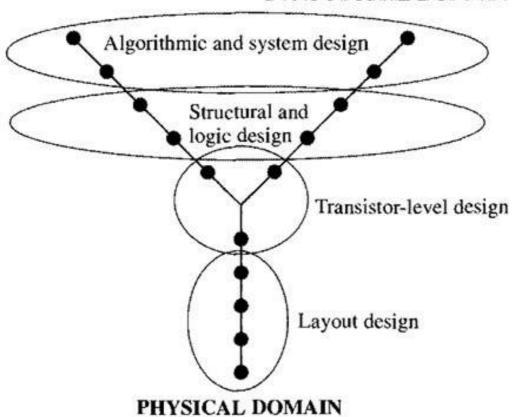
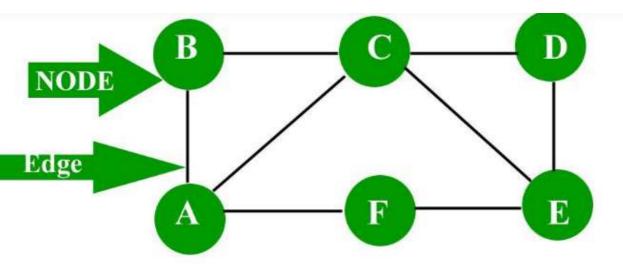
## BEHAVIORAL DOMAIN

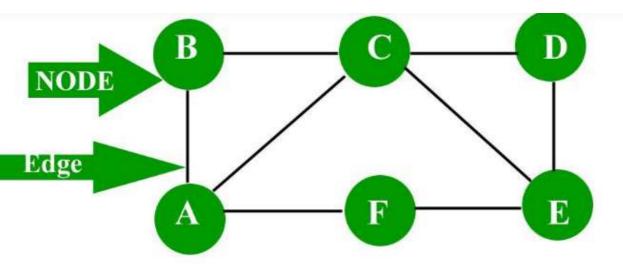
## STRUCTURAL DOMAIN



A Graph is just a way to show connections between things. It is set of edges
and vertices where each edge is associated with unordered pair of vertices.
Graph is a data structure that is defined by two components :

- 1. **Node or Vertex:** It is a point or joint between two lines like people, cities, or websites. In below diagram the nodes are A, B, C, D, E, F.
- 2. **Edge:** It is line or connection between two nodes like connections between them (friendships, roads, links). In the below diagram edges are the connecting lines in between them.





## **Basic Concepts in Graph**

Ordered Pair: Ordered pair is a connection between two nodes u and v which is identified by unique pair (u, v). The pair (u, v) is ordered because (u, v) is not same as (v, u). It is used in case of directed graph to show which vertex is directing to which vertex.

**Unordered Pair:** In this (u, v) that is identified by unique pair(u, v) can be identified as (v, u). In this the order does not matter in which they come, they are treated same. Undirected graphs are its common example.

Weighted Graph: It is a graph (directed or undirected) in which each edge is assigned some numerical value. This value is called a weight. These weights often represent costs, distances, capacities or other quantifiable relationships between vertices.

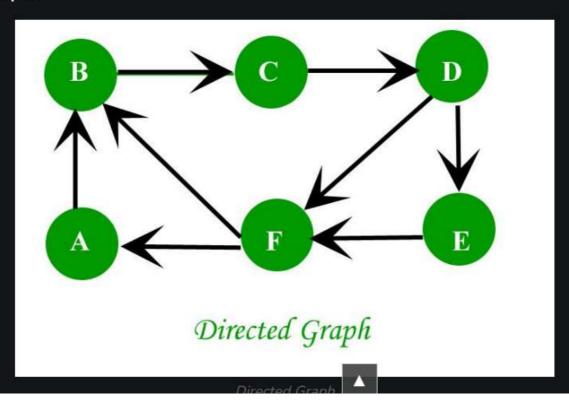
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## Terminologies in Graphs

- 1. Adjacent Node: A node 'v' is said to be adjacent node of node 'u' if and only if there exists an edge between 'u' and 'v'.
- Degree of a Node: In an undirected graph the number of edges incident on a node is the degree of the node. In case of directed graph:
  - Indegree of the node is the number of arriving edges to a node.
  - Outdegree of the node is the number of departing edges to a node.
- 3. Self-Loop: When an edge in graph connects a vertex to itself it is called self-loop. This edge starts and ends at same vertex. A self-loop is counted twice in case of degree of a node.
- 4. The sum of degree of all the vertices in a graph G is even.
- 5. Path: It is a sequence of edges which connects a sequence of distinct vertices. In this sequence of edges no vertices is repeated except in case of closed path, where only first and last vertex can be repeated.
- 6. Isolated Node: A node with degree 0 is known as isolated node. Isolated node can be found by Breadth first search(BFS). It finds its application in LAN network in finding whether a system is con ted or not.

