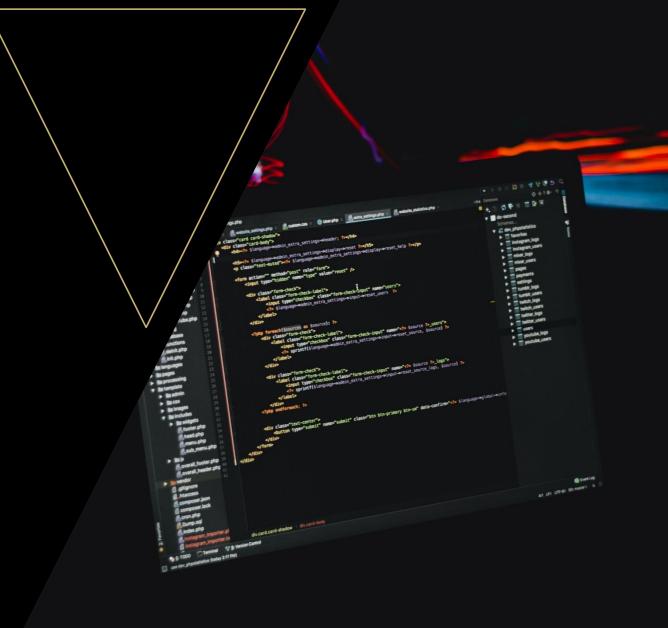
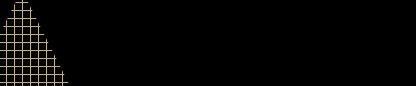


CSE 2003

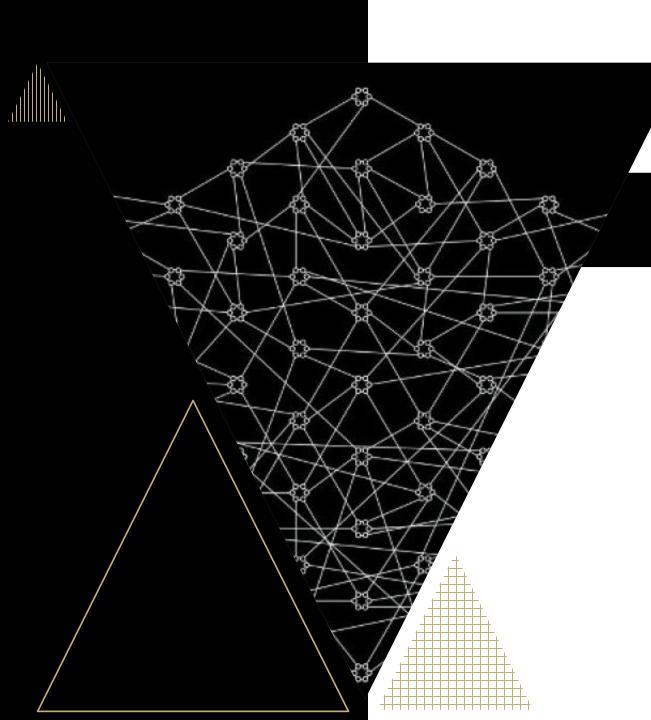
DSA ETh Digital Assignment - 1

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Group 4

Running time, Time-complexity, & Performance analysis of an algorithm



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Running Time



The running time of an algorithm for a specific input depends on the number of operations executed.

The greater the number of operations, the longer the running time of an algorithm.

It could take nanoseconds, or it could go on forever. It also depends on the input.

```
For example: function addEm(a, b) {
    var c = a + b;
    return c;
    };
```

This algorithm takes two arguments, adds them together, then returns their sum. This wouldn't take long at all. Fractions of a second. However, check this one:

```
function endless() {
     while(2 === 2) {
     console.log("aaaaaaaaah");
     };
```

This is a "while true" algorithm. It takes a condition that will always return true (2 will always equal 2) and does something over and over again ad infintium. In this case, it will continuously print "aaaaaaaah" to the console until you force it to end or your computer catches fire.



