

Ques What is Data Structure? What are its classifications?

Ans → Data Structure is a storage format that is used to store and organise data. It is a way of arranging data on a computer so that it can be accessed and updated easily. In other words, data structures refers to the way of assembling or organising data so that it can be used efficiently. It is a particular way of storing data in computer's memory so that it can be used easily. The choice of a good data structure makes it possible to perform a variety of critical operations effectively. An efficient data structure uses minimum memory space and execution time to process the structure as possible. The choice of choosing the right data structure may depend upon the size of program or time complexity of the program. The data structure only deals with the organization of data in main memory (RAM) and not in secondary memory. Hence, data structure is one of the factors that is responsible for the execution time taken by the program to load in Main memory.

CLASSIFICATIONS ➤

Data structures are classified into four categories

1) Linear and Non-Linear D.S.

In a linear d.s. the elements are arranged in a sequence so that processing of elements can be possible in linear manner. In non-linear d.s. the elements are not arranged in a sequence manner.

2) Homogeneous and Non-Homogeneous D.S.

In homogeneous D.S., all elements are of same datatype & in heterogeneous D.S., all elements are of different datatype.

3) Static and Dynamic Data Structure

The D.S. whose size is fixed is known as static D.S. and whose size is not fixed is known as Non-static data structure.

4) Primitive & Non-Primitive D.S.

Primitive D.S. is the D.S. that can not be further subdivided.

Non-Primitive D.S. is the D.S. that can be further sub-divided into various parts.

Ques Define array and its types. Also discuss the advantages and disadvantages of Arrays.

Ans → Arrays ⇒ An array is a collection of homogeneous data elements of a specific datatype that have been given a single name which are stored in contiguous memory location. All the elements of array can be accessed by using its index number enclosed in square brackets after the array name.

Ex. int a[5];

Or

a[5] = 10;

Types of Arrays:-

- 1) While declaring an array, if only one square bracket is used then the array is one-dimensional.
- 2) While declaring an array, if two square brackets are used then the array is two-dimensional.
- 3) While declaring an array, if three or more square brackets are used then the array is multidimensional.

Example := # include <stdio.h>

void main()

{

int a[10]; // 1-D

int b[10][10]; // 2-D

int c[10][10][10]; // Multi-D

{}

The index of array always starts from 0.

The array elements can be accessed by incrementing the index value by 1. We can access the array randomly and as well as sequentially.

By incrementing the value of index by 1 we can access the next block of the array. We can perform various operations on arrays like,

- 1) Searching
- 2) Sorting
- 3) Traversing
- 4) Merging
- 5) Insertion
- 6) Deletion

We can also initialize the array while we are declaring it.

Ex:- $\text{int } a[5] = \{ 1, 2, 3, 4, 5 \}$

By default, an array contains garbage value.

But if we initialize less elements than the size of array during declaration, rest of the blocks will contain the value 0.

Advantages of Array \Rightarrow

- 1) Array is the simplest kind of D.S.
- 2) Array is the most compact data structure i.e. it takes minimum possible space in memory.
- 3) Computation of vectors & matrices become easy.
- 4) Elements of array have direct access to them.
- 5) Array elements can be accessed instantly.
- 6) Arrays help programmers to refer to many values with a single name.

Disadvantages of Array \Rightarrow

- 1) An array use reference mechanism to work with memory which can cause instable behaviour of operating system.
- 2) The size of array should be known in advance i.e. before compile time.
- 3) Insertion and Deletion in case of array is very costly since a lot of data movement has to be done.
- 4) Sometimes, it's not easy to work with many

arrays.

- 5) Array holds the similar kind of information.
So it is impossible to store unrelated information
into an array.

Memory Representation of 1-D Array

Array elements are stored in continuous memory locations. Computer does not need to keep track of the address of every element of linear array, but only of the first element. The address of the First element of linear array is called Base Address.

Base	→	200	2002	2004	2006	2008	2010	2012
Address	200	10	20	30	40	50	60	70

Memory Representation of 2-D Array

In 2-D arrays, elements are arranged in rows and columns, But computer memory is linear. So, elements of 2-D array are mapped into linear memory. 2-D arrays can be stored in memory in two ways :

- 1) Column - major Order
- 2) Row - major Order

1) Column - major Order

In column - major order, the elements are stored column wise i.e. first column is stored, then second column & then third column and so on.

If we have 3×2 matrix, then elements will be stored as:-

A ₁₁
A ₂₁
A ₃₁
A ₁₂
A ₂₂
A ₃₂

2) Row-Major Order

In row-major order, the elements are stored row-wise i.e. first row is stored, then second row, then third row and so on. If we have 3×2 matrix, then the elements will be stored as:-

A ₁₁
A ₁₂
A ₂₁
A ₂₂
A ₃₁
A ₃₂

STACK

Stack is a linear d.s. that implements LIFO scheme. A stack is a simple d.s. that allows adding and removing elements only at one end. That end is called Top of the Stack i.e. items can be inserted or removed only at the top of the stack. The last item added to the stack will be first item to removed from stack.

Imagine stacking a set of books on top of each other - you can place a new book on the top of stack and you can remove the book that is currently on the top of stack. Size of stack depends upon number of elements currently present in the stack.

In D.S., we can represent stack as follows :-
Data Structure (Variable) used :-

TOP → It always address top most element of stack.

Operations

PUSH → Pushing something on the stack means "placing it on top". Push operation is used to insert any element at the TOP of stack.

During a push operation, first we check availability of space in stack. If stack has reached its limit, it is overflow condition. In such case, we can not insert the element in stack. After push operation, TOP of stack is incremented by 1.

Push in Array Implementation of Stack.

In array representation, stack is represented using a 1-D array say 'stack' with size 'n'. Here 'n' is the limit of Stack. Following steps are followed to Push an element onto stack.

- 1) Firstly, we check for overflow. If $\text{Top} = n$, then we cannot insert a new element.
- 2) Then we check whether the stack is empty. If yes, i.e. $\text{top} = \text{NULL}$, then we initialize top to 1 otherwise we simply increment top by 1 i.e. $\text{Top} = \text{Top} + 1$.
- 3) Then, we insert new element at the Top.

POP → Pop means "retrieve data" from the stack.

Pop operation of the stack is used to remove any element from top of stack. After pop operation, top element is retrieved from stack and next element becomes top of stack.

Pop in Array implementation of stack.

Following are the steps to pop an element from stack.

- 1) Firstly, we check for underflow. If the stack is empty i.e. if $\text{TOP} = \text{NULL}$ then it is underflow.
- 2) If underflow condition is false then we move to next step. In this step, we retrieve the Top element of stack.
- 3) Then we set the top pointer. If the popped element was the last element of stack i.e. top was equal to

then top is set to NULL otherwise we simply decrement top by 1. i.e. $\text{Top} = \text{Top} - 1$

Queue

A queue is a particular kind of collection in which the entries in the collection are kept in order. A queue is a data structure where insertions are made at one end and deletions are made at another end. We add elements at the end also called as rear of queue and remove element from the front.

For ex:- Any new entry is made at the back of the queue and persons at the first are served first.

Queue is an example of linear Data Structure.

