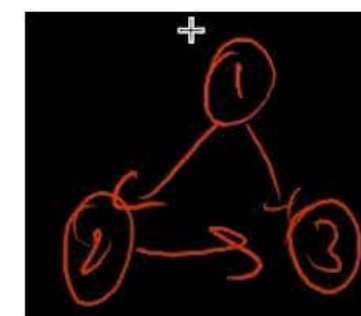


## Different approaches to solve graph questions

1. Identify bfs/dfs
2. Apply appropriate strategy to solve the problem
  - i. variation of **get connected components**
  - ii. variation of minimum spanning tree -**Kruskals algo**
  - iii. Variation of **topological sorting** (One has to come before other)
  - iv. Which uses priority queue (Variation of **Dijkstra**) -> When each step has to choose min or max among all adjacent edges
  - v. Variation of articulation point - Removal of an edge/node will break the system into n parts, find the edge/node
  - vi. Using **disjoint sets** - Questions in which we dynamically add edges to graph while solving the problem OR questions that have to check if edge or vertex belongs to a component OR Anything that involves **grouping /merging**
  - vii. Variation of hamiltonian or euler's path

Mother Vertex-> vertex through which we can reach all the other vertices of the Graph. First element of the topological sort is the mother vertex

Detection of a cycle using Disjoint Set for directed graph is not possible BUT if we know that the graph has a cycle, we can detect the edge which is causing the cycle





# Graphs

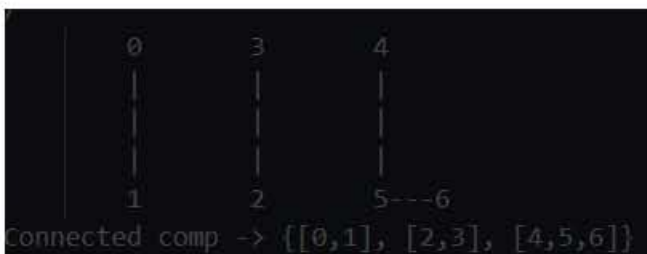
```
void DFS_Traversal(int node){
    visited[node] = 1;
    cout << node << " ";
    for(int i = 0; i < v; i++){
        if(adj[node][i] == 1 && visited[i] != 1)
            DFS_Traversal(i);
    }
}
```

DFS

```
void BFS_Traversal(int src){
    vector<int> visited (v, 0);
    queue<int> q;
    //Starting with source node
    q.push(src);
    visited[src] = 1;
    while(!q.empty()){
        int node = q.front();
        q.pop();
        cout << node << " ";
        //Visiting all connected nodes and enqueue unvisited nodes for exploration
        for(int j = 0; j < v; j++){
            if(adj[node][j] == 1 && visited[j] != 1){
                visited[j] = 1;
                q.push(j); //Enqueuing unvisited node
            }
        }
    }
    cout << endl;
}
```

T:  $O(V+E)$   
S:  $O(V)$

BFS



```
void getConnectedComponent(int src, vector<bool> &visited, vector<int> &connectedComp){
    visited[src] = true;
    connectedComp.push_back(src);
    for(int &edge : adj[src]){
        if(visited[edge] == false){
            getConnectedComponent(edge, visited, connectedComp);
        }
    }
}
```

```
vector<vector<int>> comps; //To store all connected components
vector<bool> visited(7, false);
for(int i = 0; i < 7; i++){
    if(visited[i] == false){
        vector<int> comp; //To store one single component
        g.getConnectedComponent(i, visited, comp);
        comps.push_back(comp);
    }
}
```

get Connected Components

```
//If a graph is connected, all nodes must be visited in one traversal
vector<bool> visited(7, false);
vector<int> comp;
g.getConnectedComponent(0, visited, comp);
for(bool x : visited){
    if(x == false) cout << "false" << endl;
}
cout << "true" << endl;
```

Is graph connected

```
void hamiltonmian(vector<bool> visited, int src, string psf){
    visited[src] = true;
    //Base case
    bool allVisited = true;
    for(bool x : visited){
        if(x == false){
            allVisited = false;
            break;
        }
    }
    if(allVisited){
        cout << psf + to_string(src) ;
        for(int node : adj[src]){
            if(node == psf[0] - '0'){
                cout << "*" << endl;
                return;
            }
        }
        cout << "." << endl;
        return;
    }
    //Code
    for(int node : adj[src]){
        if(visited[node] == false)
            hamiltonmian(visited, node, psf + to_string(src));
    }
    visited[src] = false;
}
```

0123456.  
0123465.  
0125643\*  
0346521\*

jayte hue visit kare  
vapas ate hue unvisit kare  
base case-All nodes visited

hamiltonian path

A hamiltonian path is such which visits all vertices without visiting any twice. A hamiltonian path becomes a cycle if there is an edge between first and last vertex.



```
bool hasCycleDFS(int node, vector<vector<int>>& adjList, vector<bool>& visited, int parent) {
    visited[node] = true;

    for (int neighbor : adjList[node]) {
        if (!visited[neighbor]) {
            if (hasCycleDFS(neighbor, adjList, visited, node))
                return true;
        } /* If we are visiting an already visited node which is not its parent => Cyclic
        else if (neighbor != parent) {
            return true;
        }
    }

    return false;
}
```

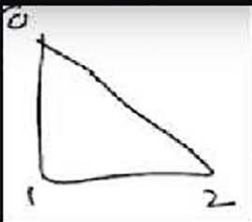
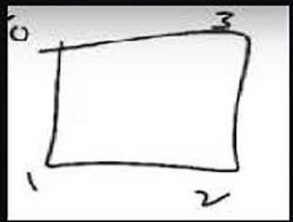
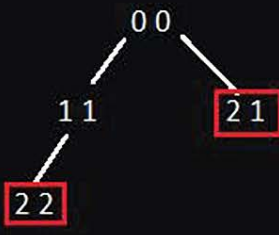
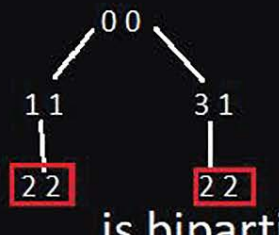
Cycle detection using  
DFS in undirected graph

The approach is the same for BFS  
we create a queue {node, parent}  
Whenever we are adding an adj node, we check if the adj  
node is visited, if the node is visited and is not parent then  
the graph is cyclic


```
bool Graph::isBipartite(int src, vector<int> &visited){
    // <node, Level at which it is visited>
    queue<pair<int, int>> q;
    q.push({src, 0});
    while(!q.empty()){
        pair<int, int> node = q.front();
        q.pop();
        //if node is already visited
        if(visited[node.first] != -1){
            //visited-> contains Level at which node was visited
            //node.second -> current level
            //If they both dont match => the node is being visited at different levels => Not bipartite
            if(node.second != visited[node.first]){
                return false;
            }
        }
        //If unvisited, mark 'visited at current level'
        visited[node.first] = node.second;
        for(int i : adj[node.first]){
            if(visited[i] == -1){
                q.push({i, node.second + 1});
            }
        }
    }
    return true;
}
```

This is true when the graph is cyclic, we  
further check if graph contains odd no of  
nodes or even no of node

If the node is visited at two different  
levels, then it has odd no of nodes  
Else it has even no of nodes

is bipartite



A graph is said to be bipartite if its vertices  
can be divided into two mutually exclusive  
and exhaustive sets such that all the edges are  
present across the sets

All non cyclic graphs --> Bipartite  
If cyclic-> Even no of Nodes -> Bipartite  
Odd no of nodes -> Not Bipartite

If there are different components in a graph  
--> All the components must be bipartite,  
for graph to be bipartite,

```
void Graph::topologicalSort(int src, vector<bool> &visited, stack<int> &s){
    visited[src] = true;
    for(int node : adj[src]){
        if(visited[node] == false)
            topologicalSort(node, visited, s);
    }
    s.push(src);
}
```

topological sort

```
vector<bool> visited(v, false);
stack<int> s;
for(int i = 0; i < v; i++){
    if(visited[i] == false){
        g.topologicalSort(i, visited, s);
    }
}
while(!s.empty()){
    cout << s.top() << endl;
    s.pop();
}
```

Topological sort--> It is a permutation of all vertices for a  
directed graph in called a topological sort if for all directed  
edges uv, u appears before v in the graph.

