

Algorithms and Data Structures

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Introduction

- Exam will be written, but you can do oral if you want
 - After lectures, around June 25
- An algorithm is a finite series of steps that solves a problem
 - In CS, an algorithm is a well defined computational procedure
- For a sorting algorithm, the input is a series of numbers and the output is an ordered series of numbers
- Algorithms are for humans, while a program is for a computer
- Algorithms are written in pseudocode, which follow specific conventions
- A problem can be solved by many algorithms
- An algorithm can be implemented in many different programs
- Properties of algorithms
 - Input for an algorithm can have 0 or more inputs
 - It always has 1 or more outputs
 - It should be clearly defined and unambiguous
 - It should terminate after a finite number of steps
 - All operations must be basic
 - * They can be solved exactly and in finite time
- The correctness of an algorithm is difficult to prove
 - I would need to try all possible inputs (!)
 - Published algorithms have a mathematical proof
- Incorrect algorithms can produce a wrong output or not produce any for some instances
 - In some cases they are still useful, if I can control their error rate
- Efficiency is related to the ability of an algorithm to be executed with available resources
- Resources are time and memory
- Time is measured in running time, not CPU time
 - CPU time is dependent on CPU (!)
 - CPU time is number of instructions divided by number of instructions per unit time
 - Running time is the number of primitive operations to be performed in proportion to the input size
- The algorithm influences time much more than hardware, we do not focus on hardware (!)
- Running time of n^2 is unacceptable for large inputs
- Using the right data structure is important for efficiency
- Decision trees are essential in CS
- An instance of a problem is a specific input for that problem
- In a while and for loop, the test is always executed once more than the body

Math background

- Finite sums
 - $\sum_{k=1}^n a_k = a_1 + a_2 + \dots + a_n$
- Infinite sums

- $\sum_{k=1}^{\infty} a_k = a_1 + a_2 + \dots$
- Sums are linear
 - $\sum_{k=1}^n (ca_k + b_k) = c \sum_{k=1}^n a_k + \sum_{k=1}^n b_k$
- The arithmetic series
 - $\sum_{k=1}^n k = 1 + 2 + \dots + n = \frac{n(n+1)}{2}$
- The quadratic arithmetic series
 - $\sum_{k=1}^n k^2 = 1 + 4 + \dots + n^2 = \frac{n(n+1)(2n+1)}{6}$
- The cubic arithmetic series
 - $\sum_{k=1}^n k^3 = 1 + 8 + \dots + n^3 = \frac{n^2(n+1)^2}{4}$
- The geometric series
 - $\sum_{k=1}^n x^k = 1 + x + x^2 \dots + x^n = \frac{x^{n+1}-1}{x-1}$
 - When x is less than 1
 - * $\sum_{k=1}^{\infty} x^k = \frac{1}{1-x}$
 - * $\sum_{k=1}^{\infty} kx^k = \frac{x}{(1-x)^2}$
- Other formulas
 - $\sum_{k=1}^n \log k \approx n \log n$
 - $\sum_{k=1}^n k^p \approx \frac{n^{p+1}}{p+1}$
- A set is a non-ordered and non repetitive collection of elements
- Sets can be infinite
- It is described as $S = \dots$
- 2 sets are equal if they contain the same elements
- The cardinality of a set $|S|$ is the number of elements it contains
- If A contains all the elements contained in B and also other elements, then B is a proper subset of A
- The power set $P(S)$ is the set of all subset of S , including the empty set and S itself
 - $|P(S)| = 2^{|S|}$
- The cartesian product of 2 sets is a set containing all possible pairs of elements
- The cartesian product of n sets is a set of n -tuples
- A tree has only one way to go from one node to the other
- A graph can have cycles
- A forest is made of many trees
- Any non-empty tree with n nodes has $n-1$ edges
 - If this is not true, we don't have a tree
- A tree is rooted if one of its nodes is distinguished as root
 - It can be defined recursively such that every non-root node of a rooted tree is itself the root of a subtree
- Tree terminology is similar to that of ancestry trees
- The depth of a node is its distance from the root (number of edges)
- The height of a node is the lenght of the longest path to a leaf
- A binary tree is an order tree with 2 subtrees wich are themselves binary

Pseudocode

- We start counters from 1 since it is easier to understand
- Bold words are reserved words like **return**
- Variables are always local to the current procedure
- We can have loops like **while** and **for**

for $i=0$ to/downto $i=4$ (by 3)

 <statement>

- We have if statements

if <condition>

 <statement>

- Comments are rendered with `//`
- No colon/semicolon at the end of lines
- Use of indented blocks
- Differentiate conditional expressions and assignments (!)
- A slice of an array is indicated as `A[3..5]`
- Attributes of objects are indicated as `object.attribute`
 - The length of an array can be indicated as `A.length`