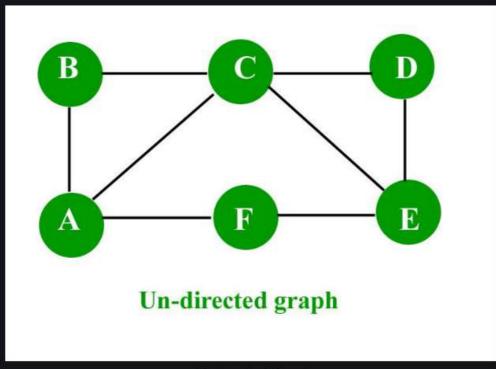
# **Directed Graph**

A graph in which the direction of the edge is defined to a particular node is a directed graph.



# **Undirected Graph**

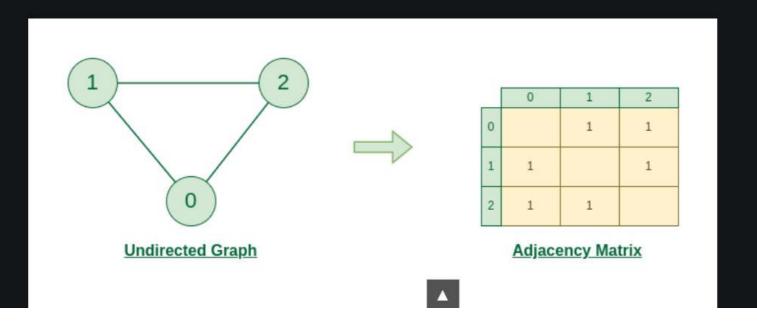
A graph in which the direction of the edge is not defined. So if an edge exists between node 'u' and 'v', then there is a path from node 'u' to 'v' and vice-versa.



Undirected Granh

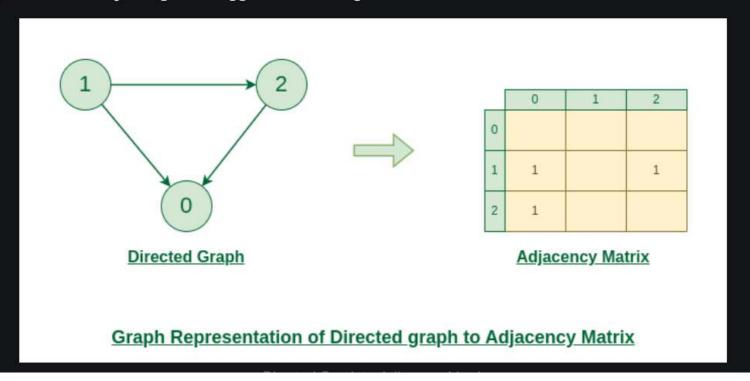
# Representation of Undirected Graph as Adjacency Matrix:

The below figure shows an undirected graph. Initially, the entire Matrix is initialized to **0**. If there is an edge from source to destination, we insert **1** to both cases (adjMat[source][destination] and adjMat[destination][source]) because we can go either way.



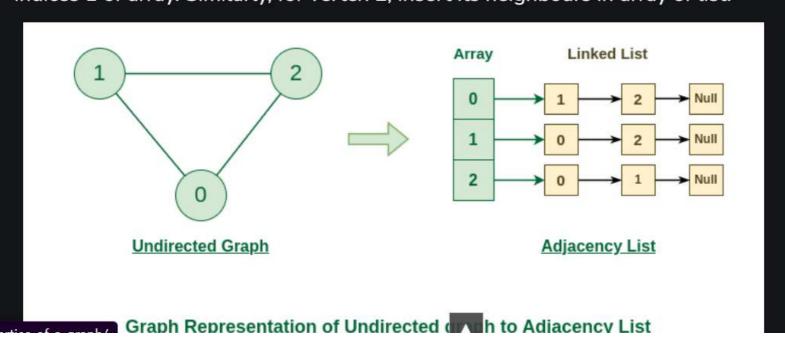
# Representation of Directed Graph as Adjacency Matrix:

The below figure shows a directed graph. Initially, the entire Matrix is initialized to **0**. If there is an edge from source to destination, we insert **1** for that particular **adjMat[source][destination]**.



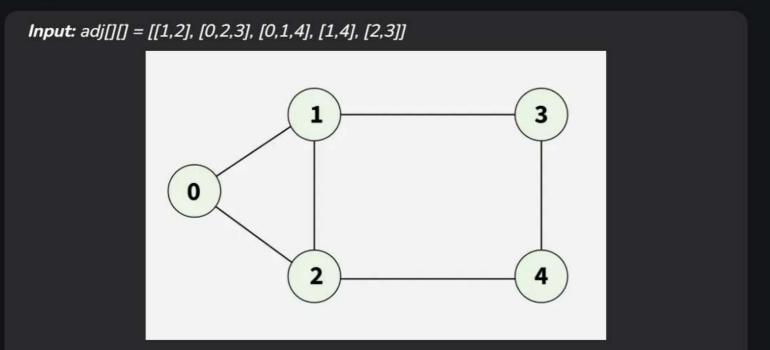
## Representation of Undirected Graph as Adjacency list:

The below undirected graph has 3 vertices. So, an array of list will be created of size 3, where each indices represent the vertices. Now, vertex 0 has two neighbours (i.e, 1 and 2). So, insert vertex 1 and 2 at indices 0 of array. Similarly, For vertex 1, it has two neighbour (i.e, 2 and 0) So, insert vertices 2 and 0 at indices 1 of array. Similarly, for vertex 2, insert its neighbours in array of list.



