

Machine Learning the WEKA directions

Week-7

WEKA IS

- ✓ Machine Learning toolkit
 - ✓ Open Source (Java)
 - ✓ Well documented + huge community
 - ✓ Provides API, command line and GUI-swing tools
 - ✓ Relatively easy to learn
-
- ✓ Runnable on a remote server so you can “dumb terminal” your laptop and keep your data in one place!

WEKA is not...

- A complete replacement for R/Matlab
- Optimized out of the box for multiple CPUs / compute farms
- An excuse to ignore the method details of a clustering/classification algorithm

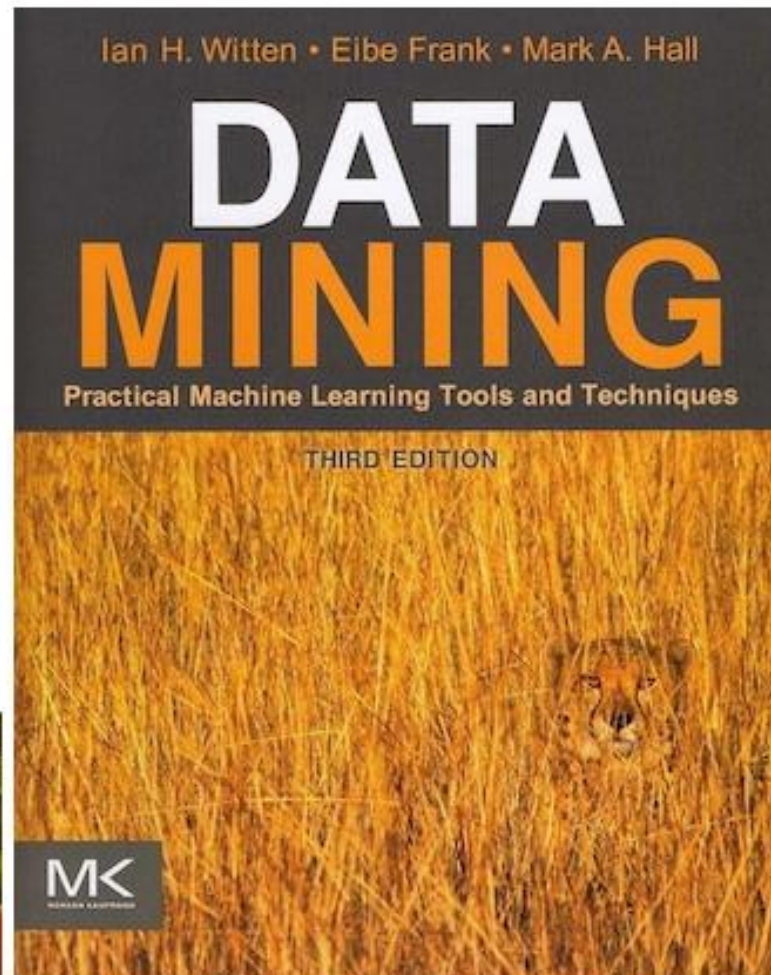
Data Mining: Practical Machine Learning Tools and Techniques (Third Edition)

Ian H. Witten, Eibe Frank,
Mark A. Hall

Morgan Kaufmann
January 2011
629 pages
Paper
ISBN
978-0-12-374856-0



Eibe Frank and Ian
Witten



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Personal WEKA dataset

- ✓ 21k+ variables
- ✓ 14k+ subjects phenotyped
- ✓ 9k+ subjects genotyped 500k Affymetrix
- ✓ 54M recorded phenotype values of widely varying types
- ✓ Even the simplest correlation matrix
 - ✓ $20k * 20k = 400M$ comparisons *before* including SNPs

WEKA basics

NOTE: GUI will be Slightly differ due to versions

Iris Example Data Set



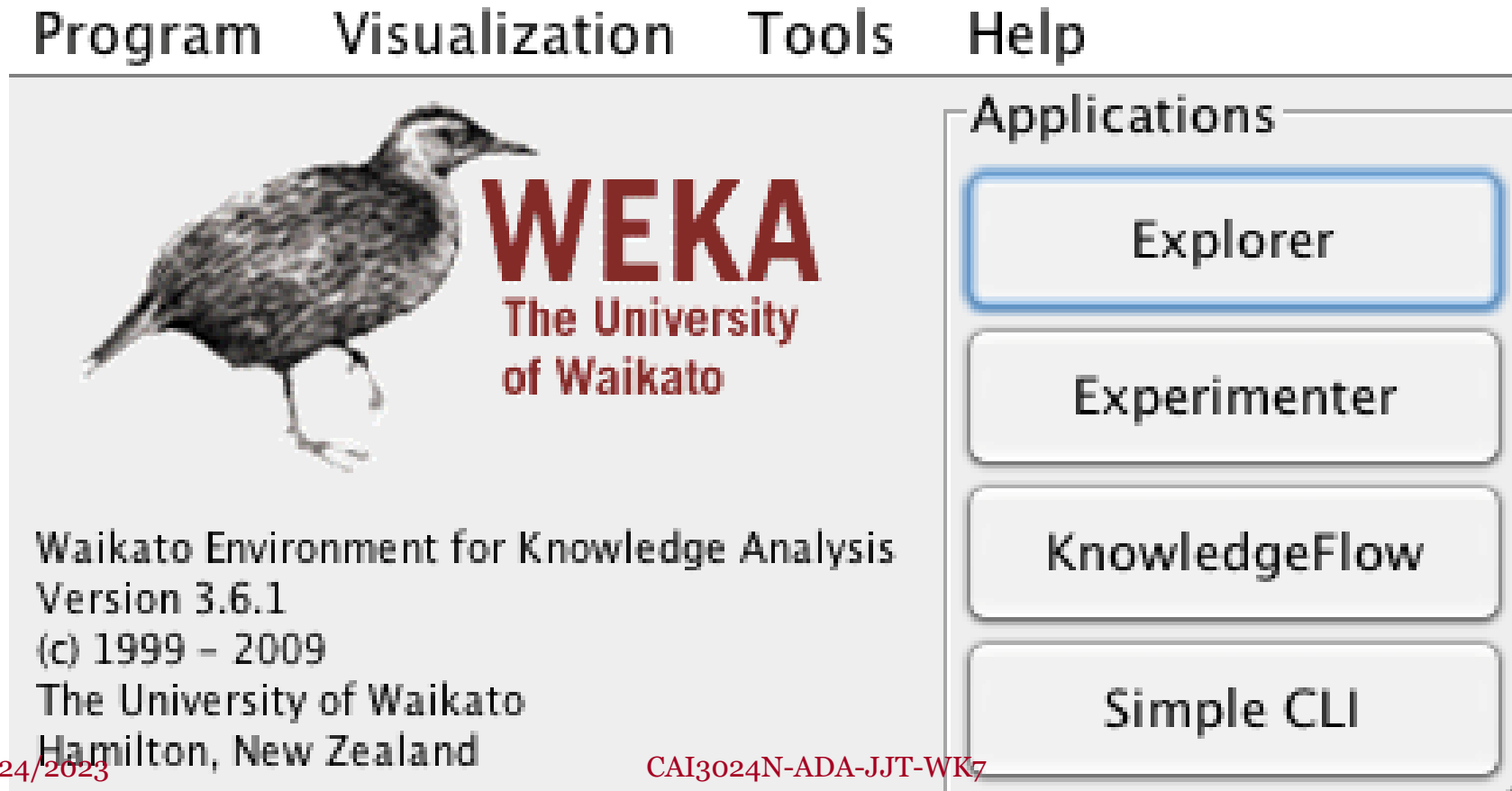
Instances

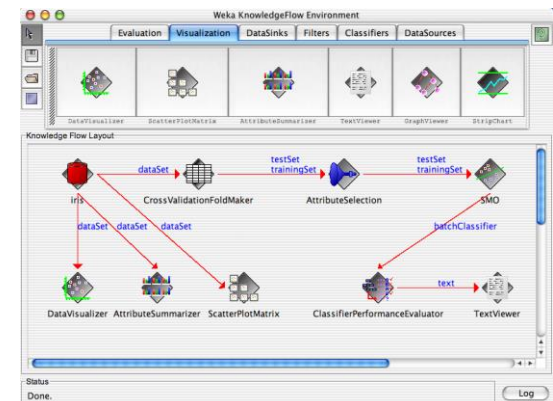
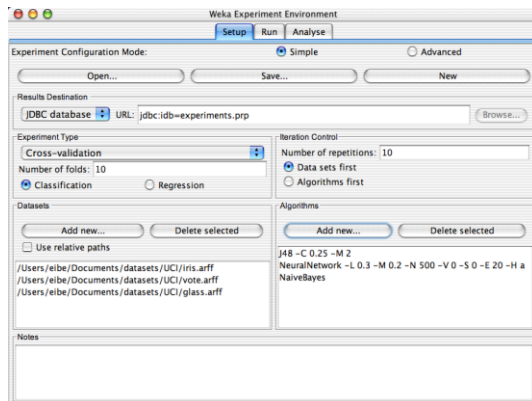
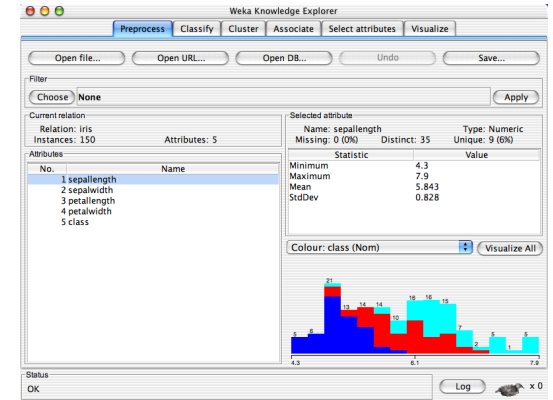
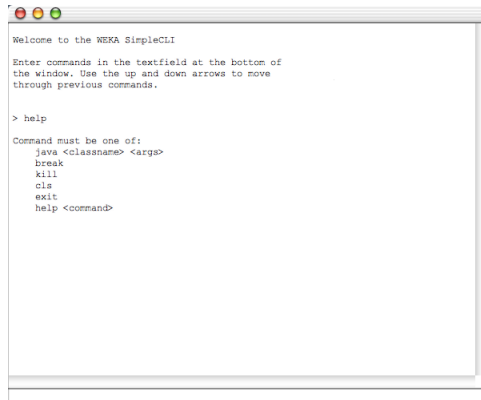
Features		Class
<u>Sepal</u>	<u>Petal</u>	<u>Species</u>
Length	Width	
Pick flower 1		1
Pick flower 2		2
Pick flower 3		3
Pick flower N		????

WEKA basics

NOTE: GUI will be Slightly differ due to versions

- API backs all functions of the CLI/GUI interfaces, can be easily used for your own project.





WEKA Explorer Tutorial Examples

❶ Preprocess

- Instance and Attribute Filters (Supervised and Unsupervised)

❷ Classify

- Bayes

❸ Cluster

- Expectation Maximization
- Hierarchical Clustering

❹ Associate

- Apriori

❺ Select Attributes

- Via clustering

Preprocess

- File: CSV, ARFF*,
- Database: direct SQL access (useful)

The screenshot shows the 'Preprocess' tab of the Weka Explorer interface. At the top, there are tabs for 'Preprocess', 'Classify', 'Cluster', 'Associate', 'Select attributes', and 'Visualize'. Below these are buttons for 'Open file...', 'Open URL...', 'Open DB...', 'Generate...', 'Undo', 'Edit...', and 'Save...'. The 'Filter' section has a 'Choose' button and a dropdown menu currently set to 'None', with an 'Apply' button. The 'Current relation' section shows 'Relation: None', 'Instances: None', and 'Attributes: None'. The 'Attributes' section has buttons for 'All', 'None', 'Invert', and 'Pattern', followed by a list box and a 'Remove' button. The 'Selected attribute' section shows 'Name: None', 'Missing: None', 'Distinct: None', 'Type: None', and 'Unique: None', with a list box and a 'Visualize All' button. The status bar at the bottom left says 'Status 10/24/2023 Welcome to the Weka Explorer'.

