Room Scheduling – Improved Solution

- [0,6], [1,4], [2,13], [3,5], [3,8], [5,7], [5,9], [6,10], [8,11], [8,12], [12,14]
- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]

-- Sorted by finish times

- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]
- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]
- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]
- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]
Greedy Algorithms

Given a set of activities \((s_i, f_i)\), we want to schedule the maximum number of non-overlapping activities.

**GREEDY-ACTIVITY-SELECTOR** \((s, f)\)

1. \(n = \text{length}[s]\)
2. \(S = \{a_1\}\)
3. \(i = 1\)
4. for \(m = 2\) to \(n\) do
5. if \(s_m\) is not before \(f_i\) then
6. \(S = S \cup \{a_m\}\)
7. \(i = m\)
8. return \(S\)
Why does it work?

- **THEOREM**
  Let $A$ be a set of activities and let $a_1$ be the activity with the earliest finish time. Then activity $a_1$ is in some maximum-sized subset of non-overlapping activities.

- **PROOF**
  Let $S'$ be a solution that does not contain $a_1$. Let $a'_1$ be the activity with the earliest finish time in $S'$. Then replacing $a'_1$ by $a_1$ gives a solution $S$ of the same size.

  Why are we allowed to replace? Why is it of the same size?

Then apply induction! How?
Greedy Algorithms – Huffman Coding

- Huffman Coding Problem

  Example: Release 29.1 of 15-Feb-2005 of TrEMBL Protein Database contains 1,614,107 sequence entries, comprising 505,947,503 amino acids. There are 20 possible amino acids. What is the minimum number of bits to store the compressed database?

  ~2.5 G bits or 300MB.

- How to improve this?

- Information: Frequencies are not the same.

<table>
<thead>
<tr>
<th>Amino Acid</th>
<th>Frequency</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ala (A)</td>
<td>7.72</td>
</tr>
<tr>
<td>Arg (R)</td>
<td>5.24</td>
</tr>
<tr>
<td>Asn (N)</td>
<td>4.28</td>
</tr>
<tr>
<td>Asp (D)</td>
<td>5.28</td>
</tr>
<tr>
<td>Cys (C)</td>
<td>1.60</td>
</tr>
<tr>
<td>Gln (Q)</td>
<td>3.91</td>
</tr>
<tr>
<td>Glu (E)</td>
<td>6.54</td>
</tr>
<tr>
<td>Gly (G)</td>
<td>6.90</td>
</tr>
<tr>
<td>His (H)</td>
<td>2.26</td>
</tr>
<tr>
<td>Ile (I)</td>
<td>5.88</td>
</tr>
<tr>
<td>Leu (L)</td>
<td>9.56</td>
</tr>
<tr>
<td>Lys (K)</td>
<td>5.96</td>
</tr>
<tr>
<td>Met (M)</td>
<td>2.36</td>
</tr>
<tr>
<td>Phe (F)</td>
<td>4.06</td>
</tr>
<tr>
<td>Pro (P)</td>
<td>4.87</td>
</tr>
<tr>
<td>Ser (S)</td>
<td>6.98</td>
</tr>
<tr>
<td>Thr (T)</td>
<td>5.52</td>
</tr>
<tr>
<td>Trp (W)</td>
<td>1.18</td>
</tr>
<tr>
<td>Tyr (Y)</td>
<td>3.13</td>
</tr>
<tr>
<td>Val (V)</td>
<td>6.66</td>
</tr>
</tbody>
</table>

- Idea: Use shorter codes for more frequent amino acids and longer codes for less frequent ones.
**Huffman Coding**

2 million characters in file.


IDEA 1: Use ASCII Code
Each need at least 8 bits,
Total = 16 M bits = 2 MB

IDEA 2: Use 4-bit Codes
Each need at least 4 bits,
Total = 8 M bits = 1 MB

IDEA 3: Use Variable Length Codes

<table>
<thead>
<tr>
<th>Length</th>
<th>A</th>
<th>C</th>
<th>G</th>
<th>T</th>
<th>N</th>
<th>Y</th>
<th>R</th>
<th>S</th>
<th>M</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>22</td>
<td>18</td>
<td>18</td>
<td>22</td>
<td>10</td>
<td>5</td>
<td>4</td>
<td>4</td>
<td>3</td>
</tr>
</tbody>
</table>

How to Decode?
Need Unique decoding!
Easy for Ideas 1 & 2.

What about Idea 3?

2 million characters in file.

Length = ?
Expected length = ?

Sum up products of frequency times the code length, i.e.,

\[(.22\times2 + .22\times2 + .18\times3 + .18\times3 + .10\times3 + .05\times5 + .04\times5 + .04\times5 + .03\times5) \times 2 \text{ M bits} = \]

\[3.24 \text{ M bits} = .4 \text{ MB}\]