Time Complexity

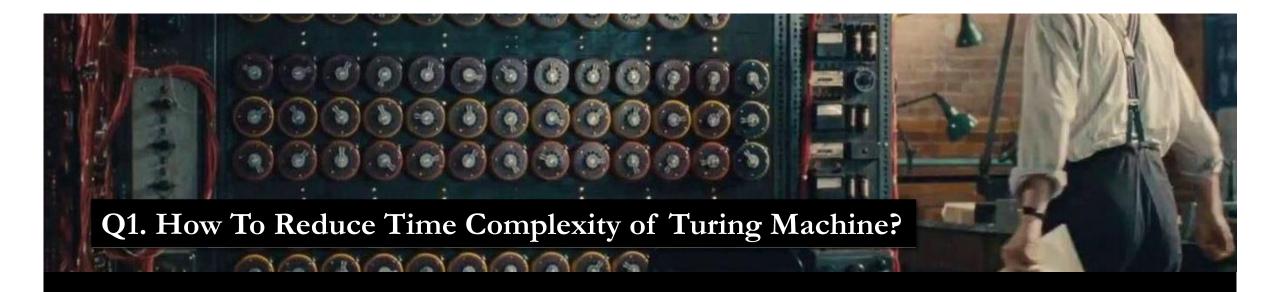
Big O Notation	Name	Example(s)
O(1)	Constant	#Odd or Even number, #Look-up table (on average)
O(log n)	Logarithmic	#Finding element on sorted array with binary search
O(n)	Linear	#Find max element in un sorted array #Duplicate elements in array with Hash Map
O(n log n)	Linearithmic	#Sorting elements in array with merge sort
$O(n^2)$	Quadratic	#Duplicate elements in array **(naïve)** #Sorting array with bubble sort
$O(n^3)$	Cubic	#3 variables equation solver
O(2 ⁿ)	Exponential	#Find all subsets
O(n!)	Factorial	#Find all permutations of a given set/string

Time complexity estimates how an algorithm performs regardless of the kind of machine it runs on. It can be determined by "counting" the number of operations performed by the code.

This is defined as a function of the input size n using Big-O notation. n indicates the size of the input, while O is the worst-case scenario growth rate function.

We use the Big-O notation to classify algorithms based on their running time or space (memory used) as the input grows. The O function is the growth rate in function of the input size n.

The notation is useful when comparing algorithms used for similar purposes.

