```
1 package Drugs;
 2 import com.sun.xml.internal.bind.v2.model.core.ID;
 3
 4 import java.io.File;
 5 import java.io.FileNotFoundException;
 6 import java.io.FileWriter;
 7 import java.util.Scanner;
 8 import java.io.IOException;
 9
10 /** this is the main function Drug bank where all
11
    * the operations are being performed upon the drug
12
    *
13
   */
14
15 public class drugBank {
16
       FileWriter writeFile;
17
       Drug[] data;
18
       BinaryNode root;
19
20
       public drugBank() {
21
           ReadData();
22
       }
23
24
       /** this class reads the data and stores it into
25
        * a file and keeps the storage as an array of
   drugs
26
27
        */
28
       public void ReadData() {
29
30
           try {
               File file = new File("dockedApproved.tab"
31
   );
32
               Scanner scanner = new Scanner(file);
               String text = scanner.nextLine();
33
34
               int counter = 0;
               while (scanner.hasNextLine()) {
35
                   text = scanner.nextLine();
36
37
                   counter++;
38
               }
39
               scanner.close();
```

```
40
                data = new Drug[counter];
41
                file = new File("dockedApproved.tab");
42
                scanner = new Scanner(file);
43
                String[] store_array;
44
                text = scanner.nextLine();
45
46
                for (int i = 0; i < data.length; i++) {
47
                    store_array = scanner.nextLine().
   split("\\t");
48
                    data[i]=new Drug(store_array[0].trim
   (), store_array[1].trim(), store_array[2].trim(),
   store_array[3].trim(),store_array[4].trim(),Double.
   parseDouble(store_array[5].trim()));
49
50
           } catch (FileNotFoundException e) {
                System.out.println("No file found!");
51
52
                e.printStackTrace();
53
           }
       }
54
55
       /** the create class just creates the drug item
56
57
        * into the tree for the bank
58
        *
59
        */
60
61
       public void create() {
           for (int i = 0; i < data.length; i++) {</pre>
62
                root = insert(root, data[i]);
63
64
           }
65
       }
66
67
       /** the insert class tries to insert the nodes
        * that are being entered or read from the file
68
69
        * to their desired positions
70
        *
71
        * @param rt
72
        * @param drug
73
        * <u>@return</u>
74
        */
75
76
       public BinaryNode insert(BinaryNode rt, Drug drug
```

```
76 ) {
 77
            if (rt == null) {
                rt = new BinaryNode(null, drug,null);
 78
 79
            } else {
                if (drug.drugBankId.compareToIgnoreCase(
 80
    rt.item.drugBankId) < 0) {</pre>
                     rt.left = insert(rt.left, drug);
 81
 82
                 }else {
 83
                     rt.right = insert(rt.right, drug);
 84
                 }
 85
 86
            return rt;
 87
        }
 88
        /** the inorder traverse basically
 89
         * makes use of this traversal to print all
 90
 91
         * nodes in ascending order of drugs
 92
 93
         */
 94
 95
        public void inOrderTraverse() {
 96
            try {
 97
                 writeFile = new FileWriter("
    dockedApprovedSorted.tab");
                traverseinorder(root, writeFile);
 98
 99
                 writeFile.flush();
100
                writeFile.close();
            } catch (IOException e) {
101
                System.out.println("error occurred");
102
103
                 e.printStackTrace();
            }
104
105
        }
106
107
108
        public void traverseinorder(BinaryNode b,
    FileWriter tem) {
            if (b == null) {
109
110
                 return;
111
            } else {
112
                 traverseinorder(b.left, tem);
113
                 try {
```

```
tem.write(b.item.drugBankId + "" + b
114
    .item.GenericName + "" + b.item.drugGroups + "" + b.
    item.score + "\n");
                 } catch (IOException e) {
115
116
                     System.out.println("error occurred
    again");
117
                     e.printStackTrace();
                 }
118
119
                 traverseinorder(b.right, tem);
120
             }
        }
121
122
123
        /** the search function just searches the
    desired
124
         * drug from the tree.
