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A T M E

College of Engineering

DEPARTMENT OF COMPUTER SCIENCE AND DESIGN



LECTURE NOTES

COURSE: ANALYSIS AND DESIGN OF ALGORITHMS

COURSE CODE: BCS401

SEMESTER: IV

COURSE COORDINATOR: Dr. Nasreen Fathima

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INSTITUTIONAL MISSION AND VISION

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- To develop academic, professional and financial alliances with the industry as well as the academia at national and transnational levels.
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Program Specific Outcomes (PSOs)

PSO1: Ability to apply skills in the field of algorithms, database design, web design, cloud computing and data analytics.

PSO2: Apply knowledge in the field of computer networks for building network and internet based applications.

MODULE I

INTRODUCTION

Contents

1. Introduction

1. 1. What is an Algorithm?
1. 2. Algorithm Specification
1. 3. Analysis Framework

2. Performance Analysis

2. 1. Space complexity
2. 2. Time complexity

3. Asymptotic Notations

3. 1. Big-Oh notation
3. 2. Omega notation
3. 3. Theta notation
3. 4. Little-oh notation

4. Mathematical analysis of Non-Recursive Algorithms

5. Mathematical analysis of Recursive Algorithms

1. INTRODUCTION

1.1 What is an Algorithm?

Informal Definition:

An Algorithm is any well-defined computational procedure that takes some value or set of values as input and produces a set of values or some value as output. Thus algorithm is a sequence of computational steps that transforms the input into the output.

Formal Definition:

*An **algorithm** is a sequence of unambiguous instructions for solving a problem, i.e., for obtaining a required output for any legitimate input in a finite amount of time.*

This definition can be illustrated by a simple diagram:

- *The reference to —instructions in the definition implies that there is something or someone capable of understanding and following the instructions given.*
- *Human beings and the computers nowadays are electronic devices being involved in performing numeric calculations.*
- *However, that although the majority of algorithms are indeed intended for eventual computer implementation, the notion of algorithm does not depend on such an assumption.*

In addition, all algorithms should satisfy the following criteria or properties.

1. **INPUT** → *Zero or more quantities are externally supplied.*
2. **OUTPUT** → *At least one quantity is produced.*
3. **DEFINITENESS** → *Each instruction is clear and unambiguous.*
4. **FINITENESS** → *If we trace out the instructions of an algorithm, then for all cases, the algorithm terminates after a finite number of steps.*
5. **EFFECTIVENESS** → *Every instruction must very basic so that it can be carried out, in principle, by a person using only pencil & paper.*

Fundamentals Of Algorithmic Problem Solving

- *Algorithms are procedural solutions to problems. These solutions are not answers but specific instructions for getting answers.*
- *he following diagram briefly illustrates the sequence of steps one typically goes through in designing and analyzing an algorithm.*
- *It includes the following sequence of steps:*
 - ✓ *Understanding the problem*
 - ✓ *Deciding on: Computational means, Exact vs. approximate problem solving, data structure(s), Algorithm design techniques.*
 - ✓ *Design an algorithm*
 - ✓ *Prove correctness Analyze the algorithm*
- **Understanding the problem**
 - ✓ *Before designing an algorithm the most important thing is to understand the problem given.*
 - ✓ *Asking questions, doing a few examples by hand, thinking about special cases, etc.*
 - ✓ *An Input to an algorithm specifies an instance of the problem the algorithm that it solves.*
 - ✓ *Important to specify exactly the range of instances the algorithm needs to handle. Else it*

will work correctly for majority of inputs but crash on some —boundaryll value.
 - ✓ *A correct algorithm is not one that works most of the time, but one that works correctly for all legitimate inputs.*
- **Ascertaining the capabilities of a computational device**
 - ✓ *After understanding need to ascertain the capabilities of the device.*
 - ✓ *The vast majority of algorithms in use today are still destined to be programmed for a computer closely resembling the von Neumann machine—a computer architecture.*
 - ✓ *Von Neumann architectures are sequential and the algorithms implemented on*

- ✓ *they are called sequential algorithms.*
- ✓ *Algorithms which designed to be executed on parallel computers called parallel algorithms.*
- ✓ *For very complex algorithms concentrate on a machine with high speed and more memory where time is critical.*
- **Choosing between exact and approximate problem solving**
 - ✓ *For exact result->exact algorithm*
 - ✓ *For approximate result->approximation algorithm.*
 - ✓ *Examples of exact algorithms: Obtaining square roots for numbers and solving non-linear equations.*
 - ✓ *An approximation algorithm can be a part of a more sophisticated algorithm that solves a problem exactly.*
- **Deciding on appropriate data structures**
 - ✓ *Algorithms may or may not demand ingenuity in representing their inputs.*
 - ✓ *Inputs are represented using various data structures.*
 - ✓ *Algorithm + data structures=program.*
- **Algorithm Design Techniques and Methods of Specifying an Algorithm**
 - ✓ *An **algorithm design technique** (or —strategy|| or —paradigm||) is a general approach to solving problems algorithmically that is applicable to a variety of problems from different areas of computing.*
 - ✓ *The different algorithm design techniques are: brute force approach, divide and conquer, greedy method, decrease and conquer, dynamic programming, transform and conquer and back tracking.*
 - ✓ *Methods of specifying an algorithm: Using natural language, in this method ambiguity problem will be there.*
 - ✓ *The next 2 options are : pseudo code and flowchart.*
 - ✓ *Pseudo code: mix of natural and programming language. More precise than NL*
 - ✓ *Flow chart: method of expressing an algorithm by collection of connected geometric shapes containing descriptions of the algorithm's steps.*
 - ✓ *This representation technique has proved to be inconvenient for large problems.*

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- ✓

The state of the art of computing has not yet reached a point where an algorithm's description—be it in a natural language or pseudo code—can be fed into an electronic computer directly. Instead, it needs to be converted into a computer program written in a particular computer language.
- ✓

Hence program as yet another way of specifying the algorithm, although it is preferable to consider it as the algorithm's implementation.
- **Proving an Algorithm's Correctness**
 - ✓

After specifying an algorithm we have to prove its correctness.
 - ✓

The correctness is to prove that the algorithm yields a required result for every legitimate input in a finite amount of time.
 - ✓

For example, the correctness of Euclid's algorithm for computing the greatest common divisor stems from the correctness of the equality $\text{gcd}(m, n) = \text{gcd}(n, m \bmod n)$, the simple observation that the second integer gets smaller on every iteration of the algorithm, and the fact that the algorithm stops when the second integer becomes 0.
 - ✓

For some algorithms, a proof of correctness is quite easy; for others, it can be quite complex.
 - ✓

A common technique for proving correctness is to use mathematical induction.
 - ✓

Proof by mathematical induction is most appropriate for proving the correctness of an algorithm.
- **Analyzing an algorithm**
 - ✓

After correctness, efficiency has to be estimated.

