COMP 401 - Project in Biology and Computer Science

# Final Report

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Terms to know: unswitched case: case where training dataset has 185 instances and testing dataset has 908 instances (original way datasets were formed); switched case: case where training dataset has 908 instances and testing dataset has 185 instances; ranked case: case where training dataset has 908 instances with attributes ranked and testing dataset has 185 instances

#### 1. Introduction

Climate change is becoming an ever-growing problem. The increase in global temperatures will lead to more drought events and some areas of the world are expected to experience reductions in their crop yields. For these areas, plant breeding programs and agricultural technology must be developed (Vello et al., 2015).

The use of different phenomics technologies in plants is a key element to improve our knowledge of the genotype-phenotype association of desired agricultural traits such as the response to water deficit (Vello et al., 2015). In the McGill Plant Phenomics Platform (MP3), research is being conducted in the hopes of better understanding plant response to water deficit, among others. The MP3 is instrumental in conducting this research. The MP3 has two main components, the 3D system and the Hts systems. The 3D system has the following cameras: 2 visible light, 2 neo-infrared, and 2 infrared, consisting of top and side views for each type of camera. This allows the phenomics lab to capture most of the characteristics of the plant. The system has a lifter with two different positions, high and low, and it can take images at single degree differences, ie 0 degrees, 1 degree, 2 degrees, etc. A conveyor belt transports the plants and the cabinets in which the plants are held have different positions that can rotate. There is a weighing system which allows for the tracking of automatic watering of the plant. The system can be programmed to keep the plant at a certain watering level or absolute weight value, eg. the plant will always receive 100 mL of water or the plant will receive water until it weighs 300 g (including the weight of the carrier). MP3 is located in the McGill greenhouse in the Stewart Biology building.

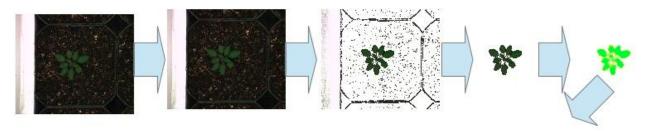
In this research, soybeans were studied because they are economically important. As the meat-packing industry is known to have very negative effects on the environment, soybeans have become even more important for the environmental sustainability of the future. Because soybean products are ingredients in many meat and dairy substitutes and soybeans are an exceptional source of essential nutrients, the shift from meat to soybeans may be essential (Arnarson et al., 2015). Learning more about soybeans would better allow us to effectively use it as a meat and dairy substitute and alleviate some negative environmental effects.

Machine learning and image analysis are important in this field so that we can better predict the effect that climate change will have on crop production and better prepare for it (Vello et al., 2015, Stavness et al., 2017). With the ability to instantly analyze different plants grown under different watering conditions, we will be able to get a better sense of what needs to be done in order to still meet the global demand for plant agriculture.

# 2. Background

# 2.1. Image Processing

All of the necessary image analysis was performed by Emilio Vello and the processed image data was stored in the MP3 soybean database. Figure 1 shows an example of the image analysis pipeline that was performed prior to this research.



**Figure 1 –** Image analysis pipeline performed prior to research and stored in arabidopsis database; From "Image Analysis Pipeline", by Emilio Vello, 2017, <a href="http://mp3.biol.mcgill.ca/mcgill\_mp3\_analysis.html">http://mp3.biol.mcgill.ca/mcgill\_mp3\_analysis.html</a>. Copyright 2017 by Emilio Vello. Reprinted with permission.

The final image in Figure 1 shows the image from which the data is taken and stored in the phenomics lab database.

# 2.2. Weka

Machine learning has an ever-growing important role in many fields, including imagebased plant phenotyping. Weka is a collection of state-of-the-art machine learning algorithms (Frank and Witten, 1999). It was developed at the University of Waikato in New Zealand and the name stands for Waikato Environment for Knowledge Analysis (Frank and Witten, 1999). The system is written in Java and runs on almost any platform. Weka provides implementations of learning algorithms that one can easily apply to a dataset. All of the algorithms take their input in ARFF format which can be read from a file or generated by a database query (Frank and Witten, 1999). For this research, the database queries were saved as ARFF files and then read in by the program. One way of using Weka is to use learned models to generate predictions on new instances, which is the way that Weka was used for this research (Frank and Witten, 1999). Besides just training and building models based on a given training dataset, Weka also has several methods for "Attribute Selection". These different "Attribute Selection" methods read the training dataset and determine which attributes in the dataset are most important in predicting the developmental stage of the plant. One of these methods ranks the attributes by level of importance and these rankings can then be used while training and building the classification models. Weka is available through an easy-to-use graphical user interface or directly in code through the Java Weka library.

#### 3. Methods

#### 3.1. Software

During this research, a 'Weka Application' Java project was developed consisting of one class and several methods, that will be used by Dr. Thomas Bureau and Mr. Emilio Vello in their phenomics lab. For simplicity of building an initial program that Mr. Vello can later build upon, the developmental stage of the plant was based on its DAS (days after seeding). The different das were split into 5 categories, each containing 10 values and the developmental stages were then assigned to one of these 5 categories (Stage 1, Stage 2, Stage 3, Stage 4, or Stage 5). For example, a plant who was 34 days after seeding was considered to be in Stage 4.

This class provides an easy way to query the image data from the phenomics lab's soybean database, write the data to an ARFF file that is understandable to Weka methods, build and train classification models provided by the Java Weka library, and test these models on a testing dataset, which also comes from the soybean database. The jdbc drivers were used to be

able to write SQL queries within the class and the Java Weka library was used to directly import Weka methods. The Weka library provided all of the machine learning classification algorithms that were trained and tested within the class. The program simply reads the training data, builds a classification model based on this data and the name of the model inputted by the user (or builds the program's default model options). The program then runs on the testing dataset and calculates a precision of the model. All of the predictions of the different classification algorithms tested were outputted to a CSV file which also contained the barcode, das, and true stage of the plant. Another CSV is produced that stores the name of each algorithm tested and their respective precision value. The precision value was simply calculated as the number of instances that the model classifies correctly divided by the total number of instances within the dataset. These precision values matched those reported by Weka when performing the training and testing of the models directly within the Weka GUI. The CSV containing the stage prediction of each plant was then analyzed in R.

Both the Java code and R analysis code can be viewed at the end of this report.

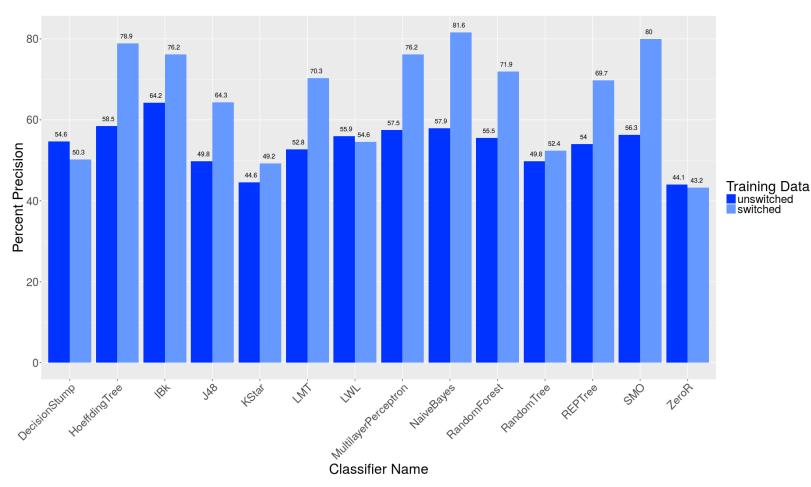
#### 3.2. Datasets and Tests

The training and testing sets were formed by querying the phenomics lab's soybean database. The public schema was used as were the dasplusev, imageev, imgobjectev, and soyidentification tables. These tables allowed the joining of the identification of the plant along with all of the important image characteristics that would be used to determine the developmental stage of the plant. The original training data only looked at the vis-side-1-0 camera view, line 1, and set 3. The original testing data also only looked at the vis-side-1-0 camera, lines 1, 2, and 3 from set 2 and lines 1 and 2 from set 3. It is important to only choose one camera for training and testing otherwise this will lead to non-comparable image views. The area of the plant from an aerial view will be calculated differently than the area of the plant from a side view. Therefore, it is important to be consistent with the type of camera when querying the data. The data selected additionally only included "well-watered" plants as opposed to also including "drought" plants. This is because the developmental stage of plants in these different conditions vary greatly. The term 'original' training data refers to the fact that this data is what was initially used to build the program (this will later on in the paper be referred to as the 'unswitched case', as described in 'terms to know' above). The testing data was later on used as the training data and vice versa to see if there would be any improvement in precision (this will later on in the paper be referred to as the 'switched case', as described in 'terms to know' above).