In short, we summarize Queue as follows:

- 1) A list structure with two access points called the front and the rear.
- 2) All insertions occur at the rear and deletions occur at the front.
- 3) Queue is of varying length.
- 4) Queue mostly stores homogeneous components.
- 5) Queue has FIFO characteristics.

Applications Of Queue

Queue has applications in day to day life as well as in computers.

1) In Real Life

- a) Queue in front of ticket window
- b) Queue for admission
- c) Queue for fee deposit
- d) Queue at a bank counter
- e) Queue waiting for bus at local bus stop.

2) In Computers

- a) Multiprogramming - Queues are used to implement multiprogramming which is very important area of operating system. Multiprogramming requires queues for every field from processor management to device management.
- b) Simulating Real life Queues - We can use queues for simulating real life simulations like banking.
- c) Message Queue - Message queue is a good way to communicate from one application to another.
- d) Playlist - Add songs to the end, play from the front of list.

Queue Operations

① Enqueue

Enqueue means insertion of new elements in the queue. In array representation of queue, we have an array 'queue' with maximum 'n' elements and two variables 'front' and 'rear' which points to the first and last element of queue. Initially, both front and rear are set to NULL. When we want to insert first element, we set both front and rear to 1. After that for each insertion, we check for overflow condition. It can be checked as

if (Rear=n) then

Print "overflow"

Return

[end if]

If no overflow, it means that space is available and we increment rear pointer as

Rear = Rear + 1

After that, we insert new item in the Queue.

② Dequeue

Dequeue operation is used to delete an element from the queue. An element can be deleted only from the front of queue. Before Dequeue, we will check for underflow condition. It means that we will check for the availability of elements. Underflow can be checked using the statement.

If (FRONT = NULL And REAR = NULL) then

Print "Underflow"

Return

[end if]

If element is available i.e. front and rear are not NULL then the front element is deleted from the queue. After the deletion, we will set the values of front and rear. If front and rear both are equal then after deletion both front and rear will set to NULL; otherwise we will increment value of front by 1. In this case, there will be no change in rear.

if (front = rear) then

- a) set front = NULL
- b) set rear = NULL

else

 front = front + 1

[end if]

Difference

Stack

- 1) The stack is based on LIFO principle.
- 2) Insertion operation is called Push operation.
- 3) Deletion operation is called Pop operation.
- 4) Push and Pop operation takes place from only one end of stack.
- 5) Only one pointer is used for performing operation.
- 6) Empty condition is checked using $\text{TOP} == -1$.
- 7) Full condition is checked using $\text{TOP} == \text{Max} - 1$.
- 8) There are no variants available for stack.

Queue

- The Queue is based on FIFO principle.
- Insertion operation is called Enqueue operation.
- Deletion operation is called Dequeue operation.
- Enqueue and Dequeue operation takes place from different end of queue.
- Two pointers are used to perform operations.
- Empty condition is checked using $\text{FRONT} == -1 \& \& \text{REAR} == -1$.
- Full condition is checked using $(\text{FRONT} == 0 \& \& \text{REAR} == \text{Max} - 1) || (\text{Front} == \text{REAR} + 1)$.
- There are three types of variants :- circular, double-ended and priority Queue.

TREES

TREE TERMINOLOGY

1. Node: It is a key component of data structure and it stores the data and address of left child and right child.
2. Edge: Connection between two nodes is known as edges.
3. Parent: The immediate predecessor of a node is called its parent. All the nodes except root node exactly have one parent in level.
4. Child: The immediate successor of node is called child.
5. Sibling: Two or more nodes having same parents are known as siblings.
6. Root: Top most node of Tree is called root of tree. Root Node has no parent.
7. Leaf: All the nodes having no children at any level are called leaf nodes.
8. Path: Path is the interconnection of nodes and edges to move from one node to another node.
9. Level: It is always an integer value which measures the distance from one node to another node in tree. Root is always at level 0 and child of root nodes are at level 1 and grandchild of root nodes are at level 2, and so on.
10. Degree: Maximum number of children that are possible for a node is known as degree of a node.
11. Successor: Left and right subtrees of tree are called

- as successors or children of nodes.
12. Ancestors and Descendant: The parents and grandparents of nodes are called the ancestors and children of a node and children of these children are called descendants.
13. Height / Depth of tree: The depth of a tree is maximum level of any node. It can be also described as the length of the longest path from the root node.
14. Complete Binary Tree: A tree is said to be complete if it has maximum number of nodes at each level except the last level. And if all the nodes at last level appear as far left as possible. A complete tree of depth H has maximum $2^H - 1$ nodes.

Properties of Binary Tree

1. Maximum number of nodes in binary tree at any level will be evaluated by 2^n . where "n" is the level at any given location.
2. Minimum number of nodes in binary Tree with height H is $H+1$.
3. If N is the number of nodes in binary tree then the number of edges E in a non-empty binary tree will be $N-1$.
4. Number of leaf nodes in a binary tree will be calculated with the formula as $N+1$ where N is the number of nodes with exactly two children.
5. For a complete Binary Tree T with N nodes, the

height is $\lceil \log_2(n+1) \rceil$.

6. The number of leaf nodes in a complete Binary Tree containing N nodes is $(N+1)/2$, where N is the number of nodes in a tree.
7. A binary tree of height H has at most $2^{h+1}-1$ elements.

II Representations of Binary Tree in Memory

Memory Representation
of Binary Tree

Array

LinkedList

i) Array Representation of Binary Tree

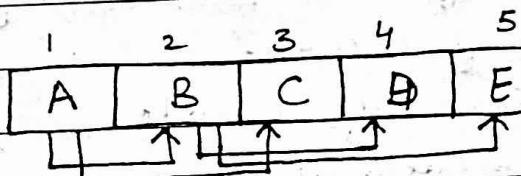
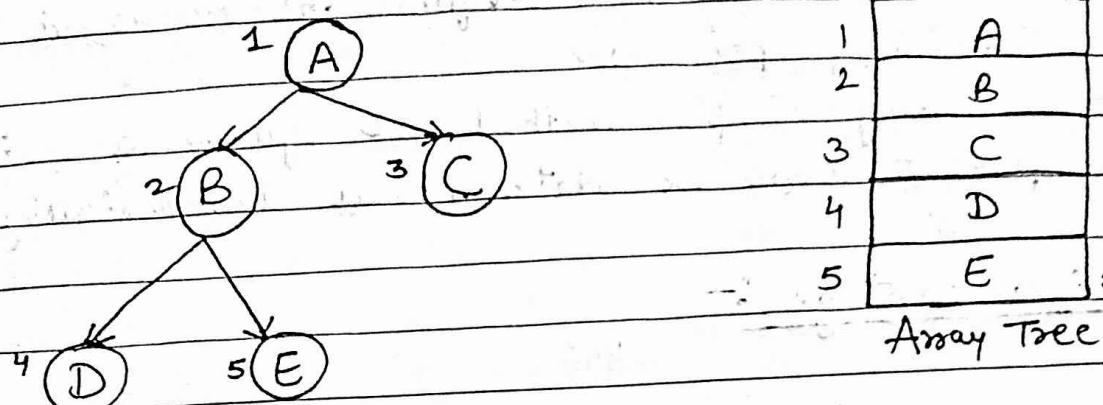
The general data structure, array representation of binary Tree follows the normal convention. If a binary tree is complete binary tree, it can be represented using an array capable of holding n elements where index position -1 in array and left and right child will store at their corresponding position calculated using $2k$ and $2k+1$ where k is the index position of parent node.