 - Time efficiency and space efficiency.
 - Time: how fast the algorithm runs?
 - Space: how much extra memory the algorithm needs?
 - ✓

Simplicity: how simpler it is compared to existing algorithms. Sometimes simpler algorithms are also more efficient than more complicated alternatives. Unfortunately, it is not always true, in which case a judicious compromise needs to be made.
 - ✓

Generality: Generality of the problem the algorithm solves and the set of inputs it accepts.

- ✓
If not satisfied with these three properties it is necessary to redesign the algorithm.
- **Code the algorithm**
 - ✓
Writing program by using programming language.
 - ✓
Selection of programming language should support the features mentioned in the design phase.
 - ✓
Program testing: If the inputs to algorithms belong to the specified sets then require no verification. But while implementing algorithms as programs to be used in actual applications, it is required to provide such verifications.
 - ✓
Documentation of the algorithm is also important.

1.2 Algorithm Specification

Algorithm can be described in three ways.

1. *Natural language like English: When this way is choosed care should be taken, we should ensure that each & every statement is definite.*
2. *Graphic representation called flowchart: This method will work well when the algorithm is small& simple.*
3. *Pseudo-code Method: In this method, we should typically describe algorithms as program, which resembles programming language constructs*

Pseudo-Code Conventions:

1. *Comments begin with // and continue until the end of line.*
2. *Bocks are indicated with matching braces {and}.*

An identifier begins with a letter. The data types of variables are not explicitly declared.

4. *Compound data types can be formed with records. Here is an example,*

data type – n data – n; node * link;

Node. Record { data type – 1 data-1; . . . }

Here link is a pointer to the record type node. Individual data items of a record can be accessed with → and period.

5. *Assignment of values to variables is done using the assignment statement.*

<Variable>:= <expression>;

6. *There are two Boolean values TRUE and FALSE.*

✓

Logical Operators AND, OR, NOT

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✓

Relational Operators <, <=, >, >=, =, !=

7. The following looping statements are employed. For, while and repeat-until

While Loop:

While < condition > do

{ <statement-1> ... <statement-n> }

For Loop:

For variable: = value-1 to value-2 step step do

{ <statement-1> ... <statement-n> }

repeat-until:

repeat <statement-1> ... <statement-n> until<condition>

8. A conditional statement has the following forms.

✓

If <condition> then <statement>

✓

If <condition> then <statement-1> Else <statement-1>

✓

Case

statement:

Case

{ : <condition-1> : <statement-1>

...

: <condition-n> : <statement-n>

: else : <statement-n+1>

}

9. Input and output are done using the instructions read & write.

10. There is only one type of procedure:

✓

Algorithm, the heading takes the form, Algorithm Name (Parameter lists)

✓

As an example, the following algorithm finds & returns the maximum of „n“ given numbers:

Algorithm Max(A,n)

/ A is an array of size n

{

Result := A[1];

```
    for i:= 2 to n
    do
        if A[i] > Result then
            Result :=A[i];
        return Result;
    }
```

In this algorithm (named Max), A & n are procedure parameters. Result & i are Local variables.

1.3 Analysis Framework

There are two kinds of efficiency:

- ◆ **Time efficiency** - indicates how fast an algorithm in question runs.
- ◆ **Space efficiency** - deals with the extra space the algorithm requires.

✓

Measuring An Input Size

- ◆ An algorithm's efficiency is investigated as a function of some parameter n indicating the algorithm's input size.
- ◆ In most cases, selecting such a parameter is quite straightforward.
- ◆ For example, it will be the size of the list for problems of sorting, searching, finding the list's smallest element, and most other problems dealing with lists.
- ◆ For the problem of evaluating a polynomial $p(x)$ of degree n , it will be the polynomial's degree or the number of its coefficients, which is larger by one than its degree.
- ◆ There are situations, of course, where the choice of a parameter indicating an input size does matter.

- **Example** - computing the product of two n -by- n matrices. There are two natural measures of size for this problem.
- The matrix order n .
- The total number of elements N in the matrices being multiplied.

◆ Since there is a simple formula relating these two measures, we can easily switch from one to the other, but the answer about an algorithm's efficiency will be qualitatively different depending on which of the two measures we use.

◆ The choice of an appropriate size metric can be influenced by operations of the algorithm in question. For example, how should we measure an input's size for a spell-

DAA module-1

checking algorithm? If the algorithm examines individual characters of its input, then we should measure the size by the number of characters; if it works by processing words, we should count their number in the input.

◆ We should make a special note about measuring size of inputs for algorithms involving properties of numbers (e.g., checking whether a given integer n is prime).

✓

Units For Measuring Run Time

◆ We can simply use some standard unit of time measurement—a second, a millisecond, and so on—to measure the running time of a program implementing the algorithm.

◆ There are obvious drawbacks to such an approach. They are

- ◆ Dependence on the speed of a particular computer
- ◆ Dependence on the quality of a program implementing the algorithm
- ◆ The compiler used in generating the machine code
- ◆ The difficulty of clocking the actual running time of the program.

◆ Since we are in need to measure algorithm efficiency, we should have a metric that does not depend on these extraneous factors.

◆ One possible approach is to count the number of times each of the algorithm's operations is executed. This approach is both difficult and unnecessary.

◆ The main objective is to identify the most important operation of the algorithm, called the

basic operation, the operation contributing the most to the total running time, and compute the number of times the basic operation is executed.

◆ As a rule, it is not difficult to identify the basic operation of an algorithm.

For e.g., The basic operation is usually the most time-consuming operation in the algorithm's innermost loop

◆ Consider the following example:

Let c_{op} be the execution time of an algorithm's basic operation on a particular computer, and let $C(n)$ be the number of times this operation needs to be executed for this algorithm. Then we can estimate the running time $T(n)$ of a program implementing this algorithm on that computer by the formula

$$T(n) \approx c_{op}C(n).$$

Total number of steps for basic operation execution, $C(n) = n$



Orders Of Growth

- ◆ A difference in running times on small inputs is not what really distinguishes efficient algorithms from inefficient ones.
- ◆ When we have to compute, for example, the greatest common divisor of two small numbers, it is not immediately clear how much more efficient Euclid's algorithm is compared to the other two algorithms discussed in previous section or even why we should care which of them is faster and by how much. It is only when we have to find the greatest common divisor of two large numbers that the difference in algorithm efficiencies becomes both clear and important.
- ◆ For large values of n , it is the function's order of growth that counts: look at Table which contains values of a few functions particularly important for analysis of algorithms.