Sorting

- Sorting is an intermediate step in many tasks in CS
- There are many sorting algorithms

Insertion sort

- It is like arranging card in order in your hand by picking one at a time
- I take 1 unsorted object at a time and I insert it in the correct position in the sorted array
 - I compare with all the objects in the sorted array, until I find the right position
- I start from the first element of the array and I don't do anything
- I take the second element, and if it is smaller than the first I swap them
- I take the third, and if it is smaller than the second I compare it with the first and I swap in the right position
- I continue like this for all the elements

Pseudocode

```

INSERTION-SORT(A)
  for j = 2 to A.length
    key = A[j]
    i = j-1
    while i > 0 and A[i] > key
      A[i+1] = A[i]
    A[i+1] = key

```

Running time

- Nearly sorted numbers can be sorted much faster with insertion sort
- The input size is the length of the array
 - $n = A.length$
- The initial **for** test is executed n times
 - It is n , not $n-1$ because even when it is false we still have to check ones (!)
 - The body of the **for** is executed $n-1$ times
- The assignment of `key` is therefore executed $n-1$ times
- The **while** test is executed $\sum_{j=2}^n t_j$ times
 - The body of the **while** is executed $\sum_{j=2}^n t_j - 1$
 - There are 2 assignments on the while body
- The final assignment after the **while** inside the **for** is executed $n-1$ times

Best case

- The array is already sorted
- I never enter the while, but I do completely the for
- This means that t_j is 1, I only do the test
- The time is linear

Worst case

- The array is in reverse sorted order
- The time is quadratic

Average case

- It is really difficult to do, we prefer to focus on the worst case

Evaluation

- For almost sorted sequences its running time is almost linear
- Can be online
 - It can sort sequences as they arrive
- In the worst and average cases it is quadratic
 - Quadratic is really bad (!)

Merge sort

- It is a divide and conquer algorithm
 - Divide a problem in subproblems of smaller size
 - Solve the subproblems recursively (conquer)
 - Combine the solution to solve the original problem
- The complicated part is the merging process
- It runs always as $n \cdot \log(n)$, there is not worst or best case
- It requires a lot of memory to store all the sub-arrays
- It is worse than insertion sort in the best case, but better in most cases
- It cannot work online (!)

Idea

- I want to sort the array A
- I split the array using the indices p,q,r such that $p \leq q < r$
- I want to produce a single sorted subarray
- I call initially on A with p=1 and r=A.length
- The index q is the one that best splits the array in 2
- For merging I always have sorted arrays to merge
 - The first element of each array is guaranteed to be the smallest one of the entire array
 - I compare the first element of the 2 arrays to be merged, and I put the smallest in the output array
 - I repeat until one of the arrays is empty
 - I finish by putting what remains of the other array in the output
 - I put an imaginary infinite at the end of any array
 - * This is so that when I finish the elements of an array, whatever remains in the other is smaller and so it is inserted in the output

Pseudocode

```

MERGESORT(A,p,r)
  if p < r
    q = (p+r)/2
    MERGESORT(A,p,q)
    MERGESORT(A,q+1,r)
    MERGE(A,p,q,r)

```

```

MERGE(A,p,q,r)
  n1 = q - p + 1
  n2 = r - q
  for i = 1 to n1
    L[i] = A[p+i-1]

```

```

for j = 1 to n2
    R[i] = A[q+j]
L[n1+1] = \infty
R[n2+1] = \infty
i = 1
j = 1
for k = p to r
    if L[i] <= R[j]
        A[k] = L[i]
        i += 1
    else A[k] = R[j]
        j += 1

```

MERGESORT(A,1,A.lenght)

Running time

- MERGE
 - Copying the elements into the subarrays takes $\Theta(n)$
 - Adding elements to the final array takes n iterations of that themselves take constant time
* $\Theta(n)$
 - In total, the merging takes $\Theta(n)$
- MERGE-SORT
 - Let $T(n)$ be the unknown running time of MERGE-SORT
 - Calculating q: $\Theta(1)$
 - Solve recursively 2 subproblems of size $n/2$: $2T(n/2)$
 - Call to MERGE: $\Theta(n)$
 - So, $T(n) = \begin{cases} \Theta(1) & \text{if } n = 1 \\ 2T(n/2) + \Theta(1) + \Theta(n) & \text{if } n > 1 \end{cases}$
 - The recursive equation can be solved and we find that $T(n) = \Theta(n \log n)$

Limiting behaviour of functions

- There are different notations to define the behaviour of functions
- The Θ (theta) notation signifies asymptotic equality
 - Formally, for a function $f(n)$ having a certain Θ notation there are 2 constant that multiplied for the Θ function are constantly greater or smaller than $f(n)$ for $n > n_0$
* This is defined as a tight bound
 - If I say that $f(n) = \Theta(g(n))$ I mean that $f(n)$ belongs to the family of functions with order of growth $g(n)$
- The O (big-O) notation indicates an upper bound for the asymptotic behaviour
 - The formal definition is similar to that of Θ , but instead of a tight bound I only search for an upper bound
* I only want a constant, not 2 (!)
- The Ω (big-Omega) notation indicates a lower bound for the function
- An important theorem: $f(n) = \Theta(g(n)) \iff f(n) = O(g(n)) \wedge f(n) = \Omega(g(n))$
- Factorials are faster than exponentials, but slower than n^n (!)