Preprocess | Classify | Cluster | Associate | Select attributes | Visualize

Open file... | Open URL... | Open DB... | Generate... | Undo | Edit... | Save...

Filter
Choose | None | Apply

Current relation
Relation: None
Instances: None
Attributes: None

Attributes
All | None | Invert | Pattern
[List Box]
Remove

Selected attribute
Name: None
Missing: None
Distinct: None
Type: None
Unique: None
[List Box]
Visualize All

Status
10/24/2023
Welcome to the Weka Explorer

WEKA “flat” files

@relation heart-disease-simplified

@attribute age numeric

@attribute sex { female, male}

@attribute chest_pain_type { typ_angina, asympt, non_anginal, atyp_angina}

@attribute cholesterol numeric

@attribute exercise_induced_angina { no, yes}

@attribute class { present, not_present}

@data

63,male,typ_angina,233,no,not_present

67,male,asympt,286,yes,present

67,male,asympt,229,yes,present

38,female,non_anginal,?,no,not_present

...



**Flat file in
ARFF format**

WEKA “flat” files

@relation heart-disease-simplified

@attribute age numeric

@attribute sex { female, male}

@attribute chest_pain_type { typ_angina, asympt, non_anginal, atyp_angina}

@attribute cholesterol numeric

@attribute exercise_induced_angina { no, yes}

@attribute class { present, not_present}

@data

63,male,typ_angina,233,no,not_present

67,male,asympt,286,yes,present

67,male,asympt,229,yes,present

38,female,non_anginal,?,no,not_present

...

numeric attribute

nominal attribute





Weka Knowledge Explorer

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

None

Apply

Current relation

Relation: None

Instances: None

Attributes: None

Selected attribute

Name: None

Missing: None

Type: None

Distinct: None

Unique: None

Attributes

Empty list for attributes

Empty list for selected attribute details



Visualize All

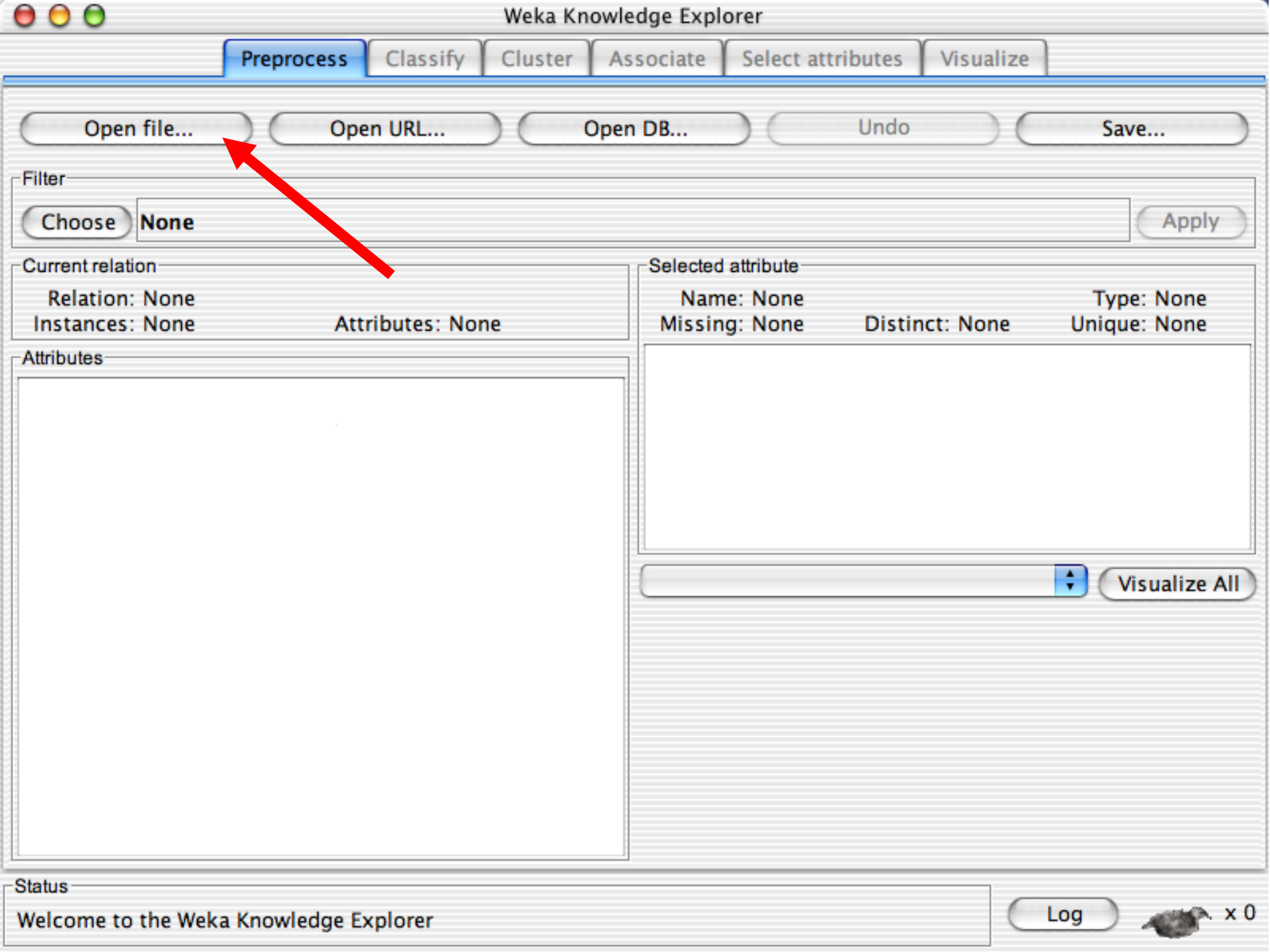
Status

Welcome to the Weka Knowledge Explorer

Log



x 0





Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

None

Apply

Current relation

Relation: iris

Instances: 150

Attributes: 5

Attributes

No.	Name
1	sepallength
2	sepalwidth
3	petallength
4	petalwidth
5	class

Selected attribute

Name: sepallength

Type: Numeric

Missing: 0 (0%)

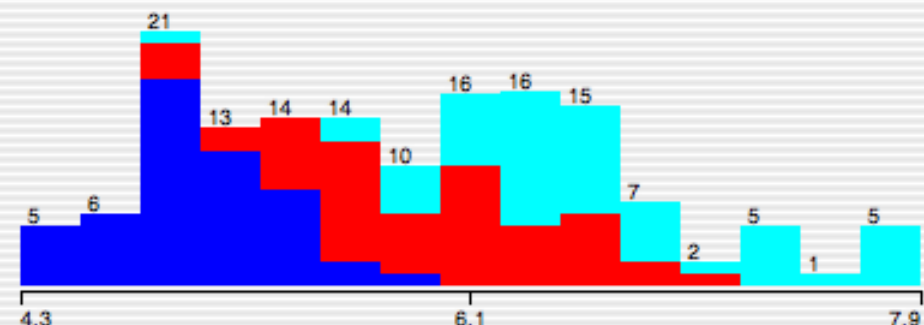
Distinct: 35

Unique: 9 (6%)

Statistic	Value
Minimum	4.3
Maximum	7.9
Mean	5.843
StdDev	0.828

Colour: class (Nom)

Visualize All

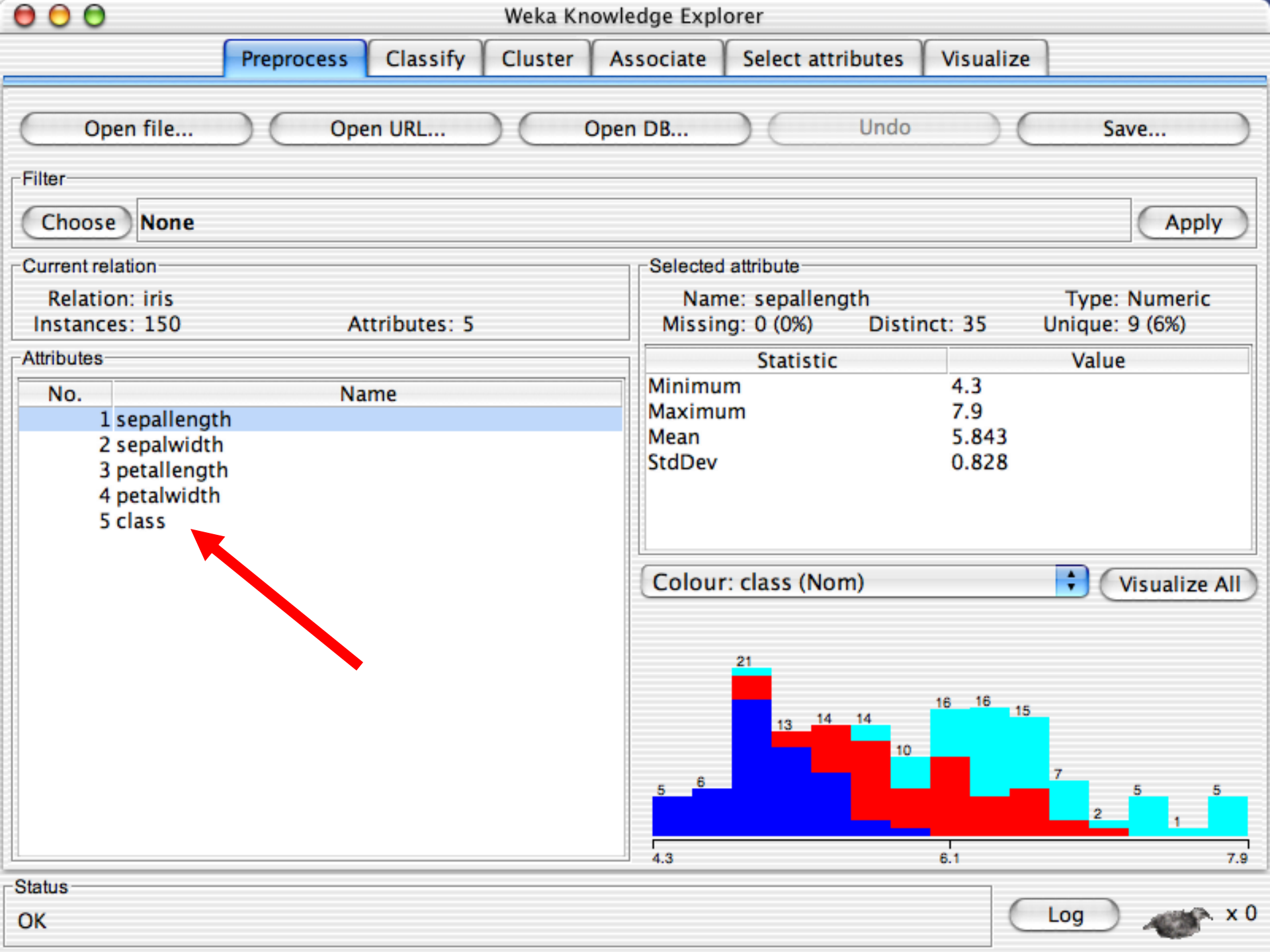


Status

OK

Log

 x 0





Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

None

Apply

Current relation

Relation: iris

Instances: 150

Attributes: 5

Attributes

No.	Name
1	sepal.length
2	sepal.width
3	petal.length
4	petal.width
5	class

Selected attribute

Name: class

Missing: 0 (0%)