Given a **undirected graph** represented by an adjacency list adj, where each adj[i] represents the list of vertices connected to vertex i. Perform a **Breadth First Search (BFS)** traversal starting from vertex 0, visiting vertices from left to right according to the adjacency list, and return a list containing the BFS traversal of the graph.

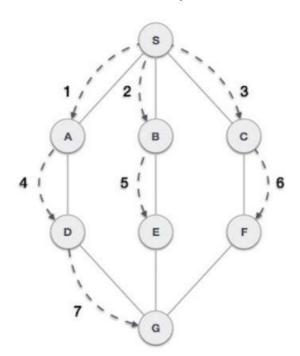
## Examples:

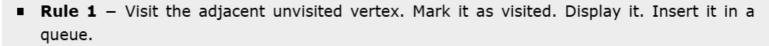


## Breadth First Search (BFS) Algorithm

Breadth First Search (BFS) algorithm traverses a graph in a breadthward motion to search a graph data structure for a node that meets a set of criteria. It uses a queue to remember the next vertex to start a search, when a dead end occurs in any iteration.

Breadth First Search (BFS) algorithm starts at the tree root and explores all nodes at the present depth prior to moving on to the nodes at the next depth level.

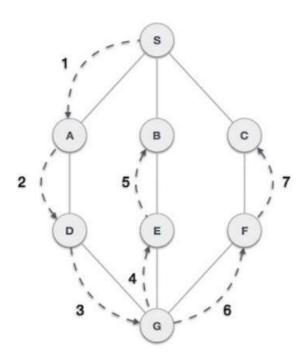




- Rule 2 If no adjacent vertex is found, remove the first vertex from the queue.
- Rule 3 Repeat Rule 1 and Rule 2 until the queue is empty.

