Disjoint Set ADT

-The implementation of disjoint-set data structure using trees, particularly with the union by rank method, offers several advantages over other data structures like arrays

Efficient Union and Find Operations

- -In a tree-based implementation, the union operation is typically faster compared to array-based implementations.
- -By linking smaller trees to larger ones based on their rank, the height of trees is kept low, resulting in faster find operations as well.

Balanced Trees

- -Union by rank ensures that the resulting trees remain balanced.
- -Balanced trees prevent degenerate cases where the tree becomes essentially a linked list, ensuring that find operations remain efficient even in worst-case scenarios.

Optimized Find Operation

- -The find operation in a tree-based disjoint-set structure has a time complexity of $O(\log n)$ in the worst case, where n is the number of elements.
- -This is because path compression is applied during the find operation, which flattens the tree structure, reducing the height of the tree.

Memory Efficiency

- -Tree-based implementations typically require less memory compared to array-based implementations, especially when dealing with large sets of elements.
- -This is because only the parent pointer, rank, and possibly some additional metadata are stored for each element, rather than allocating memory for an entire array.

Scalability

- -Tree-based implementations scale well with the number of elements in the disjoint sets.
- -Even for large sets of elements, the time complexity of union and find operations remains logarithmic, making tree-based implementations suitable for a wide range of applications.

Flexibility

- -Tree-based disjoint-set implementations offer flexibility in terms of the underlying data structure.
- -Different variations, such as using path compression or weighted union, can be applied based on the specific requirements of the application.

Overall, tree-based implementations with union by rank strike a balance between efficiency, memory usage, and scalability, making them a popular choice for implementing disjoint-set data structures in various algorithms and applications.

Algorithm 1: Creating a set

Input: capacity, nodes(array), data (integer).

Output: newnode (pointer to a node).

- 1. Create a newnode.
- 2. If capacity is lesser than or equal to data
 - a. new_capacity<-data+1and dynamically resize the internal array (nodes) to accommodate the new data.
- 3. Initialise i to 0
- 4. Repeat till i is capacity
 - a. new_nodes[i] <- nodes[i]
 - b. Increment i
- 5. Initialise i to capacity
- 6. Repeat till i is new_capacity
 - a. new_nodes[i] is NULL
- 7. Delete nodes
- 8. nodes <- new nodes
- 9. capacity <- new_capacity
- 10. If nodes[data] is NULL
 - a. Delete newnode
 - b. Return NULL
- 11. newnode->data <- data
- 12. newnode->parent <- newnode
- 13. newnode->rank <- 0
 - a. If nodes[data] is NULL
 - i. nodes[data] <- newnode</pre>
 - b. Otherwise
 - i. Delete newnode
- 14. Return newnode

Algorithm 2: Finding root of the tree by path compression

Input: capacity, nodes(array), ele (integer).

Output: root of the tree.

1. If ele lesser than 0 or ele greater than or equal to capacity or nodes[ele] is NULL

- a. Return -1
- 2. Create a pointer temp <- nodes[ele]
- 3. If temp is not temp->parent
 - a. root_data <- find(temp->parent->data)
 - b. temp->parent <- nodes[root_data]</p>
- 4. Return temp->parent->data

Algorithm 3: Merge two sets by rank

Input: nodes(array), ele_1 and ele_2 (integer).

Output: true (success) / false (failure).

- 1. root_1 <- find(ele_1)
- 2. root_2 <- find(ele_2)
- 3. If root_1 is -1 and root_2 is -1 (elements not found)
 - a. Return false
- 4. Otherwise if root 1 is -1 (ele 1 not found)
 - a. Return false
- 5. Otherwise if root_2 is -1 (ele_2 not found)
 - a. Return false
- 6. If root_1 is root_2 (both elements are in the same set)
 - a. Return false
- 7. If nodes[root_1]->rank is lesser than nodes[root_2]->rank
 - a. nodes[root 1]->parent <- nodes[root 2]</pre>
- 8. Otherwise if nodes[root_1]->rank is greater than nodes[root_2]->rank
 - a. nodes[root_2]->parent <- nodes[root_1]</pre>
- 9. Otherwise
 - a. nodes[root_2]->parent <- nodes[root_1]</pre>
 - b. nodes[root_1]-> rank++
- 10. Return true

FUNCTIONAL REQUIREMENTS:

- 1. create()
- 2. find()
- 3. merge()

TIME COMPLEXITY ANALYSIS FOR EACH ALGORITHM:

1. $\underline{\text{create()}} - \underline{\text{O(n)}}$

$$f(n) = 4n + 19$$

$$g(n) = n \text{ and } c = 10$$

n	Cost of f(n)	Cost of c.g(n)
0	19	0
1	23	10
2	27	20

3	31	30
4	35	40

Therefore n_0 = 4 for which f(n) <= c.g(n) is true for all n_0 < n . Therefore f(n) = O(n)

Analysis summary:

Best Case: O(1)Average Case: O(1)Worst Case: O(n)

Explanation:

- Best Case (O(1)): This occurs when the data value (data) being inserted is already present in the nodes array (i.e., nodes[data]!= NULL). In this case, the function simply checks for the existing value, prints an error message, and frees the allocated memory for newnode. All these operations are constant time (O(1)).
- Average Case (O(1)): Assuming a random distribution of data insertions, the average case complexity might be closer to constant time (O(1)). This means that on average, most insertions might not trigger capacity expansion and can be handled in constant time.
- Worst Case (O(n)): The worst case occurs when the data value (data) is significantly larger than the current capacity of the nodes array.