# 3.3. Approach

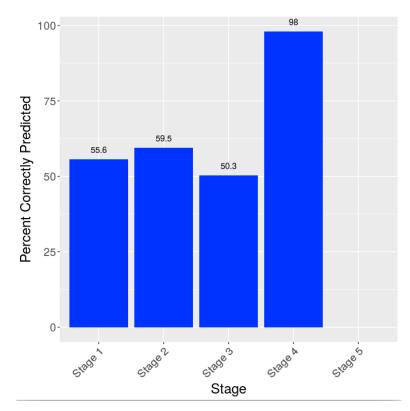
The user was left with the option of inputting which classification algorithms they want to try. This was possible due to the "AbstractClassifier" class within the Weka library. The "AbstractClassifier" class would instantiate any classifier object simply based on the name of the classifier. This prevented the necessity of importing every classifier available from Weka regardless of user input. An array list of strings containing all of the possible classification algorithms was created in order to ensure that the user input was valid. This array list is also the 'default' set of classification algorithms that the program will run if the user decides not to input any algorithms on their own. Because the datasets are fairly small, it only took a few seconds for all of the models to be built and tested. The user was also allowed to decide whether they wanted to rank the attributes in the datasets to test if there would be any improvement in precision from doing so.

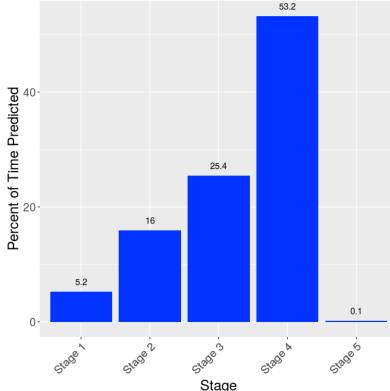
#### 4. Results



**Figure 2** - All default algorithms tested with their precisions graphed side by side with the unswitched (dark blue) and switched (light blue) case

The precision values of all of the default algorithms are visible in Figure 2. The 'unswitched' bars in the figure refer to the 'original' organization of the training and testing data as described in 'terms to know' and Methods above. The 'switched' bars therefore refer to switching which dataset was used for training versus which dataset was used for testing. The following algorithms had better precision with the 'unswitched' datasets: ZeroR, LWL, and Decision Stump. The following algorithms had better precision with the 'switched' datasets: NaiveBayes, MultilayerPerceptron, SMO, IBk, KStar, HoeffdingTree, J48, LMT, RandomForest, RandomTree, and REPTree. Therefore, 11 algorithms performed better when the larger dataset was used for training and only 3 algorithms performed better when the smaller dataset was used for training. Overall, the best classification algorithm was NaiveBayes, in the 'switched' case, with 81.6% precision.



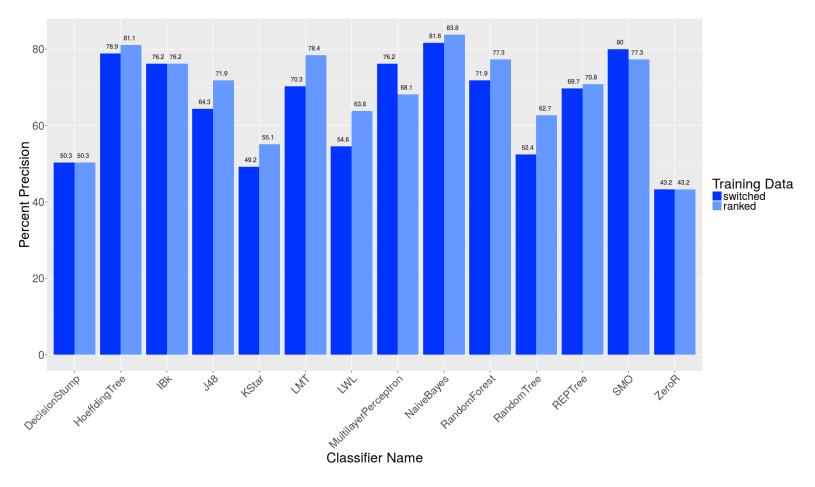


**Figure 3A** – Number of times each stage was correctly predicted as a percentage of the total number of times each stage was predicted; Sum of number of times each classifier correctly predicted the stage was calculated and divided by total number of times each stage was predicted at all; Average over all default algorithms except for ZeroR

**Figure 3B** – Frequency that each stage was predicted; Number of times each stage was predicted by each classifier divided by total number of prediction made; Average over all default algorithms except for ZeroR

If we look at just the 'switched' case, ie with the larger dataset used for training and the smaller used for testing, Figure 3A shows the number of times that each stage was correctly predicted, as a percentage of the total number of times each stage was predicted. The sum of the number of times each classifier correctly predicted the stage was calculated and then divided by the total number of times each stage was predicted at all. Note that in these calculations, the ZeroR algorithm was ignored as this algorithm simply predicts the most common stage in the training dataset. For example, in the original training set, Stage 4 was the most common stage so the ZeroR algorithm predicted Stage 4 for each instance in the testing set, therefore including this algorithm would skew the data too much. The frequency that each stage was predicted for this case was also calculated. This frequency was calculated by counting the number of times each stage was predicted by all of the algorithms and then dividing by the total number of predictions made by all of the algorithms, except ZeroR. The frequencies resulted in the following breakdown: Stage 1-5.2%, Stage 2-16%, Stage 3-25.4%, Stage 4-53.2%, Stage 5-0.1%, which is shown in Figure 3B.

When looking at the 'unswitched' case, the stage precision breakdown we saw was the following: Stage 1-63.5%, Stage 2-73%, Stage 3-29%, Stage 4-79.5%, Stage 5-0%. The frequency that each stage was predicted for this case had the following breakdown: Stage 1-7.2%, Stage 2-32.2%, Stage 3-17.1%, Stage 4-43.5%, Stage 5-0%.



**Figure 4** - All defaut algorithms tested with their precisions graphed side by side with the ranked (dark blue) and switched (light blue) case

Figure 4 shows the different precision values for the 'switched' case and the 'ranked' case. The 'ranked' case took the 'switched' data and first ranked the attributes before training and building the classification models. The same default set of models were built and tested and the precision values were compared to those achieved when the ranking algorithm was not run. The algorithms that performed better when the attributes were ranked first were: NaiveBayes, KStar, LWL, HoeffdingTree, J48, LMT, RandomForest, RandomTree, and REPTree. The algorithms that performed better when the attributes were not ranked first were: MultilayerPerceptron and SMO. The algorithms that showed no difference in precision when the attributes were ranked or unranked were: ZeroR, IBk, DecisionStump. Therefore, 9 algorithms performed better when the attributes were ranked, 2 performed worse, and 3 performed the same. The best classification algorithm after ranking the attributes was the NaiveBayes algorithm with a precision of 83.8%, a 2.2% increase in precision from NaiveBayes in just the 'switched' case with no attribute ranking.

When looking at the 'ranked' case, the stage precision breakdown we saw was the following: Stage 1 - 79.6%, Stage 2 - 81.5%, Stage 3 - 42.1%, Stage 4 - 99.5%, Stage 5 - 0%. The frequency that each stage was predicted for this case had the following breakdown: Stage 1 - 6.6%, Stage 2 - 19.8%, Stage 3 - 18.3%, Stage 4 - 55.3%, Stage 5 - 0%.