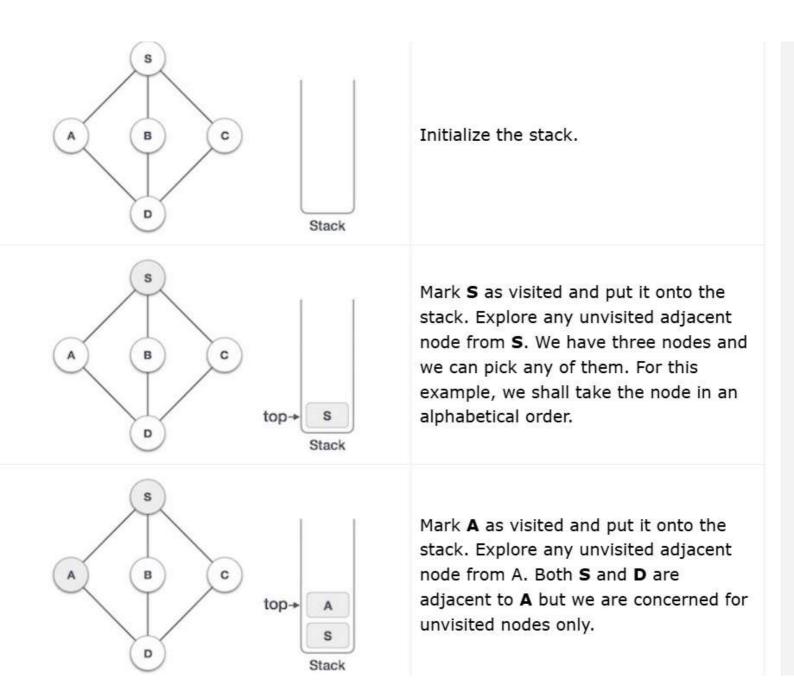
## Depth First Search (DFS) Algorithm

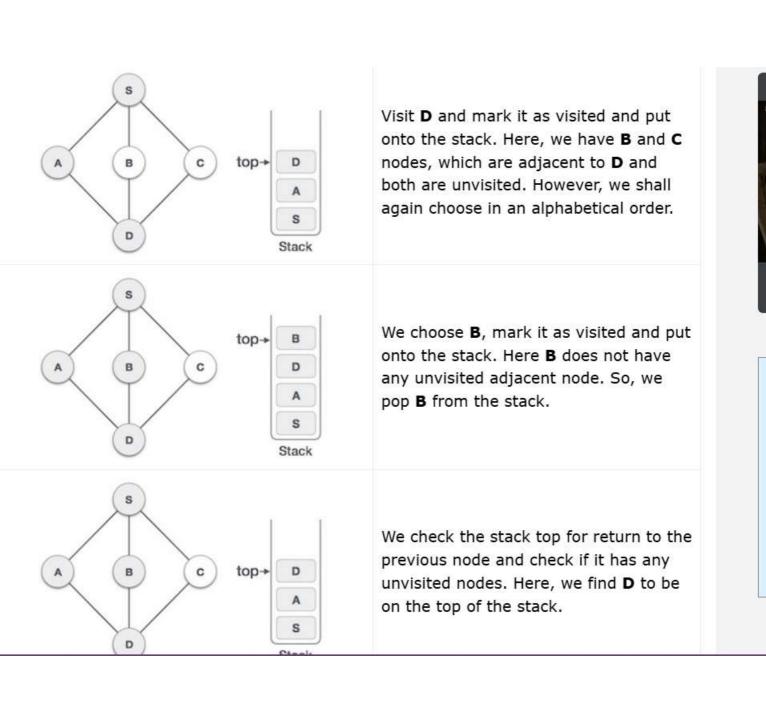
Depth First Search (DFS) algorithm is a recursive algorithm for searching all the vertices of a graph or tree data structure. This algorithm traverses a graph in a depthward motion and uses a stack to remember to get the next vertex to start a search, when a dead end occurs in any iteration.



As in the example given above, DFS algorithm traverses from S to A to D to G to E to B first, then to F and lastly to C. It employs the following rules.

- Rule 1 Visit the adjacent unvisited vertex. Mark it as visited. Display it. Push it in a stack.
- Rule 2 If no adjacent vertex is found, pop up a vertex from the stack. (It will pop up all the vertices from the stack, which do not have adjacent vertices.)
- Rule 3 Repeat Rule 1 and Rule 2 until the stack is empty.





Time Committee
Time Complexity
The time complexity of the DFS algorithm is represented in the form of $O(V + E)$ , where $V$ is the number of nodes and $E$ is the number of edges.
Space Complexity
The space complexity of the DFS algorithm is O(V).

#### **VLSI Design Automation Tools Explained in Simple Terms**

Designing a chip (VLSI) is a complex process with many steps. To make it easier, engineers use **Computer-Aided Design (CAD) tools**. These tools help automate tasks like drawing circuits, checking for errors, and optimizing performance.

Here's a breakdown of the main types of tools used in VLSI design:

#### 1. Algorithmic & System Design Tools

**Purpose:** Help define what the chip should do at a high level (like a blueprint).

- Hardware Description Languages (HDLs) Instead of writing code in C or Python, engineers use special languages like VHDL or Verilog to describe how the chip should work.
- Simulators Test if the design works correctly before building the actual chip.
- High-Level Synthesis (HLS) Converts the high-level description into a lower-level circuit design (like turning software into hardware).
- Silicon Compilers Try to automate the entire chip design process (like a software compiler, but for chips).

### 2. Structural & Logic Design Tools

Purpose: Design and optimize the logic gates and connections inside the chip.

- Schematic Editors Like a digital drawing board where engineers design circuits using logic gates (AND, OR, NOT, etc.).
- Logic Simulators Test if the logic circuits work correctly.
- Logic Synthesis Tools Automatically optimize circuits to make them faster or smaller.
  - Two-Level Logic Optimization Simplifies circuits to use fewer gates (important for fast chips).
  - Multi-Level Logic Optimization Balances speed and chip area for complex designs.
- Timing Analysis Tools Check if signals travel fast enough through the circuit (to avoid delays).

## 3. Transistor-Level Design Tools

Purpose: Work with transistors (the tiny switches that make up logic gates).

- Switch-Level Simulators Treat transistors as simple on/off switches (fast but less accurate).
- Circuit-Level Simulators Use complex math to model real electrical behavior (slow but precise).
- Circuit Extractors After drawing the chip layout, these tools identify all transistors, wires, and electrical properties.

#### 4. Layout Design 100is

**Purpose:** Place and connect all components physically on the chip.

- Placement Tools Decide where each block (like a memory unit or processor) should go on the chip.
- Routing Tools Draw the wires connecting all the blocks without overlaps.
- Floorplanning Tools Plan the chip's overall structure early in the design (like a floor plan for a house).
- Cell Compilers & Module Generators Automatically create small, reusable circuit blocks (like adders or memory cells).
- Design Rule Checkers (DRC) Ensure the layout follows manufacturing rules (like minimum wire spacing).
- Symbolic Layout & Compaction Tools Allow designers to draw circuits without worrying about exact sizes; the tool adjusts everything to fit manufacturing rules.

