

analyze of Kruskal's algorithm

1. Data Structures:

- **edge[n][3]:** This 2D array represents the edges of the graph. Each row in the array represents an edge, with:
 - `edge[i][0]`: The source vertex of the edge.
 - `edge[i][1]`: The destination vertex of the edge.
 - `edge[i][2]`: The weight of the edge.
- **parent[n]:** An array to store the parent of each vertex in the Minimum Spanning Tree. Initially, each vertex is its own parent.
- **rank[n]:** An array to store the rank of each vertex in the union-find data structure, used to optimize the union operation.

2. Algorithm Steps:

- **Sorting:** The edges are first sorted in ascending order of their weights using the `qsort` function and a custom comparator function. This ensures that edges are considered in increasing order of weight.
- **Initialization:** The `makeSet` function initializes the parent and rank arrays for each vertex, setting each vertex as its own parent and rank to 0.
- **MST Construction:**
 - For each edge in the sorted list:
 - Find the parent of the source and destination vertices using the `findParent` function.
 - If the parents are different, it means the vertices belong to different components.
 - Union the two components using the `unionSet` function.
 - Add the weight of the edge to the `minCost`.
- **Output:**
 - Print the edges included in the MST.
 - Print the total minimum cost of the MST.

3. Time Complexity:

- **Sorting:** Sorting the edges using qsort takes $O(E \log E)$ time, where E is the number of edges.
- **Initialization:** The makeSet function takes $O(V)$ time, where V is the number of vertices.
- **Union-Find:** The findParent and unionSet operations, when implemented with path compression and union by rank, have amortized time complexity of $O(\alpha(V))$, where α is the inverse Ackermann function, which grows extremely slowly. Therefore, the total time complexity of the union-find operations is $O(E \alpha(V))$.

Overall Time Complexity:

Since E (number of edges) is usually proportional to V^2 in a graph, the overall time complexity of Kruskal's algorithm is **$O(E \log E)$** or **$O(V^2 \log V)$** .

4. Space Complexity:

The space complexity of Kruskal's algorithm is $O(V + E)$ to store the graph, parent array, and rank array.

analyze for heap sort.

1. Algorithm:

The code implements the heap sort algorithm, which sorts an array in ascending order by using a binary heap data structure.

2. Key Steps:

- **Heapify:**
 - The heapify function is the core of the algorithm.
 - It ensures that the subtree rooted at a given node satisfies the heap property (parent node is greater than or equal to its children).
 - It recursively sifts down the node until the heap property is maintained.
- **Building the Heap:**
 - The heapSort function starts by building a max-heap from the given array.

- It iterates through the array from the last non-leaf node ($n/2 - 1$) to the root (0) and calls heapify for each node.
- **Sorting:**
 - The heapSort function then repeatedly extracts the maximum element (root) from the heap and swaps it with the last element of the unsorted portion.
 - The size of the heap is reduced by one after each extraction, and the heap is re-heapified to maintain the heap property.

3. Time Complexity:

- **Building the Heap:** Building the initial heap takes $O(n)$ time.
- **Extracting and Re-heapifying:** In each iteration, extracting the maximum element takes $O(1)$ time, and re-heapifying takes $O(\log n)$ time. This is repeated $n-1$ times, resulting in a total time complexity of $O(n \log n)$.

4. Space Complexity:

The space complexity of heap sort is $O(1)$ as it sorts the array in-place, without requiring any additional data structures.

5. Correctness:

The code correctly implements the heap sort algorithm and produces the expected sorted output.