```
1 package Assign_2;
 3 /** This class is an implementation of the main class
    for this Virus Tree program.
   * Micah Rose-Mighty
 5
   * 6498935
   * 2020-10-08
   * Created using IntelliJ
 8
   */
 9
10 import java.text.MessageFormat;
11
12 public class Main {
13
14
       public static void main(String[] args) {
15
16
           System.out.println("Tree Created: ");
17
           VirusTree vt = new VirusTree("
   tree_of_virus_input.txt");
18
           System.out.println();
19
20
           System.out.println("Depth of Tree: ");
21
           System.out.print(vt.depth());
22
           System.out.println();
23
           System.out.println();
24
           System.out.println("Breadth-First Traversal
25
    ");
26
           vt.bfs();
27
           System.out.println();
28
29
           System.out.println("Pre-order Traversal: ");
30
           vt.preorder();
31
           System.out.println();
32
33
           System.out.println("Post-order Traversal: ");
           vt.postorder();
34
35
           System.out.println();
36
37
           String v1 = "HCoV-OC43";
           String v2 = "Hcov-229E";
38
39
           System.out.println(MessageFormat.format("
   Distance between \{0\} and \{1\}: ", \vee1, \vee2));
40
           vt.distance(v1, v2);
```

```
\label{lem:cosc2} File - C:\Users\mbox{\sc 2P03\Assign}_2\sc \Assign} \label{lem:cosc2P03\Assign} Assign_2\sc \Assign} \label{lem:cosc2P03\Assign} In the context of the cost of the cos
    41
                                                                                                   System.out.println();
    42
                                                                                                   String v3 = "SARS-CoV";
    43
                                                                                                   String v4 = "Zika virus";
    44
                                                                                                   System.out.println(MessageFormat.format("
    45
                             Distance between \{0\} and \{1\}: ", \vee 3, \vee 4);
                                                                                                   vt.distance(v3, v4);
    46
                                                                                                   System.out.println();
    47
    48
                                                                }
    49
    50 }
    51
```

```
1 package Assign_2;
 3 /** This class is an implementation of the VirusTree
   Structure along with all the required methods.
   * Micah Rose-Mighty
 5
   * 6498935
   * 2020-10-08
   * Created using IntelliJ
 8
   */
 9
10 import java.io.File;
11 import java.io.IOException;
12 import java.text.MessageFormat;
13 import java.util.Arrays;
14 import java.util.LinkedList;
15 import java.util.Scanner;
16
17 public class VirusTree {
18
19
       private VirusTreeNode root; // tree root node
20
21
       public VirusTree() { root = null; } // empty tree
    constructor
22
23
       public VirusTree(String filename){ // Constructor
    that creates tree structure from given input file
24
           this();
25
           try(Scanner scanner = new Scanner(new File(
   filename))) {
26
               while(scanner.hasNextLine()){
27
                   String[] parts = scanner.nextLine().
   trim().split(",");
28
29
                   for(int i = parts.length-1; i>=1; i
   --){
                        if(!insert(parts[0], parts[i])){
30
31
                           throw new
   IllegalArgumentException("Can not find parent node: "
    + parts[0]);
                       }
32
                   }
33
34
                   System.out.println(MessageFormat.
   format("{0}: {1}", parts[0], String.join(" -> ",
   Arrays.copyOfRange(parts, 1, parts.length))));
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```

```
File - C:\Users\micah\Desktop\COSC 2P03\Assign_2\src\Assign_2\VirusTree.java
35
36
            }
37
            catch (IOException e) {
                throw new RuntimeException(e);
38
            }
39
        }
40
41
42
        public int depth() { // Method for getting the
   depth of the given tree using inner recursive method
   for calculation.
43
            if(root == null){
44
                return 0;
45
46
            return depthStep(root);
        }
47
48
49
        public void bfs(){ // Method for outputting the
   given tree nodes in breadth-first traversal order.
50
            if (root == null) {
51
                return;
            }
52
53
54
            LinkedList<VirusTreeNode> queue = new
   LinkedList<>();
55
            queue.addLast(root);
56
57
            while(!queue.isEmpty()){
58
                VirusTreeNode current = queue.removeFirst
   ();
59
                System.out.println(current.info);
60
61
                VirusTreeNode currChild = current.
   firstChild;
                while(currChild != null){
62
63
                     queue.addLast(currChild);
64
                     currChild = currChild.nextSibling;
                }
65
            }
66
        }
67
68
        public void preorder(){ // Method for outputting
69
   the given tree in Pre-order Traversal Order by
   calling traverse method and setting it to true.
70
            traverse(true);
```