#### 5. Verification Tools

**Purpose:** Make sure the chip works correctly before manufacturing.

- Simulation Tests the design with sample inputs (but can't check every possible case).
- Formal Verification Uses math to prove the chip will always work correctly (no guesswork).
- **Rapid Prototyping** Builds a test version using programmable chips (FPGAs) to check real-world performance.

## 6. Design Management Tools

Purpose: Keep track of all files, versions, and teamwork in big projects.

- Version Control Like "Google Docs for chip design," tracks changes and lets engineers collaborate.
- Database Systems Store huge amounts of design data efficiently.
- Framework Tools Help different CAD tools work together smoothly (like a universal adapter for software).

## 29.1. Tractable and Intractable Problems

 Let's start by reminding ourselves of some common functions, ordered by how fast they grow.

constant	O(1)
logarithmic	$O(\log n)$
linear	O(n)
n-log-n	$O(n \times \log n)$
quadratic	$O(n^2)$
cubic	$O(n^3)$
exponential	$O(k^n)$ , e.g. $O(2^n)$
factorial	O(n!)
super-exponential	e.g. $O(n^n)$

Computer Scientists divide these functions into two classes:

**Polynomial functions:** Any function that is  $O(n^k)$ , i.e. bounded from above by  $n^k$  for some constant k.

E.g. 
$$O(1)$$
,  $O(\log n)$ ,  $O(n)$ ,  $O(n \times \log n)$ ,  $O(n^2)$ ,  $O(n^3)$ 

This is really a different definition of the word 'polynomial' from the one we had in a previous lecture. Previously, we defined 'polynomial' to be any function of the form  $a_k n^k + a_{k-1} n^{k-1} + \ldots + a_1 n + a_0$ .

But here the word 'polynomial' is used to lump together functions that are bounded from above by polynomials. So,  $\log n$  and  $n \times \log n$ , which are not polynomials in our original sense, are polynomials by our alternative definition, because they are bounded from above by, e.g., n and  $n^2$  respectively.

Exponential functions: The remaining functions.

E.g. 
$$O(2^n), O(n!), O(n^n)$$

This is a real abuse of terminology. A function of the form  $k^n$  is genuinely exponential. But now some functions which are worse than polynomial but not quite exponential, such as  $O(n^{\log n})$ , are also (incorrectly) called exponential. And some functions which are worse than exponential, such as the superexponentials, e.g.  $O(n^n)$ , will also (incorrectly) be called exponential. A better word than 'exponential' would be 'super-polynomial'. But 'exponential' is what everyone uses, so it's what we'll use.

Why have we lumped functions together into these two broad classes? The next two
tables and the graph attempt to show you why.

	10	50	100	300	1000
5n	50	250	500	1500	5000
$n \times \log n$	33	282	665	2469	9966
$n^2$	100	2500	10000	90000	1 million (7 digits)
$n^3$	1000	125000	1 million (7 digits)	27 million (8 digits)	1 billion (10 digits)
$2^n$	1024	a 16-digit number	a 31-digit number	a 91-digit number	a 302-digit number
n!	3.6 million (7 digits)	a 65-digit number	a 161-digit number	a 623-digit number	unimaginably large
$n^n$	10 billion (11 digits)	an 85-digit number	a 201-digit number	a 744-digit number	unimaginably large

(The number of protons in the known universe has 79 digits.)
(The number of microseconds since the Big Bang has 24 digits.)

**Polynomial-Time Algorithm:** an algorithm whose order-of-magnitude time performance is bounded from above by a polynomial function of n, where n is the size of its inputs.

**Exponential Algorithm:** an algorithm whose order-of-magnitude time performance is not bounded from above by a polynomial function of n.

Why do we divide algorithms into these two broad classes? The next table, which
assumes that one instruction can be executed every microsecond, attempt to show
you why.

	10	20	50	100	300
$n^2$	$\frac{1}{10000}$ second	$\begin{array}{c} \frac{1}{2500} \\ \text{second} \end{array}$	$\begin{array}{c} \frac{1}{400} \\ \text{second} \end{array}$	$\frac{1}{100}$ second	$\begin{array}{c} \frac{9}{100} \\ \text{second} \end{array}$
$n^5$	$\frac{1}{10}$ second	3.2 seconds	5.2 minutes	2.8 hours	28.1 days
$2^n$	$\frac{1}{1000}$ second	1 second	35.7 years	400 trillion centuries	a 75-digit number of centuries
$n^n$	2.8 hours	3.3 trillion years	a 70-digit number of centuries	a 185-digit number of centuries	a 728-digit number of centuries

Tractable Problem: a problem that is solvable by a polynomial-time algorithm.  The upper bound is polynomial.
Intractable Problem: a problem that cannot be solved by a polynomial-time algorithm. The lower bound is exponential.
Here are examples of tractable problems (ones with known polynomial-time algorithms):
- Searching an unordered list
- Searching an ordered list
- Sorting a list
<ul> <li>Multiplication of integers (even though there's a gap)</li> </ul>

