CIS 1210—Data Structures and Algorithms—Fall 2024

BFS & DFS—Tuesday, October 15 / Wednesday, October 16

Readings

• [Lecture Notes Chapter 16: Graph Traversals: BFS and DFS](https://www.cis.upenn.edu/~cis1210/current/lectures/notes.pdf#page=97)

Review: Graph Representations

Let $G = (V, E)$, where $|V| = n$ and $|E| = m$. In other words, G contains n vertices and m edges.

Adjacency Matrix

We can represent G with an $n \times n$ matrix A, where $A[i][j] = 1$ if there is an edge (i, j) and 0 otherwise. The primary advantage of an adjacency matrix is that we can check whether or not there is an edge between two vertices in $O(1)$ time since all we have to do is index into the matrix. On the other hand, the matrix uses $\Theta(n^2)$ space, and it takes $\Theta(n)$ time to enumerate the neighbors of a vertex v since we must iterate through an entire row in the matrix.

Adjacency List

We can also represent G as an adjacency list, where we have an array A of $|V|$ lists such that $A|u|$ contains a LinkedList of vertices v, such that for every vertex v_i in $v, (u, v_i) \in E$. (Note that if we have a directed graph, we can store two LinkedLists at index u , one for u 's in-neighbors and one for u 's out-neighbors.) The primary advantage of an adjacency list is that we only use $\Theta(n+m)$ space, which is better than the $\Theta(n^2)$ space usage of an adjacency matrix especially if $m \ll n^2$. Furthermore, it only takes $O(deg(v))$ time to enumerate the neighbors of a vertex v, but on the other hand, checking whether $(u, v) \in E$ also takes $O(deg(v))$ time since we have to traverse the LinkedList. In CIS 1210, we usually use adjacency lists.

Review: BFS (Breadth First Search)

At a high-level, in BFS, we explore the graph in "layers," so we explore "wide" before exploring "deep." We begin at a node v (layer 0); then explore the children of v (layer 1); then the children of the nodes in layer 1 (which make up layer 2); and so on. In other words, we explore all nodes at layer L_i before exploring any nodes at layer L_{i+1} . Because we explore the graph in layers, as seen in the [pseudocode,](https://www.cis.upenn.edu/~cis1210/current/lectures/notes.pdf#page=99) we use a queue to implement this algorithm since it is FIFO. Note that the output of BFS is a BFS tree, and that as written, BFS is not guaranteed to visit every node in the graph.

The running time of BFS is $O(n + m)$. This is because we add and remove each node from the queue at most once, and for each node, we only examine its neighbors exactly once.

Review: DFS (Depth First Search)

In DFS, we explore "deep" before exploring "wide." We begin at some node v and examine its neighbors. When we encounter a neighbor that has not been visited yet, we visit it. Once we arrive at a node whose neighbors have all been visited, we "backtrack," via returning from recursive calls, until we reach a node that still has unvisited neighbors. Because of this, DFS can be implemented recursively as seen in this [pseu](https://www.cis.upenn.edu/~cis1210/current/lectures/notes.pdf#page=102)[docode,](https://www.cis.upenn.edu/~cis1210/current/lectures/notes.pdf#page=102) but we can also implement it iteratively by using a stack since it is LIFO. The iterative pseudocode for DFS with a stack is the BFS [pseudocode,](https://www.cis.upenn.edu/~cis1210/current/lectures/notes.pdf#page=99) except instead of a queue we use a stack, instead of dequeue we use pop, and instead of enqueue we use push. We know that DFS will visit every node in the graph, so the output of DFS is a DFS forest. Note that DFS also introduces the concept of colors and finishing times, which makes it useful for other graph applications later on.

The running time analysis for DFS is similar to that of BFS; for each vertex v , we iterate through its neighbors to yield a runtime of $O(n + m)$.

Given a DFS forest of an input graph, we can classify its edges into four types, each of which can provide important information about the graph. The four types of edges are:

- 1. Tree Edges: Edges in the DFS forest
- 2. Back Edges: Edges (u, v) where v is a ancestor of u
- 3. Forward Edges: Non-forest edges (u, v) where v is a descendant of u
- 4. Cross Edges: All remaining edges

Problems

Problem 1

Design an $O(n+m)$ algorithm to find the shortest path between nodes u and v in a connected, unweighted graph.