TOPOLOGICAL SORT IS POSSIBLE ONLY FOR DAG



# Cycle Detection

## 1. For undirected graph

```
bool hasCycleDFS(int node, vector<vector<int>>& adjList, vector<bool>& visited, int parent) {
    visited[node] = true;

    for (int neighbor : adjList[node]) {
        if (!visited[neighbor]) {
            if (hasCycleDFS(neighbor, adjList, visited, node))
                return true;
        } /* If we are visiting an already visited node which is not its parent => Cyclic
        else if (neighbor != parent) {
            return true;
        }
    }

    return false;
}
```

using DFS

```
bool hasCycleBFS(int node, vector<vector<int>>& adjList, vector<bool>& visited) {
    queue<pair<int, int>> q;
    q.push({node, -1}); // {node, parent}

    visited[node] = true;
    while (!q.empty()) {
        int currNode = q.front().first;
        int parent = q.front().second;
        q.pop();

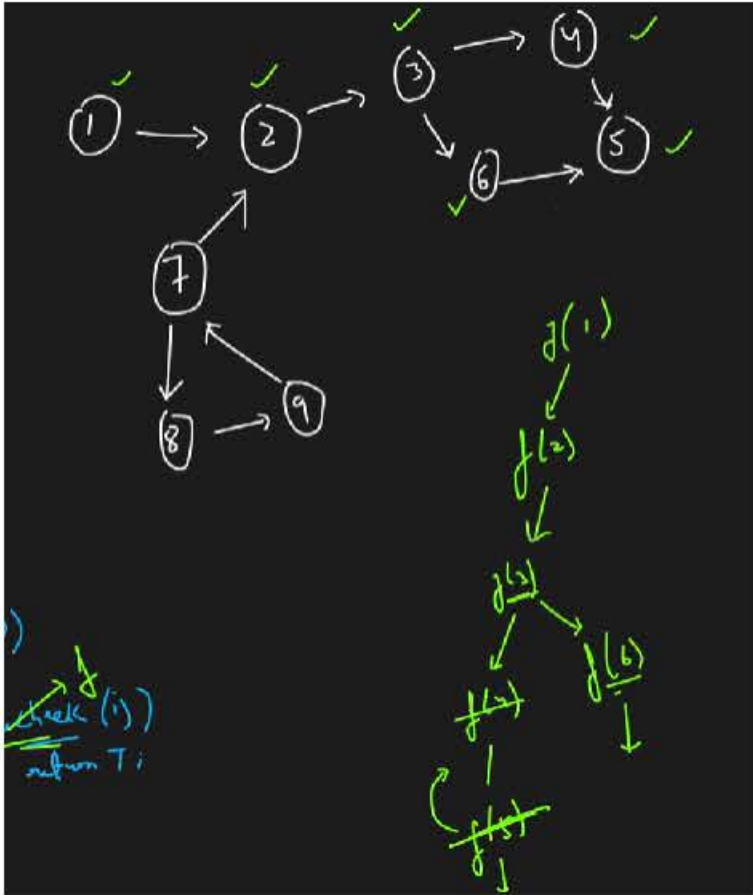
        for (int neighbor : adjList[currNode]) {
            if (!visited[neighbor]) {
                visited[neighbor] = true;
                q.push({neighbor, currNode});
            } else if (neighbor != parent) {
                return true;
            }
        }
    }

    return false;
}
```

using bfs

## 2. For directed graph

The above algorithm will not work.



Consider this example.

When we call DFS continuously we go in the path

1->2->3->4->5

Since there are no adj nodes for 5, we return

At 3 we have another adj node 6

so we go to 6

At 6 we find that 5 is an adj node which is already visited => CYCLIC

BUT ITS NOT TRUE

So we need another dfsVis array which will keep track of all nodes visited while dfs traversal

```
bool hasCycleDFS(int node, vector<vector<int>>& adjList, vector<bool>& visited, vector<bool>& dfsVisited) {
    visited[node] = true;
    dfsVisited[node] = true;

    for (int neighbor : adjList[node]) {
        if (!visited[neighbor]) {
            if (hasCycleDFS(neighbor, adjList, visited, dfsVisited))
                return true;
        } else if (dfsVisited[neighbor]) {
            return true;
        }
    }

    dfsVisited[node] = false;
    return false;
}
```

using dfs

Using BFS -> We use Khans algorithm for topological sorting. WKT topo sort is possible for only DAG. So we use the reverse approach here. If we are not able to topologically sort the nodes => CYCLIC

Approach -> Code is same as Khans algo but here we maintain a counter.

increment counter for every node popped from the queue

if cnt != n then the graph is cyclic



## Single Source Shortest Path Algos

```
void addEdge(vector<pair<int, int> > adj[], int u, int v, int wt){
    adj[u].push_back(make_pair(v, wt));
    adj[v].push_back(make_pair(u, wt));
}

void shortestPath(vector<pair<int, int> > adj[], int V, int src){
    priority_queue<iPair, vector<iPair>, greater<iPair>> pq; //Min heap

    // Create a vector for distances and initialize all distances as infinite (INF)
    vector<int> dist(V, INF);

    // Insert source itself in priority queue and initialize its distance as 0.
    pq.push(make_pair(0, src));
    dist[src] = 0;

    // Looping till all distances are not finalized
    while (!pq.empty()) {
        int u = pq.top().second;          T : O(E + V log V)
        pq.pop();                         S : O(V)

        // Get all adjacent of u.
        for (auto x : adj[u]) {
            int v = x.first;
            int weight = x.second;

            // If there is shorter path to v through u.
            if (dist[v] > dist[u] + weight) {
                // Updating distance of v
                dist[v] = dist[u] + weight;
                pq.push({dist[v], v});
            }
        }
    }
}
```

Single source shortest path to all other nodes

Dijkstra's Algo

Drawback -> Dijkstra's Algorithm may not work for negative weighted edges

### Replacing queue with priority queue in BFS -> Dijkstra's Algo

#### Time Complexity:

The time complexity of Dijkstra's algorithm will be  $O(E + V \log V)$ . This is because, we are iterating over all the edges once during the entire run of the algorithm. In each iteration, we are popping one node and pushing the unvisited neighbour nodes. Since the priority queue can contain all the vertices, the push or pop operation will be  $O(\log V)$ . Hence the total time complexity will be  $O(E) + O(V) * O(\log V) = O(E + V \log V)$ .

Note: You can argue that we might be having multiple Pairs having the same node's value. So, the maximum size of the priority queue will be not  $O(N)$  but  $O(E)$ . But, even if you replace  $\log V$  with  $\log E$  (cost of one push/pop operation), then there will be no difference in the time complexity as:  $O(E + V \log E) = O(E + V \log(V^2)) = O(E + 2V \log V) = O(E + V \log V)$  only.

