

# Algorithms and Programming Techniques

## COMP3121 UNSW

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2022T2

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# 1 Introduction - Revision

## 1.1 Rates of Growth

**The Problem** To analyse algorithms, we need a way to compare two functions representing the runtime of each algorithm. However, comparing values directly presents a few issues.

- Outliers may affect the results.
- One algorithm may be slower for a set period of time, but catch up after a while.
- The runtime of an algorithm may vary depending on the implementation or the architecture of the machine where, some instructions are faster than others.

**Asymptotic Growth** For algorithms, we often prefer to refer to them in terms of their asymptotic growth in runtime, with relation to the input size.

A function that quadruples with every extra input will always have a greater runtime than one that increases linearly with each new input, for some large enough input size.

**Big O Notation** We say  $f(n) = O(g(n))$  if there exists a positive constants  $C, N$  such that

$$0 \leq f(n) \leq Cg(n) \quad \forall n \geq N.$$

We may refer to  $g(n)$  to be the asymptotic upper bound for  $f(n)$ .

**Big Omega Notation** We say  $f(n) = \Omega(g(n))$  if there exists positive constants  $c, N$  such that

$$0 \leq cg(n) \leq f(n) \quad \forall n \geq N.$$

Then,  $g(n)$  is said to be an asymptotic lower bound for  $f(n)$ . It is useful to say that a problem is at least  $\Omega(g(n))$ .

**Landau Notation**  $f(n) = \Omega(g(n))$  if and only if  $g(n) = O(f(n))$ .

There are strict version of Big O and Big Omega notations; these are little o and little  $\omega$  respectively.

We say  $f(n) = \Theta(g(n))$  if

$$f(n) = O(g(n)) \text{ and } f(n) = \Omega(g(n)).$$

That is, both  $f$  and  $g$  have the same asymptotic growth.

**Logarithms** Logarithms are defined so that for  $a, b > 0$  where  $n \neq 1$ , let

$$n = \log_a b \Leftrightarrow a^n = b.$$

They have the following properties:

- $a^{\log_a n} = n$

- $\log_a(mn) = \log_a(m) + \log_a(n)$
- $\log_a(n^k) = k \log_a(n)$

By the change of base,

$$\log_a(x) = \frac{\log_b(x)}{\log_b(a)}.$$

As such, the denominator is constant in terms of  $x$  and so all log bases are equivalent under Big Theta notation.

## 1.2 Assumed Data Structures

**Arrays** We assume static arrays (though, it is possible to extend to dynamic arrays).

- We assume random-access in  $O(1)$
- Insert / delete  $O(n)$
- Search:  $O(n)$  -  $\log n$  if sorted

**Linked Lists** We assume the linked lists are doubly-linked since the  $2\times$  overhead is negligible.

- Accessing next / previous:  $O(1)$
- Insert / delete to head or tail:  $O(1)$
- Search:  $O(1)$ .

**Stacks** Last in, first out.

- Accessing top:  $O(1)$
- Insert / delete from top:  $O(1)$

**Queue** First in, first out.

- Access front:  $O(1)$
- Insert front:  $O(1)$
- Delete front:  $O(1)$

**Hash Tables** Store values by their hashed keys. Ideally, no two keys will hash to the same value, however this may not be guaranteed.

- Search is expected to be  $O(1)$  however, in the worst case, we expect to have to search through all the values in  $O(n)$ .
- Insertion is expected to be  $O(1)$  however, in the worst case, we expect to have to search through all the values in  $O(n)$ .
- Deletion follows the same pattern of  $O(1)$  expectation and  $O(n)$  worst case.

**Binary Search Trees** We store (comparable) keys (or key-value pairs) in a binary tree, where each node has at most two children, the left and right. The value of a node must be greater than the values of all its children in its left sub-tree and greater than those in the right sub-tree.

- In the best case, the height of the tree is  $h = \log_2 n$ . Such a tree is *balanced*. in the worst case however, the tree is a long chain of height  $n$ .
- The average search is  $\log n$ . If the tree is self-balancing then search and other operations are guaranteed to be  $\log n$ .

**Binary Heap** A max-heap is such that the value of a node is greater than or, equal to the value of all its children. A min-heap follows the same principle but in reverse.

- Finding the maximum involves finding the top item in  $O(1)$
- Deleting the top item will also require re-balancing.
- Re-balancing the heap requires  $\log n$  time.
- Insertion also requires  $\log n$  time.

## 1.3 Algorithms

**Linear Search** Given an array  $A$  of  $n$  integers. We may determine if a value is inside  $A$ , by searching through the array linearly. This occurs in  $O(n)$ .

**Sorted Arrays - Binary Search** Given a sorted array  $A$  of  $n$  integers that are sorted by value. We may determine if a value is inside  $A$ , by binary search. We pick the midpoint of the  $A$ . If the midpoint  $m$  is the value desired, we return. Otherwise, we continue to recursively search through the array by halving the search space. If the  $m$  is greater than the value we desire, then we search the right sub-array of the array; otherwise, we search the left sub-array of the array

**Decision Problems and Optimisation Problems** Decision problems are of the form

Given some parameters  $X$ , can you ...

Optimisation problems are of the form

What is the smallest  $X$  for which you can ...

**Comparison Sorting** We are given an array of  $n$  items. We may sort the array in  $O(n \log n)$  at best.

**Asymptotic Lower Bound on Comparison Sorting** There are  $n!$  permutations of the array, of which there is 1 correct sort. If we do  $k$  comparisons, then we are able to check  $2^k$  permutations of results from these comparisons and so, can at most, distinguish  $2^k$  permutation.

