INF280: Competitive programming

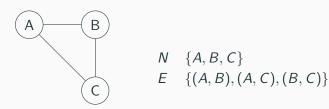
Basic graph traversals

Louis Jachiet

Introduction

You all know graphs:

- Set of nodes N
- Set of edges $E \subseteq N \times N$
- Edges can be undirected or directed, i.e., $(a, b) \neq (b, a)$



Data Structures

Several options to represent graphs:

- Adjacency matrix:
 - bool G[MAXN][MAXN];
 - G[x][y] is true if an edge between node x and y exists
 - Replace bool by int to represent weighted edges
- Adjacency list:
 - vector<int> Adj[MAXN];
 - y is in Adj[x] if an edge between node x and y exists
 - Pairs to represent weights
- Edge list:
 - vector<pair<int, int> > Edges;
 - Edges contains a pair of nodes if an edge exists between them

Nodes and edges may also be custom structs or classes

Simple Traversals

Simple Traversals

Depth-First Search

Depth-First Search

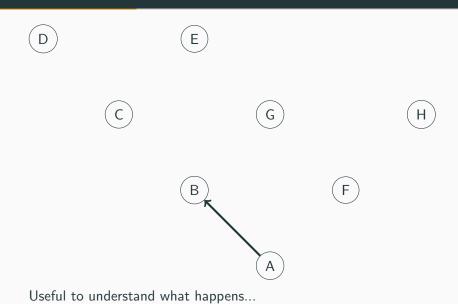
Visit each node in the graph once:

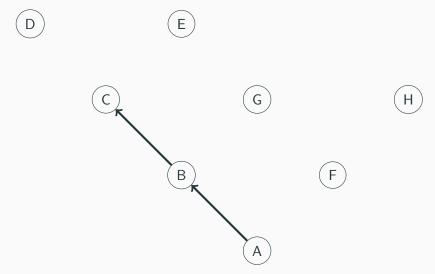
- Recursive implementation below
- Manage stack yourself for iterative version
- First visit child nodes then siblings

```
int state[ID_NODE_MAX] ;
const int NOT_VISITED = 0, IN_VISIT = 1 , VISITED = 2 ;
void dfs(int node) {
  if(state[node] == NOT_VISITED) {
    state[node] = IN_VISIT ;
    for(auto v : nxt[node])
      dfs(v);
    state[node] = VISITED ;
  }
}
```

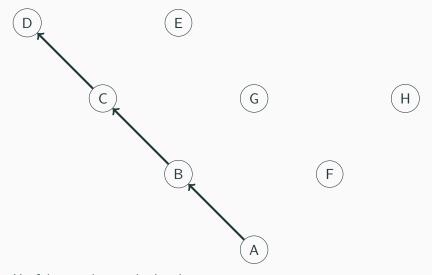
Applications of DFS

- Determine a topological order of nodes
- Detect if a cycle exists
- Check reachability between nodes
- Decompose graph into connected components
- Decompose graph in strongly connected components
- Examples: https://visualgo.net/dfsbfs

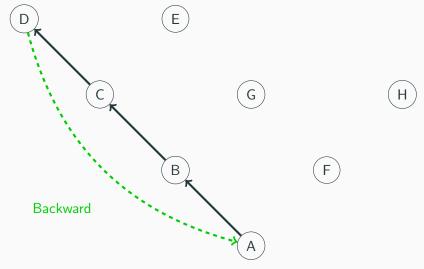




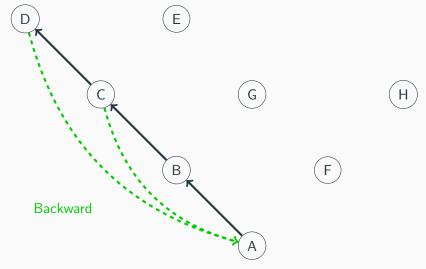
Useful to understand what happens...