#### Space Complexity:

We are taking a priority queue of Pair nodes. Hence, the space complexity will be  $O(N)$  where  $N$  = maximum Pair nodes in the queue, which is equivalent to  $O(V)$ .

```
class edge{
public:
    int u, v, wt;
    edge(int u, int v, int wt){
        this->u = u;
        this->v = v;
        this->wt = wt;
    }
};

void shortestPath(vector<edge> graph, int V, int src){
    // Create a vector for distances and initialize all distances as infinite (INF)
    vector<int> dist(V, INF);

    //Relaxing all the edges V-1 times
    for(int i = 0; i < V-1; i++){
        for(edge e : graph){
            if(dist[e.u] + e.wt < dist[e.v]){
                dist[e.v] = dist[e.u] + e.wt;
            }
        }
    }

    //Relaxing one more time to find if negative cycle is present
    bool flag = false;
    for(edge e : graph){
        if(dist[e.u] + e.wt < dist[e.v]){
            flag = true;
            cout << "Has negative weighted cycle" << endl;
            return;
        }
    }

    // Print shortest distances stored in dist[]
    printf("Vertex Distance from Source\n");
    for (int i = 0; i < V; ++i)
        printf("%d \t\t %d\n", i, dist[i]);
}
```

T :  $O(V * E)$   
S :  $O(V)$

Bellmanford Algorithm

This can be used for negative weighted graphs

#### DRAWBACK ->

Does not work for negative weighted cycle

Since this algorithm doesn't work for -ve weighted cycle, so this won't work for undirected graph with -ve weights because for an edge  $u-v$  in undirected graph it is same as  $u \rightarrow v$  and  $u \leftarrow v$ . So if there is negative weight this becomes -ve weighted cycle.

In the Bellman-Ford algorithm, we perform the relaxation step  $V-1$  times where  $V$  is the number of vertices in the graph. The reason why we need to perform  $V-1$  relaxations is that, in the worst-case scenario, the shortest path from a source vertex to a destination vertex can pass through at most  $V-1$  edges.



# Minimum Spanning Tree Algorithms

```

class Pair{
public:
    int v; //Vertex
    int av; //Acquiring vertex
    int wt; //Weight
    Pair(int wt, int v, int av){
        this->v = v;
        this->av = av;
        this->wt = wt;
    }
    Pair();
};

class Graph{
    int v;
    //vertex , weight
    list<pair<int, int>> *adj;
public:
    Graph(int vertices);
    void addEdge(int src, int dest, int wt);
    void prims(int);
};

// Add edges to the graph
void Graph::addEdge(int src, int dest, int wt){
    adj[src].push_back({dest, wt});
    adj[dest].push_back({src, wt});
}

void Graph::prims(int src){
    vector<bool> visited(v, false);

    // pair <weight, vertex, acquiring vertex>
    priority_queue<Pair, vector<Pair>, myComparator> pq;
    pq.push(Pair(0, src, -1)); // -1 -> Non existing vertex

    cout << "V AV WT" << endl << src << " - -" << endl;
    while(!pq.empty()){
        Pair p = pq.top();
        pq.pop();
        //If the node is already visited -> Continue (To avoid cycle)
        if(visited[p.v] == true) continue;

        //If unvisited, mark visited and print
        visited[p.v] = true;
        //Avoiding source node
        if(p.av != -1) cout << p.v << " " << p.av << " " << p.wt << endl;

        //Explore all unvisited adj nodes of current node
        for(pair<int, int> &neighbour : adj[p.v]){
            if(visited[neighbour.first] == false){
                pq.push(Pair(neighbour.second, neighbour.first, p.v));
            }
        }
    }
}
    
```

**T:  $O((v+e)\log v)$**   
**S:  $O(v+e)$**

T is  $V+E \log v$  because  
 Normal BFS has  $T = O(V+E)$   
 and here we are using PQ  
 => Extra  $\log v$  for each push  
 and pop operation

**Prims Algo**

Used to find the minimum spanning tree  
 Spanning tree -> Sub graph of a graph having V  
 vertices (same as original graph) and V-1 edges

No of spanning trees =  $\frac{E}{V-1} C$  - no of cycles

```

class DisjointSet {
    vector<int> rank, parent, size;
public:
    DisjointSet(int n) {
        rank.resize(n + 1, 0); //Rank of all elements is initially zero
        parent.resize(n + 1);
        size.resize(n + 1);
        for (int i = 0; i <= n; i++) {
            parent[i] = i; //Initially, element is parent of itself => Size of each component is 1
            size[i] = 1;
        }
    }

    int findUPar(int node) {
        if (node == parent[node]) return node;
        return parent[node] = findUPar(parent[node]);
    }

    void unionByRank(int u, int v) {
        int ulp_u = findUPar(u); //Ultimate parent of u
        int ulp_v = findUPar(v); //Ultimate parent of v

        if (ulp_u == ulp_v) return;
        //Whoever has a smaller rank is connected to bigger rank => Bigger rank is the parent after union
        if (rank[ulp_u] < rank[ulp_v]) {
            parent[ulp_u] = ulp_v;
        }
        else if (rank[ulp_v] < rank[ulp_u]) {
            parent[ulp_v] = ulp_u;
        }
        //If rank is same, make anyone as parent, but here we increase the rank of the parent by 1
        else {
            parent[ulp_v] = ulp_u;
            rank[ulp_u]++;
        }
    }
}

class Solution{
public:
    //Function to find sum of weights of edges of the Minimum Spanning Tree.
    int kruskalsAlgo(int V, vector<vector<int>> adj){
        //Create a list of edges ( ) vector<weight, u, v> ) from the given graph
        vector<pair<int, pair<int, int>>> edges;
        for (int i = 0; i < V; i++) {
            for (auto it : adj[i]) {
                int adjNode = it[0];
                int wt = it[1];
                int node = i;

                edges.push_back({wt, {node, adjNode}});
            }
        }
        DisjointSet ds(V);

        //Sorting edges according to weight
        sort(edges.begin(), edges.end());
        int mstWt = 0;
        cout << "U V WT" << endl;
        for (auto it : edges) {
            int wt = it.first;
            int u = it.second.first;
            int v = it.second.second;

            //If there is no cycle -> consider the edge
            if (ds.findUPar(u) != ds.findUPar(v)) { //If the two nodes belong to different components => No cycle
                mstWt += wt;
                ds.unionBySize(u, v);
                cout << u << " " << v << " " << wt << endl;
            }
        }

        return mstWt;
    }
};
    
```

**T= $O(E \log v)$**   
**S= $O(E + V)$**

Since for parent and  
 rank array  $O(v)$  space is  
 used.  $O(E)$  space is used  
 for storing edges.