a). The 0th position is ignored

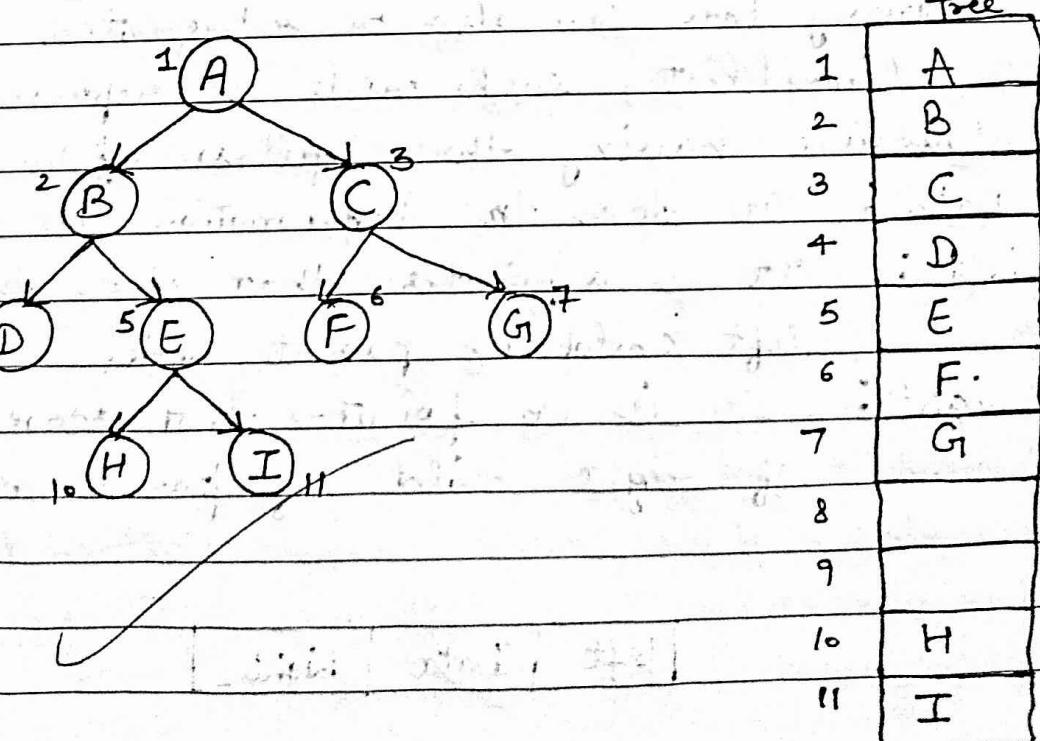
b) i) Left child : $2*k$

ii) Right child : $2*k+1$

c) To find the parent of a given element we will use $\lfloor K/2 \rfloor$. Here, K is the index of particular element.



We can see that an array representation of complete binary Tree does not lead to the waste of any storage. But if you want to represent an incomplete binary tree using array representation, then it leads to the wastage of storage. Ex:-



Advantages :-

1. This method does not require the overhead of maintaining pointers.
2. It is very simple method as given a child node, its parent can be determined immediately.

Disadvantages :-

1. The sequential representation is usually inefficient to use. In case of incomplete Binary tree, a lot of wastage of memory occurs.
2. Insertion or deletion of a node requires considerable data movement up and down the array, using an excessive amount of processing time.

ii) Linked list Representation of Binary Tree

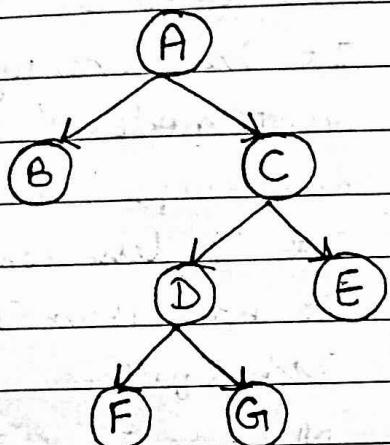
A binary tree can also be represented by linkedlist.

In linked list, each node is represented by a structure having three fields of members.

- a) Data : It stores the information or data.
- b) Left : It is a pointer that stores the address of left child of parent node.
- c) Right : It is a pointer that stores the address of right child of parent node.

Left	Data	Right
------	------	-------

Some programming languages like C, CPP support pointers, so the above mentioned concept can be implemented very easily. But the language which does not support pointer like FORTRAN 70, BASIC and then it is difficult to implement this concept. But the same concept can be implemented using three parallel linear array INFO, LEFT, RIGHT as shown below:-



	LEFT		INFO		RIGHT
1	1	1	A	1	3
2	-1	2	B	2	-1
3	4	3	C	3	5
4	6	4	D	4	7
5	-1	5	E	5	-1
6	-1	6	F	6	-1
7	-1	7	G	7	-1

Binary Tree

Advantages :-

- It is most efficient way to represent binary tree in memory.
- Memory wastage is less as compared to sequential representation.

Disadvantages :-

- Wasted memory space in -1 as shown above.
- Given a node, it is difficult to determine its parent node.
- Its implementation algorithm is more difficult.

Difference Between Graph And Tree

Graph

1. More than one path is allowed.
2. It does not have a root node.
3. Graph can have loops.
4. It is more complex.
5. It has two Traversal Techniques.
6. Here number of edges are not defined.
7. It can be directed or undirected.
8. For finding shortest path in networking graph is used.
9. Represents Data in hierarchical manner.

Tree

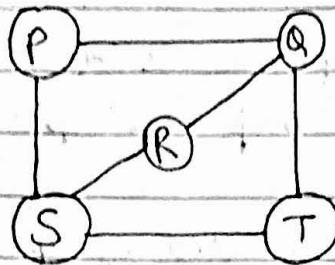
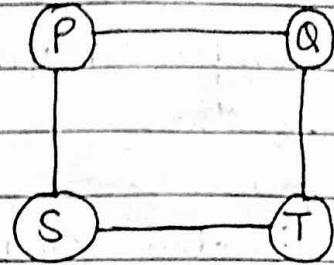
- Only one path between two vertices.
- It has exactly one root node.
- No loops are permitted.
- It is less complex.
- It has Three Traversal Techniques.
- Number of edges are $N-1$ where N is number of nodes.
- Trees are always directed.
- For game trees, decision trees, the tree is used.
- Represents data similar to a network.