Table: Values (some approximate) of several functions important for analysis of algorithms

n	$\log_2 n$	n	$n \log_2 n$	n^2	n^3	2^n	$n!$
10	3.3	10^1	$3.3 \cdot 10^1$	10^2	10^3	10^3	$3.6 \cdot 10^6$
10^2	6.6	10^2	$6.6 \cdot 10^2$	10^4	10^6	$1.3 \cdot 10^{30}$	$9.3 \cdot 10^{157}$
10^3	10	10^3	$1.0 \cdot 10^4$	10^6	10^9		
10^4	13	10^4	$1.3 \cdot 10^5$	10^8	10^{12}		
10^5	17	10^5	$1.7 \cdot 10^6$	10^{10}	10^{15}		
10^6	20	10^6	$2.0 \cdot 10^7$	10^{12}	10^{18}		



Worst-Case, Best-Case, Average Case Efficiencies

- ◆ Algorithm efficiency depends on the **input size n** . And for some algorithms efficiency not only on input size but also on the **specifics of particular or type of input**.
- ◆ We have best, worst & average case efficiencies.
Worst-case efficiency: Efficiency (number of times the basic operation will be executed) for the worst case input of size n . i.e. The algorithm runs the longest among all possible inputs of size n .
Best-case efficiency: Efficiency (number of times the basic operation will be executed)

for the best case input of size n . i.e. The algorithm runs the fastest among all possible inputs of size n . **Average-case efficiency:** Average time taken (number of times the basic operation will be executed) **to solve all the possible instances (random) of the input.**

NOTE: It is not the average of worst and best case.

◆ **Example: Sequential search.** This is a straightforward algorithm that searches for a given item (some search key K) in a list of n elements by checking successive elements of the list until either a match with the search key is found or the list is exhausted.

◆ Here is the algorithm's pseudo code, in which, for simplicity, a list is implemented as an array. (It also assumes that the second condition will not be checked if the first one, which checks that the array's index does not exceed its upper bound, fails.)

ALGORITHM *SequentialSearch*($A[0..n - 1], K$)

```
//Searches for a given value in a given array by sequential search
//Input: An array  $A[0..n - 1]$  and a search key  $K$ 
//Output: The index of the first element in  $A$  that matches  $K$ 
//         or  $-1$  if there are no matching elements
 $i \leftarrow 0$ 
while  $i < n$  and  $A[i] \neq K$  do
     $i \leftarrow i + 1$ 
if  $i < n$  return  $i$ 
else return  $-1$ 
```

Clearly, the running time of this algorithm can be quite different for the same list size n . **Worst case efficiency**

◆ The worst-case efficiency of an algorithm is its efficiency for the worst-case input of size n , which is an input (or inputs) of size n for which the algorithm runs the longest among all possible inputs of that size.

◆ In the worst case the input might be in such a way that there are no matching elements or the first matching element happens to be the last one on the list, in this case the algorithm makes the largest number of key comparisons among all possible inputs of size n :

$$C_{\text{worst}}(n) = n.$$

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- ◆ The way to determine is quite straightforward
- ◆ To analyze the algorithm to see what kind of inputs yield the largest value of the basic

operation's count $C(n)$ among all possible inputs of size n and then compute this worst-case value **$C_{\text{worst}}(n)$** .

- ◆ The worst-case analysis provides very important information about an algorithm's efficiency

by bounding its running time from above. In other words, it guarantees that for any instance of size n , the running time will not exceed **$C_{\text{worst}}(n)$** its running time on the worst-case inputs.

Best case Efficiency

- ◆ The best-case efficiency of an algorithm is its efficiency for the best-case input of size n , which is an input (or inputs) of size n for which the algorithm runs the fastest among all possible inputs of that size.
- ◆ We can analyze the best case efficiency as follows.

First, determine the kind of inputs for which the count $C(n)$ will be the smallest among all possible inputs of size n . (Note that the best case does not mean the smallest input; it means the input of size n for which the algorithm runs the fastest.)

- ◆ Then ascertain the value of $C(n)$ on these most convenient inputs.
- ◆ Example- for sequential search, best-case inputs will be lists of size n with their first element equal to a search key; accordingly, $C_{\text{best}}(n) = 1$.
- ◆ The analysis of the best-case efficiency is not nearly as important as that of the worst-case efficiency.

- ◆ But it is not completely useless. For example, there is a sorting algorithm (insertion sort) for

which the best-case inputs are already sorted arrays on which the algorithm works very fast.

- ◆ Thus, such an algorithm might well be the method of choice for applications dealing with

almost sorted arrays. And, of course, if the best-case efficiency of an algorithm is unsatisfactory, we can immediately discard it without further analysis.

Average case efficiency

- ◆ It yields the information about an algorithm about an algorithm's behaviour on a

DAA module-1

—typical and random input.

◆ To analyze the algorithm's average-case efficiency, we must make some assumptions about possible inputs of size n .

◆ The investigation of the average case efficiency is considerably more difficult than investigation of the worst case and best case efficiency.

◆ It involves dividing all instances of size n into several classes so that for each instance of the class the number of times the algorithm's basic operation is executed is the same.

◆ Then a probability distribution of inputs needs to be obtained or assumed so that the expected value of the basic operation's count can then be derived. The average number of key comparisons $C_{avg}(n)$ can be computed as follows,

◆ Let us consider again sequential search. The standard assumptions are, ◆ Let p be the probability of successful search.

◆ In the case of a successful search, the probability of the first match occurring in the i^{th} position of the list is p/n for every i , where $1 \leq i \leq n$ and the number of comparisons made by the algorithm in such a situation is obviously i .

◆ In the case of an unsuccessful search, the number of comparisons is n with the probability of such a search being $(1 - p)$. Therefore,

$$\begin{aligned} C_{avg}(n) &= \left[1 \cdot \frac{p}{n} + 2 \cdot \frac{p}{n} + \dots + i \cdot \frac{p}{n} + \dots + n \cdot \frac{p}{n} \right] + n \cdot (1 - p) \\ &= \frac{p}{n} [1 + 2 + \dots + i + \dots + n] + n(1 - p) \\ &= \frac{p}{n} \frac{n(n+1)}{2} + n(1 - p) = \frac{p(n+1)}{2} + n(1 - p). \end{aligned}$$

Example, if $p = 1$ (i.e., the search must be successful), the average number of key comparisons made by sequential search is $(n + 1) / 2$.