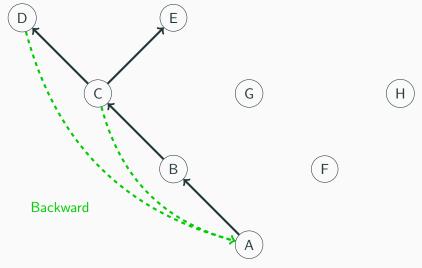
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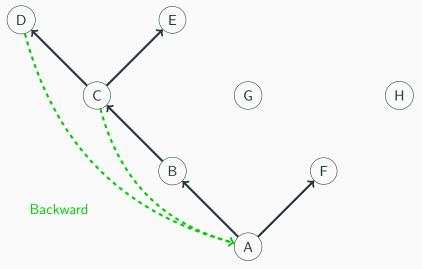
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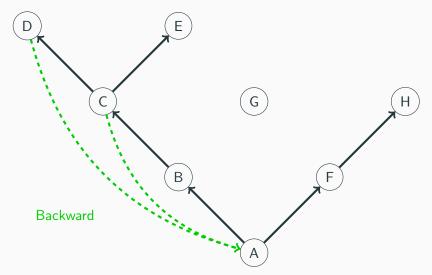
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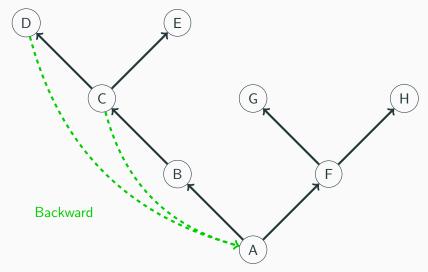
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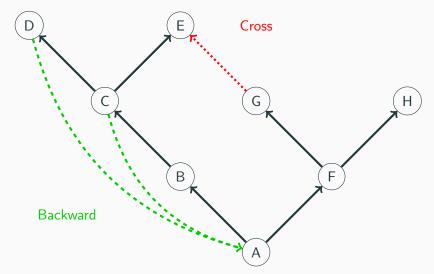
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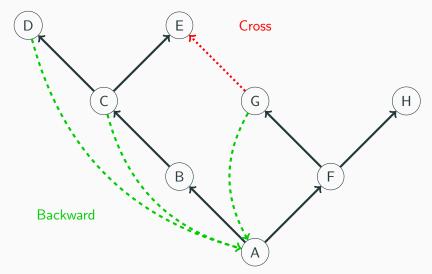
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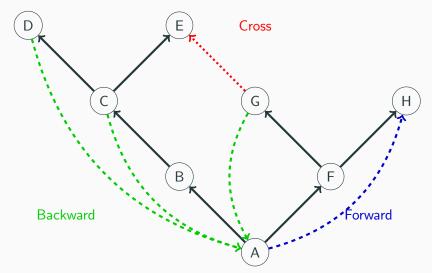
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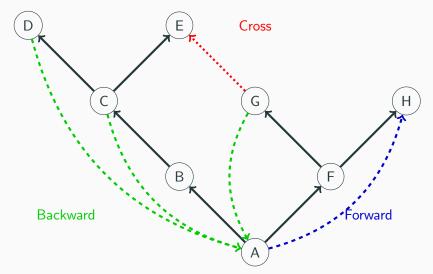
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Exercise: compute Strongly Connected Component

Simple Traversals

Breadth-First Search

Breadth-First Search

Visit each node in the graph once:

• Similar to DFS, but replaces stack by queue

```
int seen[NB_NODE_MAX] ;
void bfs(int start) {
  vector<int> todo = {start} ;
  seen[start] = true ;
  for(size_t id = 0 ; id < todo.size() ; id++)</pre>
    for(auto v : nxt[todo[id]])
      if(!seen[v]) {
        seen[v] = true;
        todo.push_back(v);
      }
```

Applications of BFS

- Shortest path search
 - Stop processing when a given node d was found
 - Minimizes number of hops, i.e., all edges have same weight or 0-1 Weights
 - Generalization follows next
- Examples: https://visualgo.net/dfsbfs

Simple Traversals

Simple Travelsais

0-1 Breadth-First Search

Breadth-First Search with edges of bounded distance

```
vector<int> nodes_at[MAX_DISTANCE];
void bfs(int start) {
  fill(dist,dist+NB_NODES_MAX,INF);
 nodes_at[0] = {start} ;
 dist[start] = 0;
  for(int cur_dist = 0 ; cur_dist < MAX_DISTANCE ; cur_dist++ )</pre>
    for(size_t id = 0 ; id < nodes_at[cur_dist].size() ; id++) {</pre>
      const int node = nodes_at[cur_dist][id] ;
      if(dist[node] == cur_dist)
        for(auto [neigh,len] : nxt[node])
          if(dist[neigh] > cur_dist+len) {
            dist[neigh] = cur_dist+len ;
            nodes_at[dist[neigh]].push_back(neigh);
```

Finding Paths

Finding Paths

Dijkstra

Dijkstra

- Dijkstra's algorithm generalizes BFS
- Constraint: all edges need to have non-negative weights
- Use a priority queue to choose which node to examine next

Finding Paths

Bellman-Ford

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Bellman-Ford DP problem: "q(n, k) is the minimal distance of n from the source node using k intermediate edges"

Bellman-Ford can also be seen as a way to solve a linear system with inequalities of the form: $x_i + c_i < y_i$