Graphs

Types of Graphs :-

1. Simple Graph - A graph is said to be simple graph if there are no ~~longer~~ loop or parallel edges present.
2. Directed graph - In a directed graph every edge of the graph is an ordered pair of vertices connected by the edge.
3. Undirected graph - In this, every edge is an unordered pair of vertices connected by the edge.
4. Complete graph - A graph is said to be complete if there is edge between each vertices.
5. Weighted graph - A graph is said to be a weighted graph if its edges are assigned some value.
6. Isomorphic graph - Two graphs are said to be isomorphic if
 - i) They have the same number of vertex.
 - ii) They have same number of edges.
 - iii) They have an equal number of vertex with a given degree.
7. Tree Graph - A connected graph which does not create cycle is said to be tree graph. In simple words, a connected graph is a tree if it does not create cycle or loops. All trees are graph but all graphs are not tree.
8. Sub Graph - A graph is said to be a sub graph if and only if it contains the vertex set V and Edges set E from another graph G . In other

In words we can say that a graph $G'_i = G_i(V', E')$ is called a subgraph of $G_i = G_i(V, E)$ if the vertices and edges of G'_i are contained in the vertices and edges of G_i .

a) Graph G_i b) Sub graph G'_i of G_i

Terminology of Graphs

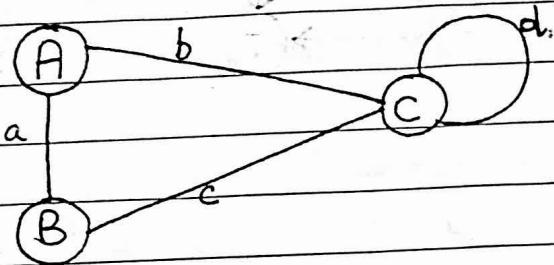
A graph G_i may be defined as a finite set V of vertices and a set E of edges -

- A graph is a non-linear data structure with cycles.
- A graph $G_i = (V, E)$ is finite non-empty set of objects.
- V called set of vertices or nodes.
- E called set of all edges or arcs.

1. Adjacent Vertices - Two vertex are called adjacent if there is an edge between them.
The degree of a vertex in an undirected graph is the number of edges associated with it.
2. Path - A path in a graph is a sequence of vertices $v_1, v_2, v_3, \dots, v_n$ such that v_i, v_{i+1} are in E for $i=1, 2, \dots, (n-1)$. The length of such a path is the number of edges on the path, which is equal

to $(n-1)$. A simple path is a path such that all vertices are distinct, except that the first and last could be the same.

3. Loops - An edge joining only one vertex is called as loop or if it has identical end points.



4. Simple Path -

- A path is said to be simple path if all the vertices on the path except the first and last are distinct.
- If first and last vertices are same, the path will be a cycle.

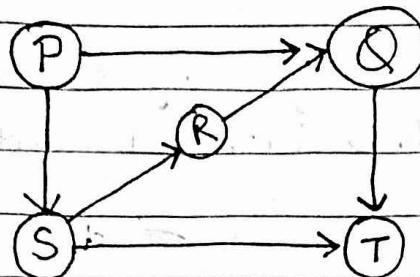
5. Degree - In the undirected graph, number of edges can be connected to the vertex to form a graph. Connected number of edges to the vertex is the degree of the vertex. If G is a graph with m edges, then

$$\sum \deg(v) = 2m$$

An edge (u, v) is counted twice in the summation above; once by its endpoint u and once by its endpoint v . Thus, the total contribution of the edges to the degrees of the vertices is twice the number of edges.

In Degree - This term is used in directed graphs, where direction is also defined with edge. Total

number of incoming edges to the vertex is known as 'In degree' of the vertex.



$$P \rightarrow 0$$

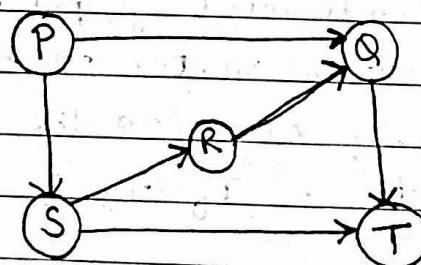
$$Q \rightarrow 2$$

$$R \rightarrow 1$$

$$S \rightarrow 1$$

$$T \rightarrow 2$$

Out-Degree - This term is also used for the directed graph. Total number of outgoing edges from the vertex is known as 'Out Degree' of the vertex.



$$P \rightarrow 2$$

$$Q \rightarrow 1$$

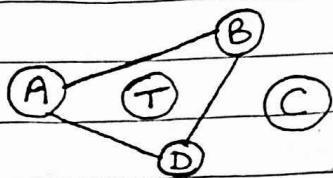
$$R \rightarrow 1$$

$$S \rightarrow 2$$

$$T \rightarrow 0$$

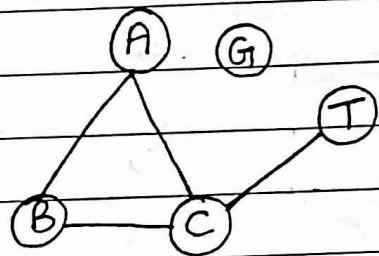
6. **Isolated - Vertex** - A vertex having no incident edge is called an isolated vertex. Or, we can

say that isolated vertices are vertices with zero degree.



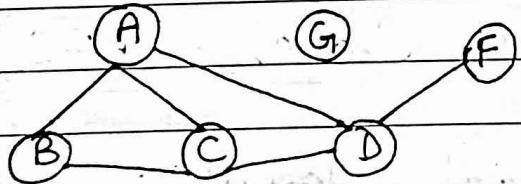
Here, isolated vertices are T and C

7. Pendent Vertex - A vertex of degree one is called pendent or end vertex.



Here Pendent vertex is T

8. Incident edge - The edge (v_i, v_j) is said to be incident if there is any edge (v_i, v_j) .

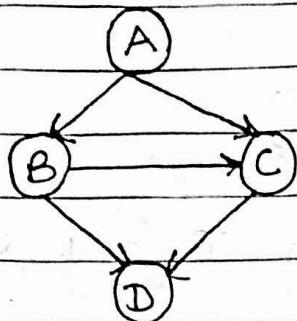


The edges incident on vertex A are (A, B) , (A, C) and (A, D) .

9. Source and Sink

- A vertex V is said to be source if it has its positive out-degree but 0 in-degree.
- A vertex V is said to be sink if it has 0 out-degree but positive in-degree.

degree. e.g.



Vertex	In-degree	Out-Degree
A	0	2
B	1	2
C	2	1
D	2	0

Here vertex A is source and vertex D is sink.

Searching Techniques

① Breadth First Search (BFS)

The BFS traversal algorithm visits each node in a graph as follows :-

- i) Firstly we examine the starting node V, then we examine all the neighbours of V, then we examine all the neighbours of the neighbours of V and so on.
- ii) We need to keep track of the neighbours of a node and we need to guarantee that no node is processed more than once.

- (iii) This can be done by using a queue to hold nodes in that are waiting to be processed.
- iv) A queue is similar to a line of people, that operates on a FIFO Principle.
- v) During the execution of algorithm each node n of Graph G_1 will be in one of the three states called as the status of n as follows:-
- a) Status = 1 (ready state): the initial state of node n
- b) Status = 2 (waiting state): the node n is in the queue waiting to be processed.
- c) Status = 3 (processed state): the node has been processed.

Algorithm :-

- 1) Initialize all the nodes to ready state. ($\text{status} = 1$).
- 2) Put the starting node in queue and change its state to waiting state ($\text{status} = 2$).
- 3) Repeat step 4 and 5 until queue is empty.
- 4) Remove the front node n of queue. Process n and change its status to processed state ($\text{status} = 3$).
- 5) Add all the neighbours of n that are in ready state ($\text{status} = 1$) to the rear of queue and change their status to waiting state ($\text{status} = 2$).
- 6) Exit.