◆ If $p = 0$ (i.e., the search must be unsuccessful), the average number of key comparisons will be n because the algorithm will inspect all n elements on all such inputs.

Recapitulation of the Analysis Framework

- 1 Both time and space efficiencies are measured as functions of the algorithm's input size.
- 2 Time efficiency is measured by counting the number of times the algorithm's basic operation is executed. Space efficiency is measured by counting the number of extra memory units consumed by the algorithm.
- 3 The efficiencies of some algorithms may differ significantly for inputs of the same size. For such algorithms, we need to distinguish between the worst-case, average-case, and best-case efficiencies.
- 4 The framework's primary interest lies in the order of growth of the algorithm's running time
(extra memory units consumed) as its input size goes to infinity.

2. PERFORMANCE ANALYSIS

2.1 Space complexity

- Total amount of computer memory required by an algorithm to complete its execution is called as **space complexity** of that algorithm.
- The Space required by an algorithm is the sum of following components
 - ✓ A **fixed** part that is independent of the input and output. This includes memory space for codes, variables, constants and so on.
 - ✓ A **variable** part that depends on the input, output and recursion stack. (We call these parameters as instance characteristics)
- Space requirement $S(P)$ of an algorithm P ,

$$S(P) = c + Sp$$

where c is a constant depends on the fixed part, Sp is the instance characteristics

Example-1: Consider following algorithm $abc()$

```
Algorithm abc(a,b,c)
{
    return a+b+++*c+(a+b-c)/(a+b) +4.0;
}
```

DAA module-1

Here fixed component depends on the size of a , b and c . Also instance characteristics $S_p=0$

Example-2: Let us consider the algorithm to find sum of array.

For the algorithm given here the problem instances are characterized by n , the number of elements to be summed. The space needed by $a[]$ depends on n . So the space complexity can be written as; $S_{sum}(n) \geq (n+3)n$ for $a[]$, One each for n , i and

```
float Sum(float a[], int n)
{
    float s = 0.0;
    for (int i=1; i<=n; i++)
        s += a[i];
    return s;
}
```

2.2 Time Complexity

- The time $T(p)$ taken by a program P is the sum of the compile time and the run time(execution time)
 - ✓ The compile time does not depend on the instance characteristics. Also we may assume that a compiled program will be run several times without recompilation .
 - ✓ The run time is denoted by t_p (instance characteristics).
- Usually, the execution time or run-time of the program is referred as its time complexity denoted by t_p (instance characteristics). This is the sum of the time taken to execute all instructions in the program.

✓

For example comment has zero steps; assignment statement has one step and in an iterative statement such as the for, while & repeat-until statements, the step

DAA module-1

count is counted for the control part of the statement

- We can determine the **steps needed by a program** to solve a particular problem instance in two ways

✓

*In the **first method** we introduce a new variable **count** to the program which is initialized to zero. We also introduce statements to increment **count** by an appropriate amount into the program. So when each time original program executes, the **count** also incremented by the step count.*

✓

Example-1: Consider the algorithm **Sum()**. After the introduction of the count the program will be as follows.

```
Algorithm Sum(a, n)
{
    s := 0.0;
    count := count + 1; // count is global; it is initially zero.
    for i := 1 to n do
    {
        count := count + 1; // For for
        s := s + a[i]; count := count + 1; // For assignment
    }
    count := count + 1; // For last time of for
    count := count + 1; // For the return
    return s;
}
```

*From the above we can estimate that invocation of **Sum()** executes total number of **2n+3** steps.*

✓

Example-2: Consider the algorithm that computes Sum of n numbers recursively

```
Algorithm RSum(a, n)
{
    count := count + 1; // For the if conditional
    if (n ≤ 0) then
    {
        count := count + 1; // For the return
        return 0.0;
    }
    else
    {
        count := count + 1; // For the addition, function
        // invocation and return
        return RSum(a, n - 1) + a[n];
    }
}
```

■

When analyzing a recursive program for its step count, we often obtain

DAA module-1

A recursive formula for the step count, for example

$$t_{RSum}(n) = \begin{cases} 2 & \text{if } n = 0 \\ 2 + t_{RSum}(n - 1) & \text{if } n > 0 \end{cases}$$

These recursive formulas are referred to as recurrence relations. One way of solving is to make repeated substitutions for each occurrence of the function t_{RSum} on the right-hand side until all such occurrences disappear

$$\begin{aligned} t_{RSum}(n) &= 2 + t_{RSum}(n - 1) \\ &= 2 + 2 + t_{RSum}(n - 2) \\ &= 2(2) + t_{RSum}(n - 2) \\ &\vdots \\ &= n(2) + t_{RSum}(0) \\ &= 2n + 2, \quad n \geq 0 \end{aligned}$$

- So, the step count for $RSum$ is $2n+2$

✓

The **second method** to determine the step count of an algorithm is to build a table in which we list the total number of steps contributed by each statement.

First determine the number of steps per execution (s/e) of the statement and the total

number of times (ie., frequency) each statement is executed.

- By combining these two quantities, the total contribution of all statements, the step count for the entire algorithm is obtained.

✓ **Example 1:** Consider the algorithm `sum()`.

Statement	s/e	frequency	total steps
<code>float Sum(float a[], int n)</code>	0	-	0
<code>{ float s = 0.0;</code>	1	1	1
<code> for (int i=1; i<=n; i++)</code>	1	$n+1$	$n+1$
<code> s += a[i];</code>	1	n	n
<code> return s;</code>	1	1	1
<code>}</code>	0	-	0
Total			$2n+3$

Table: Step table for the `Sum()` algorithm

Example 2: Consider the algorithm that computes Sum of n numbers recursively

Statement	s/e	frequency		total steps	
		$n = 0$	$n > 0$	$n = 0$	$n > 0$
1 Algorithm RSum(a, n)	0	–	–	0	0
2 {					
3 if ($n \leq 0$) then	1	1	1	1	1
4 return 0.0;	1	1	0	1	0
5 else return					
6 RSum($a, n - 1$) + $a[n]$;	$1 + x$	0	1	0	$1 + x$
7 }	0	–	–	0	0
Total				2	$2 + x$

$$x = t_{\text{RSum}}(n - 1)$$

Table: Step table for the RSum() algorithm

3. ASYMPTOTIC NOTATIONS

The efficiency analysis framework concentrates on the order of growth of an algorithm's basic operation count as the principal indicator of the algorithm's efficiency.

