

## **Same Tree Documentation**

### **Problem Statement**

Given the roots of two binary trees, p and q, write a function to check if they are the same or not. Two binary trees are considered the same if they are structurally identical and the nodes have the same value.

### **Example 1**

**Input:** p = [1,2,3], q = [1,2,3]

**Output:** true

### **Example 2**

**Input:** p = [1,2], q = [1,null,2]

**Output:** false

### **Example 3**

**Input:** p = [1,2,1], q = [1,1,2]

**Output:** false

### **Constraints**

- The number of nodes in both trees is in the range [0, 100].
- $-10^4 \leq \text{Node.val} \leq 10^4$

## **Explanation**

### **1. Base Cases:**

- If both nodes p and q are None, then the trees are identical at this branch, so return True.
- If only one of the nodes is None, the trees are not identical, so return False.
- If the values of p and q are different, the trees are not identical, so return False.

### **2. Recursive Step:**

- Recursively check if the left subtrees of p and q are the same.
- Recursively check if the right subtrees of p and q are the same.
- Both conditions must be True for the trees to be identical.

## **Complexity**

- Time Complexity:  $O(N)$ , where N is the number of nodes in the trees. Each node is visited exactly once.
- Space Complexity:  $O(N)$  in the worst case due to the recursion stack, where the trees are skewed.

This solution efficiently checks the structural and value equivalence of two binary trees using a recursive approach.