Distinct: 3

Type: Nominal

Unique: 0 (0%)

Label	Count
Iris-setosa	50
Iris-versicolor	50
Iris-virginica	50

Colour: class (Nom)

Visualize All

50



50



50



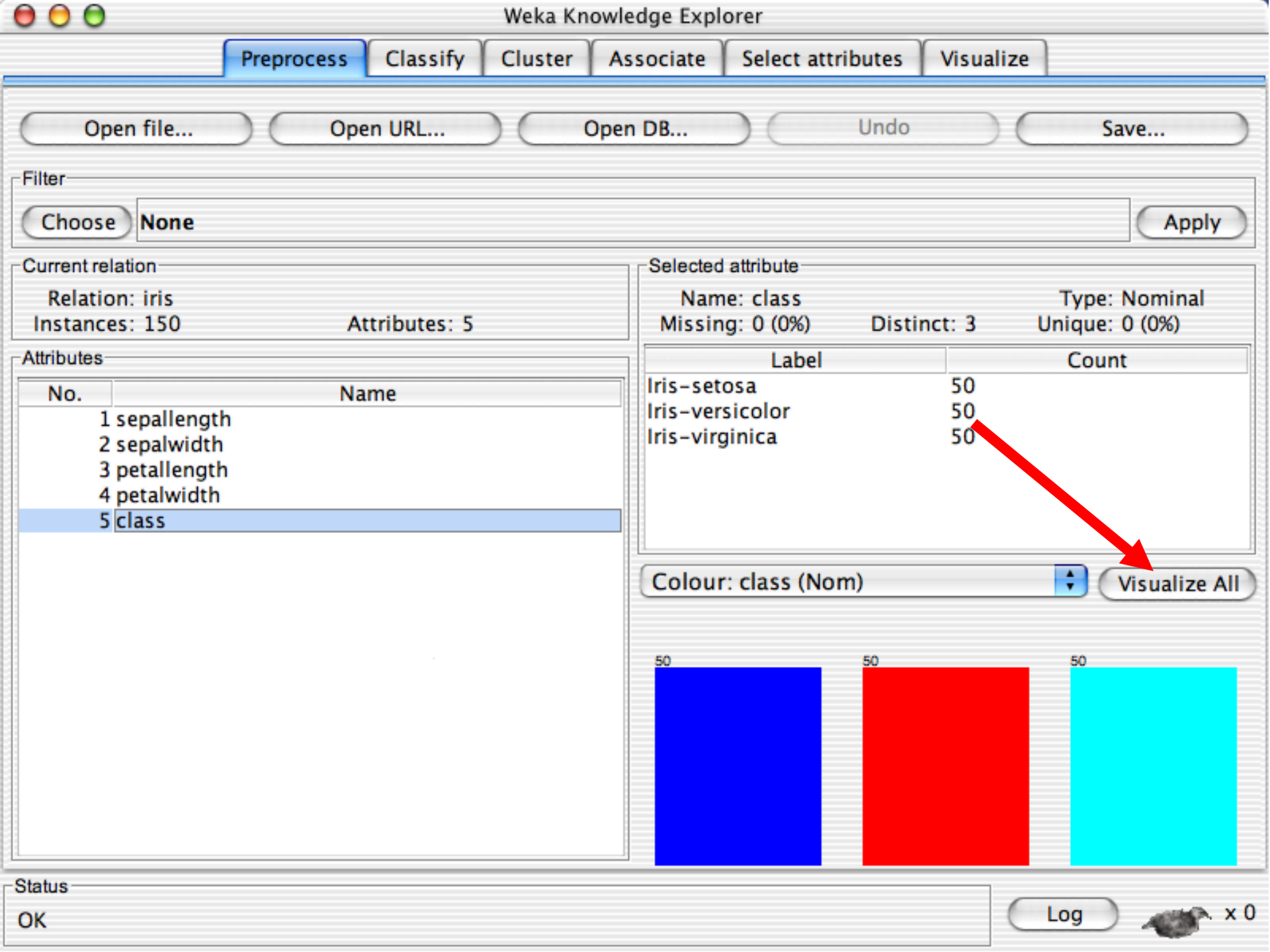
Status

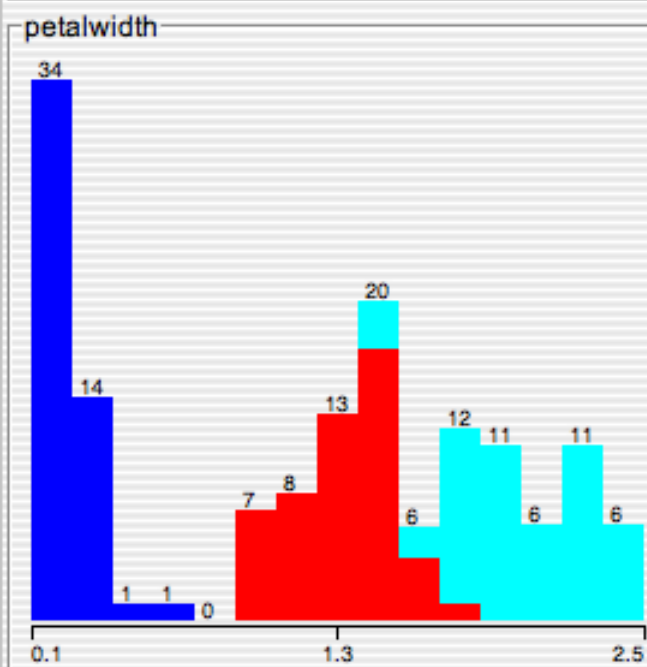
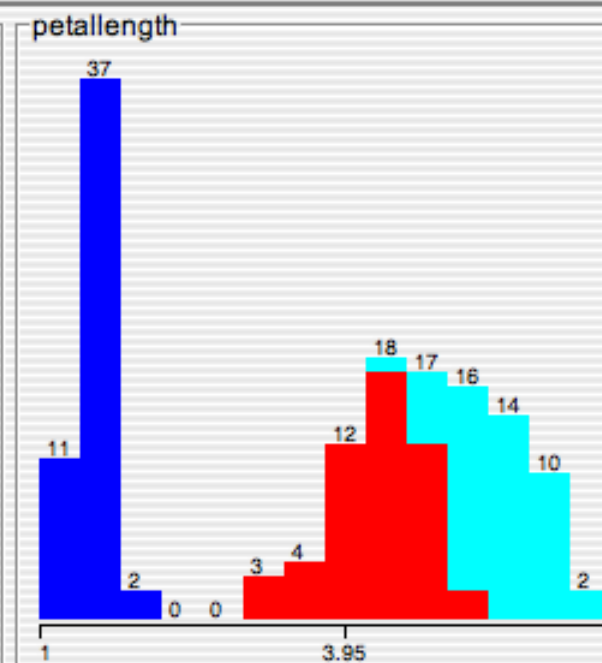
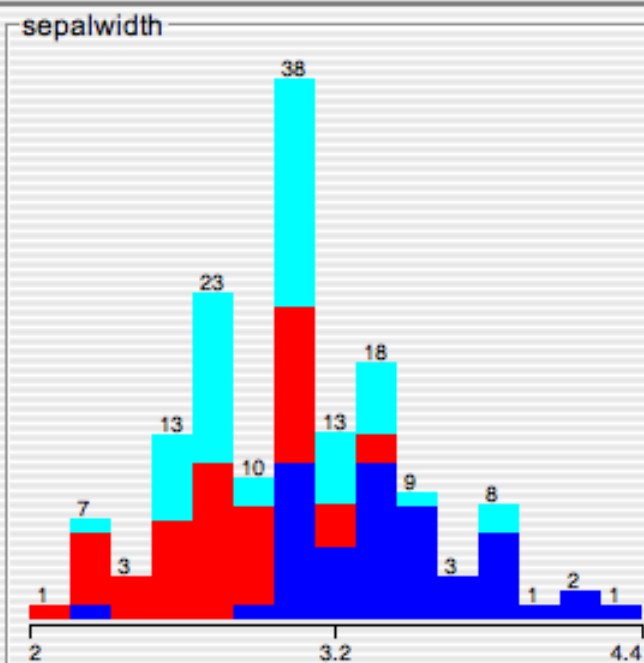
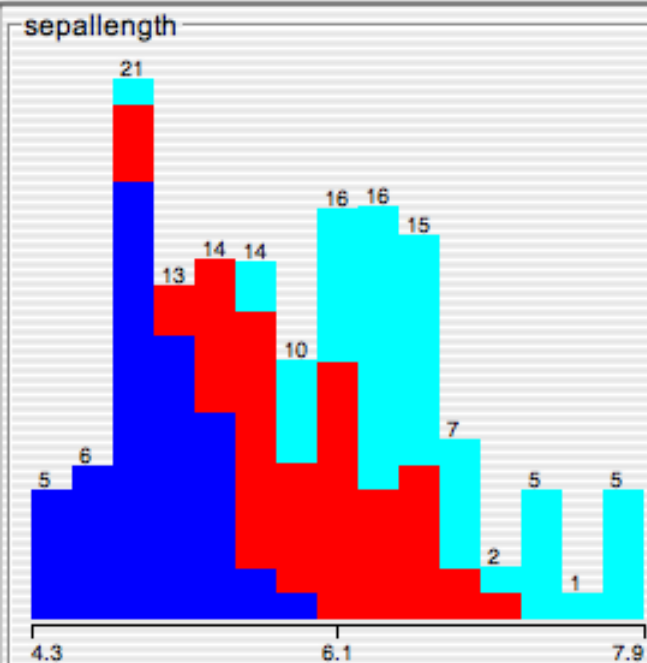
OK

Log



x 0







Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

None

Apply

Current relation

Relation: iris

Instances: 150

Attributes: 5

Attributes

No.	Name
1	sepal.length
2	sepal.width
3	petal.length
4	petal.width
5	class

Selected attribute

Name: petal.length

Type: Numeric

Missing: 0 (0%)

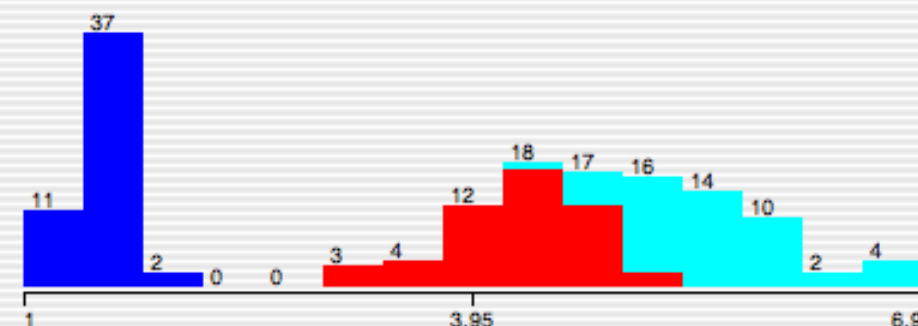
Distinct: 43

Unique: 10 (7%)

Statistic	Value
Minimum	1
Maximum	6.9
Mean	3.759
StdDev	1.764

Colour: class (Nom)