To compare and rank such orders of growth, computer scientists use three notations:

O (big oh), Ω (big omega), and

Θ (big theta).

Informal Introduction

Informally, $O(g(n))$ is the set of all functions with a lower or same order of growth as $g(n)$ (to within a constant multiple, as n goes to infinity). Thus, to give a few examples, the following assertions are all true:

$$n \in O(n^2), \quad 100n + 5 \in O(n^2), \quad \frac{1}{2}n(n - 1) \in O(n^2).$$

Indeed, the first two functions are linear and hence have a lower order of growth than $g(n) = \frac{1}{2}n^2$, while the last one is quadratic and hence has the same order of growth as n^2 . On the other hand,

$$n^3 \notin O(n^2), \quad 0.00001n^3 \notin O(n^2), \quad n^4 + n + 1 \notin O(n^2).$$

Indeed, the functions n^3 and $0.00001n^3$ are both cubic and hence have a higher order of growth than n^2 , and so has the fourth-degree polynomial $n^4 + n + 1$.

The second notation, $\Omega(g(n))$, stands for the set of all functions with a higher or same order of growth as $g(n)$ (to within a constant multiple, as n goes to infinity).

For example,

$$n^3 \in \Omega(n^2), \quad \frac{1}{2}n(n-1) \in \Omega(n^2), \quad \text{but } 100n + 5 \notin \Omega(n^2).$$

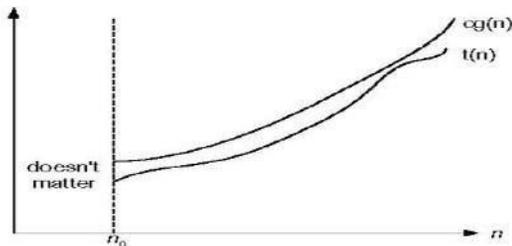
Finally, $\Theta(g(n))$ is the set of all functions that have the same order of growth as $g(n)$.

Big Oh- O notation

DEFINITION A function $t(n)$ is said to be in $O(g(n))$, denoted $t(n) \in O(g(n))$, if $t(n)$ is bounded above by some constant multiple of $g(n)$ for all large n , i.e., if there exist some positive constant c and some nonnegative integer n_0 such that

$$t(n) \leq cg(n) \quad \text{for all } n \geq n_0.$$

The definition is illustrated in the following figure.

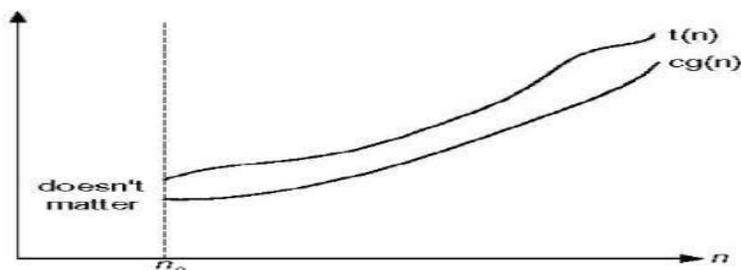


Big-oh notation: $t(n) \in O(g(n))$

Big Omega- Ω notation

DEFINITION A function $t(n)$ is said to be in $\Omega(g(n))$, denoted $t(n) \in \Omega(g(n))$, if $t(n)$ is bounded below by some positive constant multiple of $g(n)$ for all large n , i.e., if there exist some positive constant c and some nonnegative integer n_0 such that

$$t(n) \geq cg(n) \quad \text{for all } n \geq n_0.$$



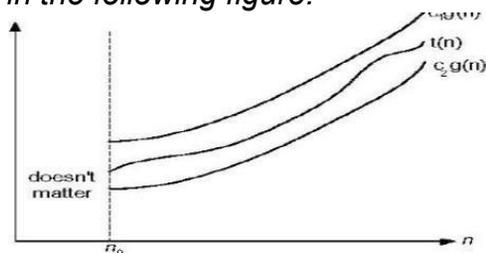
Big-omega notation: $t(n) \in \Omega(g(n))$

Big Theta- Θ notation

DEFINITION A function $t(n)$ is said to be in $\Theta(g(n))$, denoted $t(n) \in \Theta(g(n))$, if $t(n)$ is bounded both above and below by some positive constant multiples of $g(n)$ for all large n , i.e., if there exist some positive constants c_1 and c_2 and some nonnegative integer n_0 such that

$$c_2g(n) \leq t(n) \leq c_1g(n) \quad \text{for all } n \geq n_0.$$

The definition is illustrated in the following figure.



Big-theta notation: $t(n) \in \Theta(g(n))$

Examples: Refer class notes



Useful property involving the asymptotic notations

DAA module-1

Using the formal definitions of the asymptotic notations, we can prove their general properties. The following property, in particular, is useful in analyzing algorithms that comprise two consecutively executed parts

THEOREM If $t_1(n) \in O(g_1(n))$ and $t_2(n) \in O(g_2(n))$, then

$$t_1(n) + t_2(n) \in O(\max\{g_1(n), g_2(n)\}).$$

(The analogous assertions are true for the Ω and Θ notations as well.)

PROOF The proof extends to orders of growth the following simple fact about four arbitrary real numbers a_1, b_1, a_2, b_2 : if $a_1 \leq b_1$ and $a_2 \leq b_2$, then $a_1 + a_2 \leq 2 \max\{b_1, b_2\}$.

Since $t_1(n) \in O(g_1(n))$, there exist some positive constant c_1 and some non-negative integer n_1 such that

$$t_1(n) \leq c_1 g_1(n) \quad \text{for all } n \geq n_1.$$

Similarly, since $t_2(n) \in O(g_2(n))$,

$$t_2(n) \leq c_2 g_2(n) \quad \text{for all } n \geq n_2.$$

Let us denote $c_3 = \max\{c_1, c_2\}$ and consider $n \geq \max\{n_1, n_2\}$ so that we can use both inequalities. Adding them yields the following:

$$\begin{aligned} t_1(n) + t_2(n) &\leq c_1 g_1(n) + c_2 g_2(n) \\ &\leq c_3 g_1(n) + c_3 g_2(n) = c_3 [g_1(n) + g_2(n)] \\ &\leq c_3 2 \max\{g_1(n), g_2(n)\}. \end{aligned}$$

Hence, $t_1(n) + t_2(n) \in O(\max\{g_1(n), g_2(n)\})$, with the constants c and n_0 required by the O definition being $2c_3 = 2 \max\{c_1, c_2\}$ and $\max\{n_1, n_2\}$, respectively. ■