**Kruskals Algo**



# Khans Algorithm - Topological sorting using BFS

```
vector<int> khansAlgo(vector<vector<int>> graph){
    //Calculating indegree of each vertex
    vector<int> indeg(graph.size());
    for(int i = 0; i < graph.size(); i++){
        for(int x : graph[i]){
            indeg[x]++;
        }
    }

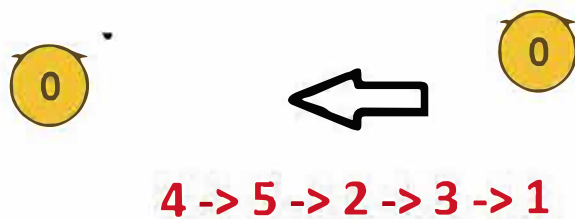
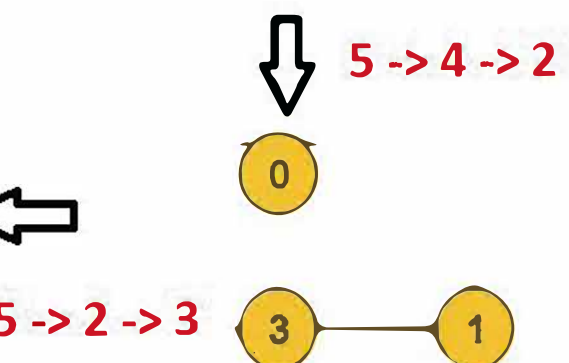
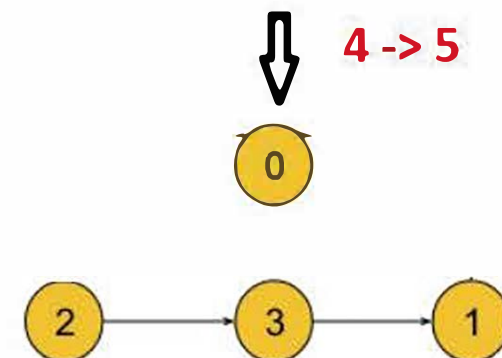
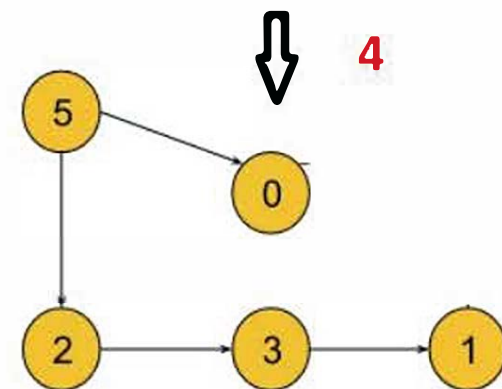
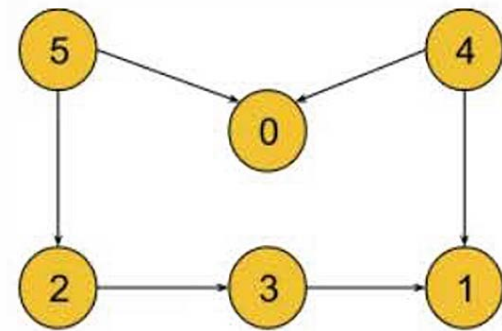
    queue<int> q;
    //Pushing all nodes with indegree 0 to queue
    for(int i = 0; i < graph.size(); i++){
        if(indeg[i] == 0) q.push(i);
    }

    int idx = 0;
    vector<int> ans(graph.size());
    while(!q.empty()){
        int node = q.front();
        q.pop();

        ans[idx++] = node;
        for(int adj : graph[node]){
            indeg[adj]--;

            if(indeg[adj] == 0) q.push(adj);
        }
    }
    return idx == graph.size() ? ans : vector<int> {-1};
}
```

T : O(V+E)  
S : O(V)

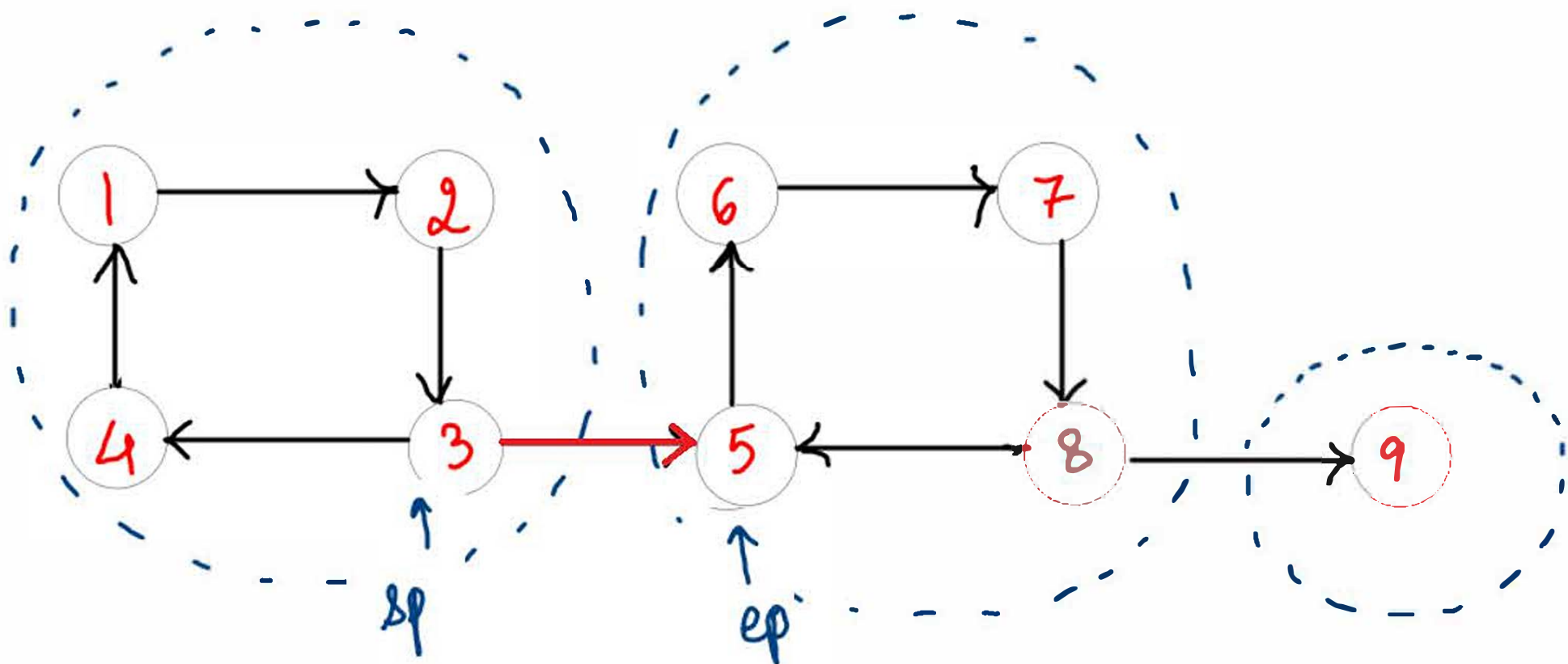


4 -> 5 -> 2 -> 3 -> 1 -> 0



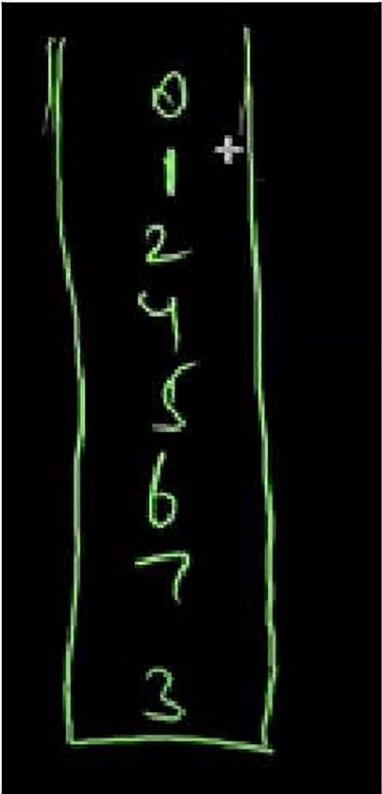
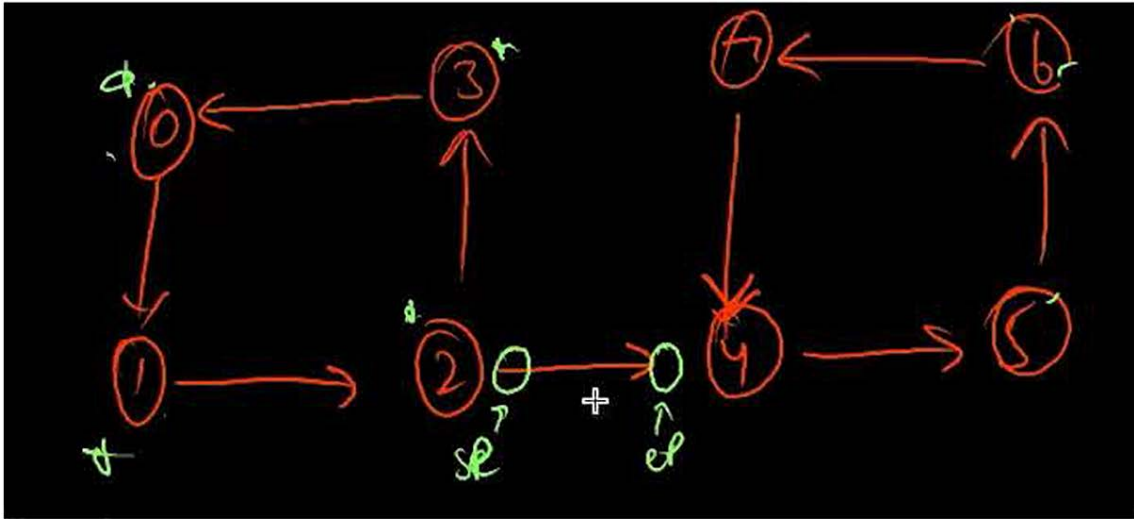
# Kosaraju Algorithm