Visualize All

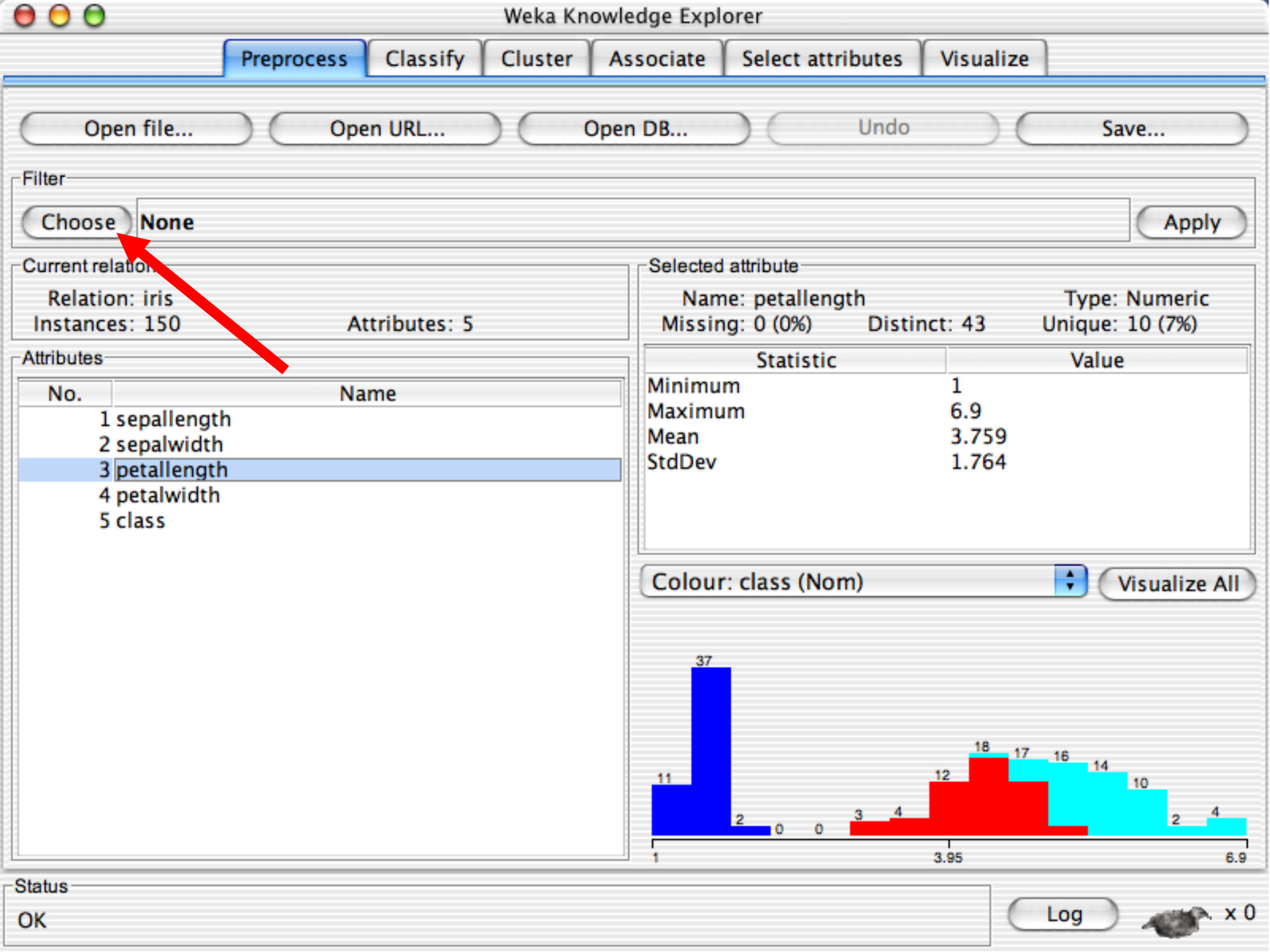


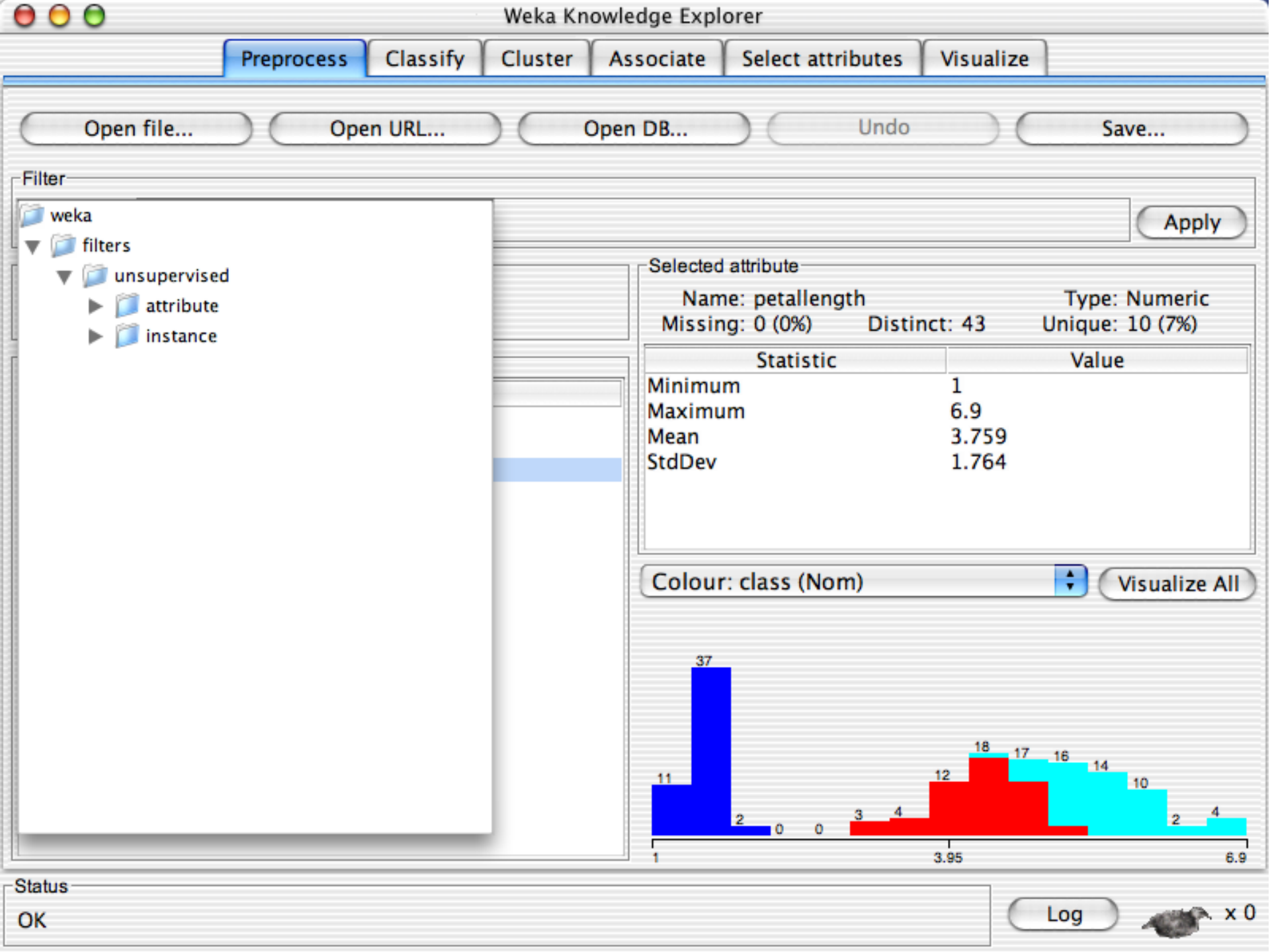
Status

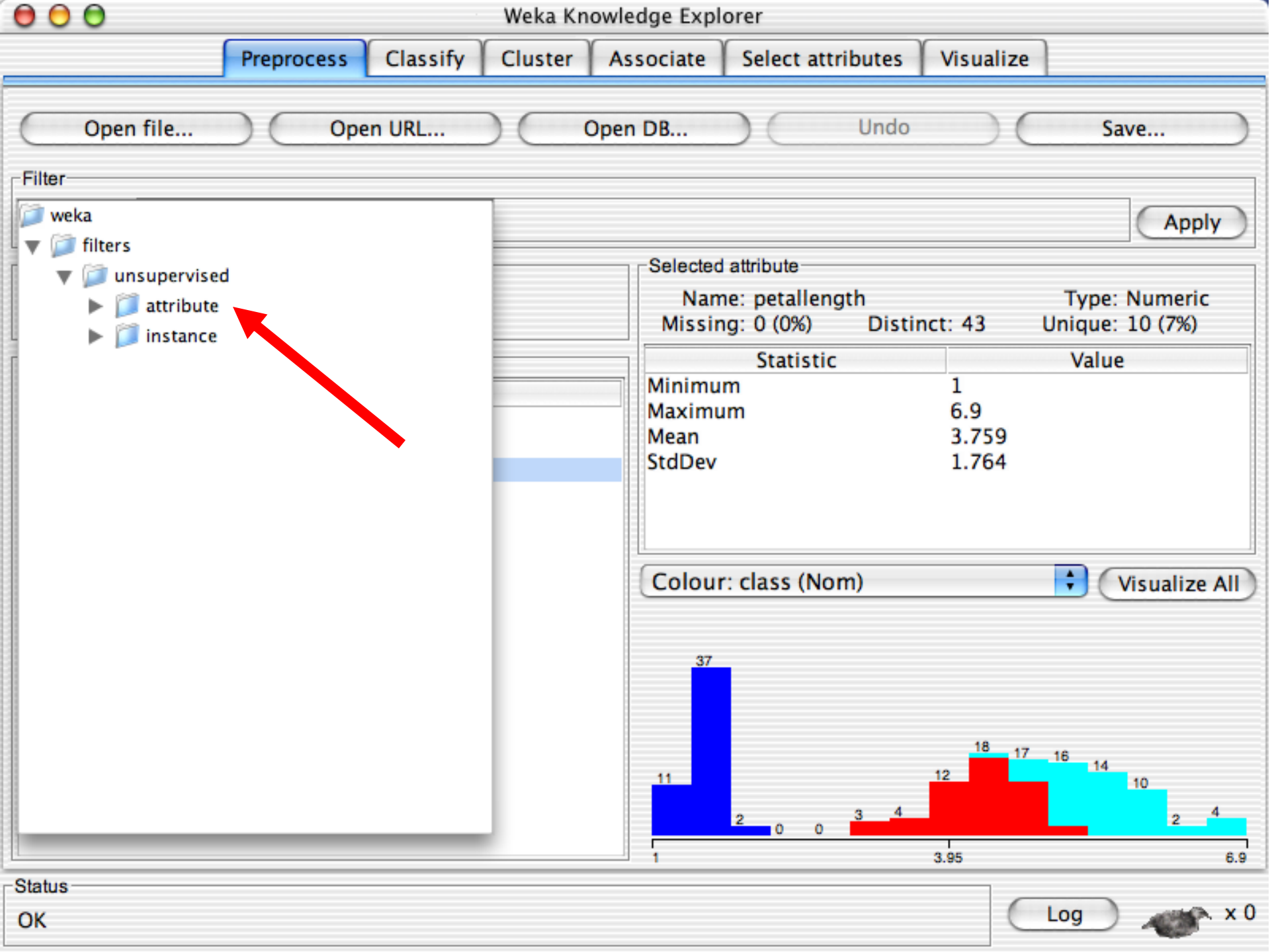
OK

Log

 x 0









Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

- weka
 - filters
 - unsupervised
 - attribute
 - Add
 - AddCluster
 - AddExpression
 - AddNoise
 - Copy
 - Discretize
 - FirstOrder
 - MakeIndicator
 - MergeTwoValues
 - NominalToBinary
 - Normalize
 - NumericToBinary
 - NumericTransform
 - Obfuscate
 - PKIDiscretize
 - Remove
 - RemoveType

