

Problem Description In this problem, we are given two binary trees represented by their root nodes root1 and root2. We need to determine if one tree can be transformed into the other tree through a series of flip operations. A flip operation consists of swapping the left and right children

of a given node. The trees are considered flip equivalent if one can be transformed into the other by doing any number of flips, possibly zero. We should return true if the trees are flip equivalent and false otherwise. Intuition

To solve this problem, we use a recursive approach that is a form of Depth-First Search (DFS). The main idea is that if two trees are

flip equivalent, then their roots must have the same value, and either both pairs of the left and right subtrees are flip equivalent, or the left subtree of one tree is flip equivalent to the right subtree of the other tree and vice versa. To implement this idea, we compare the current nodes of both trees:

2. If only one node is None or the values of the nodes differ, the trees are not flip equivalent.

1. If both nodes are None, then the trees are trivially flip equivalent.

- 3. If the nodes have the same value, we then recursively check their subtrees.
- We have two cases for recursion to check for flip equivalence:
- Case 1: Left subtree of root1 is flip equivalent to left subtree of root2 AND right subtree of root1 is flip equivalent to right subtree of root2.

returns false indicating the trees are not flip equivalent at this level.

then the trees rooted at root1 and root2 are flip equivalent at the current level.

Case 2: Left subtree of root1 is flip equivalent to right subtree of root2 AND right subtree of root1 is flip equivalent to left

- subtree of root2. If either of these cases is true, we conclude that the trees rooted at root1 and root2 are flip equivalent.
- This approach works well because it exploits the property that a tree is defined not just by its nodes and their values, but by the specific arrangement of these nodes. By checking all possible flip combinations of subtrees, we can effectively determine flip

equivalence between the two given trees.

Solution Approach The solution code implements a recursive function called dfs which stands for depth-first search. This function continuously dives deeper into the subtrees of both trees, simultaneously, for as long as it finds matching node values and structure that adhere to flip

equivalence. Below are the aspects of the implementation explained: Base Cases: The first checks in the dfs function handle the base cases. If both nodes are equal, this means that we have

subtrees equivalent.

 Recursive Checks: The dfs function then makes two critical recursive calls representing the two possible scenarios to check for equivalence:

reached equivalent leaves, or both nodes are None, which means the subtrees are empty and trivially flip equivalent. We thus

return true. If one of the nodes is None while the other isn't, or if the nodes' values are not equal, the function immediately

root1 is equivalent to the right subtree of root2. This corresponds to the situation where no flips are necessary at the current level of the trees. 2. The second scenario checks if the left subtree of root1 is equivalent to the right subtree of root2 and if the right subtree of

root1 is equivalent to the left subtree of root2. Here, it matches the case where a flip at the current node would render the

1. The first scenario checks if the left subtree of root1 is equivalent to the left subtree of root2 and if the right subtree of

• Efficient Short-Circuiting: The use of the and within each scenario creates short-circuiting behavior. This means if the first check within a scenario returns false, the second check is not performed, saving unnecessary computation.

Logical OR Operation: The or operation between the two checks is used to state that if either of these scenarios holds true,

call, allowing the algorithm to backtrace through the nodes of the trees and unwind the recursion with the correct answer. No other explicit data structures are used, which implies that the space complexity of the solution is primarily dependent on the

equivalence of the entire trees. This method effectively uses the call stack as a data structure to hold the state of each recursive

The dfs function is invoked with the root nodes of the two trees. The return value of this initial call will be the result for flip

height of the recursion tree, while the time complexity depends on the size of the binary trees being compared.

Let's take a small example of two binary trees to illustrate the solution approach: Let tree A have the following structure:

Application of the Solution Approach:

For the matched pair of nodes with value 2:

For the matched pair of nodes with value 3:

1 # Definition for a binary tree node.

self.value = value

self.right = right

return False

1. Without flipping children.

2. With flipping children.

self.left = left

class TreeNode:

class Solution:

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Example Walkthrough

We want to determine if tree B can be transformed into tree A via flip operations.

And let tree B have a structure that is a flipped version of tree A:

Step 1: We begin with the dfs function call on the root of tree A (which has the value 1), and the root of tree B (also value 1). Step 2: Since both root nodes' values are identical and not None, we check their subtrees, implementing the two scenarios described in the solution approach.

Step 4: We then check the left subtree of A (2) with the right subtree of B (2) and the right subtree of A (3) with the left subtree of B (3). Here, as we first match the values, we need to delve deeper recursively.

Recursive Step: For each matched pair (2 with 2, 3 with 3), we proceed recursively:

Both child pairs match in value; hence, recursion into them will return true.

Again, both child pairs match in value and further recursive checks would yield true.

 Compare left child of A's 2 (4) with right child of B's 2 (4). Compare right child of A's 2 (5) with left child of B's 2 (5).

Step 3: We check the left subtree of A (2) with the left subtree of B (3) and the right subtree of A (3) with the right subtree of B (2).

Since the values don't match, we proceed to the next step without making any further recursive calls in this scenario.

 Compare left child of A's 3 (6) with the right child of B's 3 (6). Compare right child of A's 3 (7) with the left child of B's 3 (7).

As all recursive checks return true, we can say that tree A is flip equivalent to tree B; hence, the initial call to dfs from the roots will

Python Solution

if not node1 or not node2 or node1.value != node2.value:

return (is_flip_equivalent(node1.left, node2.left) and \

is_flip_equivalent(node1.right, node2.right)) or \

(is_flip_equivalent(node1.left, node2.right) and \

is_flip_equivalent(node1.right, node2.left))

* Helper method to perform depth-first search to determine flip equivalence.