Bellman-Ford Algorithm

```
int from[MAX_NB_EDGES], to[MAX_NB_EDGES], weight[MAX_NB_EDGES];
int dist[MAX_PATH_LENGTH+1] [MAX_NB_NODES];
bool BellmanFord(int root) {
 fill(dist[0], dist[MAX_PATH_LENGTH], INF);
 dist[0][root] = 0:
 for(int len = 0 ; len < MAX_PATH_LENGTH ; len++)</pre>
    for (int e = 0 ; e < nb_edges ; e++)</pre>
      dist[len+1][to[e]] = min(dist[len+1][to[e]],
                              dist[len][from[e]]+weight[e]);
 // to be explained later; check for negative cycles
 return dist[MAX_PATH_LENGTH+1][target];
```

- replace dist[l][n] with dist[n] = $min_I(dist[l][n])$
- MAX_PATH_LENGTH is at most nb_nodes long

Bellman-Ford Algorithm

```
int dist[MAX_NB_NODES];
void BellmanFord(int root, int target) {
  fill(dist, dist+MAX_NB_NODES, INF);
  dist[root] = 0;
  for(int k = 0 ; k < nb_nodes - 1 ; k++) // N - 1 times
    for (int i = 0 ; i < nb_edges ; i++)
        dist[to[i]] = min(dist[to[i]], dist[from[i]]+weight[i]);
}</pre>
```

```
bool detect_negative_cycle_BellmanFord(int root, int target) {
  fill(dist, dist+MAX_NB_NODES, INF);
  dist[root] = 0;
  for(int k = 0; k < nb\_nodes - 1; k++) // N - 1 times
   for (int i = 0 ; i < nb_edges ; i++)
      dist[to[i]] = min(dist[to[i]], dist[from[i]]+weight[i]);
 // now time to check for negative cycles:
  int dist_target = dist[target]; // copy distance
  for(int k = 0 ; k < nb_nodes - 1 ; k++) // N - 1 times</pre>
    for (int i = 0 ; i < nb_edges ; i++)
      dist[to[i]] = min(dist[to[i]], dist[from[i]]+weight[i]);
  return dist[target] < dist_target ; // negative cycle?</pre>
```

Finding Paths

Floyd-Warshall

Floyd-Warshall

- Dijkstra and Bellman-Ford compute shortest paths
 - From a single source (root)
 - To all other (reachable) nodes
 - This is known as: single-source shortest path problem
- Floyd-Warshall extends this to compute the shortest paths between all pairs of nodes
- This is known as: all-pairs shortest path problem

Floyd-Warshall

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Floyd-Warshall answers the DP problem: "q(start,end,pivot): what is the shortest path between start and end going through intermediate nodes 1..pivot?"

Floyd-Warshall Algorithm

```
int dist[MAX_NB_NODES] [MAX_NB_NODES];
// We store q(start, end, pivot) in dist[start][end]
void FloydWarshall() {
  // initialize distance
  fill(dist[0], dist[MAX_NB_NODES], INF);
  for (int e = 0 ; e < nb_edges ; e++)
    dist[fr[e]][to[e]] = min(dist[fr[e]][to[e]], weight[e]);
  // now compute
  for(int pivot = 0 ; pivot < nb_nodes ; pivot++)</pre>
    for(int start = 0 ; start < nb_nodes ; start++)</pre>
      for(int end = 0 ; end < nb_nodes ; end++)</pre>
        dist[start][end] = min(dist[start][end],
                   dist[start][pivot]+dist[pivot][end]);
// WARNING, the order of the loops is important!!!
// for french speakers Pivot Début Fin => PDF algorithm
```

Finding Paths

Improvements

Keeping track of the path

We only considered the length of the path so far:

- All of the above algorithms can track the actual shortest path
- This allows to *print* each edge/node along the path
- Basic idea:
 - Introduce an array int Predecessor [MAXN]
 (Use two-dimensional array for Floyd-Warshall)
 - Updated whenever Dist[v] changes
 - Simply set to the new predecessor u

Heuristics – A* Search

Heuristics may speed-up the path search

- Bellman-Ford and Floyd-Warshall equally explore all possibilities
- Dijkstra prefers nodes with shorter distance
- Basic idea behind A* Search:
 - Extension to Dijkstra's algorithm
 - Introduce an estimation of the remaining distance
 - Prefer nodes with minimal estimated remaining distance
- Advantages
 - May converge faster than Dijkstra
 - Can be used to compute approximate solutions (trading speed for precision)

Eulerian Circuits

Eulerian Circuits

Eulerian path

Use every edge of a graph exactly once. Start and end may differ

Eulerian circuit

Use every edge exactly once. Start and end at the same node

Idea of the algorithm

If you enter a node of even degree you are sure that you can go out, decreasing the degree of unused by 2. This gives a first path/circuit. If your graph is connected, you can have remaining edges unexplored, but at least one in your current path, so you can re-explore them.

We will see more graph algorithms next week...