Apply

Selected attribute

Name: petallength

Type: Numeric

Missing: 0 (0%)

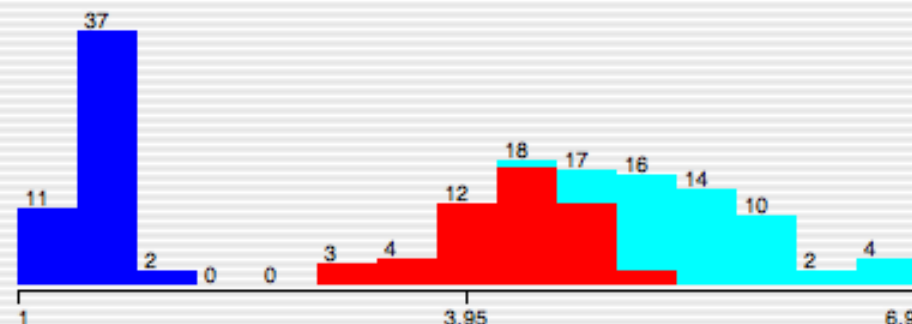
Distinct: 43

Unique: 10 (7%)

Statistic	Value
Minimum	1
Maximum	6.9
Mean	3.759
StdDev	1.764

Colour: class (Nom)

Visualize All



Status

OK

Log

x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

Discretize -B 10 -R first-last

Apply

Current relation

Relation: iris

Instances: 150

Attributes: 5

Attributes

No.	Name
1	sepal.length
2	sepal.width
3	petal.length
4	petal.width
5	class

Selected attribute

Name: petal.length

Type: Numeric

Missing: 0 (0%)

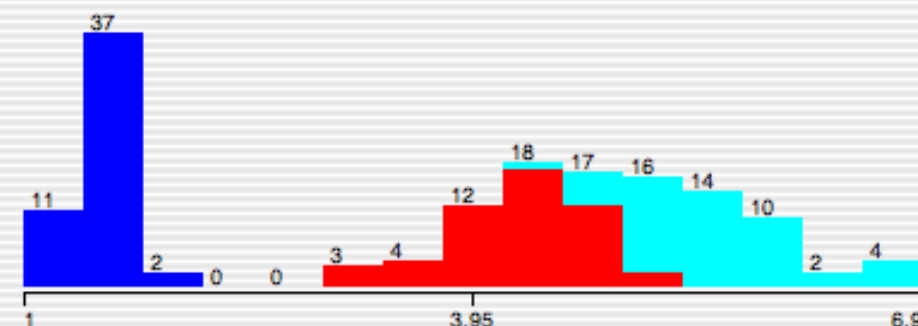
Distinct: 43

Unique: 10 (7%)

Statistic	Value
Minimum	1
Maximum	6.9
Mean	3.759
StdDev	1.764

Colour: class (Nom)

Visualize All



Status

OK

Log

x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

Discretize -B 10 -R first-last

Apply

Current relation

Relation: iris

Instances: 150

Attributes: 5

Attributes

No.	Name
1	sepal.length
2	sepal.width
3	petal.length
4	petal.width
5	class

Selected attribute

Name: petal.length

Type: Numeric

Missing: 0 (0%)

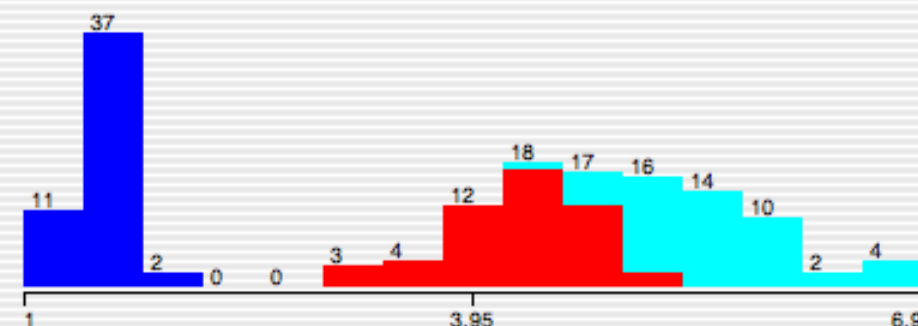
Distinct: 43

Unique: 10 (7%)

Statistic	Value
Minimum	1
Maximum	6.9
Mean	3.759
StdDev	1.764

Colour: class (Nom)

