# 1 RAM Model

## 1.1 Memory

Infinite sequence of cells, contains w bits. Every cell has an address starting at 1

## 1.2 CPU

32 registers of width w bits.

# 1.2.1 Operations

Set value to register (constant or from other register). Take two integers from other registers and store the result of; a+b, a-b,  $a \cdot b$ , a/b. Take two registers and compare them; a < b, a = b, a > b. Read and write from memory.

#### 1.3 Definitions

An algorithm is a set of atomic operations. It's cost is is the number of atomic operations. A word is a sequence of w bits

## Worst-case

Worst-case cost of an algorithm is the longest possible running time of input size n

# Dictionary search

let n be register 1, and v be register 2 register  $left \rightarrow 1$ ,  $right \rightarrow 1$ while  $left \leq right$ register  $mid \rightarrow (left + right)/2$ if the memory cell at address mid = vthen

return yes

else if memory cell at address mid > v

right = mid - 1else left = mid + 1

return no

Worst-case time:  $f_2(n) = 2 + 6 \log_2 n$ 

# 4 Big-O

We say that f(n) grows asymptotically no faster than g(n) if there is a constant  $c_1 > 0$ such that  $f(n) \leq c_1 \cdot g(n)$  and holds for all n at least a constant  $c_2$ . This is denoted by f(n) = O(g(n)). $\lim_{n \to \infty} \frac{f(n)}{g(n)} = c$  for some constant c

# 4.1 Example

 $1000\log_2 n = O(n),$  $n \neq O(10000 \log_2 n)$  $\log_{b_1} n = O(\log_{b_2} n)$  for any constants  $b_1 > 1$ and  $b_2 > 1$ . Therefore  $f(n) = 2 + 6 \log_2 n$  can be represented;  $f(n) = O(\log n)$ 

# $\mathbf{Big}$ - $\Omega$

If g(n) = O(f(n)), then  $f(n) = \Omega(g(n))$  to indicate that f(n) grows asymptotically no slower than g(n). We say that f(n) grows asymptotically no slower than g(n) if  $c_1 > 0$ such  $f(n) \ge c_1 \cdot g(n)$  for  $n > c_2$ ; denoted by  $f(n) = \Omega(g(n))$ 

# 6 Big- $\Theta$

If f(n) = O(g(n)) and  $f(n) = \Omega(g(n))$ , then  $f(n) = \Theta(g(n))$  to indicate that f(n) grows asymptotically as fast as q(n)

# Sort

# 7.1 Merge Sort

Divide the array into two parts, sort the individual arrays then combine the arrays together.  $f(n) = O(n \log n)$ .

This is the fastest sorting time possible (apart from  $O(n \log \log n)$ 

# 7.2 Counting Sort

A set S of n integers and every integer is in the range [1, U]. (all integers are distinct)

Step 1: Let A be the array storing S. Create array B of length U. Set B to zero.

Step 2: For  $i \in [1, n]$ ; Set x to A[i], Set B[x]

**Step 3:** Clear A, For  $x \in [1, U]$ ; If B[x] = 0continue, otherwise append x to A

# 7.2.1 Analysis

Step 1 and 3 take O(U) time, while Step 2 O(n) time. Therefore running time is O(n +U) = O(U).

## 8 Random

RANDOM(x, y) returns an integer between x and y chosen uniformly at random

#### Data

## 9.1 Data Structure

Data Structure describes how data is stored in memory.

#### 9.2 LinkedList

Every node stores pointers to its succeeding and preceding nodes (if they exist). The first node is called the head and last called the tail. The space required for a linkedlist is O(n) memory cells. Starting at the head node, the time to enumerate over all the integers is O(n). Time for assertion and deletion is equal to O(1)

## 9.3 Stack

The stack has two operations; Push (Inserts a new element into the stack), Pop (Removes the most recently inserted element from the stack and returns it. Since a stack is just a linkedlist, push and pop use O(1) time.

# 9.4 Queue

The queue has two operations; En-queue (Inserts a new element into the queue), Dequeue (Removes the least recently used element from the queue and returns it). Since a queue is just a linkedlist, push and pop use O(1) time.

# 10 Dynamic Arrays

## 10.1 Naive Algorithm

insert(e): Increase n by 1, initial an array A' of length n, copy all n-1 of A to A', Set A'[n]=e, Destroy A.

This takes  $O(n^2)$  time to do n insertions.

# 10.2 A Better Algorithm

insert(e): Append e to A and increase n by 1. If A is full; Create A' of length 2n, Copy A to A', Destroy A and replace with A' This takes O(n) time to do n insertions.

# 11 Hashing

The main idea of hashing is to divide the dataset S into a number m of disjoint subsets such that only one subset needs to be searched to answer any query.

# 11.1 Pre-processing

Create an array of linkedlist(L) from 1 to mand an array H of length m. Store the heads of L in H, for all  $x \in S$ ; calculate hash value (h(x)), insert x into  $L_{h(x)}$ . We will always choose m = O(n), so O(n + m) = O(n)

# 11.2 Querying

Query with value v, calculate the hash value h(v), Look for v in  $L_h(v)$ . Query time:  $O(\mid L_{h(v)}\mid)$ 

# 11.3 Hash Function

Pick a prime  $p; p \ge m, p \ge \text{any integer}$ k. Choose  $\alpha$  and  $\beta$  uniformly random from  $1, \ldots, p-1$ . Therefore:  $h(k) = 1 + (((\alpha k + \beta))$  $\mod p \pmod m$ 

# 11.3.1 Any Possible Integer

The possible integers is finite under the RAM Model. Max:  $2^w - 1$ . Therefore pexists between  $[2^w, w^{w+1}]$ .