2) Depth First Search (DFS)

The DFS traversal algorithm visits each node in a graph as follows:-

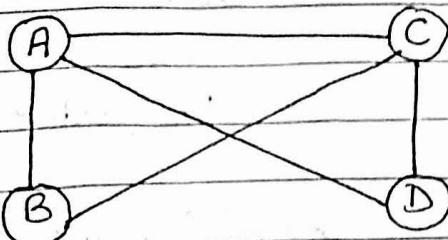
- i) First we examine the starting node V.
- ii) Then we examine each node along a path that begins at V.
- iii) After coming to the end of the path, we backtrack on this path until we can continue along another path and so on.
- iv) Eventually this method will visit every node in a connected graph.
- v) The actual traverse is done using stack. First the starting node is pushed onto the stack. Then the following process repeats:
 - 1) Pop a node off the stack.
 - 2) Traverse or print this current node.
 - 3) Push all the nodes adjacent to the current node onto the stack.
- vi) This process terminates when stack is empty.

Algorithm —

- 1) Initialize all the nodes to the ready state (Status = 1).
- 2) Push the starting node A onto stack and change its status to waiting state (Status = 2).
- 3) Repeat step 4 and 5 until stack is empty.
- 4) Pop the top node N of stack. Process N and change its status to the processed state (Status = 3).
- 5) Push onto stack all the neighbours of N that are still in the ready state (Status = 1) and change their status to the waiting state (Status = 2).
- 6) EXIT.

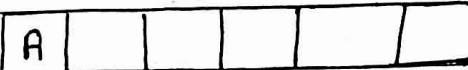
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Here is an example of BFS



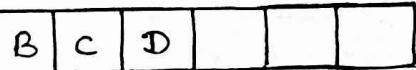
Suppose, we consider node A as a root node. ∵
Traversing would be started from node A.

Now we insert the node A in queue as shown.



Processed = NULL

- * Now we process the node A and insert the adjacent nodes of node A in queue. After processing, node A will be marked as processed.



Processed = A

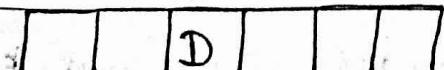
- * Now process the node B and remove it from queue and insert adjacents of node B in queue.



Processed = A, B

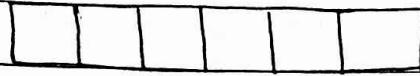
Note that, we have not to insert the processed nodes again in the queue.

- * Now process the node C and remove it from queue and add adjacents of node C in queue.



Processed = A, B, C

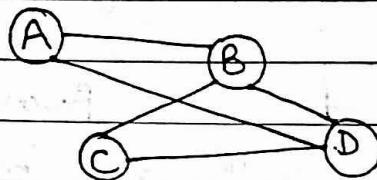
- * Now process the node **D** and remove it from queue and add its adjacents in the Queue.



Processed = A, B, C, D

We see that our queue becomes empty. So the final answer of traversal is **(A), (B), (C), (D)**.

- # Here is an example of DFS



Suppose we consider node **A** as a root node. For DFS Traversal, we use stack.

- * Push the node **A** onto the stack. Process the node **A** and pop it from stack and ~~not~~ push the adjacents of **A** onto stack.

3				3		
2				2		
1				1	D	
0	A			0	B	

After Processing →

Processed = A

- * Process the top element of stack i.e. **D** and **B** after processing pop it from stack and ~~not~~ push adjacents

of (D) onto the stack.

3		
2		
1	C	
0	B	

Processed = A, D

- * Now, Process the top element of stack i.e (C) and after processing pop it from stack and push adjacents of (C) in stack.

3		
2		
1	.	.
0	B	

Processed = A, D, C

- * Now, process the top element of stack i.e (B) and after processing pop it from stack and push its adjacents onto stack.

3	.	
2	.	
1	.	
0		

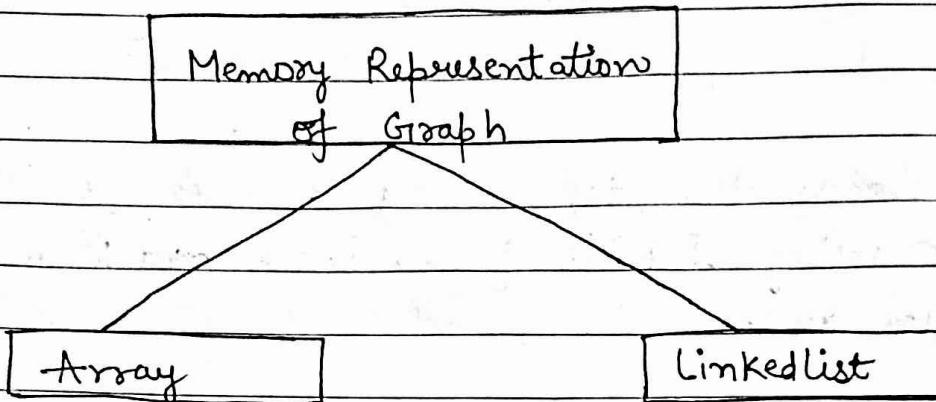
Processed = A, D, C, B

We see that our stack becomes empty. So the final result of DFS Traversal is (A, D, C, B)

Memory Representation Of Graphs

Graphs can be represented in memory by two ways

- 1) Using Array
- 2) Using linkedlist



(i) Representation of Graph in memory using Array :-

Adjacency Matrix :- Suppose G_1 is a simple directed graph with m nodes and suppose the nodes of G_1 have been ordered and are called v_1, v_2, \dots, v_m . Then the adjacency Matrix $A = (a_{ij})$ of graph G_1 is the $m \times m$ matrix defined as follows :-

$$a_{ij} = \begin{cases} 1 & \text{if } v_i \text{ is adjacent to } v_j \\ 0 & \text{otherwise} \end{cases}$$

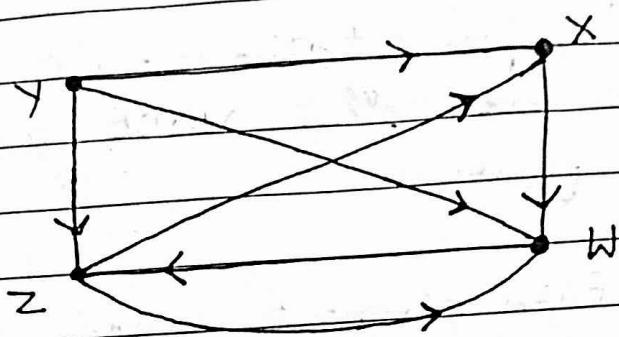
Such matrix A , which contains entries of only 0 and 1 is called bit matrix.

- ① Suppose G_1 is an undirected graph. Then the adjacency matrix A of G_1 will be a symmetric matrix i.e. one in which $a_{ij} = a_{ji}$ for every i and j . This follows from the fact that each undirected edge $[u, v]$ corresponds to the two directed edges

(u, v) and (v, u) .

(2) The above matrix representation of a graph may be extended to multigraphs. If G_1 is a multigraph then the adjacency Matrix of G_1 is the $m \times m$ matrix $A = (a_{ij})$ defined by setting a_{ij} equal to the number of edges from v_i to v_j .

