Algorithms

Karl Solomon

December 31, 2024

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Part I

Data Structures

- 1 Queue/FIFO
- 2 Stack/LIFO
- 3 Heap/Priority Queue
- 4 Tree/Graph
- 4.1 Binary Tree
 - Check if two BST are same

• Insert Element

```
void insertNode(Node* root, int data) {
   if (root == nullptr) {
      root = new Node(data);
   } else {
      if (data < root->data) {
         insertNode(root->left, data);
      } else {
         insertNode(root->right, data);
      }
}
insertNode(root->right, data);
}
```

• Delete Element

```
Node* deleteBSTNode(Node* root, int data) {
      if (root == nullptr) return root;
      if (root->data == data) {
           // Only 1 node (root)
           if (root->left == nullptr && root->right == nullptr) {
               delete root;
6
               root = nullptr;
           // Only 1 child (right)
           else if (root->left == nullptr) {
11
               Node* temp = root->right;
12
               delete root;
               root = temp;
13
14
           // Only 1 child (left)
15
           else if (root->right == nullptr) {
               Node* temp = root->left;
               delete root;
18
               root = temp;
19
           // Both children exist. Take either min from right sub-
21
              tree (implemented here) or max from left sub-tree and
              set
```

```
// to deleted node's position
22
            else {
23
                Node* temp = root->right;
24
                while (temp->left != nullptr) {
25
                     temp = temp->left;
26
27
                root -> data = temp -> data;
                root -> right = deleteBSTNode(root -> right, temp -> data);
29
30
31
       return root;
32
33
```

• Lowest Common Ancestor

```
Node* lowestCommonAncestor(Node* root, int data1, int data2) {
      // Base case
2
      if (root == nullptr) {
3
          return nullptr;
5
6
      // If both data1 and data2 are smaller than root, then LCA \,
         lies in left subtree
      if (data1 < root->data && data2 < root->data) {
           return lowestCommonAncestor(root->left, data1, data2);
10
11
      // If both data1 and data2 are greater than root, then LCA
12
         lies in right subtree
      if (data1 > root->data && data2 > root->data) {
13
14
           return lowestCommonAncestor(root->right, data1, data2);
15
16
      // If one data is on the left and the other is on the right,
17
          then root is the LCA
      return root;
18
19
```

• Determine if tree is BST

```
|bool isBST(Node* root, int min, int max) {
      if (root == nullptr) return true;
      if (root->data < min || root->data > max) return false;
      return isBST(root->left, min, root->data - 1) && isBST(root->
         right, root->data + 1, max);
5
6
  bool isBST(Node* root) {
      bool res = true;
8
      if (root == nullptr) {
9
          return res;
11
      if (root->left == nullptr && root->right == nullptr) {
          return res;
13
14
      return isBST(root, INT_MIN, INT_MAX);
15
16
```

• Traversal

- Level-Order

```
- We use a 'std::queue < Node *> ' to keep track of nodes to
      - We start by pushing the root node into the queue.
     - While the queue is not empty, we:
       - Dequeue the front node, process it by adding its data
       to the result queue.
       - Enqueue its left and right children if they exist.
6
       This process continues until all nodes have been visited,
7
        resulting in a level-order traversal of the BST.
8
  std::queue<int> traverseBSTLevelOrder(Node* root) {
       std::queue < int > result;
       if (root == nullptr) {
12
           return result;
13
14
       std::queue < Node * > nodeQueue;
16
      nodeQueue.push(root);
17
18
       while (!nodeQueue.empty()) {
19
           Node* current = nodeQueue.front();
20
           nodeQueue.pop();
21
           result.push (current -> data);
22
23
           if (current->left != nullptr) {
24
               nodeQueue.push(current->left);
25
26
           if (current->right != nullptr) {
27
               nodeQueue.push(current->right);
28
           }
30
31
       return result;
```

- In-order: Left subtree, then node, then right subtree

```
std::vector<int> bstTraverseInOrder(Node* root) {
    std::vector<int> res;
    if (root == nullptr) return res;
    res = bstTraverseInOrder(root->left);
    res.push_back(root->data);
    std::vector<int> right = bstTraverseInOrder(root->right);
    res.insert(res.end(), right.begin(), right.end());
}
```

- PreOrder: Node, then left subtree, then right subtree
- PostOrder: Left subtree, then right subtree, then Node

4.2 Graphs

Storing Graphs:

• Adjacency Matrix: 2D array NxN. [n,m] = nonZeroNum means there is an edge from node n to node m. All nodes should not include edges to self. Edges can be weighted (to suggest things like cost/distance). In directional-graphs you can have [n,m] = nonZero, but [m,n] = 0. Cons: O(N2) memory. This is especially bad for sparse graphs.

• Adjacency List: Store edges as a list of neighbor nodes. Each node contains a list of neighbors. It is possible to represent weights here with a list of std::pair<Node*, int>.