- Finding a minimum spanning tree in a graph (even though there's a gap)

- Here are examples of intractable problems (ones that have been proven to have no polynomial-time algorithm).
  - Some of them require a non-polynomial amount of output, so they clearly will take a non-polynomial amount of time, e.g.:
    - \* Towers of Hanoi: we can prove that any algorithm that solves this problem must have a worst-case running time that is at least  $2^n 1$ .
    - \* List all permutations (all possible orderings) of n numbers.
  - Others have polynomial amounts of output, but still cannot be solved in polynomial time:
    - \* For an  $n \times n$  draughts board with an arrangement of pieces, determine whether there is a winning strategy for White (i.e. a sequence of moves so that, no matter what Black does, White is guaranteed to win). We can prove that any algorithm that solves this problem must have a worst-case running time that is at least  $2^n$ .
- So you might think that problems can be neatly divided into these two classes. But
  this ignores 'gaps' between lower and upper bounds. Incredibly, there are problems
  for which the state of our knowledge is such that the gap spans this coarse division
  into tractable and intractable. So, in fact, there are three broad classes of problems:
  - Problems with known polynomial-time algorithms.
  - Problems that are provably intractable (proven to have no polynomial-time algorithm).



# **Backtracking Algorithm**

A backtracking algorithm is a problem-solving algorithm that uses a **brute force approach** for finding the desired output.

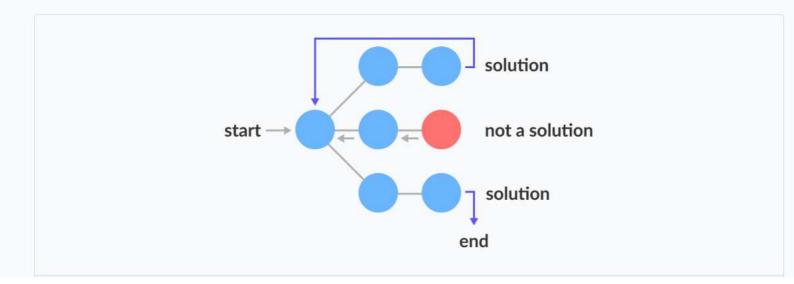
The Brute force approach tries out all the possible solutions and chooses the desired/best solutions.

The term backtracking suggests that if the current solution is not suitable, then backtrack and try other solutions. Thus, recursion is used in this approach.

This approach is used to solve problems that have multiple solutions. If you want an optimal solution, you must go for dynamic programming.

# **State Space Tree**

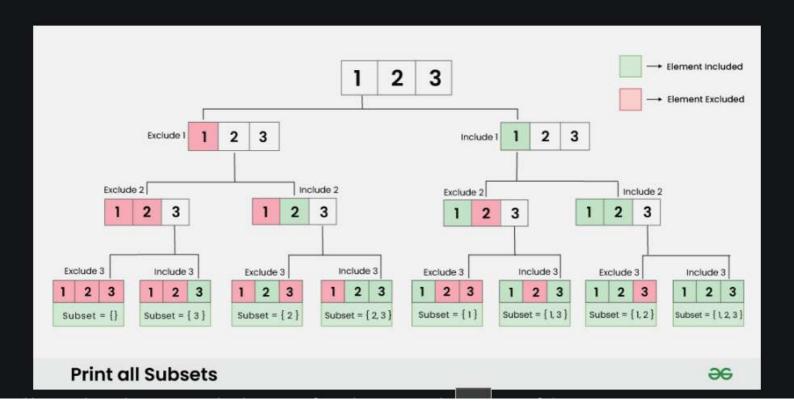
A space state tree is a tree representing all the possible states (solution or nonsolution) of the problem from the root as an initial state to the leaf as a terminal state.



# **Backtracking Algorithm**

```
Backtrack(x)
    if x is not a solution
        return false
    if x is a new solution
        add to list of solutions
    backtrack(expand x)
```

Suppose an array of size 3 having elements [1, 2, 3], the state space tree can be constructed as below:



The **branch and bound algorithm** is a technique used for solving combinatorial optimization problems. First, this algorithm breaks the given problem into multiple sub-problems and then using a bounding function, it eliminates only those solutions that cannot provide optimal solution.

**Combinatorial optimization problems** refer to those problems that involve finding the best solution from a finite set of possible solutions, such as the 0/1 knapsack problem, the travelling salesman problem and many more.

## When Branch and Bound Algorithm is used?

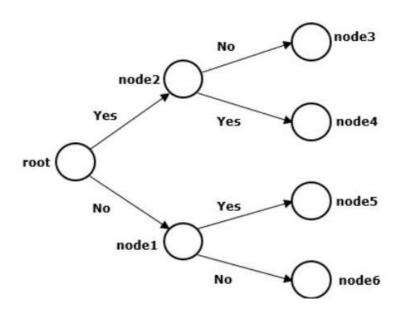
The branch and bound algorithm can be used in the following scenario -

- Whenever we encounter an optimization problem whose variables belong to a discrete set. These types of problems are known as discrete optimization problems.
- As discussed earlier, this algorithm is also used to solve combinatorial optimization problem.
- If the given problem is a mathematical optimization problem, then the branch and bound algorithm can also be applied.