Example :-



Suppose the nodes are stored in memory in a linear array DATA as follows :

DATA : X, Y, Z, W

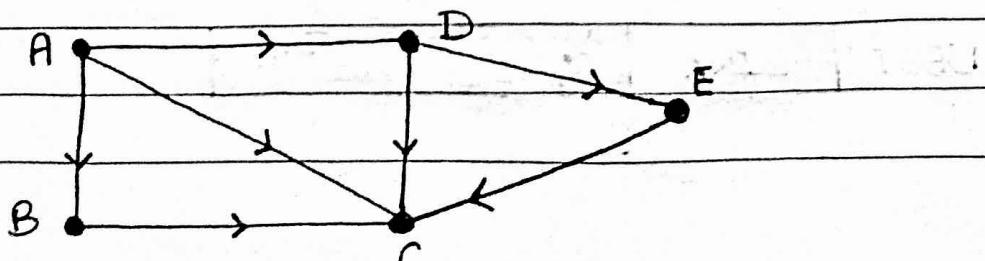
Then we assume that the ordering of the nodes in G_1 is as follows : $v_1 = X, v_2 = Y, v_3 = Z, v_4 = W$.

The adjacency Matrix A of G_1 is as follows :

$$A = \begin{bmatrix} 0 & 0 & 0 & 1 \\ 1 & 0 & 1 & 1 \\ 1 & 0 & 0 & 1 \\ 0 & 0 & 1 & 0 \end{bmatrix}$$

Note that the number of 1's in A is equal to the number of edges in G_1 .

(ii) Representation of Graph using linkedlist :-



Node	Adjacency List
A	B, C, D
B	C
C	.
D	C, E
E	C

The linked representation will contain two lists (or files) a node list NODE and an edge list EDGE, as follows.

- a) Node List → Each element in the list NODE will correspond to a node in G_1 , and it will be a record of form:

NODE	NEXT	ADJ	[Shaded Area]
------	------	-----	---------------

Here NODE will be the name or key value of node. NEXT will be a pointer to next node in the list NODE and ADJ will be a pointer to the first element in the adjacency list of the node, which is maintained in the list EDGE. The shaded area indicates that there may be other information in record such as indegree or outdegree or status of node.

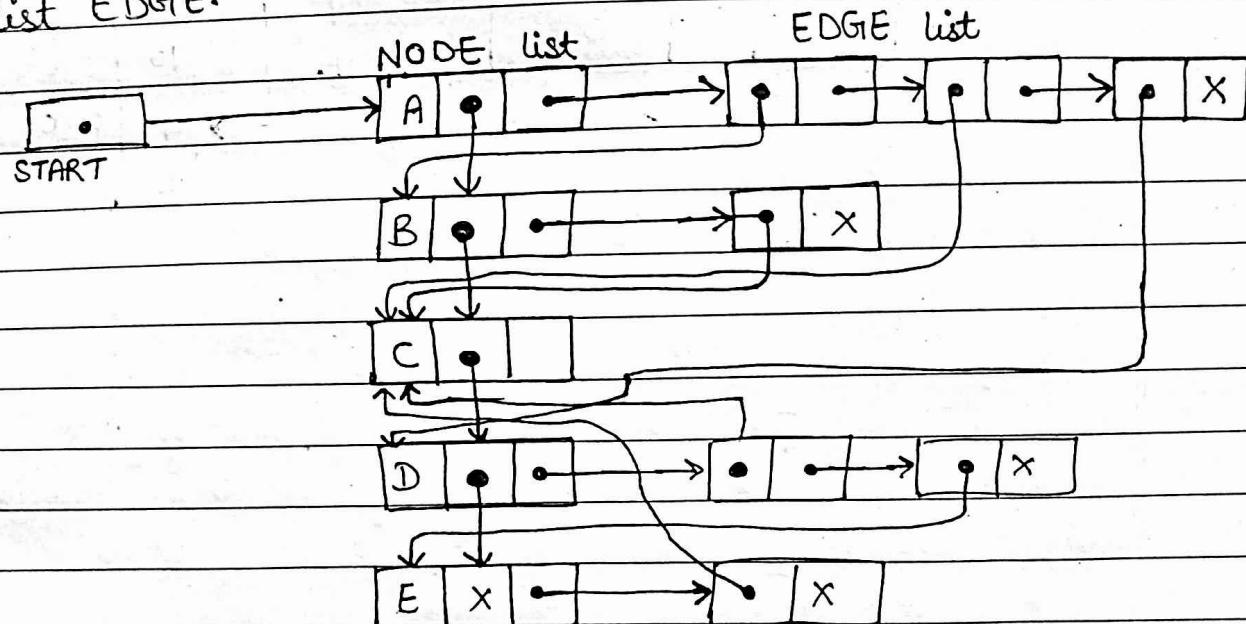
- b) Edge List → Each element in the list EDGE will correspond to an edge of G_1 will be a record of the form:

DEST	LINK	[Shaded Area]
------	------	---------------

The field DEST will point to the location in the list NODE of the destination or terminal node of edge.

The field LINK will link together the edges with the same initial node, ie: nodes in the same adjacency list.

The shaded area indicates that there may be other information in record corresponding to the edge, such as a field EDGE containing the labelled data of the edge when G is a labelled graph. We also need a pointer variable AVAILABLE for the list of available space in the list EDGE.



Schematic diagram of linked representation of G in memory.

	NODE	NEXT	ADJ	DEST	LINK
START	4	5	3	1	7
	5	C	9	2	5
	6	A	8	3	10
AVAILN	7			7 (B)	
	8			4	9 (D)
	9	E	1	5	8
	10	B	0	6	2 (C) 0
	11			7	6 (E) 0
	12			8	9
	13	D	2	9	12
	14		10	10	2 (C) 4
	15		0	11	2 (C) 0
	16			12	0

Algorithm Complexity

It is very rare to have a single algorithm for solving a problem. How efficient a particular algorithm is for solving a problem is of no concern if the amount of data to be processed is very large. We know that when a program implementing an algorithm is executed, it uses resources of the computing system such as CPU, memory, etc. The study of algorithm is necessary to determine the amount of time and storage space an algorithm may require for execution.

The complexity of an algorithm is a function $f(n)$ that gives the running time and memory space required by an algorithm for processing the input data of size n .

Space Complexity

The analysis of algorithm based on how much memory an algorithm needs to solve a particular problem is called the space complexity of an algorithm. It is expressed as a function $f(n)$ that describes the amount of memory an algorithm takes for processing n input data elements.

The space required by the algorithm when implemented not only includes instruction space but also includes data space.

The study of space complexity of an algorithm is of major concern when programs implementing algorithms are made to run on a multi-user system when storage space is shared among users. Although space complexity is important but inexpensive memory has reduced the significance of space complexity. Thus our main area of concern is the

time complexity that focuses on amount of time elapsed when processing data.

Time Complexity

The analysis of algorithm based on time of computation is called time complexity of an algorithm. The main objective of time complexity is to compare the performance of different algorithms in solving the same problem. As we are more interested in time complexity so we have to decide how to measure it. One possible approach is to implement the algorithm used for solving a particular problem using a programming language and run them. But this method has a number of pitfalls.