Graph Algorithms:

• Bellman-Ford

```
#include <iostream>
  #include <limits>
  #include <vector>
3
   * - We define a 'Graph' class with a list of 'Edge' structures,
      each containing a source, destination, and weight.
       - The 'bellmanFord' function initializes distances and
      relaxes all edges |V| - 1 times, where |V| is the number of
   * vertices.
       - It checks for negative-weight cycles by attempting one more
       relaxation.
        - If a negative-weight cycle is detected, it prints a message
9
       - Otherwise, it prints the shortest distances from the start
      vertex to all other vertices.
13
  struct Edge {
      int source, destination, weight;
14
15
16
  class Graph {
17
   public:
18
      int vertices;
19
      std::vector < Edge > edges;
20
21
      Graph(int v) : vertices(v) {}
22
      void addEdge(int source, int destination, int weight) { edges.
          push_back({source, destination, weight}); }
       // The Bellman-Ford algorithm, is used to find the shortest
         paths from a single source vertex to all other vertices
      // in a graph. The algorithm can handle graphs with negative
26
          weight edges:
      void bellmanFord(int start) {
27
           std::vector<int> distance(vertices, std::numeric_limits
28
              int >:: max());
           distance[start] = 0;
29
30
           // Relax edges |V| - 1 times
31
           for (int i = 0; i < vertices - 1; ++i) {
32
               for (const auto& edge : edges) {
33
                   if (distance[edge.source] != std::numeric_limits <</pre>
34
                      int >:: max() &&
                       distance[edge.source] + edge.weight < distance</pre>
35
                           [edge.destination]) {
                       distance[edge.destination] = distance[edge.
36
                           source] + edge.weight;
                   }
               }
40
           // Check for negative-weight cycles
41
           for (const auto& edge : edges) {
42
               if (distance[edge.source] != std::numeric_limits<int</pre>
43
                  >::max() &&
```

```
distance[edge.source] + edge.weight < distance[</pre>
44
                           edge.destination]) {
                       std::cout << "Graph contains a negative-weight
  cycle" << std::endl;</pre>
45
                       return;
46
                  }
47
             }
49
             // Print the distances
50
             for (int i = 0; i < vertices; ++i) {
51
                  std::cout << "Distance from vertex " << start << " to</pre>
                      vertex " << i << " is " << distance[i] << std::</pre>
                      endl;
        }
54
55
  };
57
   int main() {
58
        Graph g(5);
59
        g.addEdge(0, 1, -1);
        g.addEdge(0, 2, 4);
60
        g.addEdge(1, 2, 3);
61
        g.addEdge(1, 3, 2);
62
        g.addEdge(1, 4, 2);
g.addEdge(3, 2, 5);
63
64
        g.addEdge(3, 1, 1);
g.addEdge(4, 3, -3);
65
66
67
68
        g.bellmanFord(0);
69
        return 0;
70
71
```

• Dijkstra's

```
#include <iostream>
  #include <limits>
2
  #include <queue>
3
  #include <vector>
  using namespace std;
  typedef pair<int, int> Edge; // (weight, vertex)
   * In this program:
   * - We define a 'Graph' class with an adjacency list to store
      edges as pairs of (weight, vertex).
      - The 'dijkstra' function initializes distances and uses a
13
      priority queue to explore the shortest paths.
       - It updates the shortest path estimates and pushes them into
      the priority queue.
       - Finally, it prints the shortest distances from the start
      vertex to all other vertices.
16
   * /
17
18
  class Graph {
19
   public:
20
      int vertices;
21
      vector < vector < Edge >> adjList;
22
      Graph(int v) : vertices(v), adjList(v) {}
24
```

```
25
       void addEdge(int source, int destination, int weight) {
26
           adjList[source].push_back(make_pair(weight, destination));
27
           adjList[destination].push_back(make_pair(weight, source));
28
                 // For undirected graph
29
       // Dijkstra's algorithm, is used to find the shortest paths
          from a single source vertex to all other vertices
       // in a graph with non-negative edge weights:
31
       void dijkstra(int start) {
32
           vector < int > distance(vertices, numeric_limits < int > :: max())
33
           priority_queue < Edge , vector < Edge > , greater < Edge >> pq;
34
35
           distance[start] = 0;
36
37
           pq.push(make_pair(0, start));
39
           while (!pq.empty()) {
40
               int currentVertex = pq.top().second;
41
               pq.pop();
42
                for (const auto& edge : adjList[currentVertex]) {
43
                    int weight = edge.first;
44
                    int neighbor = edge.second;
45
46
                    if (distance[currentVertex] + weight < distance[</pre>
47
                        neighbor]) {
                        distance[neighbor] = distance[currentVertex] +
                             weight;
                        pq.push(make_pair(distance[neighbor], neighbor
49
                            ));
                    }
50
                }
51
           }
           // Print the distances
54
           for (int i = 0; i < vertices; ++i) {
               cout << "Distance from vertex " << start << " to</pre>
                   vertex " << i << " is " << distance[i] << endl;</pre>
57
           }
       }
58
  };
59
60
  int main() {
61
       Graph g(5);
62
63
       g.addEdge(0, 1, 10);
       g.addEdge(0, 4, 5);
64
       g.addEdge(1, 2,
                        1);
65
       g.addEdge(1, 4, 2);
       g.addEdge(2, 3, 4);
67
       g.addEdge(3, 4, 9);
68
       g.addEdge(3, 0, 7);
69
70
       g.dijkstra(0);
71
72
       return 0;
73
```

- Ford-Fulkerson
- Kruskals's

• Nearest neighbor		
• Prim's		
• DFS		
L		
• BFS		

- 5 Lists
- 5.1 Singly Linked List
- 5.2 Doubly Linked List

Part II

Types of Algorithms

- 6 Greedy
- 7 DP
- 8 DFS
- 9 BFS
- 10 Sorting
- 11 Network Flow
- 11.1 Dijkstra's