125
126
         * <a href="mailto:aparam">aparam</a> Db_id
127
         * @return
128
         */
129
130
        public BinaryNode search(String Db_id) {
131
             BinaryNode tem = root;
132
             while(tem.item.drugBankId.
    compareToIgnoreCase(Db_id) !=0){
133
                 if (tem.item.drugBankId.
    compareToIgnoreCase(Db_id) > 0) {
134
                     tem = tem.left;
135
                 } else {
136
                     tem = tem.right;
137
                 }
                 if (tem == null) {
138
                     System.out.println("No such drug
139
    found");
140
                     return null;
141
                 }
142
             }
143
144
145
             return tem;
             }
146
147
```

```
148
149
        /** the delete fucntion deletes the drug
150
         * from the file
151
         *
152
         * @param rt
153
         * @param id
154
         * @return
155
         */
156
157
            public BinaryNode delete(BinaryNode rt,
    String id) {
158
            if (rt ==
                        null) {
159
                 return rt;
160
            }
            if(id.compareToIgnoreCase(rt.item.drugBankId
161
    ) <0) {
                 rt.left = delete(rt.left, id);
162
163
            }
            else if(id.compareToIgnoreCase(rt.item.
164
    drugBankId) >0) {
165
                 rt.right = delete(rt.right, id);
166
            }
167
           else if (rt.left != null && rt.right != null
168
    ) {
169
               rt.item = find_min(rt.right).item;
               rt.right = delete(rt.right, rt.item.
170
    drugBankId);
171
172
173
174
        }
175
           else {
               rt = (rt.left != null) ? rt.left : rt.
176
    right;
177
            }
178
            return rt;
179
180
        }
181
        public BinaryNode find_min(BinaryNode rt) {
182
```

```
183
                 if (rt!=null){
184
                     while(rt.left!=null) {
185
                         rt =rt.left;
                     }
186
187
                 }
188
                 return rt;
189
        }
190
        /** the depth function calculates the depth of
191
    the
         * tree for the drugs
192
193
         *
194
         * @param DbId
195
         * @return
196
         */
197
198
        public int Depth(String DbId) {
199
            int counter = 0;
200
            BinaryNode tem = root;
            while (tem.item.drugBankId.
201
    compareToIqnoreCase(DbId) > 0){
202
                 if (tem.item.drugBankId.compareTo(DbId
    ) != 0) {
203
                     tem = tem.left;
204
                     counter = counter + 1;
205
                 } else {
206
                     tem = tem.right;
207
                     counter = counter + 1;
208
209
                 }
            }
210
211
212
213
            return counter;
214
        }
215
        public int Depth1(BinaryNode r) {
216
217
            int x;
218
            int y;
            if (r == null) {
219
220
                 return 0;
```

```
221
222
            else{
223
                x = Depth1(r.left);
                y = Depth1(r.right);
224
225
226
                if (x > y) {
227
                     return x+1;
228
                }
                else{
229
230
                     return y+1;
                }
231
232
            }
233
        }
234
235
        /** the main function is to basically tests all
         * the functions with the drug ids to see
236
    whether
237
         * the functions are properly working or not.
238
         *
239
         * @param args
240
         */
241
242
        public static void main(String[] args) {
243
            drugBank db = new drugBank();
244
245
            db.create();
246
            db.inOrderTraverse();
247
            try{
                FileWriter f = new FileWriter("Output.
248
    txt");
                int calcDepth = db.Depth("DB01050");
249
                System.out.println(("the depth is: " +
250
    calcDepth));
                f.write("the depth of the node is: " +
251
    calcDepth+"\n");
252
253
                int deepestNode = db.Depth1(db.root);
254
                System.out.println((" the depth is :" +
255
    deepestNode));
256
                f.write("the depth is: " + deepestNode+
```

```
256 "\n");
257
                 BinaryNode search = db.search("DB00316"
    );
                if (search != null) {
258
259
                     System.out.println(("the drug has
    been found"));
260
                     f.write("drug has been found");
                 }
261
                else{
262
263
                     System.out.println(("the drug was
    not found"));
264
                     f.write("the drug was not found in
                "\n");
    the file" +
265
                 }
266
                BinaryNode delete = db.delete(db.root, "
267
    DB01065");
                 if (delete != null) {
268
                     System.out.println(("the drug has
269
    been deleted"));
                     f.write("the drug has been deleted"
270
    );
271
272
                 }
273
                 else{
                     System.out.println(("the drug to be
274
    deleted does not exist"));
                     f.write("the drug was not found");
275
276
277
                 }
                 f.flush();
278
                f.close();
279
280
281
282
            catch (IOException e) {
                System.out.println("error");
283
284
            }
285
        }
286 }
287
288
```