#### 5. Discussion

From Figure 3A (the 'switched' case), we see that Stage 4 is predicted drastically more precisely than any other stage. From Figure 3B, we also see that Stage 4 was predicted the most often. Looking at a breakdown of the number of instances belonging to each stage in the dataset we have: (Stage 1: 32, Stage 2: 163, Stage 3: 280, Stage 4: 400, Stage 5: 33). The combination of Stage 4 being the most prevalent stage in the set used for training and the fact that it was predicted the most often, gives a possible reason that this stage was predicted most accurately. Additionally, Stage 4 plants are larger since they are more days after seeding. These plants may be more developed and more defined than smaller plants which have less information to be gained from them. This gives another possible reason that Stage 4 plants were predicted most precisely. Stage 5 was never predicted correctly because in this case the testing set had zero plants in Stage 5.

In the 'unswitched' case, we see again that Stage 4 was correctly predicted the most whereas Stage 3 was correctly predicted the least often (excluding Stage 5). We also see that Stage 4 was predicted the highest number of times but we also see that Stage 3 was not predicted the fewest number of times, Stage 1 was (again excluding Stage 5). Looking at a breakdown of the number of instances belonging to each stage in the dataset we have: (Stage 1: 9, Stage 2: 36, Stage 3: 60, Stage 4: 80, Stage 5: 0). We can see from this that there were no instances in the dataset that belonged to Stage 5 and this is why Stage 5 was never predicted nor correctly predicted, the classification algorithms had nothing to learn from. Similar to the 'switched' case, Stage 4 was most likely predicted most accurately due to the fact that there is more information to be gained from plants in this stage than plants in earlier stages. However, there is not yet a good explanation why Stage 3 was correctly predicted the least often instead of Stage 1.

From Figure 4 (the 'ranked' case) we see that ranking the attributes before building and training the models improved (or at least did not affect) the precision of most algorithms. For some algorithms, it is obvious why there were no improvements in precision when the attributes were ranked. The ZeroR algorithm, for example, always predicts the value that is most common in the training dataset. Therefore, it does not matter how well the data is prepared before building the model, the ZeroR algorithm will always predict the same value. Most of the algorithms experienced an improvement in precision as a result of ranking the attributes. This makes sense as the ranking tells the algorithms which attributes are most important to analyze to determine the developmental stage of the plant and therefore these attributes receive more weight. It seems reasonable that in the worst-case, this ranking would just be extra computational work that added no benefit to precision and the precision would simply be the same as if we did not rank first, which is what we saw for three of the algorithms tested. We once again saw that Stage 4 was the best predicted stage when the attributes were ranked and we can again guess that this is due to the plants in Stage 4 being larger and better defined than the plants in earlier stages. What is interesting, however, is that we see a 24% increase in precision in predicting Stage 1, a 22% increase in precision in predicting Stage 2, and an 8.2% decrease in predicting Stage 3. As just stated, the increase in precision for certain stages is most likely due to the fact that the more important attributes have more weight being assigned to them. However, there is not a good explanation yet for why Stage 3 would have experienced a decrease in precision compared to when the attributes were unranked.

#### 6. Conclusion

In this research, a program was developed as a way of automating the querying of any database from the McGill phenomics lab, reading this data, using the training data to build machine learning classification models that would predict the developmental stage of a plant based on its image data, and testing these models on a new test set of plant image data. Based on the program, the NaiveBayes algorithm is most precise in correctly predicting the developmental stage based on a new set of data. This research also showed that a larger training set leads to better models, which seems reasonable and makes sense. As we saw here, the initial training set had 185 instances while the testing set had 908 instances. When the smaller dataset is used for training, the highest precision we achieved from any algorithm was only 64.2% (from the IBk algorithm). However, when the larger dataset was used for training and the smaller dataset was used for testing, ie the 'switched' case, we were able to achieve 81.6% precision (from the NaiveBayes algorithm). We were then able to reach an even slightly higher precision by looking at the 'switched' case and ranking the attributes before building the models. With this method, we were able to achieve 83.8% precision (again from the NaiveBayes algorithm).

We saw from the R analysis that Stage 5 was predicted the least precisely both in the 'switched' and 'unswitched' cases and we safely concluded that this is due to the small number of plants that were in Stage 5 in both datasets, relative to the other stages. We saw that Stage 4 was predicted the most precisely in both the 'switched' and 'unswitched' cases. We concluded that this was due to the fact that plants in this stage were larger and more developed, thereby being more defined than smaller plants which would have less information available to learn about them.

This program provides an easy way to test many different algorithms quickly and efficiently connected to the MP3 database. A user can easily build and train models with any algorithm that exists within the Weka library. Additionally, the program also provides direct querying from the MP3 database, though it can be easily modified to query another database. A possible future opportunity is to see if different datasets lead to even higher precision. For example, in this research we only looked at the vis-side-1-0 camera but it is very possible to choose a different camera and angle to see if there would be an improvement in precision results. Though I implemented the option for the user to rank the attributes in the dataset by which attributes are most important in determining the developmental stage, I did not have the chance to implement the other attribute selection options that Weka provides. For example, Weka has different attribute selection options that will actually eliminate certain attributes and only keep the attributes that are most important in determining the developmental stage of the plant. A further step for this research would be to implement these different attribute selection options that the user can choose from to see if there would be any improvement in precision, similar to the improvement that was observed when the attribute ranking was performed.

#### References

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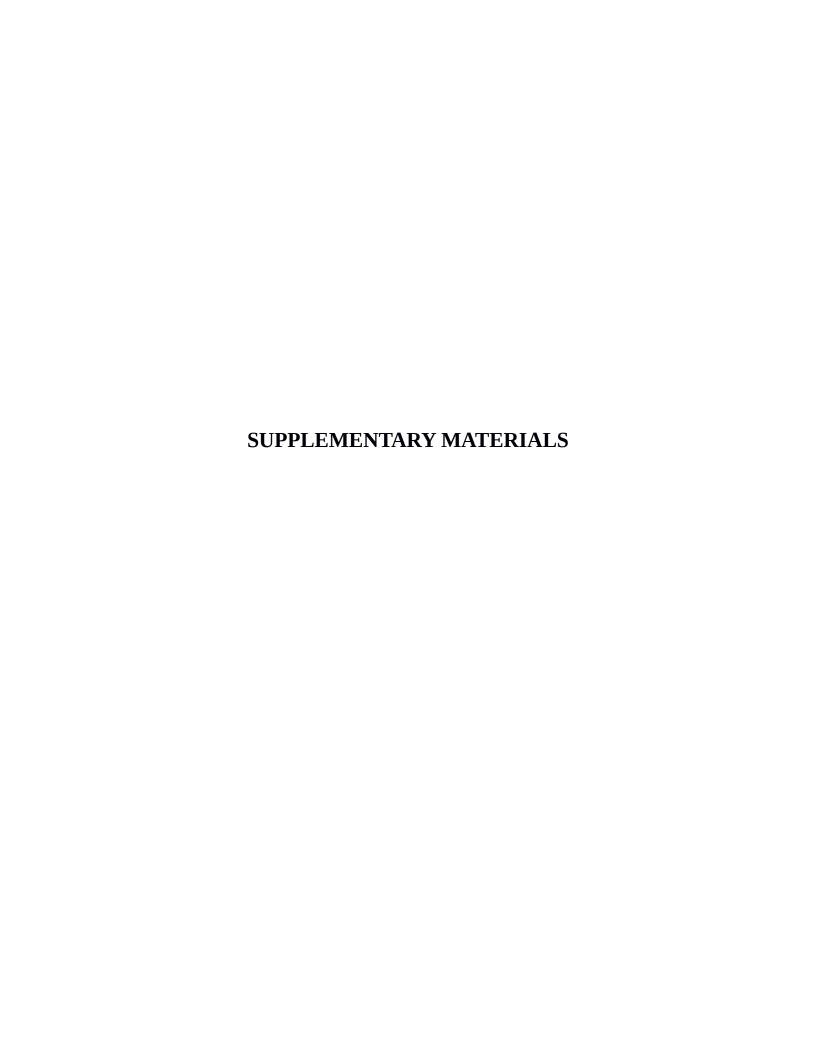
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```
1/*
 2 * @author Lea Collin
 3 */
 4 import weka.core.Instance;
 5 import weka.core.Instances;
 6 import weka.classifiers.Classifier;
 7 import weka.classifiers.AbstractClassifier;
 8 import weka.classifiers.meta.FilteredClassifier;
 9 import weka.core.converters.ArffLoader;
10 import weka.filters.unsupervised.attribute.Remove;
11 import weka.attributeSelection.*;
12 import java.io.File;
13 import java.io.FileNotFoundException;
14 import java.io.PrintWriter;
15 import java.io.IOException;
16 import java.util.ArrayList;
17 import java.util.Scanner;
18 import org.postgresql.util.PSQLException;
19 import java.sql.*;
21 public class Driver {
22
23
      static Scanner sc = new Scanner(System.in);
24
25
      public static void main(String [] args) throws Exception {
26
27
          //private database information
28
          String dbConfig = args[0];
29
30
          //ArrayList will contain all the possible algorithms that the user
  can input
          ArrayList<String> possibleClassifiers = new ArrayList<String>();
31
32
          possibleClassifiers.add("ZeroR");
33
          possibleClassifiers.add("NaiveBayes");
          possibleClassifiers.add("MultilayerPerceptron");
34
35
          possibleClassifiers.add("SMO");
36
          possibleClassifiers.add("IBk");
37
          possibleClassifiers.add("KStar");
38
          possibleClassifiers.add("LWL");
39
          possibleClassifiers.add("DecisionStump");
40
          possibleClassifiers.add("HoeffdingTree");
41
          possibleClassifiers.add("J48");
42
          possibleClassifiers.add("LMT");
          possibleClassifiers.add("RandomForest");
43
44
          possibleClassifiers.add("RandomTree");
45
          possibleClassifiers.add("REPTree");
46
47
          System.out.println("Please enter the directory name of where you
```

