

COP 4516: Competitive  
Programming and Problem Solving

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# Problems to think about!

- What is the least number of comparisons you need to sort a list of 3 elements? 4 elements? 5 elements?
- How to arrange a tennis tournament in order to find the tournament **champion** with the least number of matches? How many tennis matches are needed? How to arrange a tennis tournament in order to find the **runner up** to the champion with the least number of matches?
- How to randomize the order of a list?

# Sorting Algorithms

- SelectionSort
- InsertionSort
- BubbleSort
- ShakerSort
- MergeSort
- HeapSort
- QuickSort
- Bucket & Radix Sort
- Counting Sort

# Data Structure Evolution

- Standard operations on data structures
  - **Search**
  - **Insert**
  - **Delete**
- Linear Lists
  - Implementation: **Arrays** (**Unsorted and Sorted**)
- **Dynamic** Linear Lists
  - Implementation: **Linked Lists**
- **Dynamic** Trees
  - Implementation: **Binary Search Trees**

# Data Structures Comparison

Data Structure \ Operation	Search	Insert	Delete
Unsorted Array			
Sorted Array			
Unsorted Linked List			
Sorted Linked List			
Binary Search Trees			
Balanced Binary Search Trees			

# BST: Search

TREESearch(*node x, key k*)

▷ Search for key  $k$  in subtree rooted at node  $x$

1 **if**  $((x = \text{NIL}) \text{ or } (k = \text{key}[x]))$

2     **then return**  $x$

3 **if**  $(k < \text{key}[x])$

4     **then return** TREESearch( $\text{left}[x], k$ )

5     **else return** TREESearch( $\text{right}[x], k$ )

Time Complexity:  $O(h)$

$h$  = height of binary search tree

Not  $O(\log n)$  — Why?

# BST: Insert

TREEINSERT(*tree T, node z*)

▷ Insert node *z* in tree *T*

1  $y \leftarrow \text{NIL}$

2  $x \leftarrow \text{root}[T]$

3 **while** ( $x \neq \text{NIL}$ )

4     **do**  $y \leftarrow x$

5         **if** ( $\text{key}[z] < \text{key}[x]$ )

6             **then**  $x \leftarrow \text{left}[x]$

7             **else**  $x \leftarrow \text{right}[x]$

8  $p[z] \leftarrow y$

9 **if** ( $y = \text{NIL}$ )

10     **then**  $\text{root}[T] \leftarrow z$

11     **else if** ( $\text{key}[z] < \text{key}[y]$ )

12         **then**  $\text{left}[y] \leftarrow z$

13         **else**  $\text{right}[y] \leftarrow z$

Time Complexity:  $O(h)$

$h$  = height of binary search tree

Search for  $x$  in  $T$

Insert  $x$  as leaf in  $T$

# BST: Delete

Time Complexity:  $O(h)$

$h$  = height of binary search tree

TREEDeLETE(*tree*  $T$ , *node*  $z$ )

▷ Delete node  $z$  from tree  $T$

```
1  if ((left[z] = NIL) or (right[z] = NIL))
2    then y ← z
3    else y ← TREE-SUCCESSOR(z)
4  if (left[y] ≠ NIL)
5    then x ← left[y]
6    else x ← right[y]
7  if (x ≠ NIL)
8    then p[x] ← p[y]
9  if (p[y] = NIL)
10   then root[T] ← x
11   else if (y = left[p[y]])
12         then left[p[y]] ← x
13         else right[p[y]] ← x
14  if (y ≠ z)
15   then key[z] ← key[y]
16        cop y's satellite data into z
17  return y
```

Set  $y$  as the node to be deleted. It has at most one child, and let that child be node  $x$

If  $y$  has one child, then  $y$  is deleted and the parent pointer of  $x$  is fixed.

The child pointers of the parent of  $x$  is fixed.

The contents of node  $z$  are fixed.



# Data Structures Comparison

Data Structure \ Operation	Search	Insert	Delete
Unsorted Array	$O(n)$	$O(1)$	$O(n)$
Sorted Array	$O(\log n)$	$O(n)$	$O(n)$
Unsorted Linked List	$O(n)$	$O(1)$	$O(n)$
Sorted Linked List	$O(n)$	$O(n)$	$O(n)$
Binary Search Trees	$O(h)$	$O(h)$	$O(h)$
Balanced Binary Search Trees	$O(\log n)$	$O(\log n)$	$O(\log n)$

# Animations

- **BST:**

[http://babbage.clarku.edu/~achou/cs160/examples/bst\\_animation/BST-Example.html](http://babbage.clarku.edu/~achou/cs160/examples/bst_animation/BST-Example.html)

- **Rotations:**

[http://babbage.clarku.edu/~achou/cs160/examples/bst\\_animation/index2.html](http://babbage.clarku.edu/~achou/cs160/examples/bst_animation/index2.html)

- **RB-Trees:**

[http://babbage.clarku.edu/~achou/cs160/examples/bst\\_animation/RedBlackTree-Example.html](http://babbage.clarku.edu/~achou/cs160/examples/bst_animation/RedBlackTree-Example.html)

# Example

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

# Example

- [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.**

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

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

# Huffman Coding

2 million characters in file.

A, C, G, T, N, Y, R, S, M

**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**

Percentage  
Frequencies

**IDEA 3: Use Variable Length Codes**

A	22	11
T	22	10
C	18	011
G	18	010
N	10	001
Y	5	00011
R	4	00010
S	4	00001
M	3	00000

**How to Decode?**

Need Unique decoding!  
Easy for Ideas 1 & 2.  
What about Idea 3?

110101101110010001100000000110

110101101110010001100000000110

2 million characters in file.

Length = ?

Expected length = ?

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

$(.22 \times 2 + .22 \times 2 + .18 \times 3 + .18 \times 3 + .10 \times 3 + .05 \times 5 + .04 \times 5 + .04 \times 5 + .03 \times 5) \times 2 \text{ M bits} =$

3.24 M bits = **.4 MB**