So what does this property imply for an algorithm that comprises two consecutively executed parts? It implies that the algorithm's overall efficiency is determined by the part with a higher order of growth, i.e., its least efficient part:

Class	Name	Comments
1	<i>constant</i>	Short of best-case efficiencies, very few reasonable examples can be given since an algorithm's running time typically goes to infinity when its input size grows infinitely large.
$\log n$	<i>logarithmic</i>	Typically, a result of cutting a problem's size by a constant factor on each iteration of the algorithm (see Section 4.4). Note that a logarithmic algorithm cannot take into account all its input or even a fixed fraction of it: any algorithm that does so will have at least linear running time.
n	<i>linear</i>	Algorithms that scan a list of size n (e.g., sequential search) belong to this class.
$n \log n$	<i>linearithmic</i>	Many divide-and-conquer algorithms (see Chapter 5), including mergesort and quicksort in the average case, fall into this category.
n^2	<i>quadratic</i>	Typically, characterizes efficiency of algorithms with two embedded loops (see the next section). Elementary sorting algorithms and certain operations on $n \times n$ matrices are standard examples.
n^3	<i>cubic</i>	Typically, characterizes efficiency of algorithms with three embedded loops (see the next section). Several nontrivial algorithms from linear algebra fall into this class.
2^n	<i>exponential</i>	Typical for algorithms that generate all subsets of an n -element set. Often, the term "exponential" is used in a broader sense to include this and larger orders of growth as well.
$n!$	<i>factorial</i>	Typical for algorithms that generate all permutations of an n -element set.

fast ↓ slow	1	constant	High time efficiency low time efficiency
	$\log n$	logarithmic	
	n	linear	
	$n \log n$	$n \log n$	
	n^2	quadratic	
	n^3	cubic	
	2^n	exponential	
	$n!$	factorial	

4. MATHEMATICAL ANALYSIS OF NON-RECURSIVE ALGORITHMS

General plan for analyzing efficiency of non-recursive algorithms:

1. Decide on parameter n indicating the **input size of the algorithm**.
2. Identify algorithm's **basic operation**.
3. Check whether the number of times the basic operation is executed depends only on the input size n . If it also depends on the type of input, then investigate **worst, average, and best case efficiency** separately.
4. Set up **summation** for $C(n)$ reflecting the number of times the algorithm's basic operation is executed.
5. Simplify summation using standard formulas (**Refer class notes for formulas used for simplifying**).

Example: Finding the largest element in a given array

ALGORITHM *MaxElement*($A[0..n - 1]$)

//Determines the value of the largest element in a given array

//Input: An array $A[0..n - 1]$ of real numbers

//Output: The value of the largest element in A

maxval $\leftarrow A[0]$

for $i \leftarrow 1$ **to** $n - 1$ **do**

if $A[i] > \text{maxval}$

maxval $\leftarrow A[i]$

return *maxval*

Analysis:

DAA module-1

1. Input size: the number of elements = n (size of the array)
2. Two operations can be considered to be as basic operation i.e.,
 - a) Comparison :: $A[i] > \text{maxval}$.
 - b) Assignment :: $\text{maxval} \leftarrow A[i]$.

Here the comparison statement is considered to be the basic operation of the algorithm.

3. No best, worst, average cases- because the number of comparisons will be same for all arrays of size n and it is not dependent on type of input.

4. Let $C(n)$ denotes number of comparisons: Algorithm makes one comparison on each execution of the loop, which is repeated for each value of the loop's variable i within the bound

between 1 and $n - 1$.
$$C(n) = \sum_{i=1}^{n-1} 1$$

This is an easy sum to compute because it is nothing other than 1 repeated $n - 1$ times. Thus

$$C(n) = \sum_{i=1}^{n-1} 1 \quad \begin{array}{l} 1 + 1 + 1 + \dots + 1 \\ [(n-1) \text{ number of times}] \end{array}$$

$$C(n) = \sum_{i=1}^{n-1} 1 = n - 1 \in \Theta(n).$$

```
Algorithm UniqueElements (A[0..n-1])
//Checks whether all the elements in a given array are distinct
//Input: An array A[0..n-1]
//Output: Returns true if all the elements in A are distinct and false otherwise
for i ← 0 to n - 2 do
    for j ← i + 1 to n - 1 do
        if A[i] == A[j]
            return false
return true
```

Example: Element uniqueness problem-Checks whether the elements in the array are distinct or not.

$$C(n) = \sum_{i=1}^{n-1} 1 \quad [1 + 1 + 1 + \dots + 1 \text{ [(n-1) number of times]}]$$
$$C(n) = \sum_{i=1}^{n-1} 1 = n - 1 \in \Theta(n).$$

```
Algorithm UniqueElements (A[0..n-1])
//Checks whether all the elements in a given array are distinct
//Input: An array A[0..n-1]
//Output: Returns true if all the elements in A are distinct and false otherwise
for i ← 0 to n - 2 do
    for j ← i + 1 to n - 1 do
        if A[i] == A[j]
            return false
return true
```

Analysis

1. Input size: number of elements = n (size of the array)
2. Basic operation: Comparison
3. Best, worst, average cases exists

- Array with no equal elements
- Array with last two elements are the only pair of equal elements

4. Let $C(n)$ denotes number of comparisons in worst case: Algorithm makes one comparison for

each repetition of the innermost loop i.e., for each value of the loop's variable j between its limits $i + 1$ and $n - 1$; and this is repeated for each value of the outer loop i.e., for each value of the loop's variable i between its limits 0 and $n - 2$.

$$C(n) = \sum_{i=0}^{n-2} \sum_{j=i+1}^{n-1} 1$$

5. Simplify the summation using standard formulas as follows:

$$C(n) = \sum_{i=0}^{n-2} ((n-1) - (i+1) + 1)$$

$$C(n) = \sum_{i=0}^{n-2} (n-1-i)$$

$$C(n) = \sum_{i=0}^{n-2} (n-1) - \sum_{i=0}^{n-2} i$$

$$C(n) = (n-1) \sum_{i=0}^{n-2} 1 - \sum_{i=0}^{n-2} i$$

$$C(n) = (n-1) \sum_{i=0}^{n-2} 1 - \frac{(n-2)(n-1)}{2}$$

$$C(n) = (n-1)(n-1) - \frac{(n-2)(n-1)}{2}$$

$$C(n) = (n-1) \left((n-1) - \frac{(n-2)}{2} \right)$$

$$C(n) = (n-1) \frac{(2n-2-n+2)}{2}$$

$$C(n) = \frac{(n-1)(n)}{2}$$

$$= \frac{(n^2-n)}{2}$$

$$= \frac{(n^2)}{2} - \frac{n}{2}$$

$$C(n) \in \Theta(n^2)$$

Example 3: Given two $n \times n$ matrices A and B , find the time efficiency of the definition-based algorithm for computing their product $C = AB$. By definition, C is an $n \times n$ matrix whose elements are computed as the scalar (dot) products of the rows of matrix A and the columns of matrix B :