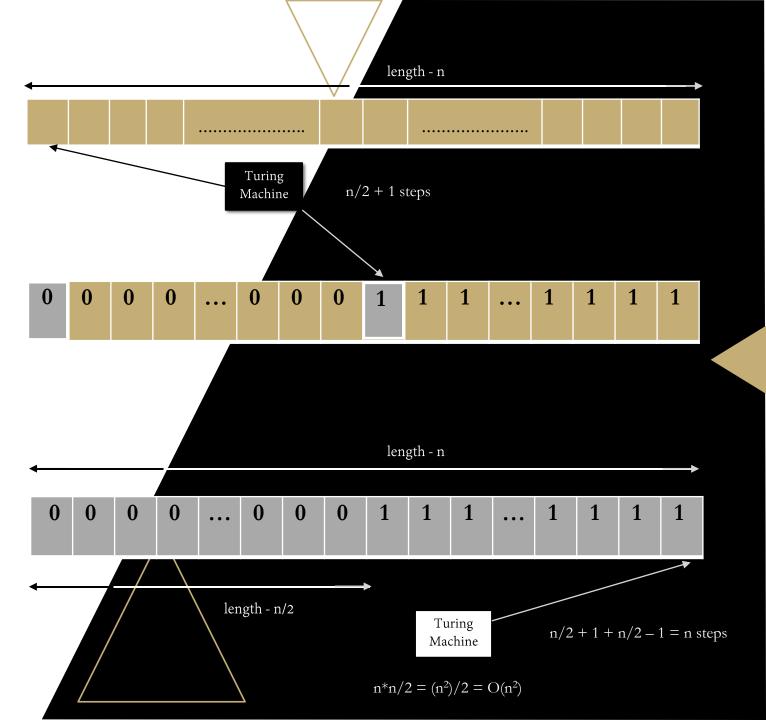


- During WW2, a brilliant mathematician named Alan Turing joined the British Military to crack the German Enigma code. He went on to become one of the founding fathers of Computer Science by inventing the TM.
- So the Turing Machine basically is used for the verification whether a string belongs to a particular language or not.

i. First, we check for the corner cases that every 0 in the string must be followed by A 1. Else we reject the string

ii. Now, in the 1st algorithm we check every zero and corresponding to that we check A 1. This takes n/2 + n/2 = n steps

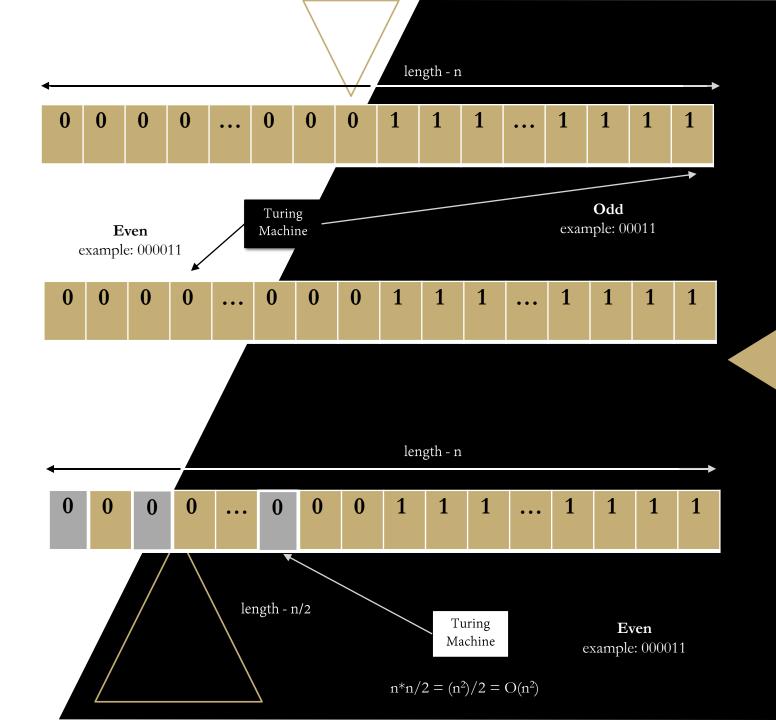
iii. So, each scan takes n steps and we repeat it for n/2 times, thereby achieving a total of n^2 time complexity in this way; represented by $O(n^2)$.



[Better Optimal Strategy]

iv. Now, here also at first we check the corner cases. Then we count the total number of O's and 1's. If it's odd, we reject it and if it's even, we accept it.

v. After this we'll cross every other 0 that we see in the input string. Same operation is performed on 1's as well.