Visualize All

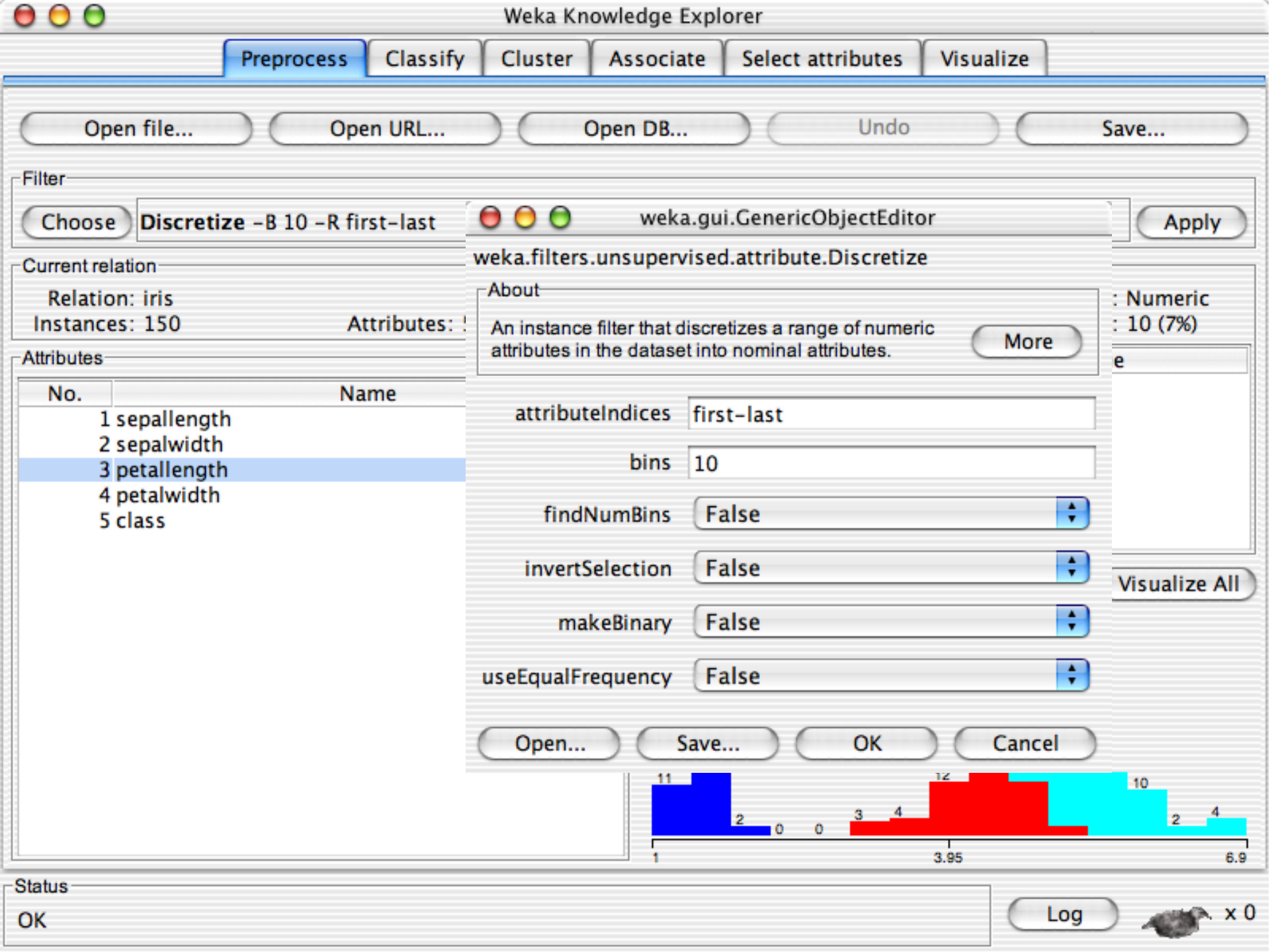


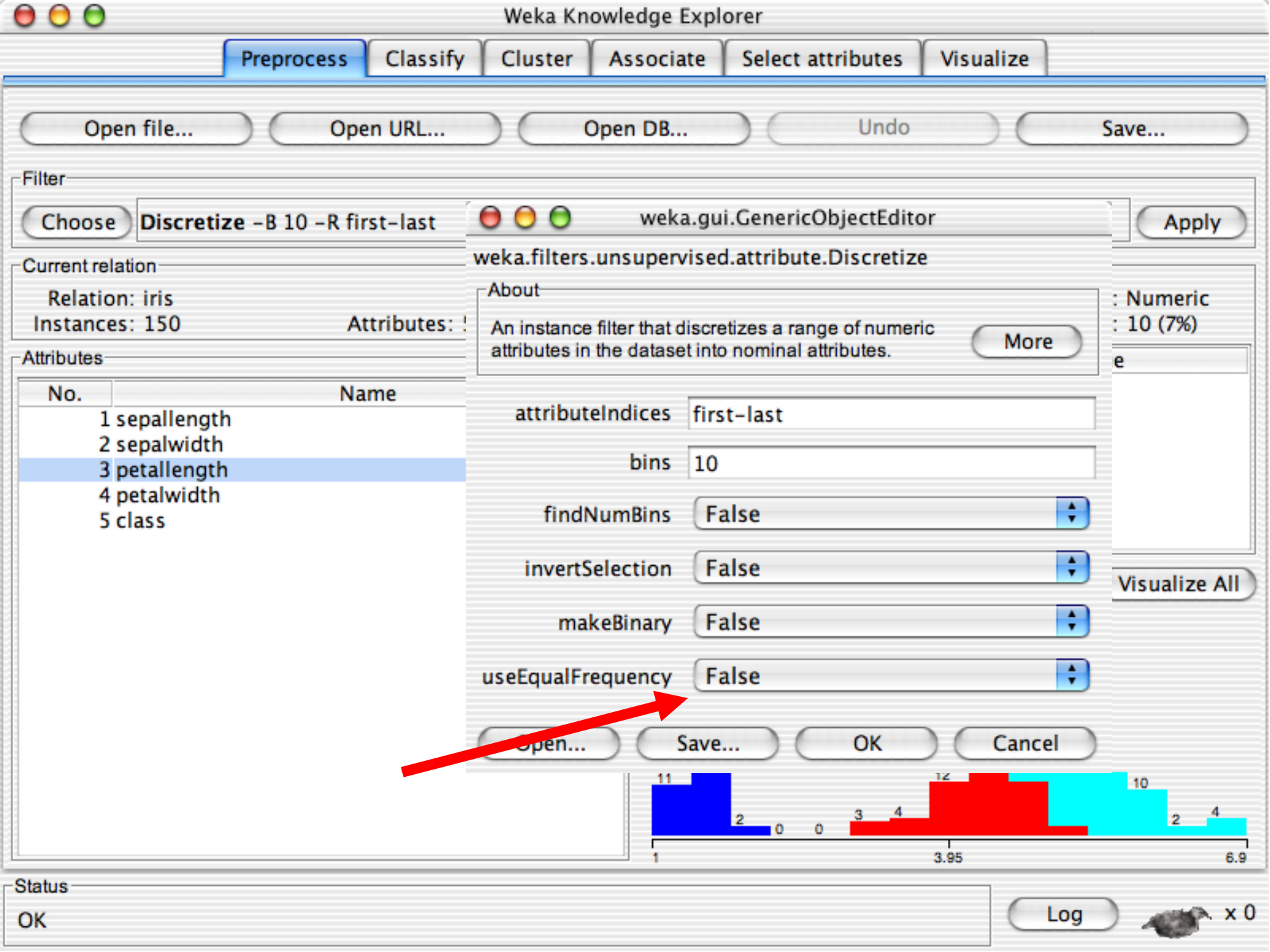
Status

OK

Log

 x 0





attributeIndices first-last

bins 10

findNumBins False

invertSelection False

makeBinary False

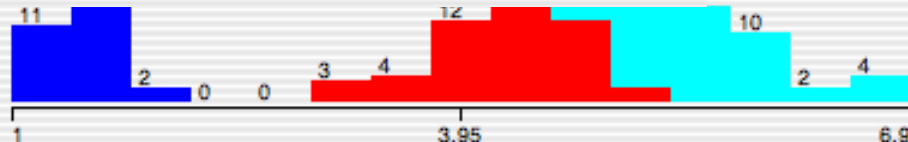
useEqualFrequency False

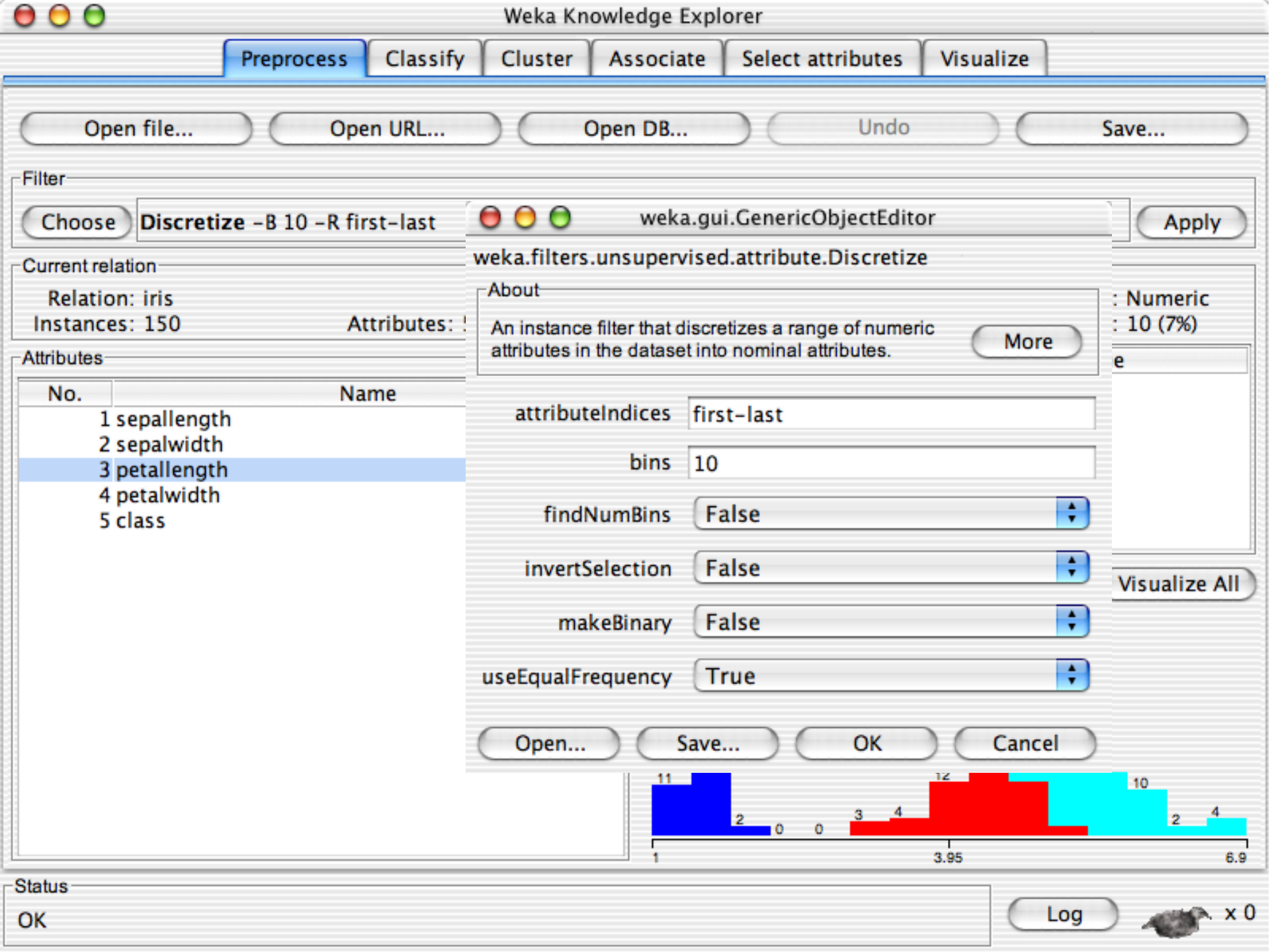
Open...

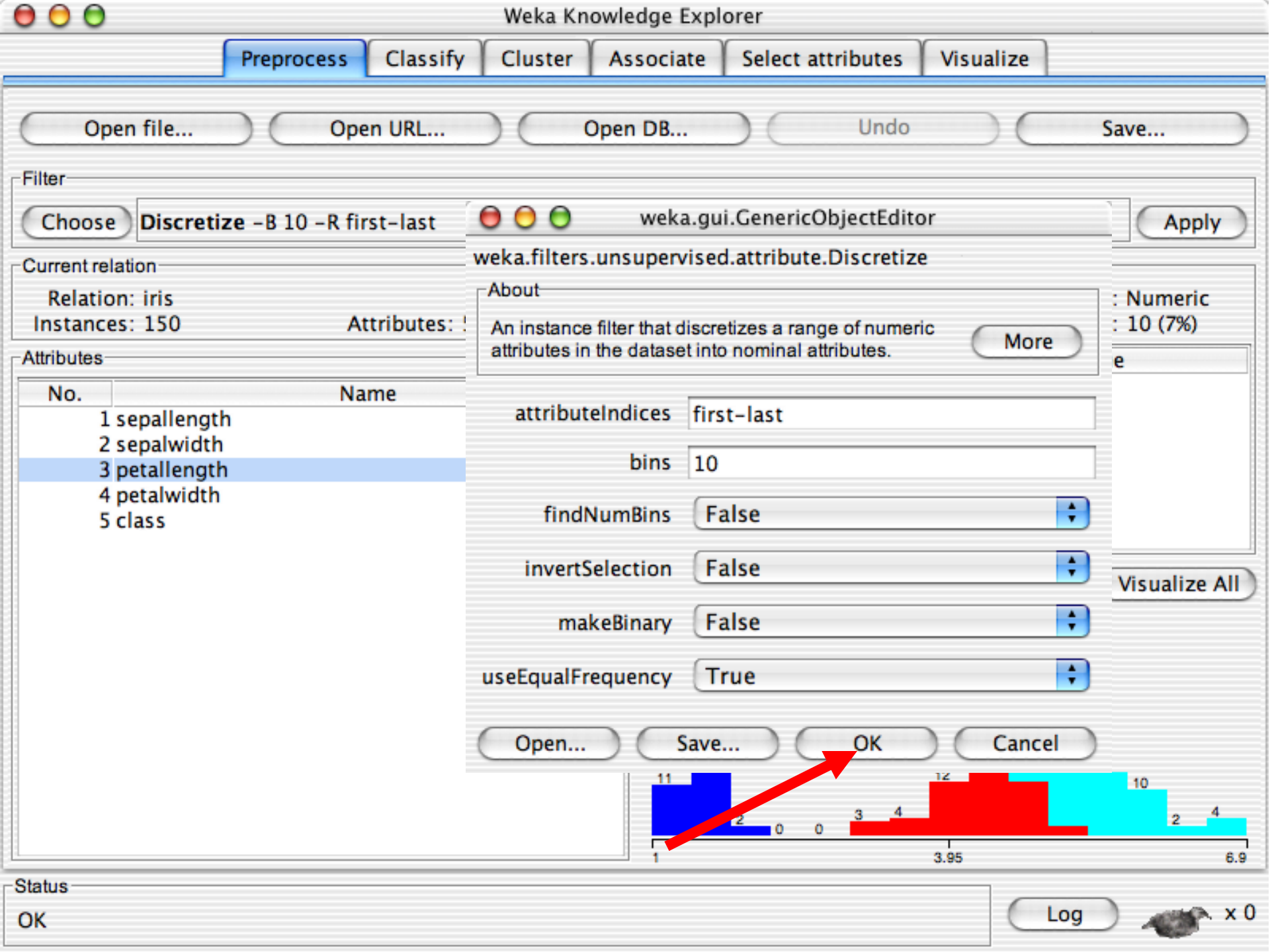
Save...

OK

Cancel









Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

Discretize -F -B 10 -R first-last

Apply

Current relation

Relation: iris

Instances: 150

Attributes: 5

Attributes

No.	Name
1	sepal.length
2	sepal.width
3	petal.length
4	petal.width
5	class

Selected attribute

Name: petal.length

Type: Numeric

Missing: 0 (0%)

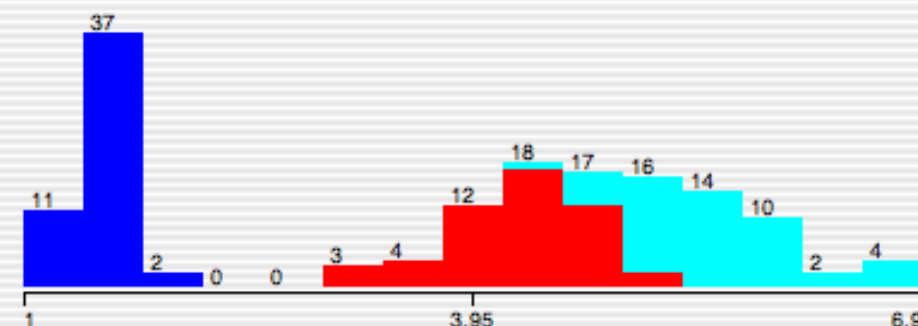
Distinct: 43

Unique: 10 (7%)

Statistic	Value
Minimum	1
Maximum	6.9
Mean	3.759
StdDev	1.764

Colour: class (Nom)