$$\begin{array}{c}
 \text{A} \\
 \left[\begin{array}{c} \text{row } i \\ \hline \square \square \square \square \square \end{array} \right] * \left[\begin{array}{c} \text{B} \\ \hline \square \\ \square \\ \square \\ \square \\ \square \\ \square \end{array} \right] = \left[\begin{array}{c} \text{C} \\ \hline \square \\ \square \\ \square \\ \square \\ \square \\ \square \end{array} \right] \\
 \text{col. } j
 \end{array}$$

where $C[i, j] = A[i, 0]B[0, j] + \dots + A[i, k]B[k, j] + \dots + A[i, n-1]B[n-1, j]$
for every pair of indices $0 \leq i, j \leq n-1$.

ALGORITHM *MatrixMultiplication*($A[0..n-1, 0..n-1]$, $B[0..n-1, 0..n-1]$)
 //Multiplies two square matrices of order n by the definition-based algorithm
 //Input: Two $n \times n$ matrices A and B
 //Output: Matrix $C = AB$
for $i \leftarrow 0$ **to** $n-1$ **do**
 for $j \leftarrow 0$ **to** $n-1$ **do**
 $C[i, j] \leftarrow 0.0$
 for $k \leftarrow 0$ **to** $n-1$ **do**
 $C[i, j] \leftarrow C[i, j] + A[i, k] * B[k, j]$
return C

Analysis:

1. We measure an input's size by matrix order n .
2. There are two arithmetical operations in the innermost loop here—multiplication and addition—that, in principle, can compete for designation as the algorithm's basic operation.
 Actually, we do not have to choose between them, because on each repetition of the innermost loop each of the two is executed exactly once. So by counting one we automatically count the other. Hence, we consider multiplication as the basic operation.
3. Let us set up a sum for the total number of multiplications $M(n)$ executed by the algorithm. (Since this count depends only on the size of the input matrices, we do not have to investigate the worst-case, average-case, and best-case efficiencies separately.)
4. Obviously, there is just one multiplication executed on each repetition of the

algorithm's

innermost loop, which is governed by the variable k ranging from the lower bound 0 to the upper bound $n - 1$. Therefore, the number of multiplications made for every pair of specific values of variables i and j is

$$\sum_{k=0}^{n-1} 1,$$
$$M(n) = \sum_{i=0}^{n-1} \sum_{j=0}^{n-1} \sum_{k=0}^{n-1} 1.$$

Now, we can compute this sum by using formula (S1) and rule (R1). Starting with the innermost sum, which is equal to n , we get

$$M(n) = \sum_{i=0}^{n-1} \sum_{j=0}^{n-1} \sum_{k=0}^{n-1} 1 = \sum_{i=0}^{n-1} \sum_{j=0}^{n-1} n = \sum_{i=0}^{n-1} n^2 = n^3.$$

If we now want to estimate the running time of the algorithm on a particular machine, we can do it by the product

$$T(n) \approx c_m M(n) = c_m n^3,$$

where c_m is the time of one multiplication on the machine in question. We would get a more accurate estimate if we took into account the time spent on the additions, too:

$$T(n) \approx c_m M(n) + c_a A(n) = c_m n^3 + c_a n^3 = (c_m + c_a) n^3,$$

Example 4: The following algorithm finds the number of binary digits in the binary representation of a positive decimal integer

ALGORITHM *Binary*(n)

//Input: A positive decimal integer n

//Output: The number of binary digits in n 's binary representation

$count \leftarrow 1$

while $n > 1$ **do**

$count \leftarrow count + 1$

$n \leftarrow \lfloor n/2 \rfloor$

return $count$

✓

First, notice that the most frequently executed operation here is not inside the **while** loop but rather the comparison $n > 1$ that determines whether the loop's body will be executed.

✓

Since the number of times the comparison will be executed is larger than the number of repetitions of the loop's body by exactly 1, the choice is not that important.

✓

A more significant feature of this example is the fact that the loop variable takes on only a few values between its lower and upper limits; therefore, we have to use an alternative way

of computing the number of times the loop is executed. Since the value of n is about halved on each repetition of the loop, the answer should be about $\log_2 n$.

✓

The exact formula for the number of times the comparison $n > 1$ will be executed is actually

$$\lfloor \log_2 n \rfloor + 1$$

the number of bits in the binary representation of n .

5. MATHEMATICAL ANALYSIS(TIME EFFICIENCY) OF RECURSIVE ALGORITHMS

General plan for analyzing efficiency of recursive algorithms:

1. Decide on parameter n indicating **input size**

2. Identify algorithm's **basic operation**

3. Check whether the number of times the basic operation is executed depends only on the input size n . If it also depends on the type of input, investigate **worst, average, and best case efficiency** separately.

4. Set up **recurrence relation**, with an appropriate initial condition, for the number of times the

algorithm's basic operation is executed.

5. **Solve** the recurrence.

Example 1: Factorial function

DAA module-1

Definition: $n! = 1 * 2 * \dots * (n-1) * n$ for $n \geq 1$ and $0! = 1$

Recursive definition of $n!$: $F(n) = F(n-1) * n$ for $n \geq 1$ and

$$F(0) = 1$$

ALGORITHM Factorial (n)

//Computes $n!$ recursively

//Input: A nonnegative integer n

//Output: The value of $n!$

if $n = 0$

return 1

else

return Factorial ($n - 1$) * n

Analysis:

1. Input size: given number = n
2. Basic operation: multiplication
3. No best, worst, average cases.
4. Let $M(n)$ denotes number of multiplications

$$M(n) = M(n-1) + 1 \quad \text{for } n > 0$$

$$M(0) = 0 \quad \text{initial condition}$$

Where: $M(n-1)$: to compute Factorial ($n-1$)

1 : to multiply Factorial ($n-1$) by n

5. Solve the recurrence relation using backward substitution method:

$$M(n) = M(n-1) + 1 \quad \text{substitute } M(n-1) = M(n-2) + 1$$

$$= (M(n-2) + 1) + 1 \quad \text{substitute } M(n-2) = M(n-3) + 1$$

$$= (M(n-3) + 1) + 2 \quad \text{substitute } M(n-3) = M(n-4) + 1$$

...