## How does Branch and Bound Algorithm work?

The branch and bound algorithm works by exploring the search space of the problem in a systematic way. It uses a tree structure (state space tree) to represent the solutions and their extensions. Each node in the tree is part of the partial solution, and each edge corresponds to an extension of this solution by adding or removing an element. The root node represents the empty solution.

The algorithm starts with the **root node** and moves towards its **children nodes**. At each level, it evaluates whether a child node satisfies the constraints of the problem to achieve a feasible solution. This process is repeated until a **leaf node** is reached, which represents a complete solution.



### Searching techniques in Branch and Bound

There are different approaches to implementing the branch and bound algorithm. The implementation depends on how to generate the children nodes and how to search the next node to expand. Some of the common searching techniques are –

- **Breadth-first search** It maintains a queue of nodes to expand, which means this searching technique uses First in First out order to search next node.
- Least cost search This searching technique works by computing bound value of each node. The algorithm selects the node with the lowest bound value to expand next.
- **Depth-first search** It maintains a stack of nodes to expand, which means this searching technique uses Last in First out order to search the next node.

# Types of Branch and Bound Solutions

The branch and bound algorithm can produce two types of solutions. They are as follows –

- Variable size solution This type of solution is represented by a subset which is the optimal solution to the given problem.
- Fixed-size solution This type of solution is represented by 0s and 1s.

# **Dynamic Programming**

Dynamic Programming is a technique in computer programming that helps to efficiently solve a class of problems that have overlapping subproblems and optimal substructure property.

If any problem can be divided into subproblems, which in turn are divided into smaller subproblems, and if there are overlapping among these subproblems, then the solutions to these subproblems can be saved for future reference. In this way, efficiency of the CPU can be enhanced. This method of solving a solution is referred to as dynamic programming.

Such problems involve repeatedly calculating the value of the same subproblems to find the optimum solution.

# **Dynamic Programming Example**

Let's find the fibonacci sequence upto 5th term. A fibonacci series is the sequence of numbers in which each number is the sum of the two preceding ones. For example, 0,1,1, 2, 3. Here, each number is the sum of the two preceding numbers.

#### **Algorithm**

```
Let n be the number of terms.

1. If n = 0, return 0.

2. If n = 1, return 1.

3. Else, return the sum of two preceding numbers.
```

We are calculating the fibonacci sequence up to the 5th term.

- 1. The first term is 0.
- 2. The second term is 1.
- 3. The third term is sum of 0 (from step 1) and 1(from step 2), which is 1.
- 4. The fourth term is the sum of the third term (from step 3) and second term (from step 2) i.e. 1 + 1 = 2.
- 5. The fifth term is the sum of the fourth term (from step 4) and third term (from step 3) i.e. 2 + 1 = 3.

# **How Dynamic Programming Works**

Dynamic programming works by storing the result of subproblems so that when their solutions are required, they are at hand and we do not need to recalculate them.

This technique of storing the value of subproblems is called memoization. By saving the values in the array, we save time for computations of sub-problems we have already come across.

```
var m = map(0 → 0, 1 → 1)
function fib(n)
  if key n is not in map m
    m[n] = fib(n - 1) + fib(n - 2)
  return m[n]
```

Dynamic programming by memoization is a top-down approach to dynamic programming. By reversing the direction in which the algorithm works i.e. by starting from the base case and working towards the solution, we can also implement dynamic programming in a bottom-up manner.

```
function fib(n)
  if n = 0
    return 0
  else
    var prevFib = 0, currFib = 1
    repeat n - 1 times
       var newFib = prevFib + currFib
       prevFib = currFib
       currFib = return currentFib
```

#### What is Linear Programming?

**Linear programming** or **Linear optimization** is a technique that helps us to find the optimum solution for a given problem, an optimum solution is a solution that is the best possible outcome of a given particular problem.

In simple terms, it is the method to find out how to do something in the best possible way. With limited resources, you need to do the optimum utilization of resources and achieve the best possible result in a particular objective such as least cost, highest margin, or least time.

The situation that requires a search for the best values of the variables subject to certain constraints is where we use linear programming problems. These situations cannot be handled by the usual calculus and numerical techniques.

#### **Linear Programming Definition**

Linear programming is the technique used for optimizing a particular scenario. Using linear programming provides us with the best possible outcome in a given situation. It uses all the available resources in a manner such that they produce the optimum result.