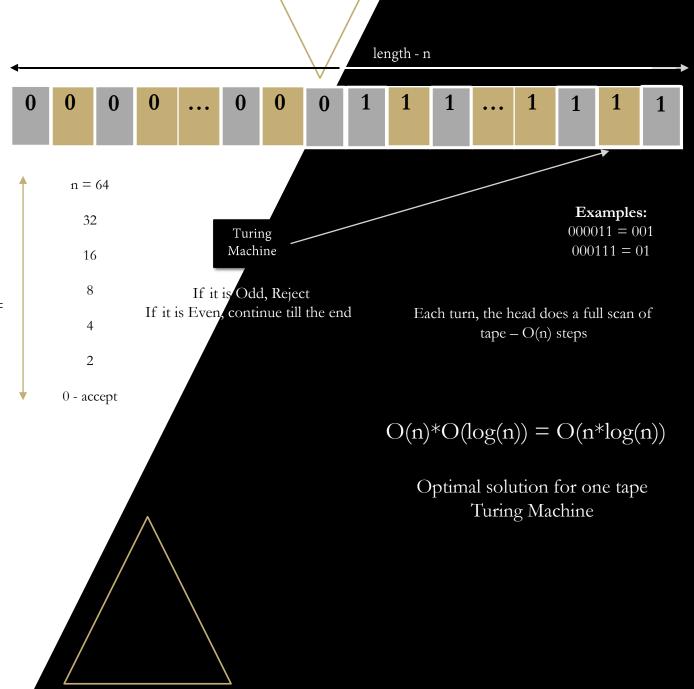


vi. After performing the above operation, we'll count the remaining 0's and 1's. Now if that sum is zero, we'll accept it and if it's odd, we reject it. And the same operation is performed until we are left with nothing.

$$log2(n) + 1 =$$
 $log2(64) + 1 =$
 7

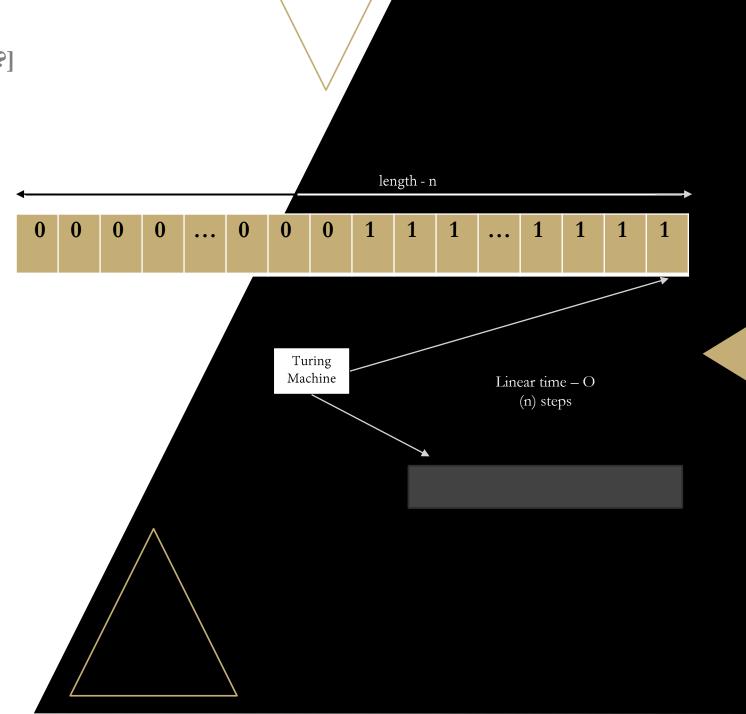
vii. So in this way, in each repetition we'll deal with half of the actual number of bits.

