

## Homework 3

### 1. Run four models:

#### 1. J48 with 10-fold CV (or some decision tree algorithm)

=== Run information ===

Scheme: weka.classifiers.trees.J48 -C 0.25 -M 2

Relation: HW3\_1

Instances: 8416

Attributes: 23

- Class
- cap-shape
- cap-surface
- cap-color
- ruises
- odor
- gill-attachment
- gill-spacing
- gill-size
- gill-color
- stalk-shape
- stalk-root
- stalk-surface-above-ring
- stalk-surface-below-ring
- stalk-color-above-ring
- stalk-color-below-ring
- veil-type
- veil-color
- ring-number
- ring-type
- spore-print-color
- population
- habitat

Test mode: 10-fold cross-validation

=== Classifier model (full training set) ===

J48 pruned tree

-----

odor = ALMOND: EDIBLE (400.0)

odor = ANISE: EDIBLE (400.0)

odor = NONE

| spore-print-color = PURPLE: EDIBLE (0.0)

| spore-print-color = BROWN: EDIBLE (1472.0)

### Homework 3

```
| spore-print-color = BLACK: EDIBLE (1424.0)
| spore-print-color = CHOCOLATE: EDIBLE (48.0)
| spore-print-color = GREEN: POISONOUS (72.0)
| spore-print-color = WHITE
| | gill-size = NARROW
| | | gill-spacing = CROWDED
| | | | population = SEVERAL: EDIBLE (72.0)
| | | | population = SCATTERED: EDIBLE (0.0)
| | | | population = NUMEROUS: EDIBLE (0.0)
| | | | population = SOLITARY: EDIBLE (0.0)
| | | | population = ABUNDANT: EDIBLE (0.0)
| | | | population = CLUSTERED: POISONOUS (16.0)
| | | gill-spacing = CLOSE: POISONOUS (32.0)
| | gill-size = BROAD: EDIBLE (528.0)
| spore-print-color = YELLOW: EDIBLE (48.0)
| spore-print-color = ORANGE: EDIBLE (48.0)
| spore-print-color = BUFF: EDIBLE (48.0)
odor = PUNGENT: POISONOUS (256.0)
odor = CREOSOTE: POISONOUS (192.0)
odor = FOUL: POISONOUS (2160.0)
odor = FISHY: POISONOUS (576.0)
odor = SPICY: POISONOUS (576.0)
odor = MUSTY: POISONOUS (48.0)
```

Number of Leaves : 24

Size of the tree : 29

Time taken to build model: 0.02 seconds

=== Stratified cross-validation ===

=== Summary ===

Correctly Classified Instances	8416	100	%
Incorrectly Classified Instances	0	0	%
Kappa statistic	1		
Mean absolute error	0		
Root mean squared error	0		
Relative absolute error	0	%	
Root relative squared error	0	%	
Total Number of Instances	8416		

=== Detailed Accuracy By Class ===

## Homework 3

	TP Rate	FP Rate	Precision	Recall	F-Measure	MCC	ROC Area	PRC Area	Class
	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	EDIBLE
	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	POISONOUS
Weighted Avg.	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	

=== Confusion Matrix ===

```
a  b <-- classified as
4488  0 |  a = EDIBLE
  0 3928 |  b = POISONOUS
```

## 2. RandomForests with 10-fold CV

=== Run information ===

Scheme: weka.classifiers.trees.RandomForest -P 100 -I 100 -num-slots 1 -K 0 -M 1.0 -V 0.001  
-S 1

Relation: HW3\_1

Instances: 8416

Attributes: 23

- Class
- cap-shape
- cap-surface
- cap-color
- ruises
- odor
- gill-attachment
- gill-spacing
- gill-size
- gill-color
- stalk-shape
- stalk-root
- stalk-surface-above-ring
- stalk-surface-below-ring
- stalk-color-above-ring
- stalk-color-below-ring
- veil-type
- veil-color
- ring-number
- ring-type
- spore-print-color
- population