```
would like to store all program outputs:");
48
49
          //add control to check for valid directory
50
          String outputDir = setOutputDirectory();
51
52
          System.out.println();
53
          System.out.println("Please enter the name of the file you'd like to
  store the TRAINING data. Please end the file name in '.arff'");
          String trainingFile = setFileName(".arff", outputDir);
54
55
          //String trainingFile = outputDir + "trainingData.arff";
56
          System.out.println();
57
58
          System.out.println("Please enter the name of the file you'd like to
  store the TESTING data. Please end the file name in '.arff'");
59
          String testingFile = setFileName(".arff", outputDir);
60
          System.out.println();
61
62
          //try to connect to database given username and password, user
  prompted to enter username and password again if connection is unsuccessful
63
          boolean successfulConnection = false;
          while(!successfulConnection) {
64
65
                  System.out.println("Please enter your database username:");
66
                  String dbUsr = sc.next();
67
                  System.out.println("Password:");
                  String dbPwd = sc.next();
68
69
                  sc.nextLine();
70
              try {
                  connectToDatabase(dbUsr, dbPwd, dbConfig, testingFile,
71
  trainingFile);
72
73
                  successfulConnection = true:
74
              }catch (PSQLException s){
                  System.out.println("Username or password was incorrect.
75
  Please try again.");
76
              }
77
          }
78
79
          boolean rankAttributes = false:
80
          boolean isValid = false;
81
          while(!isValid) {
              System.out.println("Would you like to rank the attributes? (Y/n),
82
  (Could lead to better results)");
83
              String answer = sc.next();
84
              if(answer.equals("Y") || answer.equals("y")) {
85
                  isValid = true;
86
                  rankAttributes = true;
87
              }
88
              else if(answer.equals("N") || answer.equals("n")){
```

```
89
                    isValid = true;
 90
                    rankAttributes = false;
 91
               }
 92
               else {
 93
                    System.out.println("Could not understand input. Please enter
   a name again. \n");
 94
               }
 95
           }
 96
 97
           //getting user input for classifier names, checking if input is valid
           String [] classifiers = null;
 98
 99
           boolean validClassifier = false;
100
           while(!validClassifier){
101
102
                System.out.println("Do you want to run the default set of
   classification algorithms? (Y/n)");
103
               String answer = sc.next();
104
               if(answer.equals("Y") || answer.equals("y")) {
105
                    classifiers = new String[possibleClassifiers.size()];
106
                    for(int i = 0; i < classifiers.length; i++) {</pre>
107
                        classifiers[i] = possibleClassifiers.get(i);
108
109
                    validClassifier = true;
110
               else if(answer.equals("N") || answer.equals("n")){
111
                    System.out.println("Please enter the names of the classifiers
112
   you'd like to test, separated by a single space.");
113
                    sc.nextLine();
114
                    String classifierInput = sc.nextLine():
115
                    classifiers = classifierInput.split("\\s+");
116
117
                    for(int i = 0; i < classifiers.length; i++) {</pre>
                        if(!possibleClassifiers.contains(classifiers[i])) {
118
119
                            System.out.println(classifiers[i] + " is not a valid
   classifier name.");
120
                            System.out.println("Please try again.");
121
                            System.out.println();
122
                        }
123
124
                        if(i == classifiers.length - 1 &&
   possibleClassifiers.contains(classifiers[i])) {
125
                            validClassifier = true;
126
                        }
127
                    }
128
               }
129
               else {
130
                    System.out.println("Could not understand input.");
131
               }
```

```
132
133
           }
134
135
           System.out.println();
           System.out.println("Please enter the name of the file you would like
136
   to store all of the predictions. Please end the file in '.csv'");
137
           String predictionFile = setFileName(".csv", outputDir);
138
139
           System.out.println();
140
           System.out.println("Finally, please enter the name of the file you
   would like to store the precision of each algorithm you are testing. "
                   + "Please end the file in '.csv'");
141
           String precisionFile = setFileName(".csv", outputDir);
142
143
           System.out.println();
144
145
           sc.close():
146
147
           //reading the files and getting all the instances of each one
148
           Instances instancesTrain = fileReader(trainingFile);
149
           Instances instancesTest = fileReader(testingFile);
150
151
           //what attribute do we want to predict
152
           String classAttribute = "Stage";
153
           //these are the attributes to not be included in the classifier, das
154
   is a giveaway of the stage, barcode is irrelevant
           String [] attributesToRemove = {"barcode", "das"};
155
156
           //returns the indices of the attributes to be ignored
157
           String indicesToRemove = removeAttribute(instancesTrain,
   attributesToRemove);
158
159
           double maxPrecision = 0.0;
160
           String bestMethod = "";
161
162
163
           //run all the different classifiers
164
           Double [] precisions = predict(instancesTrain, instancesTest,
   classifiers, indicesToRemove, classAttribute, predictionFile,
   rankAttributes);
165
           //writing precision values to a csv
166
                   File output = new File(precisionFile);
167
168
                   PrintWriter pw = new PrintWriter(output);
169
                   StringBuilder sb = new StringBuilder();
170
                   sb.append("Algorithm");
171
                   sb.append(',');
172
                   sb.append("Precision");
173
                   sb.append("\n");
```

```
174
175
           //find best algorithm and write to file
176
           for(int i = 0; i < classifiers.length; i++) {</pre>
177
                if(precisions[i] > maxPrecision) {
178
                   maxPrecision = precisions[i];
179
                    bestMethod = classifiers[i];
180
               }
181
182
           //simply printing out for the user which algorithm was best and its
   precision
183
           System.out.println("Best Algorithm: " + bestMethod + " with
   precision: " + maxPrecision);
184
185
           pw.write(sb.toString());
186
           pw.close();
187
       }
188
189
       public static boolean validArff(String file) {
190
           //string must contain and end in .arff to be a valid arff file
191
           return (file.contains(".arff") && file.indexOf(".arff") ==
   file.length() - 5);
192
193
194
       public static boolean validCsv(String file) {
           return (file.contains(".csv") && file.index0f(".csv") == file.length
195
   () - 4);
196
       }
197
198
       public static boolean fileExists(String file) {
           File newFile = new File(file);
199
200
           return newFile.exists();
201
       }
202
203
       public static String setOutputDirectory() {
           String dirName = "";
204
           boolean isValid = false;
205
206
           while(!isValid) {
207
               dirName = sc.next();
208
               File dir = new File(dirName);
209
               if(dir.exists() && (dirName.charAt(dirName.length()-1) == '/')) {
210
211
                    isValid = true;
212
               }
               else {
213
214
                    System.out.println("Sorry that was an invalid directory.
   Please try again.");
215
               }
           }
216
```