1. Time is wasted in implementing each of the algorithm as ^{only} finally one of these will be used.
2. The machine on which the program runs might influence its running time as time complexity in such a situation depends on a number of factors as follows:-
 - RAM capacity of computer.
 - Operating system used in computer.
 - Quality of code generated by the compiler.

A better method is to employ mathematical model that analyzes algorithm independently of specific implementations, computers or data. For ex:- In searching algs, the key operations are the number of comparisons made and in sorting algorithms the key operations are swapping and comparison. In sorting algorithms, time complexity will be small if the number of elements are to be processed are small compared to when large number of elements are to be processed.

in former case the total number of comparisons made will be smaller than the later case. Thus, the time complexity of an algorithm is a function $f(n)$ that describes the efficiency of an algorithm in terms of size n of input data. As time complexity is dependent on the key operations rather than all operations so we usually express complexity function $f(n)$ as proportional to input size (n) rather than exact function. Now let us consider a few segments of algorithm performing some key operations where we have to measure the time complexity.

1. A segment of algorithm that involves simple loop.

```
for ( i=1 ; i<501 ; i++ )  
    { :statement(s);  
    }
```

In above segment, we need to know that how many times the body of loop is executed. It is 500 times. As Time complexity is directly proportional to the number of iterations so for loop factor M , it can be generally expressed as $f(n) \propto n$.

2. Now, let us consider another loop

```
for ( i=1 ; i<501 ; i=i+2 )  
    { :statements;  
    }
```

In above case, the number of iteration is half the loop factor as step size is 2. So the time complexity of this loop is proportional to half the loop factor.

$$f(n) \propto n/2$$

3. Now let us consider an example in which loop counter is multiplied or divide each iteration.

```
for (i=1; i<=500; i=i*2)
{
    Statement(s);
}
```

```
for (i = 500; i >= 1; i = i/2)
{
    statements;
}
```

In above two segments of algorithm, we can not say that the body of loop iterates 500 times in each case as the number of times we iterate is governed by loop counter variable i which does not change in a linear fashion. In first case, it changes by a multiple of 2 and in the second case it changes by dividing it by 2.

For multiply

$\begin{matrix} \text{Iteration} \\ 2 \\ \leq 500 \end{matrix}$

For Divide

$\begin{matrix} 500/2 \\ \text{Iteration} \\ >= 1 \end{matrix}$

Expressing it in term of logarithms, we can say that there are $\log_2 500$ i.e. 9 iterations. So, for loop factor n , time complexity is proportional to $\log_2 n$.

i.e. $f(n) \propto \log_2 n$

4. Now let us consider an example of nested loop where we need to determine how many times each loop iterates. In nested loop, the total no. of iterations is the product of number of iterations in the inner loop and the outer loop.

```
a) for (i=1; i<=20; i++),
{
    for (j=1; j<=20; j++)
    {
        statements;
    }
}
```

If both loop iterates n times then the time complexity which is proportional to no. of iterations i.e.

$$f(n) \propto n^2$$

b) $\text{for } (i=1; i \leq 20; i++)$
 { $\text{for } (j=1; j \leq 20; j = j \times 2)$
 { Statement(s);
 } . . .
 }

Here the outer loop iterates 20 times and inner loop iterates $\log_2 20$ times as loop factor J changes by multiple of 2
 $\therefore f(n) \propto n \log n$

c) $\text{for } (i=1; i \leq 20; i++)$
 { $\text{for } (j=1; j \leq i; j++)$
 { Statement(s);
 } . . .
 }

As in previous case, the outer loop iterates 20 times but in this case the number of times inner loop iterates is dependent on the value of outer loop counter variable I .

if $I=1$, the body of loop iterates only one

if $I=2$, " " " " twice.

!

and so on.

\therefore The number of times the body of inner loop iterates is $1+2+3+\dots+20 = 210$ which is total no. of iterations performed by the nested loops.

Since, the number of times the body of inner loop outer loop iterates is 20 so the number of times inner loop iterates is $210/20 = 21/2$ which is same as the number of iterations (20) plus 1 divided by 2 i.e. $(20+1)/2$.

As we know that in a nested loop,

Total number of iterations = Number of iterations of Outer loop
X Number of iterations of inner loop.

In general, if loop factor is 'n' then time Complexity which is proportional to the number of iteration is
 $f(n) \propto n \cdot (n+1)/2$.

Difference between

Array

1. An array is a grouping of data elements of equivalent datatype.
2. The memory is assigned at compile time.
3. It stores data in a contiguous memory location.
4. The elements are not dependent on each other.
5. It is faster to access the element in an array.
6. Memory utilization is ineffective.
7. In case of array, memory size is fixed and is not possible to change it during run time.

Linklist

- A linkedlist is a group of entities called a node. It has two segments [Data + Address].
- The memory is assigned at run time.
- It stores data randomly in the memory.
- The elements are dependent on each other.
- In this, process of accessing the elements takes time.
- Memory utilization is effective.

In linklist, the placement of element is allocated during the run time.

Asymptotic Notations

Asymptotic notations are the mathematical notations used to describe the running time of an algorithm when the input tends towards a particular value or a limiting value.

For Ex:- In bubble sort, when the input array is already sorted, the time taken by the algorithm is linear i.e. the best case.

But when the input array is in reverse condition, the algorithm takes the maximum time to sort the elements i.e. the worst case.

When the input array is neither sorted nor in reverse order, then it takes average time. These durations are denoted using asymptotic notations.

There are mainly three asymptotic notations :-

- 1) Big O notation
- 2) Omega notation
- 3) Theta notation

1. Big O notation \Rightarrow Big O notation represents upper bound of the running time of an algorithm. Thus, it gives the worst case complexity of an algorithm.

~~graph~~ $O(g(n)) = \{ f(n) : \text{there exists positive constant } C \text{ and } n_0 \text{ such that } 0 \leq f(n) \leq Cg(n) \text{ for all } n \geq n_0 \}$

~~Ex~~ The above expression can be described as a function $f(n)$ belongs to the set $O(g(n))$ if there exist a positive constant C such that it lies between O and $Cg(n)$,

for sufficiently large n .

2. Omega (Ω) notation \Rightarrow Omega notation represents the lower bound of the running time of an algorithm. Thus, it provides the best case complexity of an algorithm.

$$\Omega(g(n)) = \{ f(n) : \text{there exist positive constants } c \text{ and } n_0 \text{ such that } 0 \leq cg(n) \leq f(n) \text{ for all } n \geq n_0 \}$$

The above expression can be described as function $f(n)$ belongs to the set $\Omega(g(n))$ if there exist a positive constant c such that it lies above $cg(n)$, for sufficiently large n .