## Homework 3

habitat

Test mode: 10-fold cross-validation

=== Classifier model (full training set) ===

RandomForest

Bagging with 100 iterations and base learner

weka.classifiers.trees.RandomTree -K 0 -M 1.0 -V 0.001 -S 1 -do-not-check-capabilities

Time taken to build model: 0.21 seconds

=== Stratified cross-validation ===

=== Summary ===

Correctly Classified Instances	8416	100	%
Incorrectly Classified Instances	0	0	%
Kappa statistic	1		
Mean absolute error	0.0004		
Root mean squared error	0.003		
Relative absolute error	0.0704	%	
Root relative squared error	0.6078	%	
Total Number of Instances	8416		

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	MCC	ROC Area	PRC Area	Class
	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	EDIBLE
	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	POISONOUS
Weighted Avg.	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	

=== Confusion Matrix ===

a	b	<-- classified as
4488	0	a = EDIBLE
0	3928	b = POISONOUS

### 3. OneR with 10-fold CV (or some non-pruning rule algorithm)

=== Run information ===

Scheme: weka.classifiers.rules.OneR -B 6

## Homework 3

Relation: HW3\_1

Instances: 8416

Attributes: 23

Class

cap-shape

cap-surface

cap-color

ruises

odor

gill-attachment

gill-spacing

gill-size

gill-color

stalk-shape

stalk-root

stalk-surface-above-ring

stalk-surface-below-ring

stalk-color-above-ring

stalk-color-below-ring

veil-type

veil-color

ring-number

ring-type

spore-print-color

population

habitat

Test mode: 10-fold cross-validation

=== Classifier model (full training set) ===

odor:

ALMOND -> EDIBLE

ANISE -> EDIBLE

NONE -> EDIBLE

PUNGENT -> POISONOUS

CREOSOTE -> POISONOUS

FOUL -> POISONOUS

FISHY -> POISONOUS

SPICY -> POISONOUS

MUSTY -> POISONOUS

(8296/8416 instances correct)

Time taken to build model: 0.03 seconds

## Homework 3

=== Stratified cross-validation ===

=== Summary ===

Correctly Classified Instances	8296	98.5741 %
Incorrectly Classified Instances	120	1.4259 %
Kappa statistic	0.9713	
Mean absolute error	0.0143	
Root mean squared error	0.1194	
Relative absolute error	2.8644 %	
Root relative squared error	23.9349 %	
Total Number of Instances	8416	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	MCC	ROC Area	PRC Area	Class
	1.000	0.031	0.974	1.000	0.987	0.972	0.985	0.974	EDIBLE
	0.969	0.000	1.000	0.969	0.984	0.972	0.985	0.984	POISONOUS
Weighted Avg.	0.986	0.016	0.986	0.986	0.986	0.972	0.985	0.979	

=== Confusion Matrix ===

```
a  b  <-- classified as
4488  0 |  a = EDIBLE
120 3808 |  b = POISONOUS
```

### 4. JRip with 10-fold CV (or some rule-pruning algorithm)

=== Run information ===

Scheme: weka.classifiers.rules.JRip -F 3 -N 2.0 -O 2 -S 1  
Relation: HW3\_1  
Instances: 8416  
Attributes: 23  
Class  
cap-shape  
cap-surface  
cap-color  
ruises  
odor  
gill-attachment  
gill-spacing  
gill-size  
gill-color

## Homework 3

stalk-shape  
stalk-root  
stalk-surface-above-ring  
stalk-surface-below-ring  
stalk-color-above-ring  
stalk-color-below-ring  
veil-type  
veil-color  
ring-number  
ring-type  
spore-print-color  
population  
habitat

Test mode: 10-fold cross-validation

=== Classifier model (full training set) ===

JRIP rules:

=====

(odor = FOUL) => Class=POISONOUS (2160.0/0.0)  
(gill-size = NARROW) and (gill-color = BUFF) => Class=POISONOUS (1152.0/0.0)  
(gill-size = NARROW) and (odor = PUNGENT) => Class=POISONOUS (256.0/0.0)  
(odor = CREOSOTE) => Class=POISONOUS (192.0/0.0)  
( spore-print-color = GREEN) => Class=POISONOUS (72.0/0.0)  
( stalk-surface-below-ring = SCALY) and (stalk-surface-above-ring = SILKY) => Class=POISONOUS  
(80.0/0.0)  
(stalk-color-above-ring = YELLOW) => Class=POISONOUS (8.0/0.0)  
(population = CLUSTERED) and (cap-color = WHITE) => Class=POISONOUS (8.0/0.0)  
=> Class=EDIBLE (4488.0/0.0)

Number of Rules : 9

Time taken to build model: 0.18 seconds

=== Stratified cross-validation ===

=== Summary ===

Correctly Classified Instances	8416	100	%
Incorrectly Classified Instances	0	0	%
Kappa statistic	1		
Mean absolute error	0		
Root mean squared error	0		

## Homework 3

Relative absolute error            0 %  
 Root relative squared error       0 %  
 Total Number of Instances       8416

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	MCC	ROC Area	PRC Area	Class
	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	EDIBLE
	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	POISONOUS
Weighted Avg.	1.000	0.000	1.000	1.000	1.000	1.000	1.000	1.000	

=== Confusion Matrix ===

```

a   b  <-- classified as
4488  0 |  a = EDIBLE
  0 3928 |  b = POISONOUS
    
```

**2. Compare and contrast the models: time to build, size, accuracy/error measure, confusion matrix, etc. Discuss your results.**

### 1. J48 with RandomForest

	J48	RandomForests
<b>Time to build</b>	0.02 seconds	0.21 seconds
<b>Size</b>	29	-
<b>Accuracy/Error measure</b>	100 %	100 %
<b>Confusion matrix</b>	<pre> a   b  &lt;-- classified as 4488  0    a = EDIBLE   0 3928    b = POISONOUS           </pre>	<pre> a   b  &lt;-- classified as 4488  0    a = EDIBLE   0 3928    b = POISONOUS           </pre>

J48 and RandomForest are both from the Trees model. J48 takes much less time than RandomForest to build with the cross validation of 10.

### 2. OneR with JRip

	OneR	JRip
<b>Time to build</b>	0.03 seconds	0.18 seconds
<b>Size</b>	-	-
<b>Accuracy/Error measure</b>	98.5741 %	100 %
<b>Confusion matrix</b>	<pre> a   b  &lt;-- classified as 4488  0    a = EDIBLE 120 3808    b = POISONOUS           </pre>	<pre> a   b  &lt;-- classified as 4488  0    a = EDIBLE   0 3928    b = POISONOUS           </pre>



## Homework 3

OneR and JRip are both from the Rules model. In comparison of JRip, OneR takes much less time to build with the cross validation of 10.

### 3. Trees with rules

J48 and RandomForest are models of Trees. OneR and JRip are models of Rules. First comparison of J48, which is a model of Trees with OneR, which is a model of Rules. In this comparison, the time to build for J48 is less than the time taken for OneR, when cross-validation folds is 10. The accuracy for J48 is 100% while the accuracy for OneR is 98.5741%. Second comparison of RandomForest, which is a model of Trees with JRip, which is a model of Rules. In this comparison, the time to build for JRip is less than the time taken for RandomForest, when cross-validation folds is 10.

### 3. Examine the results of your models. Which is more understandable?

**How would you present these results to the user? Which model do you prefer, and why?**

There are parts of two models which I examined i.e., Trees and Rules. For the tree-based, it forces the consideration of all possible outcomes of a decision and traces each path to a conclusion. For the rule-based, it is sometimes problematic because those who want to comply with rules are not always sure of everything they need to look at. In the tree-based, the accuracy will be 100% but, in the rule-based, it is not sure the accuracy to be 100%. These are the reason I will prefer the Trees model. The graph is a way to present these results to the user. We can use matplotlib from which we can display scatter plot, bar plot, etc. In the Tree model, I would prefer a J48 model, as the time taken to build this model is 0.02 seconds which is less than all the examined models. Also, the accuracy of this model is 100%.