```
217
218
           return dirName;
219
       }
220
221
       public static String setFileName(String fileType, String outputDir) {
           String outputFile = "";
222
223
           boolean isValid = false:
224
           while(!isValid) {
225
                outputFile = outputDir + sc.next();
226
                if(fileType.equals(".arff")) {
227
                    if(validArff(outputFile) == isValid) {
228
                        System.out.println("Sorry, the file you entered does not
   end in '.arff'. Please try again.");
229
                        continue;
230
                    }
231
               if(fileType.equals(".csv")) {
232
233
                    if(validCsv(outputFile) == isValid) {
234
                        System.out.println("Sorry, the file you entered does not
   end in '.csv'. Please try again.");
235
                        continue;
236
                    }
237
               if(fileExists(outputFile)) {
238
                    System.out.println("This file already exists in this
239
   directory. Do you want to overwrite it? (Y/n)?");
240
                    String answer = sc.next();
241
                    if(answer.equals("Y") || answer.equals("y")) {
242
                        isValid = true:
243
                    else if(answer.equals("N") || answer.equals("n")){
244
245
                        System.out.println("Please enter another name.");
246
                    }
247
                    else {
                        System.out.println("Could not understand input. Please
248
   enter a name again.");
249
                    }
250
               }
251
               else {
252
                   isValid = true;
253
               }
254
           }
255
           return outputFile;
256
       }
257
258
       public static Instances fileReader(String input) throws IOException {
259
260
           File inputFile = new File(input);
```

```
261
           ArffLoader atf = new ArffLoader();
262
           atf.setFile(inputFile);
263
           Instances data = atf.getDataSet();
264
265
           return data;
266
267
       }
268
269
       public static String removeAttribute(Instances data, String []
   attributes) throws Exception{
270
271
           String [] options = new String[2];
272
           options[0] = "-R";
273
274
           String indices = "";
275
           //getting all the indices of attributes we want to later remove (or
276
   ignore)
277
           for(int i = 0; i < attributes.length; i++) {</pre>
278
                int index = (data.attribute(attributes[i])).index() + 1;
279
               indices += index;
               if(i != attributes.length-1) {
280
281
                    indices += ",";
282
               }
283
           }
284
285
           return indices;
286
       }
287
288
       public static Double [] predict(Instances train, Instances test, String
   [] classifierNames,
           String indicesToRemove, String classAttribute, String outputFile,
   boolean rankAttributes) throws Exception {
290
           //removing attributes we don't want to include such as das and
291
   barcode
292
                    Remove rm = new Remove();
293
                    rm.setAttributeIndices(indicesToRemove);
294
295
           //running an attribute selector if user chose to rank attributes
296
                    if(rankAttributes) {
                        Instances [] reduced = attributeSelector(train, test);
297
298
299
                        train = reduced[0]:
300
301
                        test = reduced[1];
302
                    }
303
```

```
304
           //set the Class (what we want to predict)
305
           test.setClass(test.attribute(classAttribute));
306
307
           //setting the train class index to be the same as the testing class
   index
           train.setClassIndex(test.classIndex());
308
309
310
           //going to make an array of classifiers
311
           Classifier [] classifiers = new Classifier [classifierNames.length];
312
313
           //need somewhere to store the precision of each classifier +
   initializing the array
           Double precision [] = new Double [classifiers.length];
314
           for(int i = 0; i < precision.length; i++) {</pre>
315
316
                precision[i] = 0.0;
317
           }
318
319
           //creating filtered versions of each classifier (removing das and
   barcode)
320
           for(int i = 0; i < classifiers.length; i++) {</pre>
                Classifier temp = AbstractClassifier.forName(classifierNames[i],
321
   null):
322
                FilteredClassifier fc = new FilteredClassifier();
323
                fc.setFilter(rm);
324
325
                fc.setClassifier(temp);
326
               classifiers[i] = fc;
327
328
           }
329
330
           //building the models for each classifier
           for(int i = 0; i < classifiers.length; i++) {</pre>
331
                classifiers[i].buildClassifier(train);
332
333
           }
334
335
           //writing the header to the output csv
336
           File output = new File(outputFile);
337
           PrintWriter pw = new PrintWriter(output);
338
           StringBuilder sb = new StringBuilder();
339
           sb.append("Barcode");
340
           sb.append(',');
341
           sb.append("Das");
342
           sb.append(",");
           sb.append("Actual");
343
344
           sb.append(',');
345
           //making the names of the classifiers part of the header, will
346
   indicate the stage that classifier has predicted for a particular plant
```

```
347
           for(int i = 0; i < classifierNames.length; i++) {</pre>
348
                sb.append(classifierNames[i]);
349
               if(i != classifierNames.length - 1) {
                    sb.append(",");
350
351
               }
352
353
           sb.append("\n");
354
355
           double numInst = test.numInstances();
356
           //actually running the classifier
357
358
           for(int i = 0; i < numInst; i++){</pre>
359
360
               Instance current = test.instance(i);
361
               //will set the Stage of this 'temp' instance to be the predicted
362
   value to then compare it to the actual value of 'current'
               Instance temp = (Instance)current.copy();
363
364
               //attributes are given as array positions, getting the string
365
   value
366
               String actualVal = current.stringValue(test.classIndex());
367
368
                sb.append((int) current.value(test.attribute("barcode")));
369
               sb.append(',');
370
                sb.append((int) current.value(test.attribute("das")));
371
                sb.append(",");
                sb.append(actualVal);
372
373
               sb.append(',');
374
375
               for(int j = 0; j < classifiers.length; j++) {</pre>
               //getting the predicted value of the class attribute of this
376
   instance
377
               double predicted = classifiers[j].classifyInstance(test.instance
   (i));
378
379
               //setting this value to the temp class attribute
380
                temp.setValue(test.classIndex(), predicted);
381
382
               //getting the string value
               String predictedVal = temp.stringValue(temp.classIndex());
383
384
385
               //comparing predicted with actual value
386
               if(predictedVal.equals(actualVal)) {
387
                    precision[j]++;
388
               }
389
390
               sb.append(predictedVal);
```

```
391
392
               if(i != classifiers.length - 1) {
393
                   sb.append(",");
394
               }
395
396
               else if( j == classifiers.length - 1) {
397
                   sb.append('\n'):
398
               }
399
400
              }
           }
401
402
403
           for(int i = 0; i < precision.length; i++) {</pre>
404
               precision[i] = 100*precision[i]/numInst;
405
           }
406
407
           sb.append('\n');
408
           pw.write(sb.toString());
409
           pw.close();
410
411
           return precision;
       }
412
413
414
       private static void connectToDatabase(String usrDB, String passwordDB,
   String conDB, String trainName, String testName) throws SQLException,
   FileNotFoundException {
415
416
           File trainingOutput = new File(trainName);
417
           PrintWriter trainingPw = new PrintWriter(trainingOutput);
418
           StringBuilder sb = new StringBuilder();
419
420
           File testingOutput = new File(testName);
421
           PrintWriter testingPw = new PrintWriter(testingOutput);
422
423
           //creating the header for the arff file
           sb.append("@relation databasetraining" + "\n" + "\n" + "@attribute
424
   barcode numeric" + "\n" + "@attribute area numeric" + "\n" + "@attribute
   perimeter numeric" + "\n" +
425
           "@attribute circularity numeric" + "\n" + "@attribute compactness
   numeric" + "\n" + "@attribute major numeric" + "\n" +
           "@attribute minor numeric" + "\n" + "@attribute eccentricity numeric"
426
   + "\n" + "@attribute hisgreypeak numeric" + "\n" +
           "@attribute qlgrey numeric" + "\n" + "@attribute q2grey numeric" +
427
   "\n" + "@attribute q3grey numeric" + "\n" +
           "@attribute glr numeric" + "\n" + "@attribute g2r numeric" + "\n" +
428
   "@attribute g3r numeric" + "\n" + "@attribute g1g numeric" + "\n" +
           "@attribute q2g numeric" + "\n" + "@attribute q3g numeric" + "\n" +
   "@attribute q1b numeric" + "\n" + "@attribute q2b numeric" + "\n" +
```