Designing algorithms

- A recurrence equation describes a function in terms of its value on a smaller input
- An example: analysis of a divide and conquer algorithm

- $T(n)$ is the running time of the algorithm on input n
- Dividing takes $D(n) = \Theta(1)$ time
- Conquer takes the same T on a smaller input, so $aT(n/b)$
 - * I need to solve a subproblems in with an input size reduced of a factor b
- Combining the solutions takes $C(n) = \Theta(n)$ time
- So we have that $T(n) = \begin{cases} c & n = 1 \\ 2T(n/2) + c + cn & n > 1 \end{cases}$
- Solving recurrence equations: the iteration method
 - If I have $T(n) = T(n/2) + c \implies T(n/2) = T(n/4) + c$ and so on
 - This implies that $T(n) = c + c + T(n/4)$
 - If I continue this until I get to the base case $T(1)$
 - I can write therefore $T(n) = c * k + T(n/2^k)$
 - The base case will be when $n = 2^k$ and therefore $k = \log n$
 - So I get that $T(n) = c * \log n + T(n/2^{\log n}) = c * \log n + T(1)$
 - This means that $T(n) = \Theta(\log n)$
- Solving recurrence equations: the recursion tree method
 - I convert the equation into a tree and sum up the cost of each node
 - Let's try with $T(n) = 2T(n/2) + n^2$
 - The root has a cost n^2 and 2 children of cost $T(n/2)$ each
 - I continue to expand until the base case
 - This gives us a $O(n^2)$, since I cannot determine a tight bound

Heapsort

- Normally, the numbers to be sorted are a key that is paired to other data, forming a record
 - A record is composed of a key and satellite data
- When I want to sort, are all keys unique?
- Its running time is $\Theta n \log n$
- It is based on a data structure called heap
 - An heap is a nearly complete binary tree
 - * All nodes are binary except for possibly the last level
 - * If the last level is not full, it is filled from left to right
 - It follows the heap property: the value of a parent must be greter that that of a children
 - * This is a max heap, there are also min heaps where the property is opposite
 - The size of an heap is the number of nodes
- We can represent an heap with an array
 - The first element is the root
 - The children of the root are the second and third element
 - The fourth and fifth element are the children of the second, and so on
 - The children of node $A[i]$ are nodes $A[2i]$ and $A[2i+1]$
 - The parent of $A[i]$ is $A[\lfloor i/2 \rfloor]$
 - The maximum element is always the root
- How to maintain the max heap property (MAX-HEAPIFY)
 - I recursively explore the tree
 - If I find a parent smaller than its child, I swap them and continue
 - I assume that there is only one violation
 - The running time is $O(\log n)$, or linear to heap size ($O(h)$)

MAX-HEAPIFY(A, i)

```

1, r = 2i, 2i+1
if 1 <= A.heap-size and A[1]>A[i]
    largest = 1
else largest = i

```

```

if r <= A.heap-size and A[r]>A[i]
    largest = r
else largest = i
if largest != i
    exchange A[i] and A[largest]
    MAX-HEAPIFY(A,largest)

```

- Now we start from a random array and we want to make it a max heap
 - Note that $A[(\lfloor n/2 \rfloor + 1) \dots n]$ are leaves

BUIL-MAX-HEAP(A)

```

A.heap-size = A.lenght
for i = floor(A.lenght/2) downto 1
    MAX-HEAPIFY(A,i)

```

- This operations has an loose upper boundf of $O(n \log n)$
 - I do $n/2$ times a $O(\log n)$ operation
- However, the argument to MAX-HEAPIFY is almost never n (!)
 - In the first step it is 1, then 2 and so on
- The worst case running time of MAX-HEAPIFY is $O(\log i)$ where i is the value of the for loop in BUILD-MAX-HEAP
 - We obtain $O(n)$
- The next step is to actually sort the array
 - We swap the root with the last element and decrease the heap-size by 1
 - We call MAX-HEAPIFY on the root to rebuild the max-heap property
 - We repeat until the heap-size is 1
 - The array is sorted (!)

HEAPSORT(A)

```

BUILD-MAX-HEAP(A)
for i = A.lenght down to 1
    exchange A[1] with A[-1]
    A.heap-size = A.heap-size - 1
    MAX-HEAPIFY(A,1)

```

- The total running time is $O(n \log n)$