You will be given a graph with nodes and M directed edges. You have to find the number of Strongly Connected Components in the graph. First, understand what exactly is meant by Strongly Connected Components. A strongly connected component is a portion of the graph in which there is a path from each vertex to another vertex. It is only applicable on a directed graph. Also, all the strongly connected components of a graph are cyclic. But it does not mean that each cycle in a graph will be a strongly connected component. It is possible that multiple cycles can form one strongly connected component. Let’s take a look at the graph given below:



If we apply normal DFS to this graph, it will give us the answer for the number of connected components as 1. This is because there is an edge between 3 and 5. But, since we can’t go back from 5 to 3, it contradicts the statement for strongly connected components, and so, it should not be a part of the connected component, and form a separate strongly connected component. This is why we need a new algorithm to figure out the number of strongly connected components in a graph. So, through this, we figure out that the **edge between 3 and 5 is a problematic edge**

Our focus is now on the problematic edge so let us consider the graph

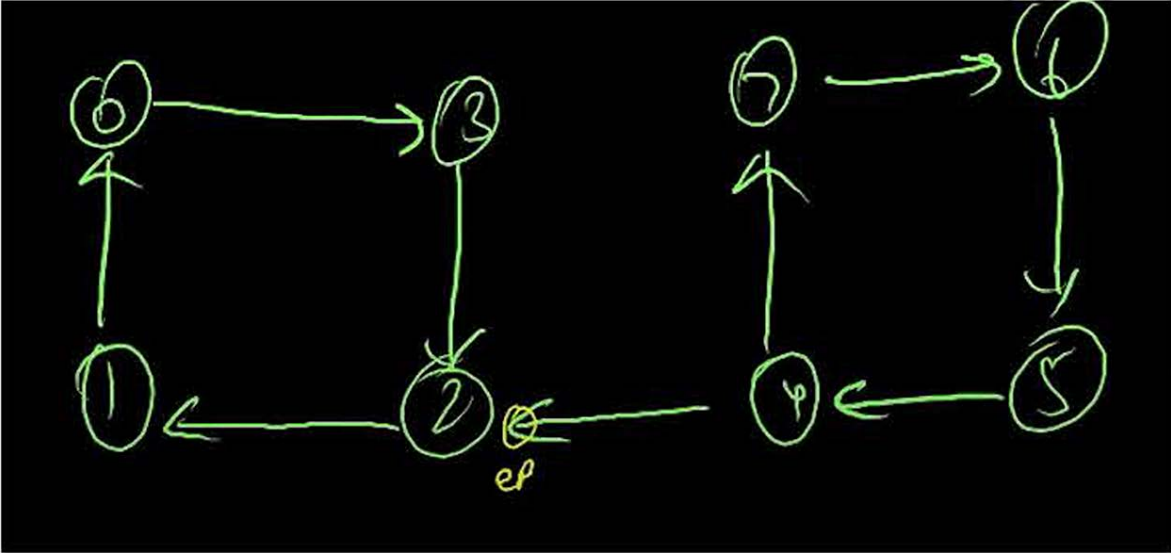


The problematic edge is from 2 -> 4 here, 2 -> starting point , 4 -> End point of the problematic edge

The solution to the above problem is that we have to run DFS on the component at endpoint of the problematic edge.

But, the problem is, how will we figure out the problematic edge? Therefore, Kosaraju algorithm was introduced. In this, we have used a simple trick.

1. First we impliment a random order dfs and add the nodes into a stack while backtracking. The element on the top of the stack will always be one of the elements from the side of the starting point of the problematic edge
2. Now, our next step will be to reverse all the edges of the graph



This will not cause any change in the strongly connected components as they will still be cyclic and remain connected to each other. The only major difference will be seen in the problematic edge. As now, the starting and ending points will be reversed.

How will this help us ? As we have seen earlier, by running random DFS we found an element from the side of the strting point of the problematic edge. Now after reversing the edges, the element on the side of starting point has become an element on the ending point of problematic edge which is what we needed

3. Now, apply DFS to the reversed graph in the order arranged in the stack

```
void dfs(int node, vector<vector<int>> &graph, vector<bool> &visited, stack<int> &s){
    visited[node] = true;
    for(int adj : graph[node]){
        if(!visited[adj]) dfs(adj, graph, visited, s);
    }
    s.push(node);
}

void revDfs(int node, vector<vector<int>> &graph, vector<bool> &visited){
    visited[node] = true;
    for(int adj : graph[node]){
        if(!visited[adj]) revDfs(adj, graph, visited);
    }
}

int kosarajuAlgp(vector<vector<int>> graph, int v){
    stack<int> s;
    vector<bool> visited(v, false);
    //1. Random DFS
    for(int i = 0; i < v; i++){
        if(!visited[i]){
            dfs(i, graph, visited, s);
        }
    }
    //2. Reversing all edges
    vector<vector<int>> revGraph(v);
    for(int i = 0; i < v; i++){
        visited[i] = false;
        for(int adj : graph[i]){
            revGraph[adj].push_back(i);
        }
    }
    // 3. DFS for reversed graph in the order of stack which we filled earlier
    int ans = 0;
    while(!s.empty()){
        int node = s.top();
        s.pop();

        if(!visited[node]){
            revDfs(node, revGraph, visited);
            ans++;
        }
    }
    return ans;
}
```



# Articulation Point

Articulation Point

disc []  
low []

par -> parent of curr node  
discovery -> Time at which curr node is discovered  
low -> Ignoring the path from src->curr, which is the node with minimum discovery we are able to touch

adj node can be  
1. Parent  
2. 2. Visited but not parent  
3. 3. not visited and not parent

if ( par ) {  
continue;  
}  
else if ( vis == true ) {  
low[u] = math.min(low[u], disc[v])  
}  
else {  
dfs ( nbr );  
low[u] = math.min(low[u], low[v]);  
if ( low[v] >= disc[u] ) {  
ap[u] = true;  
}

