

1. What is the running time of DFS if the graph is given as an adjacency list and adjacency matrix? Justify your running time.

Solution:

If the number of nodes in the graph is V and the number of edges in the graph is E , the running time are:

Adjacency list: $O(V + E)$

Adjacency matrix: $O(V^2)$

In DFS, we will visit every node once and visit those nodes are adjacent to them. In both case, we first need $O(V)$ time to visit all nodes.

If we use adjacency list, we only need to visit every edge in each list, which is $O(E)$. So, the total running time is $O(V) + O(E) = O(V + E)$.

If we use adjacency matrix, although there are only E edges, once we visit a node (i.e. the i^{th} node), we will visit the element in the i^{th} row of the adjacency matrix. In total, we will visit every element in the adjacency matrix (no matter it is 0 or 1), which is V^2 . So the total running time will be $O(V^2) + O(V) = O(V^2)$.

2. Write a method that takes any two nodes u and v in a tree T , and quickly determines if the node u in the tree is a *descendant* or *ancestor* of node v

Solution:

We can simply run DFS on a tree T and set the start node as the root and record the discover time and finish time of every node. Because T is a tree, the number of edges is $n - 1$. So, the total running time is $O(V + E) = O(n + n - 1) = O(n)$. As we have proved in the textbook, given any 2 nodes u, v in the DFS tree, if $u.d < v.d < v.f < u.f$, u is the ancestor of v . Else if $v.d < u.d < u.f < v.f$, u is the descendant of v . Because we only need $O(n)$ time to complete DFS and store the discover time and finish time. Other operations are in constant time ($O(1)$), so the total running time is $O(n)$. More specifically, we need $O(n)$ time to initialize and $O(1)$ time to determine the relationship between 2 nodes.

3. (u (v (y (x x) y) v) u) (w (z z) w)

4.

Alg: find_subsets:

procedure subsets(nums: array):

var combinations : array

dfs(nums, 0, [], combinations)

return combinations

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precedure dfs(nums: array, index: integer, combination: array, combinations, array):
    combinations.append(combination.copy)// insert the copy of combination to combinations
begin
    for i := index to nums.length:
        combination.append(nums[i])
        dfs(nums, i + 1, combination, combinations)
        combination.pop()
end;
```