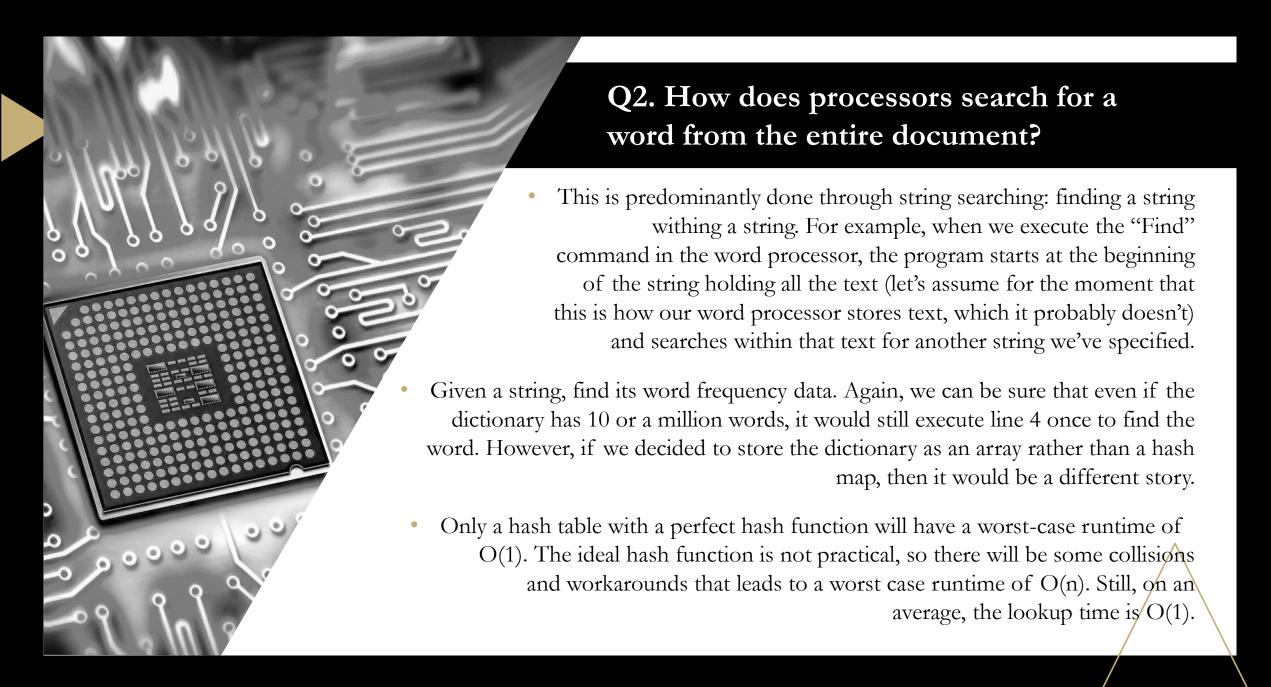
viii. So the number of times we need to repeat this log2(n) and in each turn the head does a full scan of tape that is O(n) steps. So the resultant reduced time complexity is obtained as O(n*log2(n)).



[What if we use a Two-Tape Turing Machine?]

ix. So in yet another algorithm if we employ a stack and pop in all zeroes and pop out them on occurrence of A 1 one by one time complexity in this case will be O(n)





Rabin-Karp example:

LLLLLLLM

Pattern: LLLLLM

Let Hash Value of "LLLLLL" = 0;

And Hash Value of "LLLLLM" = 1;

[How can we optimise and do better?]

Using Rabin-Karp String Search

- Michael O. Rabin, a professor at Harvard, along with his colleague Richard Karp devised a method from using hashing to do string search in O(m+n), as opposed to O(mn). In other words, in linear time as opposed to quadratic time, a nice speedup.
- The Rabin-Karp algorithm uses a technique called fingerprinting.

LLLLLM ---- 0! = 1 (One Comparison)

LLLLLM ----- 0! = 1 (One Comparison)