① min -> disc(v)  
② ap counting array  
③ actual src

low of node 6 will disc of node 3 ie 6

A vertex is said to be an articulation point in a graph if removal of the vertex and associated edges disconnects the graph. So, the removal of articulation points increases the number of connected components in a graph.

```
class Pair{
public:
    int par;
    int disc;
    int low;
    bool isAP;
    Pair(){
        par = 0;
        disc = 0;
        low = 0;
        isAP = false;
    }
};

void dfs(vector<vector<int>> &graph, int u, vector<bool> &visited, vector<Pair> &ap, int disc){
    ap[u].disc = ap[u].low = disc;
    int cnt = 0; //This variable is used to keep the count of no of dfs calls we make. This is useful

    visited[u] = true;
    for(int v : graph[u]){
        //If nbr is parent => nbr already is present in the path from src to current node
        if(v == ap[u].par) continue;
        //If nbr is not parent but is visited then the edge curr->nbr is not along the path from src to curr node => low can be updated
        else if(visited[v]){
            ap[u].low = min(ap[u].low, ap[v].disc);
        }
        //else it is an unvisited node => DFS
        else{
            ap[v].par = u;
            dfs(graph, v, visited, ap, disc+1);
            cnt++;

            ap[u].low = min(ap[u].low, ap[v].low);
            //Parent[u] = -1 => it is actual source.
            if(ap[u].par == -1 && cnt >= 2){
                ap[u].isAP = true;
            }
            else if(ap[v].low >= ap[u].disc){
                ap[u].isAP = true;
            }
        }
    }
}

void articulationPoint(vector<vector<int>> &graph, int v){
    vector<bool> visited(v, false);
    vector<Pair> ap(v);
    ap[0].par = -1;

    dfs(graph, 0, visited, ap, 1);
    for(int i = 0; i < v; i++){
        if(ap[i].isAP) cout << i+1 << " ";
    }
}
```

For an edge  $u \rightarrow v$   
if  $low(v) \geq disc(u)$   
then U is articulation point

What does the above condition mean -> If condition is true if low (child) is less than discovery (parent) this means that the whole graph below it is not able to reach parent or any of the nodes above parent

**THIS CONDITION WILL NOT WORK FOR ROOT IE ACTUAL SOURCE**

Articulation Point

disc []  
low []

par -> parent of curr node  
discovery -> Time at which curr node is discovered  
low -> Ignoring the path from src->curr, which is the node with minimum discovery we are able to touch

adj node can be  
1. Parent  
2. 2. Visited but not parent  
3. 3. not visited and not parent

if ( par ) {  
continue;  
}  
else if ( vis == true ) {  
low[u] = math.min(low[u], disc[v])  
}  
else {  
dfs ( nbr );  
low[u] = math.min(low[u], low[v]);  
if ( low[v] >= disc[u] ) {  
ap[u] = true;  
}

① min -> disc(v)  
② ap counting array  
③ actual src

low of node 6 will disc of node 3 ie 6

**FOR ROOT OR SOURCE NODE -> IF THERE ARE MORE THAN ONE DFS CALLS FOR SRC NODE THEN SRC NODE IS ALSO AN ARTICULATION POINT**

for edge 4->3  
disc(4) = 5  
low(3) = 6  
=> low(child) >= disc(parent)  
Iska matlab ye hain ki disc=5 ie node 4 ko ye pata chalgaya ki mere badh ek aisa banda hain jiski reach mere se pehle valo tak nahi he  
=> node 4 is an articulation point



# EULRIAN PATH

An Eulerian path is a path in a graph that visits every edge exactly once, allowing for the possibility of revisiting vertices. In other words, it is a sequence of edges that traverses the entire graph without lifting the pen from the paper (or without removing the finger from the touchscreen).

It is same as hamiltonian path but here it is edges instead of vertices

## 1. For Undirected Graph

An undirected graph with all nodes having **even degree** has a eularian circuit



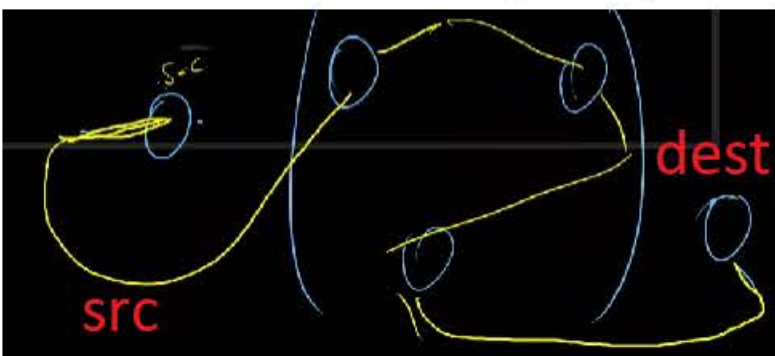
We can visualise the graph like this. Each node has an even degree  $\Rightarrow$  Each node has an entry point and exit point of an edge  $\Rightarrow$  We can easily make a eularian circuit



Node with even degree has an entry point but no exit point  $\Rightarrow$  we will be stuck inside

## For eularian path

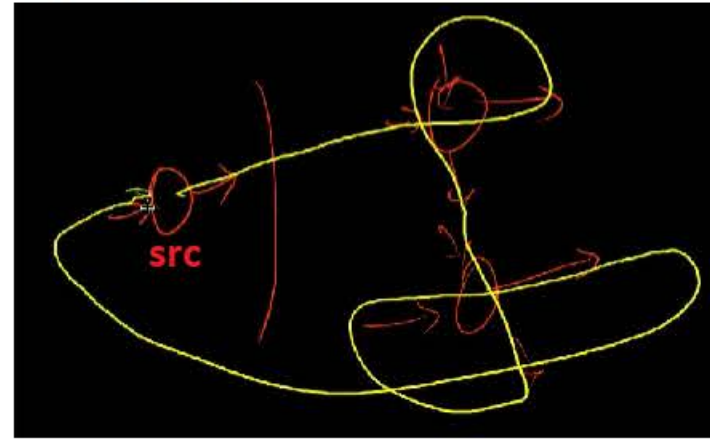
(n-2) nodes  $\rightarrow$  should have even degree  
2 nodes  $\rightarrow$  odd degree (src and dest)



We want the inner nodes to have an even degree which will force it to have entry and exit point  
Src  $\rightarrow$  odd deg  $\Rightarrow$  1 exit point  
dest  $\rightarrow$  1 entry point

## 2. For directed graph

**Eularian circuit**  $\rightarrow$  Indegree = Outdegree for all nodes



## Eularian path

(n-2) nodes  $\rightarrow$  indegree = outdegree

2 nodes  $\rightarrow$  special condition

src  $\rightarrow$  Outdeg should be 1 more than its indeg

dest  $\rightarrow$  Indeg should be 1 more than its outdeg



**Implimentation**  $\rightarrow$  Just run DFS until all the edges are visited  
**Node**  $\rightarrow$  Addition of a node into path  
should be in post order

Eularian path  $\rightarrow$  AN edge can be visited exactly once and a vertex can be visited any no of times  
Hamiltonian path  $\rightarrow$  An edge can be visited any no of times but a vertex can be visited exactly once



1. A common approach for variation of get connected components

```
bool isValid(int i,int j,int m,int n){
    return !(i == m || j == n || j < 0 || i < 0);

}

vector<pair<int, int>> dir = {{1,0},{0,1},{0,-1},{-1,0}}; //Direction
vector<vector<int>> updateMatrix(vector<vector<int>>& matrix) {
    queue<pair<int,int>> q;
    int m = matrix.size();
    int n = matrix[0].size();
    //creating a dist matrix and marking all 1's as unexplored
    vector<vector<int>> dis(m,vector<int>(n,-1));
    //Pushing all 0's to queue so that we can explore nearest 1 for each of them
    for(int i = 0; i < m; i++){
        for(int j = 0; j < n; j++){
            if(matrix[i][j] == 0){
                q.push({i,j});
                dis[i][j] = 0;
            }
        }
    }
    while(!q.empty()){
        pair<int,int> curr = q.front();
        q.pop();

        int r = curr.first;
        int c = curr.second;
        for(auto &x : dir){
            int nr = r + x.first; //next row
            int nc = c + x.second; //next col
            //If nr and nc are valid indices and unvisited
            if(isValid(nr, nc, m, n) && dis[nr][nc] == -1){
                q.push({nr, nc});
                dis[nr][nc] = dis[r][c] + 1;
            }
        }
    }
    return dis;
}
```

nearest 0 to all 1's

- 1. Declare a direction array
- 2. Push all elements of a component onto queue
- 3. BFS :
  - For all 4 dirctrions
  - 3.1. Calc nr and nc
  - 3.2. If nr and nc are valid do required WORK

2. Application of topological sort

Pepcoding offers total of n courses labelled from 0 to n-1. Some courses may have prerequisites. you have been given m pairs ai,bi. where 1 pair means you must take the course bi before the course ai. Given the total number of courses numCourses and a list of the prerequisite pairs, return the ordering of courses you should take to finish all courses. If it is impossible to finish all courses print -1.