* @return true if the subtrees rooted at the given nodes are flip equivalent, false otherwise

// If one of the nodes is null, or the values are not equal, they are not flip equivalent

// Recursively check for flip equivalence for both children without flipping and with flipping

boolean checkWithoutFlip = isFlipEquiv(node1.left, node2.left) && isFlipEquiv(node1.right, node2.right);

boolean checkWithFlip = isFlipEquiv(node1.left, node2.right) && isFlipEquiv(node1.right, node2.left);

* @param nodel the current node being compared in the first tree

* @param node2 the current node being compared in the second tree

// Both nodes are null, they are flip equivalent (base case)

if (node1 == null || node2 == null || node1.val != node2.val) {

// Return true if either possibility resulted in flip equivalence

private boolean isFlipEquiv(TreeNode node1, TreeNode node2) {

// Check both possibilities: not flipped and flipped

bool isFlipEquivalent(TreeNode* root1, TreeNode* root2) {

if (root1 == root2) return true;

// If both nodes are null, they are flip equivalent.

// Check if children are flip equivalent in two ways:

if (!root1 || !root2 || root1->val != root2->val) return false;

// 1. Without flipping (left with left and right with right)

// 2. With flipping (left with right and right with left)

return checkWithoutFlip;

if (node1 == null && node2 == null) {

return true;

return false;

Recursively check if subtrees are flip equivalent:

return true, validating the flip equivalence of the two binary trees.

def __init__(self, value=0, left=None, right=None):

Helper function to perform depth-first search.

def is_flip_equivalent(node1, node2): # If both nodes are the same, or both are None, trees are flip equivalent. if not node1 and not node2: return True # If one of the nodes is None or values are not equal, trees not flip equivalent.

def flipEquiv(self, root1: Optional[TreeNode], root2: Optional[TreeNode]) -> bool:

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# Initiate the depth-first search from the root nodes of both trees.
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           return is_flip_equivalent(root1, root2)
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Java Solution
2 class TreeNode {
       int val;
       TreeNode left;
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1 // Definition for a binary tree node.
                           // Value of the node
                           // Reference to the left child
       TreeNode right;
                           // Reference to the right child
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       // Constructors
       TreeNode() {}
       TreeNode(int val) { this.val = val; }
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       TreeNode(int val, TreeNode left, TreeNode right) {
           this.val = val;
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           this.left = left;
           this.right = right;
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15 }
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   public class Solution {
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       /**
        * Determines if two binary trees are flip equivalent.
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        * Flip equivalent binary trees are trees that are
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        * the same when either flipped or not flipped at any level of their descendants.
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        * @param root1 the root of the first binary tree
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        * @param root2 the root of the second binary tree
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        * @return true if the binary trees are flip equivalent, false otherwise
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       public boolean flipEquiv(TreeNode root1, TreeNode root2) {
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            return isFlipEquiv(root1, root2);
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57 }
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C++ Solution
   // Definition for a binary tree node.
2 struct TreeNode {
       int val;
                          // The value of the node.
       TreeNode *left;
                          // Pointer to the left child.
       TreeNode *right;
                         // Pointer to the right child.
       TreeNode() : val(0), left(nullptr), right(nullptr) {}
       TreeNode(int x) : val(x), left(nullptr), right(nullptr) {}
       TreeNode(int x, TreeNode *left, TreeNode *right) : val(x), left(left), right(right) {}
9 };
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11 class Solution {
12 public:
       // Check whether two binary trees are flip equivalent.
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       bool flipEquiv(TreeNode* root1, TreeNode* root2) {
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           return isFlipEquivalent(root1, root2);
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       // Helper function to check recursively if two trees are flip equivalent.
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// If one of the nodes is null or the values don't match, they aren't flip equivalent.

return (isFlipEquivalent(root1->left, root2->left) && isFlipEquivalent(root1->right, root2->right)) ||

(isFlipEquivalent(root1->left, root2->right) && isFlipEquivalent(root1->right, root2->left));

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Typescript Solution
1 interface TreeNode {
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val: number;
                             // The value of the node.
     left: TreeNode | null; // Pointer to the left child.
     right: TreeNode | null; // Pointer to the right child.
   // Check whether two binary trees are flip equivalent.
   function flipEquiv(root1: TreeNode | null, root2: TreeNode | null): boolean {
     return isFlipEquivalent(root1, root2);
10 }
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   // Helper function to check recursively if two trees are flip equivalent.
   function isFlipEquivalent(root1: TreeNode | null, root2: TreeNode | null): boolean {
     // If both nodes are null, they are flip equivalent.
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     if (root1 === root2) return true;
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     // If one of the nodes is null or the values don't match, they aren't flip equivalent.
18
     if (!root1 || !root2 || root1.val !== root2.val) return false;
19
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     // Check if children are flip equivalent in two ways:
     // 1. Without flipping (left with left and right with right)
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     // 2. With flipping (left with right and right with left)
     return (isFlipEquivalent(root1.left, root2.left) && isFlipEquivalent(root1.right, root2.right)) ||
23
            (isFlipEquivalent(root1.left, root2.right) && isFlipEquivalent(root1.right, root2.left));
24
25 }
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Time and Space Complexity
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to two more recursive calls.

The time complexity is O(N) where N is the smaller number of nodes in either root1 or root2. This is because the recursion stops at the leaf nodes or when a mismatch is found. In the worst case, every node in the smaller tree will be visited. The space complexity is also O(N) due to the recursion stack. In the worst case, the recursion goes as deep as the height of the tree,

The code defines a recursive function dfs which checks if two binary trees are flip equivalent. The dfs function is called for each

corresponding pair of nodes in the two trees. At each step, the code performs constant time operations before potentially making up

which can be O(N) in the case of a skewed tree (a tree in which every internal node has only one child). For a balanced tree, the space complexity will be 0(log N) because the height of the tree would be logarithmic relative to the number of nodes.