Visualize All

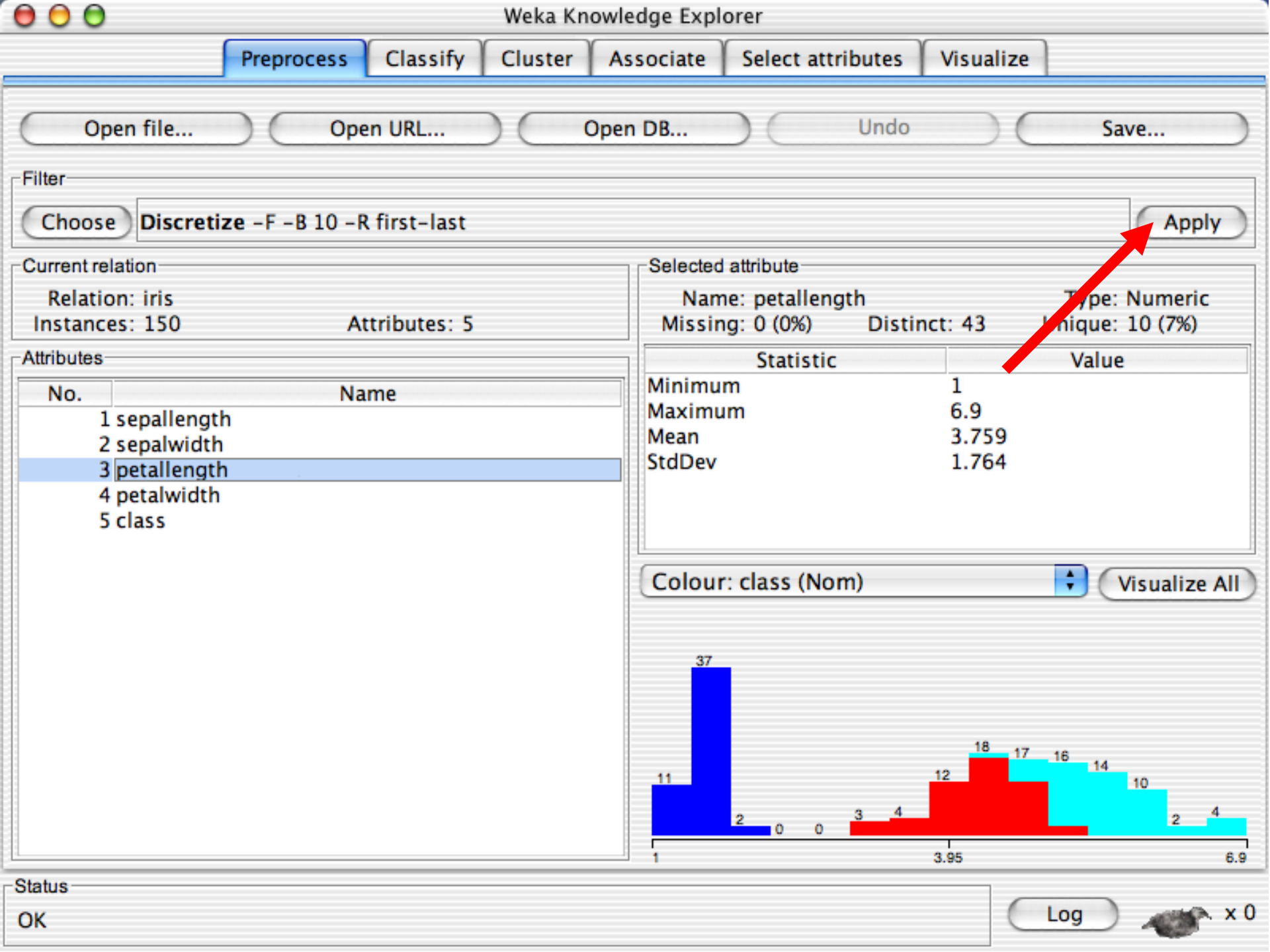


Status

OK

Log

 x 0





Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Open file...

Open URL...

Open DB...

Undo

Save...

Filter

Choose

Discretize -F -B 10 -R first-last

Apply

Current relation

Relation: iris-weka.filters.unsupervised.attribute.Disc...

Instances: 150

Attributes: 5

Attributes

No.	Name
1	sepalength
2	sepalwidth
3	petallength
4	petalwidth
5	class

Selected attribute

Name: petallength

Type: Nominal

Missing: 0 (0%)

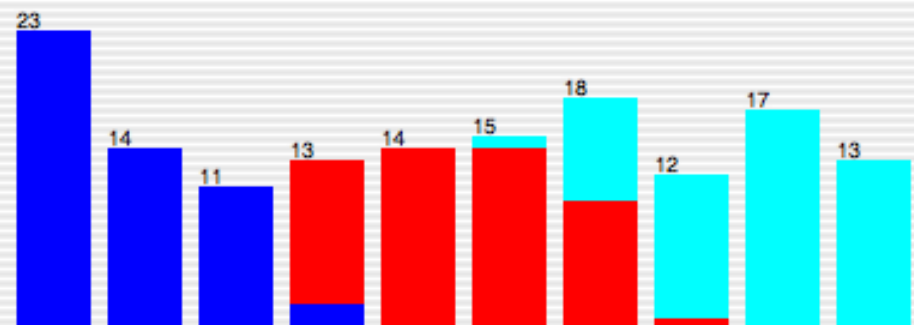
Distinct: 10

Unique: 0 (0%)

Label	Count
'(-inf-1.45]'	23
'(1.45-1.55]'	14
'(1.55-1.8]'	11
'(1.8-3.95]'	13
'(3.95-4.35]'	14
'(4.35-4.65]'	15
'(4.65-5.05]'	18

Colour: class (Nom)

Visualize All



Status

OK

Log

 x 0

Preprocess: filters

- Choosing a filter
 - Supervised vs. Unsupervised
 - Attribute vs. Instance
- Supervised filters “require a class attribute” ;
unsupervised filters do not
- “ meta-filters ” can filter results from clustering and
classification steps

Preprocess: filter examples

	Instance	Attribute
UNsupervised	Resample	Discretize
Supervised	Resample	Discretize

Preprocess: instance filter example

● Resampling

➤ Unsupervised

- ✦ random % of the dataset

➤ Supervised

- ✦ takes the *class distribution* into account when generating a random sample
- ✦ Can add bias towards a specific class value
- ✦ Can specify maximum spread for rare/common class values

Preprocess: attribute filter example

❶ Discretize

➤ Unsupervised

- K-Interval : simplest, can ensure small bin sizes
- Proportional K-Interval : optimized for classification (Naïve Bayes)

➤ Supervised

- Entropy based
 - state of the art
 - computationally expensive
- see *Chapter 7 of Data Mining by I. H. Witten and E. Frank*

Preprocess: attribute filter favorites

- Finding and Discarding variables
 - RemoveUseless : cut using variation threshold
- Datatype Transforms
 - NumericToNominal & NominalToBinary
 - StringToWord : NLP
- Value transforms
 - Normalize
 - ReplaceMissingValues : with mean value from training data
 - AddExpression : any math expression (think R)

WEKA Explorer Tutorial Examples

● Preprocess

- Instance and Attribute Filters (Supervised and Unsupervised)

● Classify

- ZeroR
- Bayes

● Cluster

- Expectation Maximization
- Hierarchical Clustering

● Associate

- Apriori

● Select Attributes

- Via clustering



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

ZeroR

Test options

☐ Use training set☐ Supplied test set

Set...

☒ Cross-validation Folds 10☐ Percentage split % 66

More options...

(Nom) class



Start

Stop

Result list (right-click for options)

Classifier output

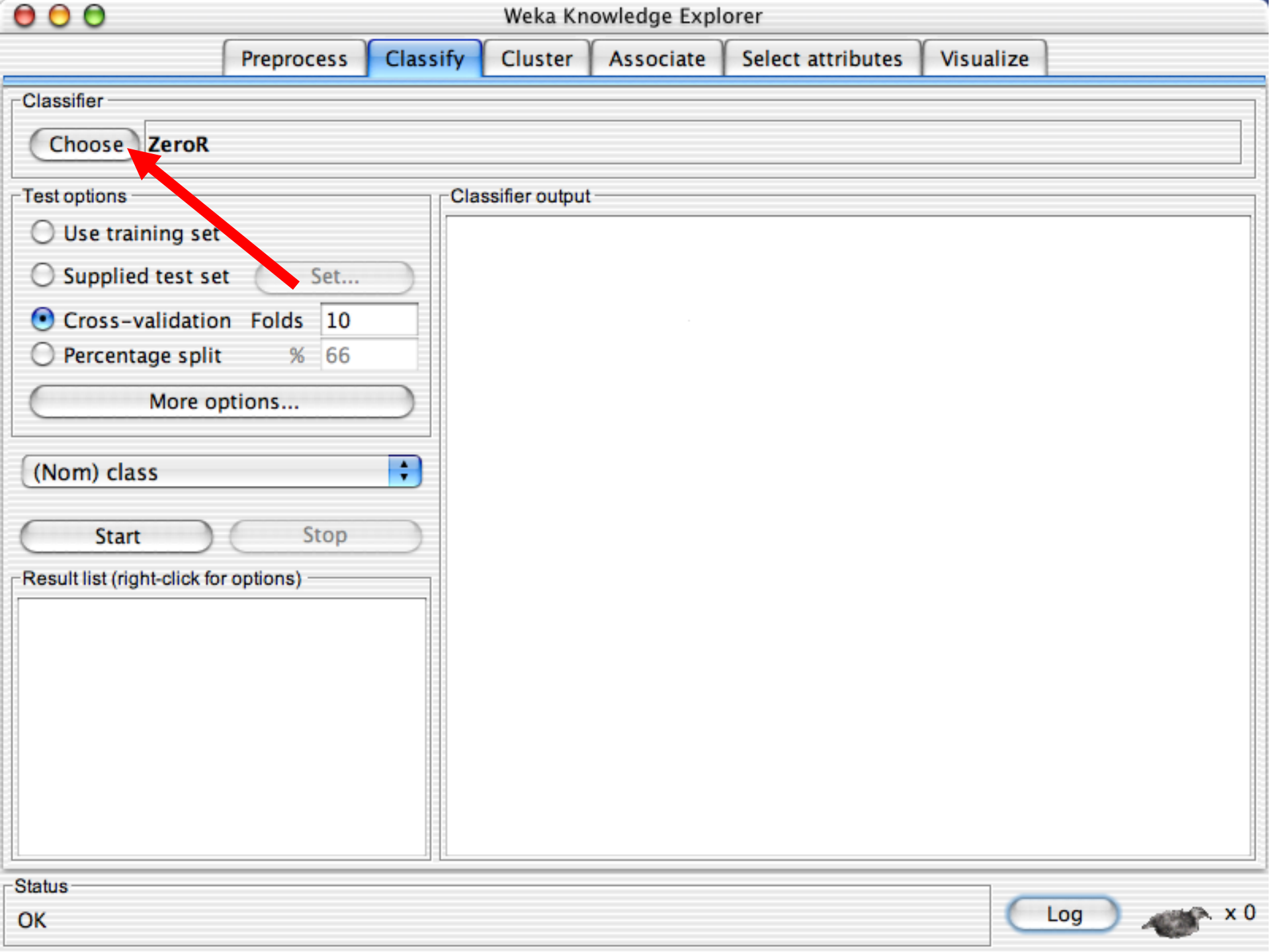
Status

OK

Log



x 0





Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

- weka
 - classifiers
 - bayes
 - functions
 - lazy
 - meta
 - misc
 - trees
 - adtree
 - DecisionStump
 - Id3
 - j48
 - J48
 - lmt
 - m5
 - RandomForest
 - RandomTree
 - REPTree
 - UserClassifier
 - rules

Classifier output

Status

OK

Log



x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☒ Cross-validation Folds 10☐ Percentage split % 66

More options...