Other example -> Alien Dictionary

3. Application of MST

```
int kruskalsAlgo(int V, vector<pair<int, pair<int, int>>> edges){
    DisjointSet ds(V);

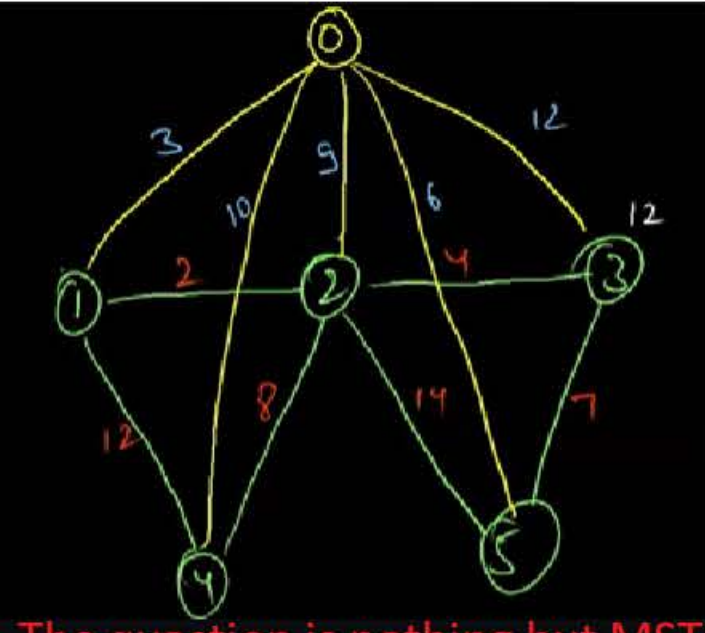
    //Sorting edges according to weight
    sort(edges.begin(), edges.end());
    int mstWt = 0;
    int pipeCnt = 0;
    for (auto edge : edges) {
        int wt = edge.first;
        int u = edge.second.first;
        int v = edge.second.second;

        //If there is no cycle -> consider the edge
        if (ds.findUPar(u) != ds.findUPar(v)) { //If the two nodes belong to different comp
            mstWt += wt;
            ds.unionBySize(u, v);
        }
    }

    return mstWt;
}

int minCostToSupplyWater(int n, vector<int>&wells, vector<vector<int>>& pipes){
    //vector<wt, u, v>
    vector<pair<int, pair<int, int>>> edges;
    //We introduce a dummy node 0 which acts like well and digging well in ith polt = edge between 0 and i
    for(int i = 1; i <= n; i++){
        edges.push_back({wells[i-1], {0, i}});
    }
    for(vector<int> &edge : pipes){
        edges.push_back({ edge[2], {edge[0], edge[1]}});
    }

    return kruskalsAlgo(n+1, edges);
}
```



The question is nothing but MST where site acts as a vertex and pipe acts as an edge. Since there must be atleast 1 well, we create an extra dummy node 0 which acts like well and digging well in ith polt = edge between 0 and i

optimum WaterDistribution

There are n construction sites in a town. We want to supply water for all the construction sites by building wells and laying pipes. For each site i, we can either build a well inside edge directly with cost wells[i-1], or pipe in water from another well to edge. The costs to lay pipes between sites are given by the 2d array cost, where each row of cost contains 3 numbers ai,bi and wi where wi is the cost to connect ai to bi. connections are bidirectional. Return the minimum total cost to supply water to all the construction sites.

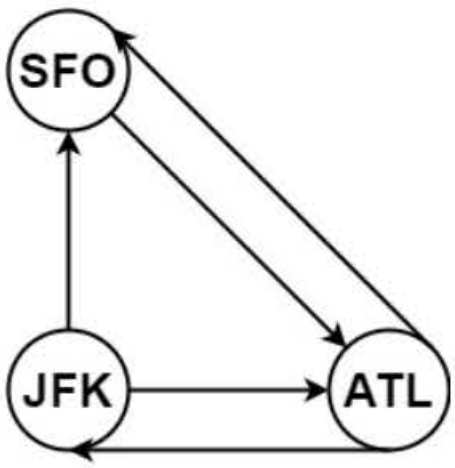
7. Eularian Path

You are given a list of airline tickets where tickets[i] = [from<sub>i</sub>, to<sub>i</sub>] represent the departure and the arrival airports of one flight. Reconstruct the itinerary in order and return it.

All of the tickets belong to a man who departs from "JFK", thus, the itinerary must begin with "JFK". If there are multiple valid itineraries, you should return the itinerary that has the smallest lexical order when read as a single string.

- For example, the itinerary ["JFK", "LGA"] has a smaller lexical order than ["JFK", "LGB"].

You may assume all tickets form at least one valid itinerary. You must use all the tickets once and only once.



**Input:** tickets = [["JFK","SFO"],["JFK","ATL"],["SFO","ATL"],["ATL","JFK"],["ATL","SFO"]]  
**Output:** ["JFK","ATL","JFK","SFO","ATL","SFO"]  
**Explanation:** Another possible reconstruction is ["JFK","SFO","ATL","JFK","ATL","SFO"] but it is larger in lexical order.

```
//map <string, priorityQueue<string>> -> PQ because we are asked to choose the cities in Lexicographical order
unordered_map<string, multiset<string>> graph;
vector<string> ans;
void dfs(string vtex){
    multiset<string> &edges = graph[vtex];
    while (!edges.empty()){
        string to = *edges.begin();
        edges.erase(edges.begin()); //Marking the edge as visited. By doing so, we need not check if this edge is visited or not in further iteration coz the edge doesnt exist now
        dfs(to);
    }
    ans.push_back(vtex);
}

vector<string> findItinerary(vector<vector<string>>> tickets) {
    for (vector<string> &e : tickets) graph[e[0]].insert(e[1]);
    dfs("JFK");
    reverse(ans.begin(), ans.end());
    return ans;
}
```

4. Application of Dijkstra

```
bool isValid(int i, int j, int n){
    return !(i < 0 || j < 0 || i == n || j == n);
}

int swimInWater(vector<vector<int>>& grid) {
    int n = grid.size();
    priority_queue<myPair, vector<myPair>, compare> pq;
    vector<vector<bool>> visited (n, vector<bool>(n, false));
    vector<pair<int, int>> dir = {{0, 1}, {1, 0}, {0, -1}, {-1, 0}};

    pq.push(myPair(0, 0, grid[0][0]));
    visited[0][0] = true;
    while(!pq.empty()){
        myPair node = pq.top();
        pq.pop();

        if(node.i == n-1 && node.j == n-1){
            return node.maxt;
        }
        for(auto p : dir){
            int ni = node.i + p.first;
            int nj = node.j + p.second;

            if(!isValid(ni, nj, n) || visited[ni][nj]) continue;

            pq.push(myPair(ni, nj, max(node.maxt, grid[ni][nj])));
            visited[ni][nj] = true;
        }
    }
    return -1;
}
```

You are given an n x n integer matrix grid where each value grid[i][j] represents the elevation at that point (i, j).