```
430
           "@attribute q3b numeric" + "\n" + "@attribute das numeric" + "\n"
   +"@attribute Stage {'Stage 1','Stage 2','Stage 3','Stage 4', 'Stage 5'}" +
   "\n" + "\n" + "@data" + "\n");
431
432
           trainingPw.write(sb.toString());
433
           testingPw.write(sb.toString());
434
435
           try {
436
               Class.forName("org.postgresql.Driver");
437
               Connection conn = DriverManager.getConnection(conDB, usrDB,
   passwordDB);
438
439
               String trainingSql = "SELECT s.barcode, o.area, "
440
                       + "o.perimeter, o.circularity, o.compactness, "
441
                       + "o.major, o.minor, o.eccentricity, o.hisgreypeak, "
442
                       + "o.q1grey, o.q2grey, o.q3grey, "
443
                       + "o.q1r, o.q2r, o.q3r,
444
                       + "o.q1g, o.q2g, o.q3g,
445
                       + "o.q1b, o.q2b, o.q3b, d.das, "
446
                       + "CASE WHEN ( d.das <= 10 ) THEN 'Stage 1' "
447
                       + "WHEN ( d.das > 10 AND d.das <= 20 ) THEN 'Stage 2' "
                       + "WHEN ( d.das > 20 AND d.das <= 30 ) THEN 'Stage 3' "
448
449
                       + "WHEN ( d.das > 30 AND d.das <= 40 ) THEN 'Stage 4' "
                       + "WHEN ( d.das > 40 AND d.das <= 50 ) THEN 'Stage 5' "
450
451
                       + "ELSE 'Stage 6' END Stage "
                       + "FROM imageev AS i, imgobjectev AS o, soyidentification
452
   AS s, dasplusev AS d "
453
                       + "WHERE i.assayid = o.assayid "
454
                       + "AND i.imgid = o.imgid "
455
                       + "AND s.barcode = ( CAST( i.barcode AS INTEGER ) ) "
456
                       + "AND i.assayid = d.assayid AND i.fdate = d.fdate "
457
                       + "AND i.set = d.set "
                       + "AND s.line = 1 "
458
459
                       + "AND i.camera = 'vis-side-1-0' "
                       + "AND i.set = '3'";
460
461
462
               String testingSql = "SELECT s.barcode, o.area, "
463
                       + "o.perimeter, o.circularity, o.compactness, "
464
                        + "o.major, o.minor, o.eccentricity, o.hisgreypeak, "
465
                       + "o.q1grey, o.q2grey, o.q3grey, '
466
                       + "o.q1r, o.q2r, o.q3r,
                       + "o.q1g, o.q2g, o.q3g, "
467
468
                       + "o.q1b, o.q2b, o.q3b, d.das, "
469
                       + "CASE WHEN ( d.das <= 10 ) THEN 'Stage 1' "
                       + "WHEN ( d.das > 10 AND d.das <= 20 ) THEN 'Stage 2' "
470
                       + "WHEN ( d.das > 20 AND d.das <= 30 ) THEN 'Stage 3' "
471
                       + "WHEN ( d.das > 30 AND d.das <= 40 ) THEN 'Stage 4' "
472
473
                       + "WHEN ( d.das > 40 AND d.das <= 50 ) THEN 'Stage 5' "
```

```
474
                        + "ELSE 'Stage 6' END Stage "
475
                        + "FROM imageev AS i, imgobjectev AS o, soyidentification
   AS s, dasplusev AS d "
                        + "WHERE i.assayid = o.assayid "
476
477
                        + "AND i.imgid = o.imgid "
478
                        + "AND s.barcode = ( CAST( i.barcode AS INTEGER ) ) "
479
                        + "AND i.assavid = d.assavid "
480
                        + "AND i.fdate = d.fdate "
                       + "AND i.set = d.set "
481
                       + "AND ( s.line = 1 OR s.line = 2 OR s.line = 3 ) "
482
                        + "AND i.camera = 'vis-side-1-0' "
483
                       + "AND i.set = '2' "
484
                       + "UNION "
485
486
                       + "SELECT s.barcode, o.area, o.perimeter, o.circularity,
                        + "o.compactness, o.major, o.minor, o.eccentricity,
487
   o.hisgreypeak, "
488
                        + "o.qlgrey, o.q2grey, o.q3grey, "
489
                        + "o.q1r, o.q2r, o.q3r,"
                        + " o.q1g, o.q2g, o.q3g,
490
491
                        + "o.q1b, o.q2b, o.q3b, d.das, "
492
                       + "CASE WHEN ( d.das <= 10 ) THEN 'Stage 1' "
493
                        + "WHEN ( d.das > 10 AND d.das <= 20 ) THEN 'Stage 2' "
                        + "WHEN ( d.das > 20 AND d.das <= 30 ) THEN 'Stage 3' "
494
                        + "WHEN ( d.das > 30 AND d.das <= 40 ) THEN 'Stage 4' "
495
                       + "WHEN ( d.das > 40 AND d.das <= 50 ) THEN 'Stage 5' "
496
497
                        + "ELSE 'Stage 6' END Stage "
498
                        + "FROM imageev AS i, imgobjectev AS o, soyidentification
   AS s, dasplusev AS d "
499
                        + "WHERE i.assayid = o.assayid "
500
                       + "AND i.imgid = o.imgid "
501
                        + "AND s.barcode = ( CAST( i.barcode AS INTEGER ) ) "
502
                        + "AND i.assayid = d.assayid "
                        + "AND i.fdate = d.fdate "
503
                       + "AND i.set = d.set "
504
505
                       + "AND (s.line = 2 OR s.line = 3 ) "
506
                       + "AND i.camera = 'vis-side-1-0' "
507
                        + "AND i.set = '3' ";
508
509
               PreparedStatement trainingPs = conn.prepareStatement
   (trainingSql);
510
               ResultSet trainingSet = trainingPs.executeQuery();
               while(trainingSet.next()) {
511
512
                   Double barcode = trainingSet.getDouble("barcode");
513
                   Double area = trainingSet.getDouble("area");
                   Double perimeter = trainingSet.getDouble("perimeter");
514
515
                   Double circularity = trainingSet.getDouble("circularity");
                   Double compactness = trainingSet.getDouble("compactness");
516
```

```
517
                    Double major = trainingSet.getDouble("major");
518
                    Double minor = trainingSet.getDouble("minor");
519
                    Double eccentricity = trainingSet.getDouble("eccentricity");
520
                    Double hisgreypeak = trainingSet.getDouble("hisgreypeak");
                    Double q1grey = trainingSet.getDouble("q1grey");
521
522
                    Double q2grey = trainingSet.getDouble("q2grey");
523
                    Double g3grey = trainingSet.getDouble("g3grey");
524
                    Double q1r = trainingSet.getDouble("q1r");
                    Double q2r = trainingSet.getDouble("q2r");
525
526
                    Double q3r = trainingSet.getDouble("q3r");
527
                    Double q1g = trainingSet.getDouble("q1g");
528
                    Double q2g = trainingSet.getDouble("q2g");
                    Double q3g = trainingSet.getDouble("q3g");
529
                    Double q1b = trainingSet.getDouble("q1b");
530
531
                    Double g2b = trainingSet.getDouble("g2b");
532
                    Double q3b = trainingSet.getDouble("q3b");
                    Double das = trainingSet.getDouble("das");
533
534
                    String stage = trainingSet.getString("Stage");
535
536
                    //writing all the different attributes from the guery to the
   output arff file
                    trainingPw.write(barcode + "," + area + ", " + perimeter + ",
537
   " + circularity + ", " + compactness + ", " + major + ", " + minor + ", " +
   eccentricity
                            + ", " + hisgreypeak + ", " + qlgrey + ", " + q2grey
538
   + ", " + q3grey + ", " + q1r + ", " + q2r + ", " + q3r