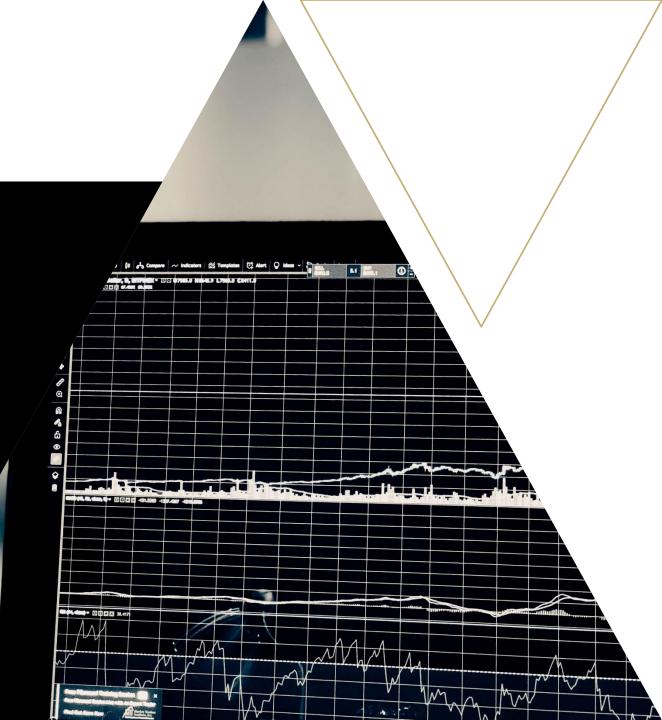
...

LLLLLM ----- 1 = 1

Hash Value is matching so compare elements one by one. (6+1 Comparisons)

Performance Analysis

Let's take an example. If we want to go from city A to city B, there can be many ways of doing this. We can go by flight, using a bus, a train and also by bicycle. Depending on the availability and convenience, we choose the one which suits us. Similarly, in computer science, there are multiple algorithms to solve a problem. When we have more than one algorithm, we need to select the most efficient one. Performance analysis helps us in this regard.

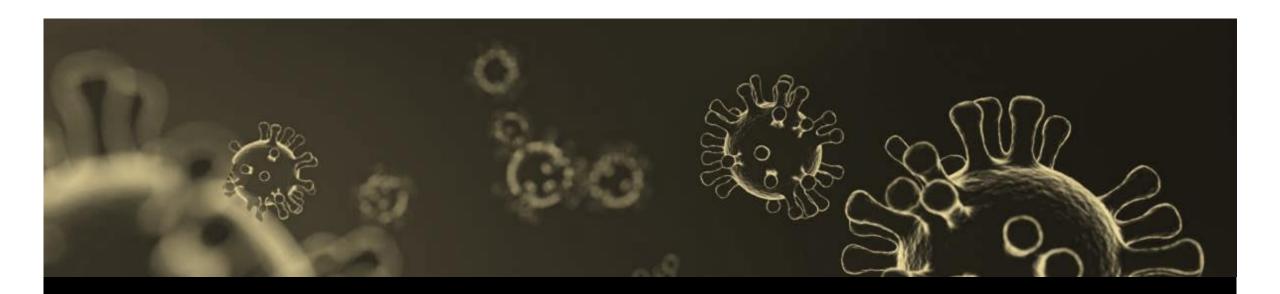




• The present outbreak of coronavirus acute respiratory disease called covid-19 has resulted in a major epidemic. The main reason why coronavirus is a major problem is because its spread can be modelled by a tree.

Before the world took lockdown measures, estimates stated that each infected person was infecting between two to four other people. This number is called R₀, a mathematical denotation that indicates how contagious an infectious disease is. For instance, if a disease has an R₀ of 15, a person who has the disease will transmit it to an average of 15 other people.



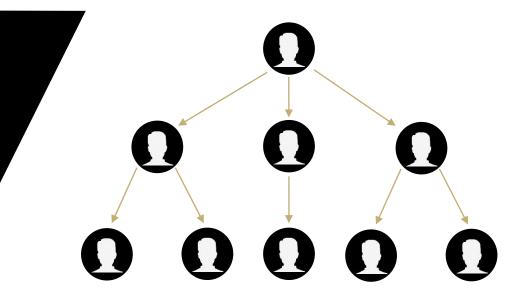


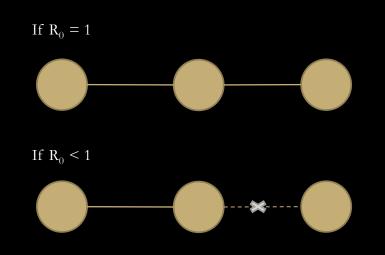
Importantly, a disease's R_0 value only applies when everyone in a population is completely vulnerable to the disease, as in the case of covid-19 where no one has been vaccinated, on one has had the disease before and there's no way to control the spread. In our model, R_0 is the average number of children each node in the tree has. This means, each model in our tree has (on average) between two and four children.