The rain starts to fall. At time t, the depth of the water everywhere is t. You can swim from a square to another 4-directionally adjacent square if and only if the elevation of both squares individually are at most t. You can swim infinite distances in zero time. Of course, you must stay within the boundaries of the grid during your swim.

Return the least time until you can reach the bottom right square (n - 1, n - 1) if you start at the top left square (0, 0).

This is an application of dijkstra algorithm, because at each step we have to choose a minimum between the adjascent nodes

0	1	2	3	4
24	23	22	21	5
12	13	14	15	16
11	17	18	19	20
10	9	8	7	6

op -> 16

5. Articulation Point

There are n servers numbered from 0 to n - 1 connected by undirected server-to-server connections forming a network where connections[i] = [a<sub>i</sub>, b<sub>i</sub>] represents a connection between servers a<sub>i</sub> and b<sub>i</sub>. Any server can reach other servers directly or indirectly through the network.

A critical connection is a connection that, if removed, will make some servers unable to reach some other server.

Return all critical connections in the network in any order.

```
void dfs(vector<vector<int>> &graph, int u, vector<bool> &visited, vector<Pair> &ap, int disc, vector<vector<int>>& ans){
    visited[u] = true;
    ap[u].disc = disc;
    ap[u].low = disc;
    int cnt = 0;

    for(int v : graph[u]){
        //if nbr is parent => nbr already is present in the path from src to current node
        if(v == ap[u].par) continue;
        //if nbr is not parent but is visited then the edge curr->nbr is not along the path from src to curr node => low can be updated
        else if(visited[v]){
            ap[u].low = min(ap[u].low, ap[v].disc);
        }
        //else it is an unvisited nod -> DFS
        else{
            ap[v].par = u;
            dfs(graph, v, visited, ap, disc+1, ans);

            ap[u].low = min(ap[u].low, ap[v].low);
            if(ap[u].par == -1){
                cnt++;
                if(cnt >= 2){
                    ans.push_back({u, v});
                }
            }
            else if(ap[v].low >= ap[u].disc){
                ans.push_back({u, v});
            }
        }
    }
}
```

This question is direct application of articulaion point but with a slight modification. Here ther have asked for edge instead of node. So when we find an articulation point, we add an edge between u and v

Critical Connection

6. DSU

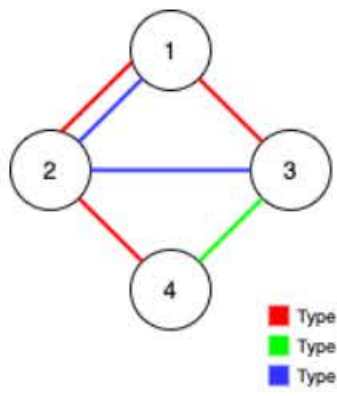
Alice and Bob have an undirected graph of n nodes and three types of edges:

- Type 1: Can be traversed by Alice only.
- Type 2: Can be traversed by Bob only.
- Type 3: Can be traversed by both Alice and Bob.

Given an array edges where edges[i] = [type<sub>i</sub>, u<sub>i</sub>, v<sub>i</sub>] represents a bidirectional edge of type type<sub>i</sub> between nodes u<sub>i</sub> and v<sub>i</sub>, find the maximum number of edges you can remove so that after removing the edges, the graph can still be fully traversed by both Alice and Bob. The graph is fully traversed by Alice and Bob if starting from any node, they can reach all other nodes.

Return the maximum number of edges you can remove, or return -1 if Alice and Bob cannot fully traverse the graph.

Example 1:



**Input:** n = 4, edges = [[3,1,2],[3,2,3],[1,1,3],[1,2,4],[1,1,2],[2,3,4]]  
**Output:** 2  
**Explanation:** If we remove the 2 edges [1,1,2] and [1,1,3]. The graph will still be fully traversable by Alice and Bob. Removing any additional edge will not make it so. So the maximum number of edges we can remove is 2.

Approach is very simple, we keep two Disjoint sets for alice and bob, out main goal is to keep both alice and bob's graph connected.

- Process the type 3 edges first- union in both alice and bob dsu
- Run bfs algorithm on all the edges -> if the nodes of an edge are already connected ie thay both have the same ultimate parent then there is no need of that edge -> decrement no of edges

TO MAKE DSU OF A 2D MATRIX, WE MAKE USE OF CELL NO  
CNO = R \* N + C



# HATKE questions

## 01 BFS

```
void addEdge(vector<vector<pair<int, int>>> &graph, int u, int v){
    graph[u].push_back({v, 0});
    graph[v].push_back({u, 1}); //Reverse edge with cost 1
}

// Approach-> Just add a reverse edge with cost 1 and apply path from
// src to dest using BFS and add cost at each step
int solution(vector<vector<pair<int, int>>> &graph, int src, int dest){
    list<pair<int, int>> q;
    q.push_back({src, 0});

    vector<bool> visited(graph.size(), false);
    while(q.size()){
        pair<int, int> node = q.front();
        q.pop_front();

        if(node.first == dest) return node.second;

        visited[node.first] = true;
        for(pair<int, int> &adj : graph[node.first]){
            if(visited[adj.first]) continue;

            if(adj.second == 0) q.push_front({adj.first, node.second});
            else q.push_back({adj.first, node.second + 1});
        }
    }
    return -1;
}
```

You are given a directed graph , src and destinationn  
You have to find the minimum number of edges you have to reverse in order to have atleast one path from vertex 1 to N, where the vertices are numbered from src to dest

## Bus routes

```
int numBusesToDestination(vector<vector<int>>& routes, int source, int target) {
    if(source == target) return 0;
    //map <stop, busses>
    unordered_map <int, vector<int>> mp;
    for(int i = 0; i < routes.size(); i++){
        for(int j = 0; j < routes[i].size(); j++){
            mp[routes[i][j]].push_back(i);
        }
    }
    unordered_set<int> bv; //To check if bus is already boarded once -> BUS VISITED
    unordered_set<int> bsv; //To check if the bus stop is already visited -> BUS STOP VISITED
    queue <int> q;
    q.push(source);
    int count = 0;
    while(!q.empty()){
        int size = q.size();
        while(size--){
            int stop = q.front();
            q.pop();
            for(int bus : mp[stop]){
                if(bv.find(bus) != bv.end()) continue; //If bus is already boarded => continue
                for(int busStop : routes[bus]){
                    if(bsv.find(busStop) != bsv.end()) continue; //If bus stop is already visited => Continue
                    if(busStop == target) return count+1; //Because current bus is not counted
                    bsv.insert(busStop);

                    q.push(busStop);
                }
                bv.insert(bus);
            }
        }
        count++;
    }
    return -1;
}
```

leetcode hard- Bus routes

You are given an array routes representing bus routes where routes[i] is a bus route that the i<sup>th</sup> bus repeats forever.

- For example, if routes[0] = [1, 5, 7], this means that the 0<sup>th</sup> bus travels in the sequence 1 -> 5 -> 7 -> 1 -> 5 -> 7 -> 1 -> ... forever.

You will start at the bus stop source (You are not on any bus initially), and you want to go to the bus stop target. You can travel between bus stops by buses only.

Return the least number of buses you must take to travel from source to target. Return -1 if it is not possible.

### Example 1:

**Input:** routes = [[1,2,7],[3,6,7]], source = 1, target = 6

**Output:** 2

**Explanation:** The best strategy is take the first bus to the bus stop 7, then take the second bus to the bus stop 6.