+ ", " + q1g + ", " + q2g + ", " + q3g + ", " + q1b +

", " + q2b + ", " + q3b + "," + das + "," + "'" + stage + "'" + "\n");
539
540
541
                trainingSet.close();
542
                trainingPw.close();
543
544
                PreparedStatement testingPs = conn.prepareStatement(testingSql);
545
                ResultSet testingSet = testingPs.executeQuery();
546
                while(testingSet.next()) {
547
                    Double barcode = testingSet.getDouble("barcode");
548
                    Double area = testingSet.getDouble("area");
549
                    Double perimeter = testingSet.getDouble("perimeter");
550
                    Double circularity = testingSet.getDouble("circularity");
551
                    Double compactness = testingSet.getDouble("compactness");
                    Double major = testingSet.getDouble("major");
552
553
                    Double minor = testingSet.getDouble("minor");
554
                    Double eccentricity = testingSet.getDouble("eccentricity");
555
                    Double hisgreypeak = testingSet.getDouble("hisgreypeak");
556
                    Double glgrey = testingSet.getDouble("glgrey");
                    Double q2grey = testingSet.getDouble("q2grey");
557
558
                    Double q3grey = testingSet.getDouble("q3grey");
                    Double g1r = testingSet.getDouble("g1r");
559
```

```
560
                    Double g2r = testingSet.getDouble("g2r");
561
                    Double g3r = testingSet.getDouble("g3r");
                    Double q1g = testingSet.getDouble("q1g");
562
563
                    Double g2g = testingSet.getDouble("g2g");
564
                    Double q3g = testingSet.getDouble("q3g");
565
                    Double q1b = testingSet.getDouble("q1b");
566
                    Double a2b = testingSet.getDouble("a2b");
567
                    Double q3b = testingSet.getDouble("q3b");
568
                    Double das = testingSet.getDouble("das");
569
                    String stage = testingSet.getString("Stage");
570
                    testingPw.write(barcode + "," + area + ", " + perimeter + ",
571
    + circularity + ", " + compactness + ", " + major + ", " + minor + ", " +
   eccentricity
                         + ", " + hisgreypeak + ", " + qlgrey + ", " + q2grey
" + qlr + ", " + q2r + ", " + q3r
572
   + ", " + q3grey + ",
      + ", " + q1g + ", " + q2g + ", " + q3g + ", " + q1b + " + q2b + ", " + q3b + "," + das + "," + "'" + stage + "'" + "\n");
573
574
575
                testingSet.close();
576
                testingPw.close();
577
578
                conn.close();
            }
579
            catch (ClassNotFoundException e) {
580
581
582
                System.out.println("Improper database connection set-up.");
583
                e.printStackTrace();
584
585
            }
586
       }
587
588 public static Instances [] attributeSelector(Instances train, Instances
   test) throws Exception {
589
590
            AttributeSelection selector = new AttributeSelection();
591
            Ranker search = new Ranker();
592
            OneRAttributeEval eval = new OneRAttributeEval();
593
            selector.setEvaluator(eval);
594
            selector.setSearch(search);
595
            selector.SelectAttributes(train);
596
            //rankedAttributes gives the ranking of the attributes along with
597
   their weights
598
            //selected Attributes just gives the order of the ranking
599
            Instances trainTemp = selector.reduceDimensionality(train);
600
            Instances trainTest = selector.reduceDimensionality(test);
601
```

```
602
603
604
605
606
606
607
608}
Instances [] reduced = {trainTemp, trainTest};

return reduced;
```

```
library(ggplot2)
library(reshape2)
setwd("~/Documents/U4/Comp401/output")
predictions <- read.csv("originalPredictions.csv", header = TRUE)</pre>
unswitched <- rep(0, length(predictions) - 3)</pre>
classifier.name <- colnames(predictions[4:length(predictions)])</pre>
actual.stage <- as.vector(predictions[, "Actual"])</pre>
n <- length(actual.stage)</pre>
for(i in 4:length(predictions)){
     v <- as.vector(predictions[,i])</pre>
     unswitched[i-3] <- round(100*length(which(actual.stage == v))/n, 3)
### RESULTS WHEN TRAINING DATA AND TESTING DATA ARE SWITCHED
switchedPredictions <- read.csv("switchedPredictions.csv", header = TRUE)</pre>
switched <- rep(0, length(switchedPredictions) - 3)</pre>
classifier.name.s <- colnames(switchedPredictions[4:length(switchedPredictions)])</pre>
actual.stage.s <- as.vector(switchedPredictions[, "Actual"])</pre>
sn <- length(actual.stage.s)</pre>
for(i in 4:length(switchedPredictions)){
  sv <- as.vector(switchedPredictions[,i])</pre>
  switched[i-3] \leftarrow round(100*length(which(actual.stage.s == sv))/sn, 3)
allPrecisions <- data.frame(classifier.name, unswitched, switched)
allPrecisions<- melt(allPrecisions, id.vars = "classifier.name")</pre>
precision.plot <- ggplot(allPrecisions, aes(y = value, x = classifier.name, fill = variable))
precision.plot <- precision.plot + geom_bar(stat = "identity", position = "dodge")
precision.plot <- precision.plot + xlab("Classifier Name") + ylab("Percent Precision")
precision.plot <- precision.plot + guides(fill=guide_legend(title = "Training Data"))</pre>
precision.plot <- precision.plot + scale_fill_manual(values = c("#0033FF", "#6699FF"))</pre>
precision.plot <- precision.plot + geom_text(aes(label = round(value, 1)), position = position_dodge(0.75), vjust = -1)</pre>
precision.plot <- precision.plot + theme(axis.text.x = element_text(angle = 45, hjust = 1), text = element_text(size = 1)
24))
precision.plot
### Finding out which algorithms performed better between switched data
###BETTER UNSWITCHED
classifier.name[which(unswitched > switched)]
###BETTER SWITCHED
classifier.name[which(unswitched < switched)]</pre>
### RESULTS WHEN ATTRIBUTES ARE RANKED FIRST
rankedPredictions <- read.csv("RankedPredictions.csv", header = TRUE)</pre>
ranked <- rep(0, length(rankedPredictions) - 3)</pre>
classifier.name.r <- colnames(rankedPredictions[4:length(rankedPredictions)])</pre>
actual.stage.r <- as.vector(rankedPredictions[, "Actual"])</pre>
rn <- length(actual.stage.r)</pre>
for(i in 4:length(rankedPredictions)){
  rv <- as.vector(rankedPredictions[,i])</pre>
  ranked[i-3] \leftarrow round(100*length(which(actual.stage.r == rv))/rn, 3)
allPrecisions <- data.frame(classifier.name.r, switched, ranked)
allPrecisions<- melt(allPrecisions, id.vars = "classifier.name.r")</pre>
precision.plot <- ggplot(allPrecisions, aes(y = value, x = classifier.name.r, fill = variable)) precision.plot <- precision.plot + geom_bar(stat = "identity", position = "dodge") precision.plot <- precision.plot + xlab("Classifier Name") + ylab("Percent Precision")
precision.plot <- precision.plot + guides(fill=guide_legend(title = "Training Data"))</pre>
precision.plot <- precision.plot + scale_fill_manual(values = c("#0033FF", "#6699FF"))</pre>
precision.plot <- precision.plot + geom\_text(aes(label = round(value, 1)), position = position\_dodge(0.75), vjust = -1)
precision.plot <- precision.plot + theme(axis.text.x = element text(angle = 45, hjust = 1), text = element text(size =
24))
precision.plot
###BETTER RANKED
classifier.name.r[which(ranked > switched)]
###NO DIFFERENCE
classifier.name.r[which(ranked == switched)]
###BETTER UNRANKED
classifier.name.r[which(ranked < switched)]</pre>
###LOOKING AT THE STAGE PREDICTION BREAKDOWN WHEN THE ATTRIBUTES ARE RANKED
predictions <- read.csv("RankedPredictions.csv", header = TRUE)</pre>
classifier.name <- colnames(predictions[5:length(predictions)])</pre>
actual.stage <- as.vector(predictions[, "Actual"])</pre>
n <- length(actual.stage)</pre>
stages <- c("Stage 1", "Stage 2", "Stage 3", "Stage 4", "Stage 5")
numStages <- rep(0, length(stages))</pre>
for (i in 1:length(stages)){
  numStages[i] <- length(which(predictions[,"Actual"] == stages[i]))</pre>
```