2. find() - O(log n)

$$f(n) = \log n + 5$$

$$g(n) = \log n \text{ and } c = 6$$

n	Cost of f(n)	Cost of c.g(n)
0	-	-
1	5	0
2	5.301	1.806
3	5.477	2.862
4	5.602	3.612
5	5.698	4.193
6	5.778	4.668
7	5.845	5.070
8	5.903	5.418
9	5.954	5.725
10	6	6
11	6.041	6.248

Therefore n_0 = 10 for which f(n) <= c.g(n) is true for all n_0 < n . Therefore f(n) = $O(\log n)$

Analysis summary:

- Best Case: O(1)

Average Case: O(log n)

- Worst Case: O(h)

Explanation:

- Best Case (O(1)): This occurs when the element (ele) is the root node itself (i.e., temp->parent == temp). In this case, the function directly returns temp->parent->data which is temp->data, avoiding the recursive call. All operations involved are constant time (O(1)).
- Average Case (O(log n)): Assuming balanced trees (like AVL trees), the average depth of a node is logarithmic in the number of nodes n (depth = log(n)). Therefore, the find function would typically need to make a logarithmic number of recursive calls (one for each level) to reach the root from a given node. Each call itself is constant time (O(1)). This results in an average-case time complexity of O(log n).
- Worst Case (O(h)): The worst case occurs when the tree is skewed, meaning all nodes lean to one side. In this scenario, the function might need to traverse all the way up to the root from the given node, resulting in h recursive calls (where h is the height of the tree). Each call takes constant time (O(1)). This leads to a worst-case time complexity of O(h).

3. merge() - O(log n)

$$f(n) = 2 \log n + 11$$

 $g(n) = \log n \text{ and } c = 12$

n	Cost of f(n)	Cost of c.g(n)
0	-	-
1	11	0
2	11.602	3.612
3	11.954	5.725
4	12.204	7.224
5	12.397	8.387
6	12.556	9.337
7	12.690	10.141
8	12.806	10.837
9	12.908	11.450
10	13	12
11	13.082	12.496
12	13.158	12.950
13	13.227	13.367
14	13.292	13.753

Therefore n_0 = 13 for which f(n) <= c.g(n) is true for all n_0 < n . Therefore f(n) = $O(\log n)$

<u>Analysis summary:</u>

Best Case: O(1)

- Average Case: O(log n)

- Worst Case: O(h)

Explanation:

- Best Case (O(1)): If the elements are already roots (single-node trees), merging them is very fast (constant time).

- Average Case (O(log n)): Assuming balanced trees (like AVL trees), finding the roots of elements takes a logarithmic amount of time on average. Merging based on rank is still constant time. This leads to an overall average complexity of O(log n).
- Worst Case (O(h)): In the worst case, the elements might belong to very unbalanced (skewed) trees. Finding the root in such trees can take linear time (proportional to the height h of the tree). This makes merging slower in this scenario.

APPLICATIONS:

Disjoint-set ADT (Union-Find data structure) finds various applications in computer science due to its efficient management of disjoint sets (non-overlapping sets). Here are some key areas where it's used:

1. Kruskal's Algorithm:

- This algorithm finds the minimum spanning tree (MST) of a weighted undirected graph. It works by iteratively processing edges in ascending order of weight. Disjoint-set is used to efficiently:
- Track connected components: Initially, each vertex represents a separate component. When an edge connects vertices from different components, the find operation helps determine which components to merge using the unionoperation.
- Avoid cycles: The find operation ensures that merging edges wouldn't create cycles in the MST.

2. Image Segmentation:

- In image processing, disjoint-set helps identify connected regions of similar pixels (e.g., objects in an image). Here's how:
- Initially, each pixel represents a separate set.
- Adjacent pixels with similar properties (e.g., color) are considered for merging.

- The find operation determines if adjacent pixels belong to the same region.
- If not, the union operation merges them into a single set, representing a connected object.

3. Percolation Simulation:

- This technique models systems with connected and disconnected regions, like studying fluid flow through porous materials. Disjoint-set can be used to:
- Track connected clusters: Initially, each site in the grid is a separate set.
- When a site becomes open (fluid can flow through), it's connected to its open neighbors using the union operation.
- The find operation helps determine if two sites belong to the same connected cluster, indicating fluid flow between them.

4. Network Connectivity:

- In network management, disjoint-set can be used to:
- Track connected components in a network: Each device or router can initially represent a separate set.
- When a connection is established between devices, the union operation merges their sets, representing a connected network segment.
- The find operation helps determine if two devices are part of the same network segment.

5. Equivalence Checking:

- This involves determining if two expressions or objects are equivalent based on certain rules. Disjoint-set can represent equivalence classes, and the union operation merges equivalent objects.

Additional Applications:

- Social network analysis: Grouping users based on interactions.
- Code optimization: Identifying common subexpressions in compiler optimization.
- Game development: Tracking connected entities in a game world.