```
File - C:\Users\micah\Desktop\COSC 2P03\Assign_2\src\Assign_2\VirusTree.java
 71
 72
 73
         public void postorder(){ // Method for
    outputting the given tree in Post-order Traversal
    Order by calling traverse method and setting it to
    false.
 74
             traverse(false);
 75
         }
 76
 77
        public void distance(String info1, String info2
    ){ //Method for outputting the distance between two
    nodes in the given tree.
             if(root == null) {
 78
 79
                 throw new IllegalArgumentException("Can'
    t find node: " + info1);
             }
 80
 81
 82
             LinkedList<String> path1 = getPath(root,
    info1, new LinkedList<>());
             if(path1 == null){
 83
                 throw new IllegalArgumentException("Can'
 84
    t find node: " + info1);
 85
             }
 86
 87
             LinkedList<String> path2 = getPath(root,
    info2, new LinkedList<>());
             if(path2 == null){
 88
                 throw new IllegalArgumentException("Can'
 89
    t find node: " + info2);
 90
             }
 91
 92
             for (int i = path1.size() - 1; i>=0; i--) {
    // For loop to find last common ancestor starting
    from last node in first path and if found in second
    path then that is common ancestor.
 93
                 String pathInfo = path1.get(i);
 94
                 int index = path2.index0f(pathInfo);
 95
                 if(index >= 0) {
                     int distance = Math.max(path1.size
 96
    () - 1 - i, path2.size() - 1 - index);
                     System.out.println(MessageFormat.
 97
    format("The distance between {0} and {1} is {2}.
    They have common ancestor {3}.", info1, info2,
    distance, pathInfo));
```

```
98
                    return;
 99
                }
100
            }
101
            throw new IllegalArgumentException(
    MessageFormat.format("There does not exist a common
    ancestor node for {0} and {1}", info1, info2));
102
103
104
        private LinkedList<String> getPath(VirusTreeNode
     currentNode, String info, LinkedList<String>
    currentList) { // Private recursive method for
    searching a path to a given node.
105
            String currentInfo = currentNode.info;
            currentList.add(currentInfo);
106
            if(currentInfo.equals(info)) {
107
                return currentList;
108
            }
109
110
111
            VirusTreeNode currChild = currentNode.
    firstChild;
112
            while (currChild != null){
113
                LinkedList<String> result = getPath(
    currChild, info, currentList);
114
                if(result != null) {
115
                    return result;
116
117
                currChild = currChild.nextSibling;
118
119
            currentList.removeLast();
120
            return null;
        }
121
122
        private void traverse(boolean preorder) { //
123
    Inner traverse method the outputs Pre-order
    Traversal if true and Post-order Traversal if false.
            if (root == null) {
124
125
                return;
126
            traverseStep(root, preorder);
127
        }
128
129
130
        private void traverseStep(VirusTreeNode
    currentNode, boolean preorder) { // Inner recursive
    traverse order method that is based on the given
```

```
130 node.
131
132
            if (preorder) {
                System.out.println(currentNode.info);
133
            }
134
135
136
            VirusTreeNode currChild = currentNode.
    firstChild;
137
            while(currChild != null) {
138
                traverseStep(currChild, preorder);
139
                currChild = currChild.nextSibling;
140
            }
141
142
            if (!preorder) {
143
                System.out.println(currentNode.info);
            }
144
        }
145
146
147
        private int depthStep(VirusTreeNode currentNode
    ) { // Inner recursive method for calculating depth.
148
            int max = 0;
149
150
            VirusTreeNode currChild = currentNode.
    firstChild;
151
            while (currChild != null) {
152
                max = Math.max(max, 1 + depthStep(
    currChild));
153
                currChild = currChild.nextSibling;
154
155
            return max;
156
        }
157
158
        private boolean insert(String parentInfo, String
     childInfo) { // Private method for adding parent-
    child pair into the given tree. Returns true if pair
     is added successfully and false otherwise.
159
            if (root == null) {
160
                root = new VirusTreeNode(parentInfo);
161
162
                root.firstChild = new VirusTreeNode(
    childInfo);
163
                root.nextSibling = null;
                root.firstChild.firstChild = null;
164
                root.firstChild.nextSibling = null;
165
```

191

192 } 193 }

```
1 package Assign_2;
3 /** This class is an implementation of the VirusTree
  Node.
4 * Micah Rose-Mighty
 5 * 6498935
 6 * 2020-10-08
 7 * Created using IntelliJ
   */
 9
10
11 public class VirusTreeNode {
12
       String info;
13
14
      VirusTreeNode firstChild;
15
16
      VirusTreeNode nextSibling;
17
       VirusTreeNode(String info){ this.info = info; }
18
  //Node data constructor
19 }
20
```

- 1 Virus, DNA Virus, RNA Virus
- 2 RNA Virus, Flaviviridae, Filoviridae, Coronaviridae
- 3 Flaviviridae, Flavivirus, Hepacivirus
- 4 Filoviridae, Cuevavirus, Dianlovirus, Ebolavirus, Marburgvirus
- 5 Coronaviridae, Alphacoronavirus, Betacoronavirus, Deltacoronavirus, Gammacoronavirus, Alphaletovirus
- 6 Flavivirus, Zika virus, Yellow fever virus, West Nile virus
- 7 Hepacivirus, GBV-B, hepatitis C virus
- 8 Ebolavirus, Ebola virus, Tai Forest virus, Bombali virus
- 9 Marburgvirus, Marburg virus
- 10 Alphacoronavirus, Hcov-229E, Hcov-NL63
- 11 Betacoronavirus, HCoV-OC43, HCoV-HKU1
- 12 Betacoronavirus, SARS-CoV
- 13 Betacoronavirus, MERS-CoV
- 14 DNA Virus, Adenoviridae, Poxviridae
- 15 Adenoviridae, Mastadenovirus
- 16 Poxviridae, Orthopoxvirus
- 17 Mastadenovirus, Canine hepatitis virus
- 18 Orthopoxvirus, Cowpox virus, Smallpox virus