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1) (70 pts) Consider all the leaves of a binary tree, from left to right order, the values of those leaves form a **leaf value sequence**. For example, in the given trees below, the leaf value sequence is (6, 7, 4, 9, 8). Two binary trees are considered **leaf-similar** if their leaf value sequence is the same. Return **true** if and only if the two given trees with head nodes **root1** and **root2** are leaf-similar.

2) (30 pts) Write one test case and time and space complexity.

Logic: ① Initialize a ~~map~~ unordered\_map.

② For the first tree, find leaf nodes and increment their count in the map.

③ For the second tree, find leaf nodes and decrement their count in the map.

④ Traverse through the map & return true if map value for all key elements are zero, else return false.

T.C 1

→ Both similar  
Return true

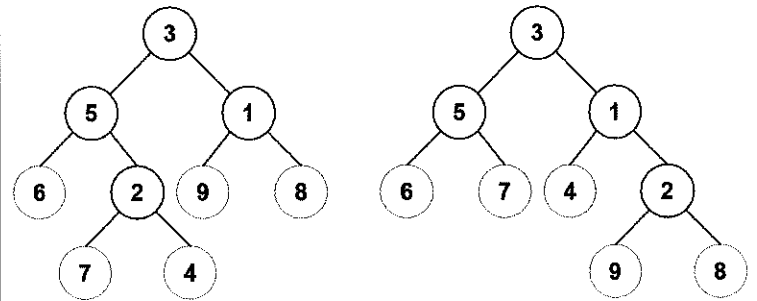
T.C 2

→ Different  
return false

Time complexity:  $O(N)$

Space complexity:  $O(N)$

```
void findLeafNodes(Node* root, unordered_map<int, int> um, int value) {
    if (!root) return;
    if (root->left == NULL && root->right == NULL) {
        um[root->data] += value;
    }
    findLeafNodes(root->left);
    findLeafNodes(root->right);
}
```



Input: root1 = [3,5,1,6,2,9,8,null,null,7,4], root2 =

[3,5,1,6,7,4,2,null,null,null,null,null,9,8]

Output: true

bool isLeafSimilar(Node\* root1, Node\* root2)

```
{ unordered_map<int, int> um;
  findLeafNodes(root1, um, 1);
  findLeafNodes(root2, um, -1);
```

```
for (auto ele : um)
{ if (ele.second != 0)
  { return false;
  }
}
return true;
```

```
findLeafNodes (Node* root,
  unordered_map<int, int> um,
  int value.
```