```
}
percent.correct <- rep(0, length(stages))</pre>
for(i in 1:length(stages)){
  indices <- which(predictions[,"Actual"] == stages[i])</pre>
  for(j in 5:length(predictions)){
    percent.correct[i] <- percent.correct[i] + length(which(predictions[indices, j] == stages[i]))</pre>
### percent.correct is an AVERAGE of the percent time a stage is predicted correctly
percent.correct <- 100*(round(percent.correct/(numStages*(length(classifier.name)-1)),3))</pre>
percent.correct <- data.frame(stages, percent.correct)</pre>
stage.plot <- ggplot(percent.correct, aes(y = percent.correct, x = stages))
stage.plot <- stage.plot + geom_bar(stat = "identity", fill = "#0033FF")</pre>
stage.plot <- stage.plot + xlab("Stage") + ylab("Percent Correctly Predicted") + ggtitle("Average Precision per Stage")
stage.plot <- stage.plot + geom text(aes(label = round(percent.correct, 1)), position = position dodge(0.75), vjust =
-1)
stage.plot
###What percentage of the time was stage 1 predicted?
times.predicted <- rep(0, length(stages))</pre>
for(i in 1:length(stages)){
  for(j in 5:length(predictions)){
    times.predicted[i] <- times.predicted[i] + length(which(predictions[,j] == stages[i]))</pre>
}
times.predicted <- 100*(round(times.predicted/(sum(times.predicted)),5))</pre>
times.predicted <- data.frame(stages, times.predicted)</pre>
times.predicted.plot <- ggplot(times.predicted, aes(y = times.predicted, x = stages))
times.predicted.plot <- times.predicted.plot + geom_bar(stat = "identity", fill = "#0033FF")
times.predicted.plot <- times.predicted.plot + xlab("Stage") + ylab("Percent of Time Predicted") + ggtitle("Percentage
Predicted per Stage")
times.predicted.plot <- times.predicted.plot + geom text(aes(label = round(times.predicted, 1)), position =
position dodge(0.75), vjust = -1)
times.predicted.plot
### WORKING WITH JUST THE SWITCHED DATA SINCE THIS LED TO OVERALL BETTER RESULTS
predictions <- read.csv("switchedPredictions.csv", header = TRUE)
classifier.name <- colnames(predictions[5:length(predictions)])</pre>
actual.stage <- as.vector(predictions[, "Actual"])</pre>
n <- length(actual.stage)</pre>
stages <- c("Stage 1", "Stage 2", "Stage 3", "Stage 4", "Stage 5")
numStages <- rep(0, length(stages))</pre>
for (i in 1:length(stages)){
  numStages[i] <- length(which(predictions[,"Actual"] == stages[i]))</pre>
percent.correct <- rep(0, length(stages))</pre>
for(i in 1:length(stages)){
  indices <- which(predictions[,"Actual"] == stages[i])</pre>
  for(j in 5:length(predictions)){
   percent.correct[i] <- percent.correct[i] + length(which(predictions[indices, j] == stages[i]))</pre>
  }
### percent.correct is an AVERAGE of the percent time a stage is predicted correctly
percent.correct <- 100*(round(percent.correct/(numStages*(length(classifier.name)-1)),3))</pre>
percent.correct <- data.frame(stages, percent.correct)</pre>
stage.plot <- ggplot(percent.correct, aes(y = percent.correct, x = stages))
stage.plot <- stage.plot + geom_bar(stat = "identity", fill = "#0033FF")</pre>
stage.plot <- stage.plot + xlab("Stage") + ylab("Percent Correctly Predicted")</pre>
stage.plot <- stage.plot + geom_text(aes(label = round(percent.correct, 1)), position = position_dodge(0.75), vjust =</pre>
stage.plot <- stage.plot + theme(axis.text.x = element text(angle = 45, hjust = 1), text = element text(size = 18))
stage.plot
###What percentage of the time was stage 1 predicted?
predictions <- read.csv("switchedPredictions.csv", header = TRUE)</pre>
times.predicted <- rep(0, length(stages))</pre>
for(i in 1:length(stages)){
  for(j in 5:length(predictions)){
    times.predicted[i] <- times.predicted[i] + length(which(predictions[,j] == stages[i]))</pre>
  }
times.predicted <- 100*(round(times.predicted/(sum(times.predicted)),5))</pre>
times.predicted <- data.frame(stages, times.predicted)</pre>
times.predicted.plot \leftarrow ggplot(times.predicted, aes(y = times.predicted, x = stages))
times.predicted.plot <- times.predicted.plot + geom_bar(stat = "identity", fill = "#0033FF")
times.predicted.plot <- times.predicted.plot + xlab("Stage") + ylab("Percent of Time Predicted")</pre>
times.predicted.plot <- times.predicted.plot + geom text(aes(label = round(times.predicted, 1)), position =
position dodge(0.75), vjust = -1)
```

```
times.predicted.plot <- times.predicted.plot + theme(axis.text.x = element text(angle = 45, hjust = 1), text =
element text(size = 18))
times.predicted.plot
###How many of each stage was there in the testing set?
trainingSet <- read.csv("DatabaseTesting.csv", header = TRUE)</pre>
stages.training <- rep(0, length(stages))</pre>
for (i in 1:length(stages)){
 stages.training[i] <- length(which(trainingSet[,"Stage"] == stages[i]))</pre>
####WORKING WITH THE ORIGINAL DATA, COPIED CODE BASICALLY FROM ABOVE
predictions <- read.csv("originalPredictions.csv", header = TRUE)</pre>
classifier.name <- colnames(predictions[5:length(predictions)])</pre>
actual.stage <- as.vector(predictions[, "Actual"])</pre>
n <- length(actual.stage)</pre>
stages <- c("Stage 1", "Stage 2", "Stage 3", "Stage 4", "Stage 5")
numStages <- rep(0, length(stages))</pre>
for (i in 1:length(stages)){
  numStages[i] <- length(which(predictions[,"Actual"] == stages[i]))</pre>
percent.correct <- rep(0, length(stages))</pre>
for(i in 1:length(stages)){
  indices <- which(predictions[,"Actual"] == stages[i])</pre>
  for(j in 5:length(predictions)){
    percent.correct[i] <- percent.correct[i] + length(which(predictions[indices, j] == stages[i]))</pre>
### percent.correct is an AVERAGE of the percent time a stage is predicted correctly
percent.correct <- 100*(round(percent.correct/(numStages*(length(classifier.name)-1)),3))</pre>
percent.correct <- data.frame(stages, percent.correct)</pre>
stage.plot <- ggplot(percent.correct, aes(y = percent.correct, x = stages))
stage.plot <- stage.plot + geom_bar(stat = "identity", fill = "#0033FF")</pre>
stage.plot <- stage.plot + xlab("Stage") + ylab("Percent Correctly Predicted") + ggtitle("Average Precision per Stage")</pre>
stage.plot <- stage.plot + geom_text(aes(label = round(percent.correct, 1)), position = position_dodge(0.75), vjust =</pre>
-1)
stage.plot
###What percentage of the time was stage 1 predicted?
times.predicted <- rep(0, length(stages))</pre>
for(i in 1:length(stages)){
  for(j in 5:length(predictions)){
    times.predicted[i] <- times.predicted[i] + length(which(predictions[,j] == stages[i]))
times.predicted <- 100*(round(times.predicted/(sum(times.predicted)),5))</pre>
times.predicted <- data.frame(stages, times.predicted)</pre>
times.predicted.plot <- ggplot(times.predicted, aes(y = times.predicted, x = stages))
times.predicted.plot <- times.predicted.plot + geom_bar(stat = "identity", fill = "#0033FF")
times.predicted.plot <- times.predicted.plot + xlab("Stage") + ylab("Percent of Time Predicted") + ggtitle("Percentage
Predicted per Stage")
times.predicted.plot <- times.predicted.plot + geom_text(aes(label = round(times.predicted, 1)), position =</pre>
position dodge(0.75), vjust = -1)
times.predicted.plot
```