The general formula for the above pattern for some i is as follows:

$$= M(n-i) + i$$

By taking the advantage of the initial condition given i.e., $M(0)=0$, we now substitute $i=n$ in the patterns formula to get the ultimate result of the backward substitutions.

$$= M(n-n) + n$$

$$= M(0) + n$$

$$M(n) = n$$

$$M(n) \in \Theta(n)$$

The number of multiplications to compute the factorial of n is n where the time complexity is linear.

Example 2: Tower of Hanoi Problem

In this puzzle, we have n disks of different sizes that can slide onto any of three pegs. Initially, all the disks are on the first peg in order of size, the largest on the bottom and the smallest on top. The goal is to move all the disks to the third peg, using the second one as an auxiliary, if necessary. We can move only one disk at a time, and it is forbidden to place a larger disk on top of a smaller one.

The problem has an elegant recursive solution, which is illustrated in Figure.

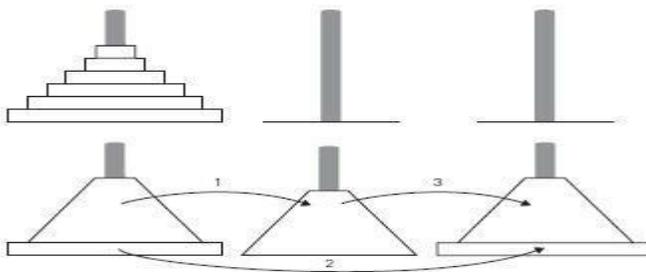
✓

To move $n > 1$ disks from peg 1 to peg 3 (with peg 2 as auxiliary):

- we first move recursively $n - 1$ disks from peg 1 to peg 2 (with peg 3 as auxiliary).
- Then move the largest disk i.e., n th disk directly from peg 1 to peg 3.
- And, finally, move recursively $n - 1$ disks from peg 2 to peg 3 (using peg 1 as auxiliary).

✓

Of course, if $n = 1$, we simply move the single disk directly from the source peg to the destination peg.



Analysis:

Let us apply the general plan outlined above to the Tower of Hanoi problem.

DAA module-1

1. The number of disks n is the obvious choice for the input's size indicator.
2. Moving one disk as the algorithm's basic operation.
3. Clearly, the number of moves $M(n)$ depends on n only, and we get the following recurrence equation for it:

$$M(n) = M(n-1) + 1 + M(n-1) \text{ for } n > 1,$$

$$M(n) = 2M(n-1) + 1, M(1) = 1 \text{ and}$$

The initial condition is:

$$M(1) = 1 \text{ for } n=1$$

5. Solve the recurrence relation using backward substitution method

$$M(n) = 2M(n-1) + 1$$

$$\text{sub. } M(n-1) = 2M(n-2) + 1$$

$$2(2M(n-2) + 1) + 1 = 2^2 M(n-2) + 2^1 + 2^0$$

$$\text{sub. } M(n-2) = 2M(n-3) + 1$$

$$= 2^2 (2M(n-3) + 1) + 2^1 + 2^0$$

$$= 2^3 M(n-3) + 2^2 + 2^1 + 2^0$$

$$= \dots$$

Since the initial condition is specified for $n = 1$, which is achieved for $i = n - 1$, we get the following formula for the solution to recurrence

$$M(n) = 2^{n-1} M(n - (n - 1)) + 2^{n-1} - 1$$

$$= 2^{n-1} M(1) + 2^{n-1} - 1 = 2^{n-1} + 2^{n-1} - 1 = 2^n - 1.$$

Thus, we have an exponential algorithm, which will run for an unimaginably long time even for moderate values of n . This is not due to the fact that this particular algorithm is poor; in fact, it is not difficult to prove that this is the most efficient algorithm possible for this problem. It is the problem's intrinsic difficulty that makes it so computationally hard.

EXAMPLE 3 A recursive version of the algorithm that computes number of binary digits in n 's binary representation.

ALGORITHM *BinRec*(n)

//Input: A positive decimal integer n

//Output: The number of binary digits in n 's binary representation

if $n = 1$ return 1

else return *BinRec*($\lfloor n/2 \rfloor$) + 1

✓

Let us set up a recurrence and an initial condition for the number of additions $A(n)$ made by the algorithm.

✓

The number of additions made in computing $\lfloor n/2 \rfloor$, plus one more addition is made by the algorithm to increase the returned value by 1. This leads to the recurrence

✓

$A(n) = A(\lfloor n/2 \rfloor) + 1$

✓

$A(n) = A(\lfloor n/2 \rfloor) + 1 \quad \text{for } n > 1.$

✓

Since the recursive calls end when n is equal to 1 and there are no additions made then, the initial condition is

$$A(1) = 0. \text{ for } n=1$$

✓

The presence of $\lfloor n/2 \rfloor$ in the function's argument makes the method of backward substitutions stumble on values of n that are not powers of 2.

✓

Therefore, the standard approach to solving such a recurrence is to solve it only for $n = 2^k$ and then take advantage of the theorem called the **smoothness rule**, which claims that under

very broad assumptions the order of growth observed for $n = 2^k$ gives a correct answer about the order of growth for all values of n . (Alternatively, after getting a solution for powers of

✓

2, we can sometimes fine-tune this solution to get a formula valid for an arbitrary n .)

Applying the same to the recurrence, for $n = 2^k$ which takes the form

$$A(2^k) = A(2^{k-1}) + 1 \quad \text{for } k > 0,$$

$$A(2^0) = 0.$$

Now backward substitutions encounter no problems:

$$\begin{aligned} A(2^k) &= A(2^{k-1}) + 1 && \text{substitute } A(2^{k-1}) = A(2^{k-2}) + 1 \\ &= [A(2^{k-2}) + 1] + 1 = A(2^{k-2}) + 2 && \text{substitute } A(2^{k-2}) = A(2^{k-3}) + 1 \\ &= [A(2^{k-3}) + 1] + 2 = A(2^{k-3}) + 3 && \dots \\ &\dots && \\ &= A(2^{k-i}) + i && \\ &\dots && \\ &= A(2^{k-k}) + k. \end{aligned}$$

Thus, we end up with

$$A(2^k) = A(1) + k = k,$$

or, after returning to the original variable $n = 2^k$ and hence $k = \log_2 n$,

$$A(n) = \log_2 n \in \Theta(\log n).$$

into a collection of disjoint subsets.

- ✓ After being initialized as a collection of n one-element subsets, the collection is subjected to a sequence of intermixed union and search operations. This problem is called the **set union problem**.