3. Theta (Θ) notation \Rightarrow Theta notation encloses the function from above and below. Since it represents the upper and the lower bound of the running time of an algorithm, it is used for analyzing the average-case complexity of an algorithm.

$$\Theta(g(n)) = \{ f(n) : \text{there exist +ve constants } C_1, C_2 \text{ and } n_0 \text{ such that } 0 \leq C_1 g(n) \leq f(n) \leq C_2 g(n) \text{ for all } n \geq n_0 \}$$

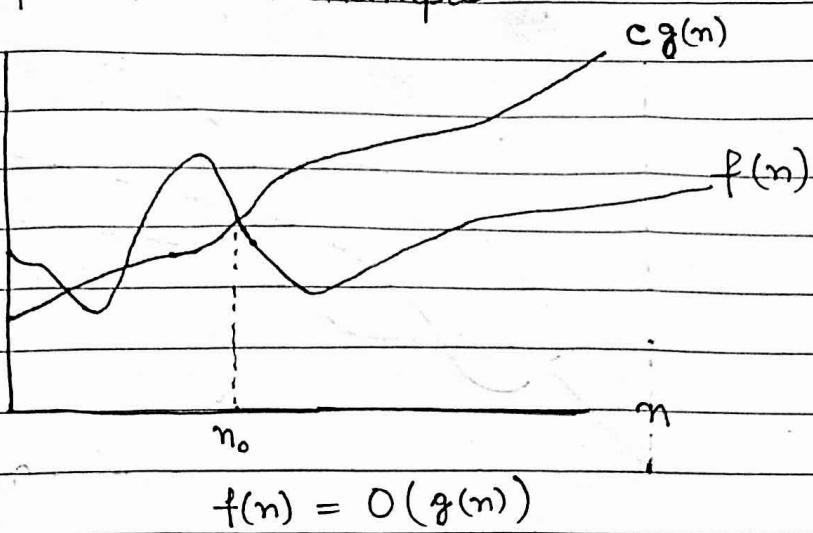
The above expression can be described as a function $f(n)$ belongs to the set $\Theta(g(n))$ if there exist +ve constants C_1 and C_2 such that it can be sandwiched between $C_1 g(n)$ and $C_2 g(n)$, for sufficiently large n .

	Best	Average	Worst
Linear Search	$O(1)$	$O(n)$	$O(n)$
Binary Search	$O(1)$	$O(\log_2(n))$	$O(\log_2(n))$
Bubble Sort	$O(n^2)$	$O(n^2)$	$O(n^2)$
Selection Sort	$O(n^2)$	$O(n^2)$	$O(n^2)$
Insertion Sort	$O(n)$	$O(n^2)$	$O(n^2)$
Quick sort	$O(n \log n)$	$O(n \log n)$	$O(n^2)$
Merge sort	$O(n \log n)$	$O(n \log n)$	$O(n \log n)$

for
for

for
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* Big O Graph with example



Ex:- $f(n) = n^2 + n + 1$
 $= O(n^2)$

let $1 \leq n \leq n^2$

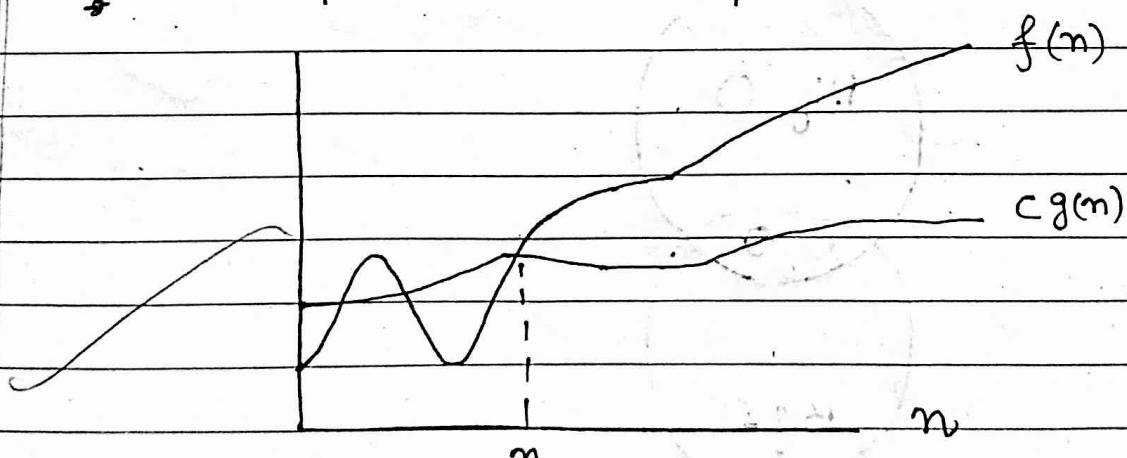
$f(n) \leq n^2 + 2n^2 + n^2$

$f(n) \leq 4n^2$

$\therefore O(n^2)$

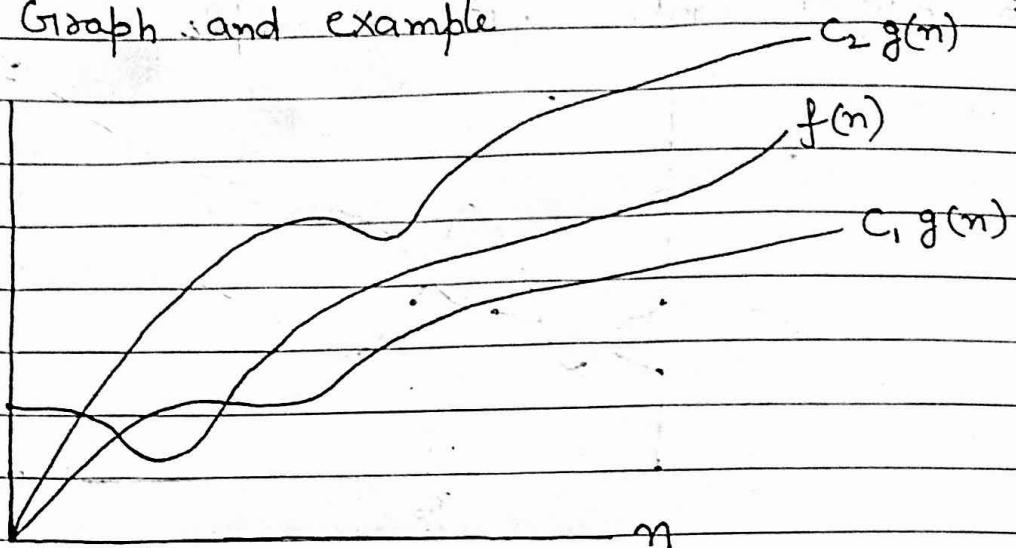
which is of form $c.g(n)$

* ~~Ω~~ Graph with example



$f(n) = \Omega(g(n))$

Ex:- $f(n) = n^2 + 2n + 1$
 $f(n) \geq 1 \cdot n^2$
 $\therefore \Omega(n^2)$

* Theta (Θ) Graph and example

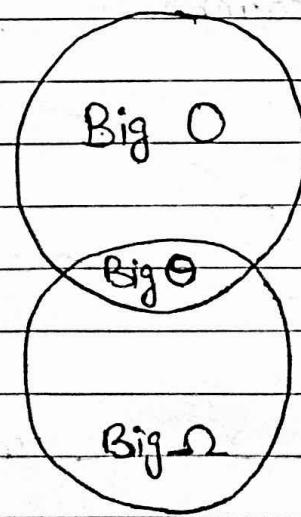
$$f(n) = \Theta(g(n))$$

$$\underline{1 \cdot n^2} \leq f(n) \leq \overline{4 \cdot n^2}$$

Lower Bound

Upper Bound

Big O



~~Surj~~
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