Thus, we need to perform comparisons such that  $2^k \geq n!$ . That is,  $k \geq \log_2(n!)$ . Therefore,

$$k \geq \frac{n}{2} \log_2 \left( \frac{n}{2} \right)$$

## 1.4 Non-Comparison Sort

**Counting Sort** Create another array  $B$  of size  $k$  where  $k$  is the maximum size of an element. We iterate through the array and count the number of times each element occurs, iterating the index of  $B$  that is equivalent to the value of our current element.

The time complexity is  $O(n + k)$  where  $O(k)$  space is required.

**Bucket Sort** Distribute the items into buckets  $1, \dots, k$  such that each bucket contains a range of items such that all items in bucket 1 are less than all the items in bucket 2 and so on. We may sort within that bucket with any algorithm and then concatenate all the buckets together.

**Radix Sort** We assume an array  $A$  of  $n$  keys, of  $k$  symbols.

We bucket the keys by their first symbol. Then, within each bucket we also classify those items by the next symbol recursively. This happens in  $O(nk)$ .

**LSD Radix Sort** Sort all keys by their last symbol, then sort all keys by their second last symbol, and so on. The sorting algorithm at each stage must be stable.

The time is  $O(nk)$  with space complexity of  $O(n + k)$ .

## 1.5 Graphs

# 2 Algorithm Analysis

## 2.1 Stable Matching Problem

**Hospital's Stable Matching Problem** Suppose there exists  $n$  doctors and hospital. Each hospital wants to hire exactly one new doctor and the hospital ranks the priority of the doctors they would prefer and each doctor also lists their preferences for the hospitals they'd like to go to.

For a stable match, we would like to create an allocation where no group would be happy to trades their allocations with each other.

**Naive Solution** The naive method computes all possible allocations and then picks one that is stable.

This runs in  $O(n!)$  time which is not preferable.

### Fun Fact about Factorials - Sterling's Approximation

$$n! \approx \left(\frac{n}{e}\right)^n$$

### Gale - Shapely Algorithm - Assumptions

- Produces pairs in stages, with possible revisions.
- A hospital which has not been paired, will be called *free*.
- Hospitals will offer jobs to doctors, who will decide whether to accept a job or not. Doctors may renege.
- All hospitals start off as free.

**Gale - Shapely - Solution** While there exists a free hospital which has not offered jobs to all doctors, pick a free  $h$  and have it offer a job to the highest doctor  $d$  on its list to whom they have not offered a job yet.

If no one has offered  $d$  a job yet, they will always accept the pair  $h, d$ .

Otherwise, if they're already on the pair  $h', d$  then if  $h$  is a higher preference than  $h'$  the doctor will renege  $h'$  for  $h$  to form  $(h, d)$ . Otherwise, the hospital will seek another person to offer a job.

**Proving Termination in  $n^2$**  In all rounds of the Gale-Shapely algorithm, the hospital will offer a job to one doctor. They may offer a job to each doctor only once. Thus, the hospital may only make  $n$  offers. There are  $n$  hospitals each making  $n$  offers so, the algorithm must terminate in  $n^2$  rounds or less.

**Proving Correctness of Gale-Shapely** If the while loop has terminated with  $h$  still free, that implies that  $h$  has offered a job to everyone. Since the only way to get declined is if another, more appealing hospital has offered a job to other doctors.

If there are no more doctors available, then  $n$  other hospitals must have offered a job to  $n$  doctors. However this would imply there are  $n+1$  hospitals in total, which is a contradiction.

Hospitals will be happy as they start with their highest preference first and, doctors will move up to accept their most preferred offer. Thus, there cannot exist a circumstance where both a hospital and doctor would like to swap.

## 3 Divide-and-Conquer Method

### 3.1 Counting Inversions

**Brute Force** Brute force will take  $O(n^2)$  time. Any method of looking at entries one by one will have at worst, a quadratic time complexity since there can be  $\frac{n^2}{2}$  inversions.

**Divide and Conquer Method** We can split the array into two equal parts  $A_{hi}$  and  $A_{lo}$ .

Then, for all of the elements in  $A_{hi}$  and the total inversions is the number of elements in  $A_{lo}$  that are less than the elements of  $A_{hi}$ .

We may sort both  $A_{hi}$  and  $A_{lo}$ . Then, we seek to merge the arrays together. Every time we pull an element from  $A$ , we add the number of elements remaining in  $A_{lo}$  to the total number of inversions as, all of those elements are greater than our current element from  $A_{hi}$  but to the left of our current element.

**Time Complexity of Divide and Conquer Method** The divide and conquer method will have the same time complexity as merge sort and thus runs in  $\Theta(n)$ .

### 3.2 The Master Theorem

**Setup Master Theorem** Let  $a \geq 1$  be an integer and  $b > 1$  be a real number,  $f(n) > 0$  be a non-decreasing function defined on the positive integers. Then,  $T(n)$  is the solution of the recurrence

$$T(n) = aT\left(\frac{n}{b}\right) + f(n).$$

Then, we define the critical exponent  $C^* = \log_b(a)$  and the critical polynomial  $n^{C^*}$ .

#### Master Theorem

1. If  $f(n) = O(n^{C^*})$  for some  $\epsilon > 0$  then,  $T(n) = \Theta(n^{C^*})$ .
2. If  $f(n) = \Theta(n^{C^*})$  then,  $T(n) = \Theta(n^{C^*}) \log n$ .
3. If  $f(n) = \Omega(n^{C^*+\epsilon})$  for some  $\epsilon > 0$  and, for some  $c < 1$ , and some  $n_0$ ,

$$af\left(\frac{n}{b}\right) \leq cf(n)$$

holds for all  $n > n_0$  then,  $T(n) = \Theta(f(n))$ .

If the conditions above do not hold then, the master theorem is not applicable.