#### **Components of Linear Programming**

The basic components of a linear programming (LP) problem are:

- Decision Variables: Variables you want to determine to achieve the optimal solution.
- Objective Function: Mathematical equation that represents the goal you want to achieve
- Constraints: Limitations or restrictions that your decision variables must follow.
- Non-Negativity Restrictions: In some real-world scenarios, decision variables cannot be negative

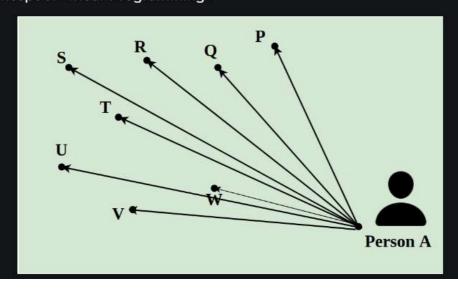
#### **Additional Characteristics of Linear Programming**

- Finiteness: The number of decision variables and constraints in an LP problem are finite.
- Linearity: The objective function and all constraints must be linear functions of the decision variables. It means the degree of variables should be one.

### **Linear Programming Examples**

We can understand the situations in which Linear programming is applied with the help of the example discussed below,

Suppose a delivery man has to deliver 8 packets in a day to the different locations of a city. He has to pick all the packets from A and has to deliver them to points P, Q, R, S, T, U, V, and W. The distance between them is indicated using the lines as shown in the image below. The shortest path followed by the delivery man is calculated using the concept of Linear Programming.



#### **Linear Programming Problems**

**Linear Programming Problems (LPP)** involve optimizing a linear function to find the optimal value solution for the function. The optimal value can be either the maximum value or the minimum value.

In LPP, the linear functions are called **objective functions**. An objective function can have multiple variables, which are subjected to conditions and have to satisfy the **linear constraints**.

#### Types of Linear Programming Problems

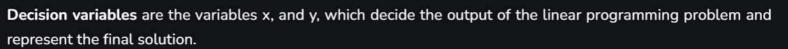
There are many different linear programming problems(LPP) but we will deal with three major linear programming problems in this article.

#### **Manufacturing Problems**

Manufacturing problems are a problem that deals with the number of units that should be produced or sold to maximize profits when each product requires fixed manpower, machine hours, and raw materials.

#### **Diet Problems**

It is used to calculate the number of different kinds of constituents to be included in the diet to get the minimum cost, subject to the availability of food and their prices.



The **objective function**, generally represented by Z, is the linear function that needs to be optimized according to the given condition to get the final solution.

The **restrictions** imposed on decision variables that limit their values are called constraints.

Now, the general formula of a linear programming problem is,

Objective Function: Z = ax + by

Constraints:  $cx + dy \ge e$ ,  $px + qy \le r$ 

**Non-Negative restrictions:**  $x \ge 0$ ,  $y \ge 0$ 

In the above condition x, and y are the decision variables.

#### What is Simulated Annealing?

Simulated Annealing is an optimization algorithm designed to search for an optimal or near-optimal solution in a large solution space. The name and concept are derived from the process of annealing in metallurgy, where a material is heated and then slowly cooled to remove defects and achieve a stable crystalline structure. In Simulated Annealing, the "heat" corresponds to the degree of randomness in the search process, which decreases over time (cooling schedule) to refine the solution. The method is widely used in combinatorial optimization, where problems often have numerous local optima that standard techniques like gradient descent might get stuck in. Simulated Annealing excels in escaping these local minima by introducing controlled randomness in its search, allowing for a more thorough exploration of the solution space.

#### **How Simulated Annealing Works**

The algorithm starts with an initial solution and a high "temperature," which gradually decreases over time. Here's a step-by-step breakdown of how the algorithm works:

- Initialization: Begin with an initial solution So and an initial temperature To. he temperature controls how likely the algorithm is to accept worse solutions as it explores the search space.
- Neighborhood Search: At each step, a new solution S' is generated by making a small change (or perturbation) to the current solution S.
- Objective Function Evaluation: The new solution S' is evaluated using the objective function. If S' provides a better solution than S, it is accepted as the new solution.
- Acceptance Probability: If S' is worse than S, it may still be accepted with a probability based on the
  temperature and the difference in objective function value. The acceptance probability is given by:

$$P(\mathsf{accept}) = e^{-\frac{\Delta E}{T}}$$

- Cooling Schedule: After each iteration, the temperature is decreased according to a predefined cooling schedule, which determines how quickly the algorithm converges. Common cooling schedules include linear, exponential, or logarithmic cooling.
- Termination: The algorithm continues until the system reaches a low temperature (i.e., no more significant improvements are found), or a predetermined number of iterations is reached.

#### Cooling Schedule and Its Importance

The cooling schedule plays a crucial role in the performance of Simulated Annealing. If the temperature decreases too quickly, the algorithm might converge prematurely to a suboptimal solution (local optimum). On the other hand, if the cooling is too slow, the algorithm may take an excessively long time to find the optimal solution. Hence, finding the right balance between exploration (high temperature) and exploitation (low temperature) is essential.