# 11.4 Timing

Space: O(n), Preprocessing time: O(n), Query time: O(1) in expectation

## 12 Week 3 - Extra

When using 'Direction 1: Constant Finding' setting  $c_1$ , always set it to match the coefficent on the LHS so that you can cancel.

When trying to get a contradiction, try and isolate an  $x \cdot c_1$  on the RHS, where  $x \in \mathbb{Z}$ , such that an expression that contains n is  $\leq x \cdot c_1$ 

Make judicious use of the max function when adding functions together If  $f_1(n) + f_2(n) \leqslant c_1 \cdot g_1(n) + c'_1 \cdot g_2(n) \leqslant$  $max\{c_1, c_1'\} \cdot (g_1(n) + g_2(n)), \text{ for all }$  $n \geqslant \max\{c_2, c_2'\}.$ 

#### **13** The Master Theorem

## 13.1 Theorem 1

$$n+\frac{n}{c}+\frac{n}{c^2}+\ldots+\frac{n}{c^h}=O(n)$$

# 13.2 Theorem 2

Let f(n) be a function that returns a positive value for every integer n > 0. We know:

$$f(1) \le c_1$$
  
 $f(n) \le \alpha \cdot f(\lceil n/\beta \rceil) + c_2 \cdot n^{\gamma} \text{ for } n \ge 2$ 

where  $\alpha, \beta, \gamma, c_1$  and  $c_2$  are positive constants. Then:

- If  $log_b \alpha < \gamma$  then  $f(n) = O(n^{\gamma})$
- If  $log_b \alpha = \gamma$  then  $f(n) = O(n^{\gamma} \cdot log(n))$
- If  $loq_b \alpha > \gamma$  then  $f(n) = O(n^{log_\beta(a)})$

# SE Set 3

Find out how many times a recurrence takes to terminate, and then proceed to eyeball the time complexity

#### Hierarchy **15**

$$O(1) \leqslant O(\log(n)) \leqslant O(n^c)$$
  
$$\leqslant O(n) \leqslant O(n^2)$$
  
$$\leqslant O(n^c) \leqslant O(c^n)$$

#### 16 Trees

An undirected graph if a pair of (V, E)where:

- V is a set of elements, eac of which is called a node
- E is a set of pairs u, v such that:
  - -u and v are distint nodes;
  - If (u, v) is in E, then (v, u) is also in E - we say that there is an edge between u and v.

A node may also be called a vertex. We will refer to V as the vertex set or the node set of the graph, and E the edge set.

For example, there is a graph (V, E) wher  $\{a, b, c, d, e\}$ E $\{(a,b),(b,a),(b,c),(c,b),(a,d),(d,a),(b,d),(d,b),(c,e),(e,c)\} \text{is at the level immediately below } u$ The number of edges equals |E|/2 =

Let G = (V, E) be an undirected graph. A path in G is a sequence of nodes  $(v_1, v_2, ..., v_k)$  such that

• For every  $i \in [1, k-1]$ , there is an edge between  $v_i$  and  $v_{i+1}$ 

A cycle in G is a path  $(v_1, v_2, ..., v_k)$  such that

k ≥ 4

10/2 = 5

- $\bullet v_1 = v_k$
- $v_1, v_2, ..., v_k$  are distinct

An undirected graph G = (V, E) is connected if, for any two distinct vertices u and v, G has a path from u to v.

A tree is connected undirected graph contains no cycles.

A tree with n nodes has n-1 edges.

Given any tree T and anarbitrary node r, we can allocate a level to each node as

- ullet r is the root of T this is leel 0 of the
- $\bullet$  All the nodes that are n edge away from r constitute level n of the tree

The number of levels is called the height of the tree. We say that T has been rooted once a root has been designated.

Consider a tree T that has been rooted. Let u and v be two nodes in T. We say that u is the parent of v if:

- - There is an edge below u and v

Accordingly, we say that v is a child of u.

Let u and v be two nodes in T. We say that u is the ancestor of v if one of the following holds:

- $\bullet \ v = u$
- u is the parent of an v
- u s the parent of an ancestor of v

Accordingly, we say that v is a decendant of u.

In particular, if  $u \neq v$ , we say that u is a proper ancestor of v, and likewise, v is a proper descendant of u.

Let u be a node in a rooted tree T. The subtree of u is the part of T that is "at or below" u.

In a rooted tree, a node is a leaf node if it has no children; otherwise it is an internal node.

Let T be a rooted tree where every internal node has at least 2 child nodes. If m is the number of leaf nodes, then the number of internal nodes is at least m-1.

A k-ary tree is a rooted tree where every internal node has at most k child nodes.

A 2-ary tree is called a binary tree.

Consider a binary tree with height h. Its level  $n \ (0 \le n \le h-1)$  is full if it contains  $2^n$ .

A binary tree of height h is complete if:

- At Level h-1, the leaf nodes are "as far left as possible"

A complete tree with  $n \ge 2$  nodes has height  $O(\log n)$ 