A total of 3 possibilities exist for the transmission or decline of a disease, depending on its R_0 value:

Cases	Values of R ₀	Condition of disease
1	$R_0 < 1$	Will eventually die out
2	$R_0 = 1$	Disease will stay alive and stable but there won't be any outbreak
3	$R_0 > 1$	May be an epidemic or outbreak

As this is a tree structure, by analysing it we know that this is going to get very large very quickly. The early objective of health organizations worldwide was to reduce R_0 to around one (or less). If $R_0 = 1$, then each leaf node in our tree now becomes the head of a linked list. Each person is infecting exactly one other person, in the same way, that a (singly) linked list has a reference to the next node in the list.





If R_0 < 1, then at some point a person will infect no-one else, and the line of infection (for the leaf) is broken. We can model that in code by having the node point to a null reference, meaning it is the ultimate node in the linked list.

One way to "solve" the coronavirus situation is to change the behaviour of the virus so that it can be modelled by a collection of (eventually finite) linked lists, rather than a tree. And hence, by analysing the structure of the virus we can potentially stop its spread.

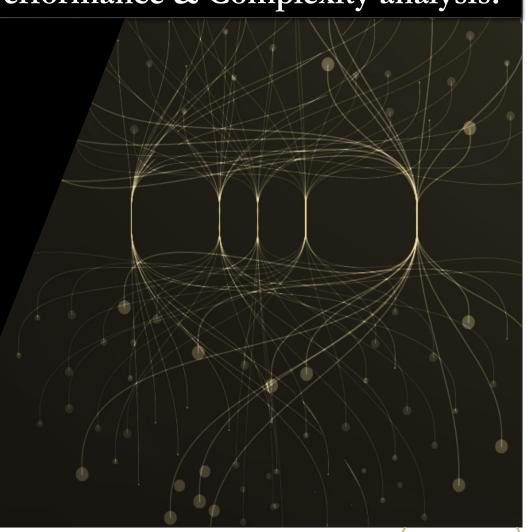


It is sometimes best to write algorithms from first principles rather than to modify a large computer program already written. The way to do this would be to go back to the original problem. Develop a paradigm for analysis. It may be along a similar line to what has been done before, or it may be a totally new approach. However, to make a good comparison, we should understand the existing algorithm thoroughly. That is one reason why performance analysis of existing algorithms is important. To understand the existing algorithm, we may need to analyse it piece by piece. It should be kept simple. If we've more than one degree of freedom, we cannot know what variable caused a certain result. So break the algorithm into components with one degree of freedom, and then see what each does. Performing the analysis of the algorithm, it is possible to improve on it and develop an efficient method, thereby creating our own version.

Q5. What is the real-life difference between Performance & Complexity analysis?

The only way to know for sure what solution is best for a problem is to measure it. However, measuring performance is actually quite a hard thing to do; there are many things that go into making sure we measure what we want to measure. Outside noise introduced by the OS, other processors, random lags, compiler optimizations, etc. It's impossible to know all the factors that influence the performance of your algorithms, after all even a random burst of electromagnetic radiation from the sun might fry your computer and your algorithm will never finish performing its task. This is where performance measurements are so much different from complexity analysis.

Obviously no one is expected to predict solar flares, or other random events that we have no control over. But when talking about performance the factors that the developer can consistently account for are very important to ensure the maximum performance of our software. One such very important concern is hardware constraints. When considering which algorithm and data structure to choose, from a performance analysis standpoint it is very important to take into account things like on what machine will this code run in and optimize it for the environment it will be used in.



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