Solution

Algorithm: Run BFS starting from u; for each node x that is visited, maintain a "parent" pointer to its parent node (the node we visit x from). Once we visit v , stop and backtrack through these parent pointers until we reach u, outputting this path as the shortest path. For example, by backtracking, we could see that v's parent was d, d's parent was c, and c's parent was u, so the path outputted would be $u-c-d-v$.

Proof of Correctness: The correctness of this algorithm follows from the lemma that BFS outputs the shortest paths in an unweighted graph. That is, for a vertex v in layer L_i , i represents the length of the shortest path from the source u to v. We prove this lemma via strong induction on i :

Base Case: If $i = 0$, then by design, L_0 only contains the source vertex u. The shortest path from u to u has length 0, so the claim holds.

Induction Hypothesis: Assume that for all vertices v such that v is in layer L_k , where $k < i$, that k is the length of the shortest path from u to v .

Induction Step: We want to show that the claim holds for some arbitrary vertex v in layer L_{k+1} . By design, v was discovered from another vertex w, such that there is an edge between w and v and w is in layer L_k . By IH, we know that the shortest path from source u to vertex w has length k, so by adding the edge from w to v, we get a path of length $k+1$ from source u to vertex v. We want to show that this path is a shortest path from u to v. Assume for the sake of contradiction that some other path $u - \cdots - w' - v$ is the true shortest path, meaning that it must have length $l \leq k$ and that w' is reachable from u by a shortest path with length less than or equal to $k-1$. Thus, we know that vertex w' is in some layer L_{l-1} , where $l-1 \leq k-1$. By design, this means that w' was removed from the BFS queue before w , implying that when we processed w' , we would have discovered v. However, this means we would have added v to layer L_l , and since $l \leq k$, we have reached a contradiction because we stated that v was an arbitrary vertex in layer L_{k+1} . Therefore, we have shown that the length of the shortest path from u to v is in fact $k + 1$, completing our Induction Step and thus our proof.

From the lemma, we know that BFS correctly computes the shortest paths from the source in an unweighted graph. In our algorithm, note that once we visit v , we stop and backtrack through its parent pointers until we reach u, so we just backtrack through the layers to yield the shortest path from u to v. Hence, outputting this path correctly yields the shortest path from u to v .

Runtime Analysis: Running BFS takes $O(m+n)$ time, and to output the shortest path, we backtrack no more than $O(n)$ times since the longest path in a graph has $n-1$ edges. Therefore, our algorithm runs in $O(m+n)$ time.

Problem 2

The CIS Department wants to assign prerequisites to their n classes. It has a list of m prerequisite pairings, such as CIS 110 being a prerequisite for CIS 120. Given the list of prerequisite pairings, design an algorithm that determines if this list of pairings is a valid list of prerequisites. For example, (CIS 110, CIS 120), (CIS 120, CIS 121), (CIS 121, CIS 110) is not a valid list of prerequisites. What is the runtime of your algorithm?

Solution

Algorithm: Construct a graph G , where each of the vertices is one of the n classes in the CIS department and there is a directed edge (u, v) if u is a prerequisite for v. Run DFS on G. If the DFS traversal finds a back edge during its execution, return that this list is invalid. Otherwise, return that this is list is valid.

Proof of Correctness: If u is a prerequisite for v, then by definition, we know that students must take course u before taking course v. Directing an edge from u to v thus translates this exact information and nothing more, so G sufficiently translates all of the information given to us. Observe that a list of prerequisites is valid iff G does not contain a cycle. Now, we want to show that G has a cycle iff a DFS traversal finds a back edge:

 (\Rightarrow) If G has some cycle C, then let v be the first vertex on this cycle discovered by DFS and let (u, v) be the edge directly preceding v on C . When v is discovered, we know that all nodes in C are still undiscovered and colored white. By definition, v is an ancestor of u in the DFS forest, so we know (u, v) is a back edge.

 (\Leftarrow) If there is a back edge (u, v) , then by definition, we know that there is a path from v to u made of tree edges. Concatenating this path from v to u with the back edge (u, v) creates a cycle.

Therefore, since G has a cycle iff a DFS traversal finds a back edge, we know that our algorithm is correct.

Runtime Analysis: Constructing G as an adjacency list takes $O(n + m)$ time. Running DFS on G and determining if there are any back edges takes $O(n + m)$ time as well, so this algorithm runs in $O(n + m)$ time.