# **LECTURE 3**

### **BASIC GRAPH REDUCTION**

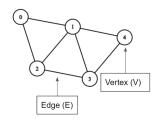
CS200

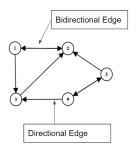
Feb 10, 2023



# **Definition**

## Our View of A Graph

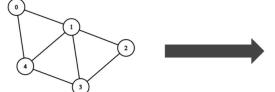




## **Code Representation**

Can represent graph in two ways.

### 1. Adjacency Matrix



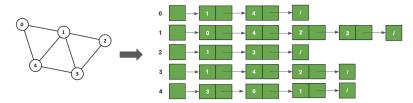
Node	0	1	2	3	4
0	0	1	1	0	0
1	1	0	1	1	1
2	1	1	0	1	0
3	0	1	1	0	1
4	0	1	0	1	0

Adding a node and graph traversal are  $O(V^2)$ .

 $M_{u,v} = 1$  iff u has a directed edge to v

### **Code Representation**

### 2. Adjacency List (List of LinkedLists)



Complexity:  $\bigcirc$  ( $\lor$  +  $\lor$ )

Definition 00000

## Adjacency List in C++

```
vector<int> adj[10001];
int main() {
    int n, m;
    cin >> n >> m; // read in vertices and edges
    for (int i = 0; i < m; i++) { // m edges follows</pre>
        int u, v;
        cin >> u >> v; // read in edge between u and v
        adj[u].push back(v);
        adj[v].push_back(u); // if undirected, must add both ways
```

# **DFS & BFS**

## Depth-First Search (DFS)

- Explore as deep as possible, backtrack once branch is fully explored.
- Implement recursively or with a stack.

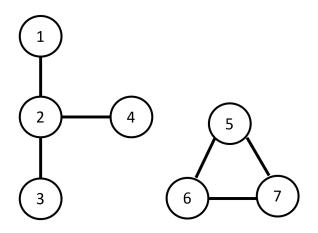
```
2 vector<int> adj[1001];
 vector<int> vis[1001];
 void DFS(int curr){
      vis[curr] = 1; // mark it visited
      for(int &next : adj[curr]){
          if(!vis[next]){ // if has not visited
              DFS(next); // step in
```

## **Breadth-First Search (BFS)**

- Explore as wide as possible, look through all neighbors before moving on.
- Implement using a queue.

```
vector<int> adj[10001];
vector<int> vis(10001);
int main() {
   int source = 1; // start at node 1
   queue<int> a:
   q.push(source); // start BFS with source
   vis[source] = 1; // mark as visited
   while(!q.empty()) {
        int curr = q.front();
        q.pop();
        for (int i = 0; i < adj[curr].size(); i++) {</pre>
            int next = adj[curr][i];
            if (!vis[next]) {
                vis[next] = 1:
                q.push(next);
```

## **Aside - Connected Components**



Is this a graph?

# **Graph Reduction**

### **Molecule Interactions**

**Problem**: A pharmaceutical company has two sets of molecules and wants to test a hypothesis: whether or not each molecule only interacts with molecules from the other group. Output "Yes" if the hypothesis holds (there are no intra-interactions) or "No" otherwise.

**Input**: A single row containing n, the # of molecules, and m, the # of interactions. m rows will follow, each containing two numbers representing an interaction between molecules.

## **Example**

### Example 1

Input:

42

12

3 4





Output: Yes

### Example 2

Input:

33

Graph Reduction 0000000

12

23

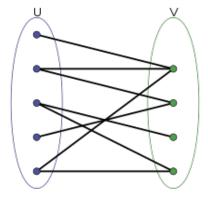
31



Output: No

### **Observation**

We can separate the two groups, circle them, and map each interaction. If we find that no two molecules of the same group have an interaction, we know the answer! This is a test of bipartiteness.



**Theorem**: A graph is bipartite iff it is two-colorable.

### **Prime Path**

Given two 4-digit prime numbers x and y, what is the minimum amount of steps needed to change x into y? You can modify one digit each time to any digit of your choice, but each intermediate number also has to be prime.

**Input**: 1033 8179

Output: 6

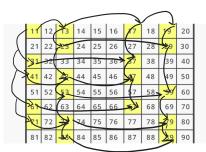
**Explanation**: 1033  $\rightarrow$  1733  $\rightarrow$  3739  $\rightarrow$  3779  $\rightarrow$ 

 $8779 \rightarrow 8179$ 

### **Prime Path**

Note: This is an example with 2-digit numbers

- Need to do two things first:
  - Find all prime numbers
  - Identify all those that are 1 change away



### **Prime Path**

### Explore with BFS to find the shortest path



# A Little More to Think About

## Algorithmic Use of BFS/DFS

What are some of the other ways we could use/modify BFS/DFS?

- Dijkstra
- Bellman-Ford
- Double-Ended Parallel