(Nom) class



Start

Stop

Result list (right-click for options)

Classifier output

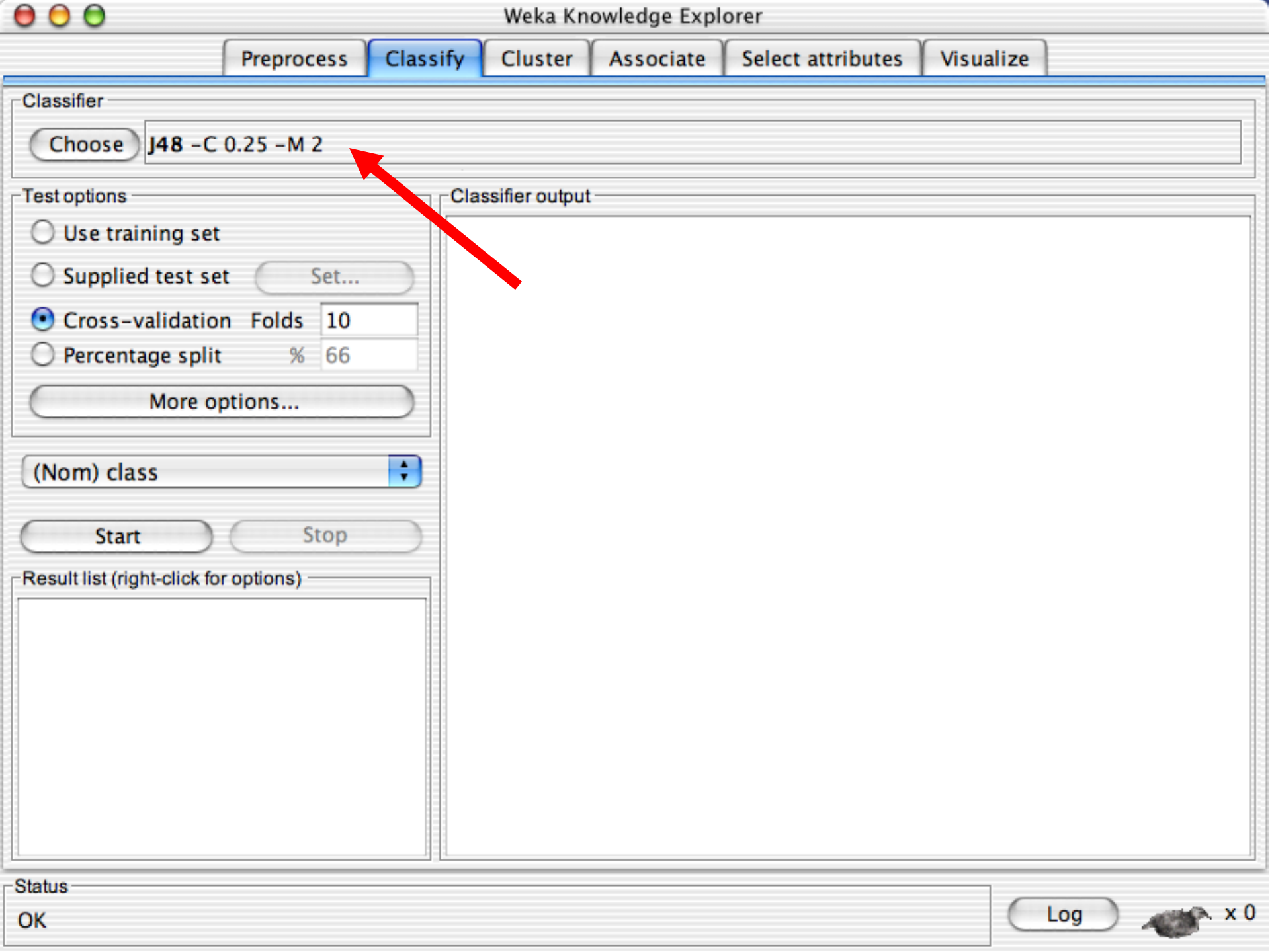
Status

OK

Log



x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set

☐ Supplied test set

Set...

☒ Cross-validation Folds 10

☐ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

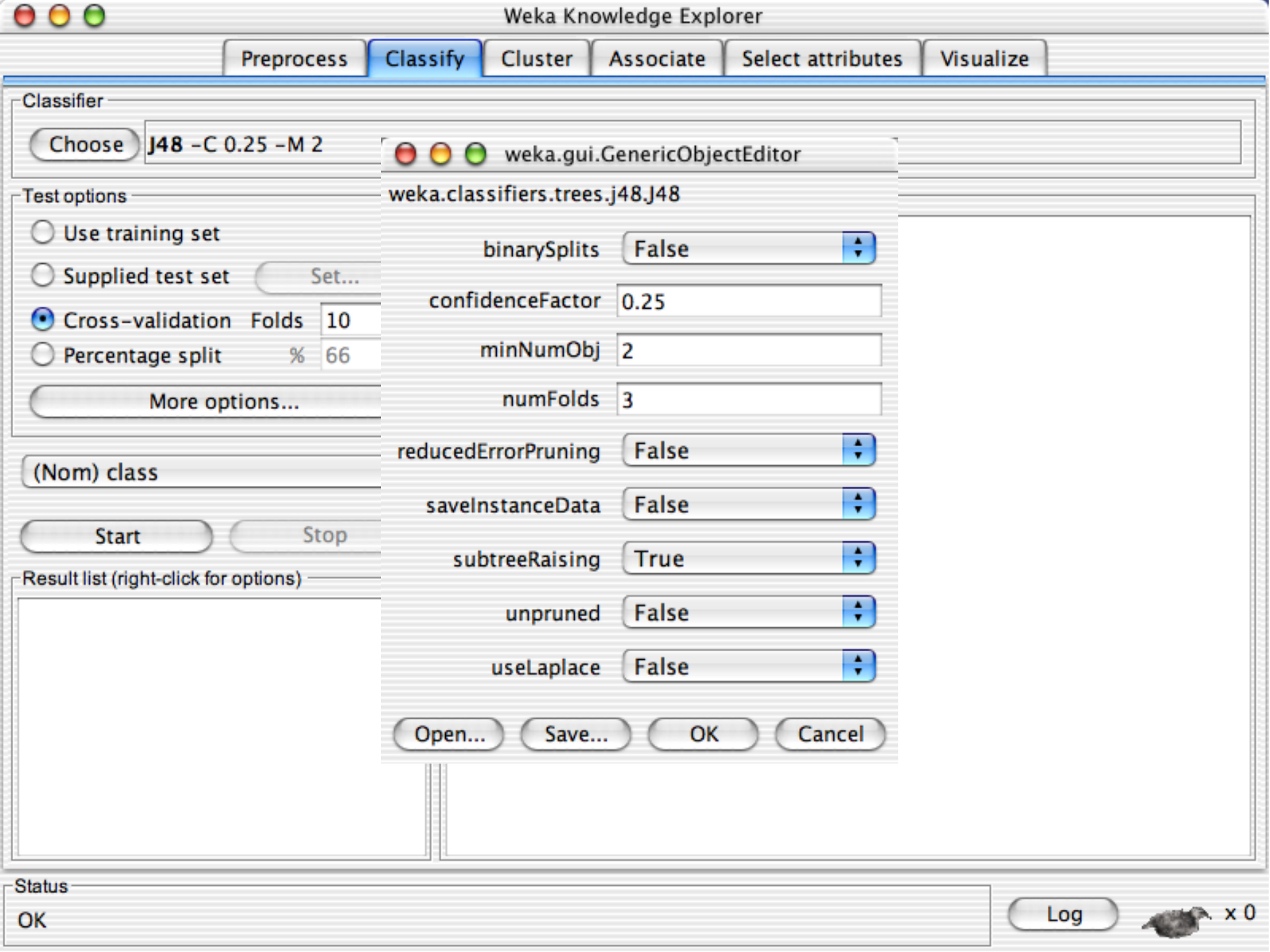
Classifier output

Status

OK

Log

x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☒ Cross-validation Folds 10☐ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)



weka.gui.GenericObjectEditor

weka.classifiers.trees.j48.J48

binarySplits False

confidenceFactor 0.25

minNumObj 2

numFolds 3

reducedErrorPruning False

saveInstanceData False

subtreeRaising True

unpruned False

useLaplace False

Open...

Save...

OK

Cancel

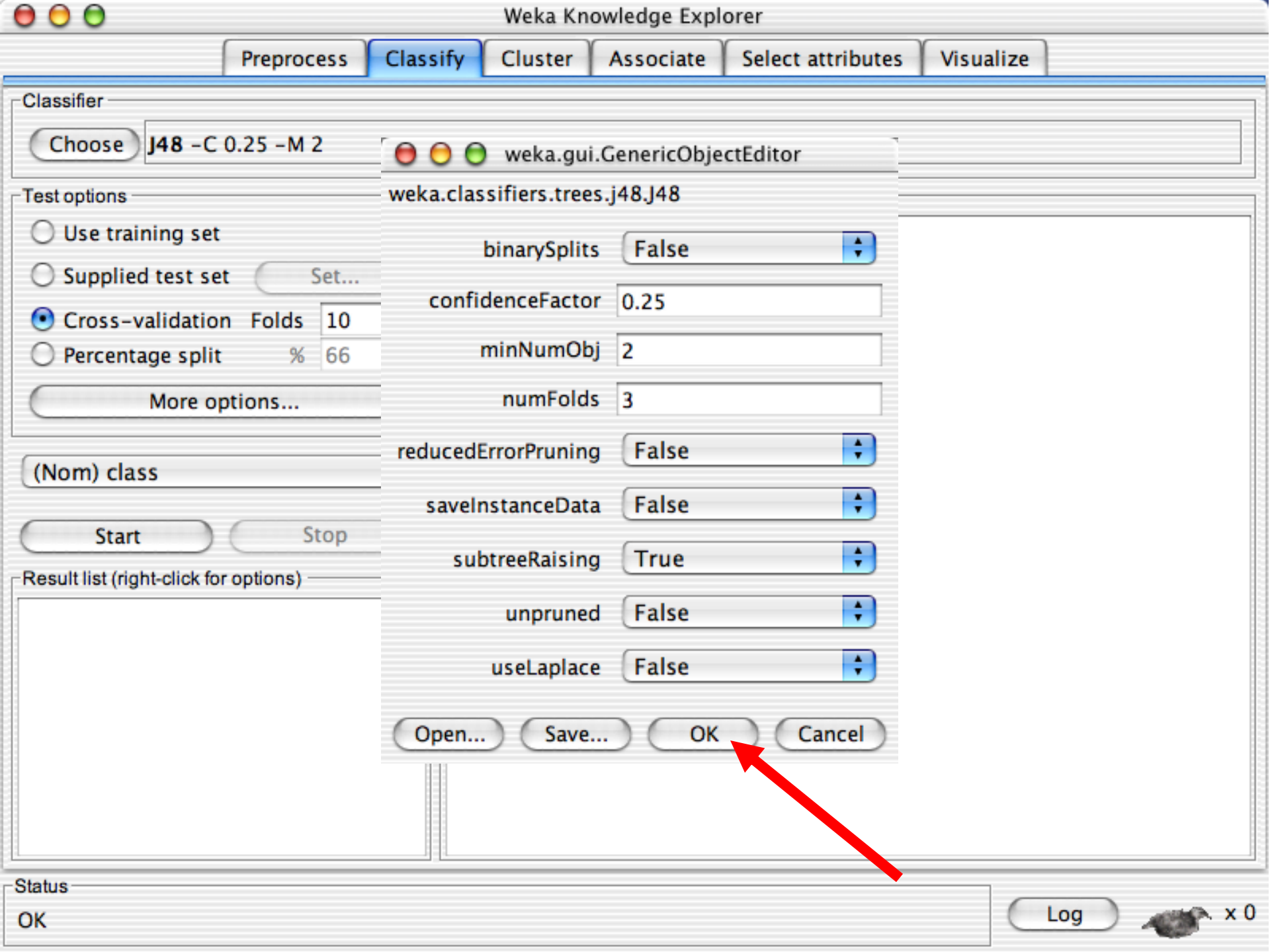
Status

OK

Log



x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set Set...☒ Cross-validation Folds 10☐ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

weka.gui.GenericObjectEditor

weka.classifiers.trees.j48.J48

binarySplits False

confidenceFactor 0.25

minNumObj 2

numFolds 3

reducedErrorPruning False

saveInstanceData False

subtreeRaising True

unpruned False

useLaplace False

Open...

Save...

OK

Cancel

Status

OK

Log

x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☒ Cross-validation Folds 10☐ Percentage split % 66

More options...

(Nom) class



Start

Stop

Result list (right-click for options)

Classifier output

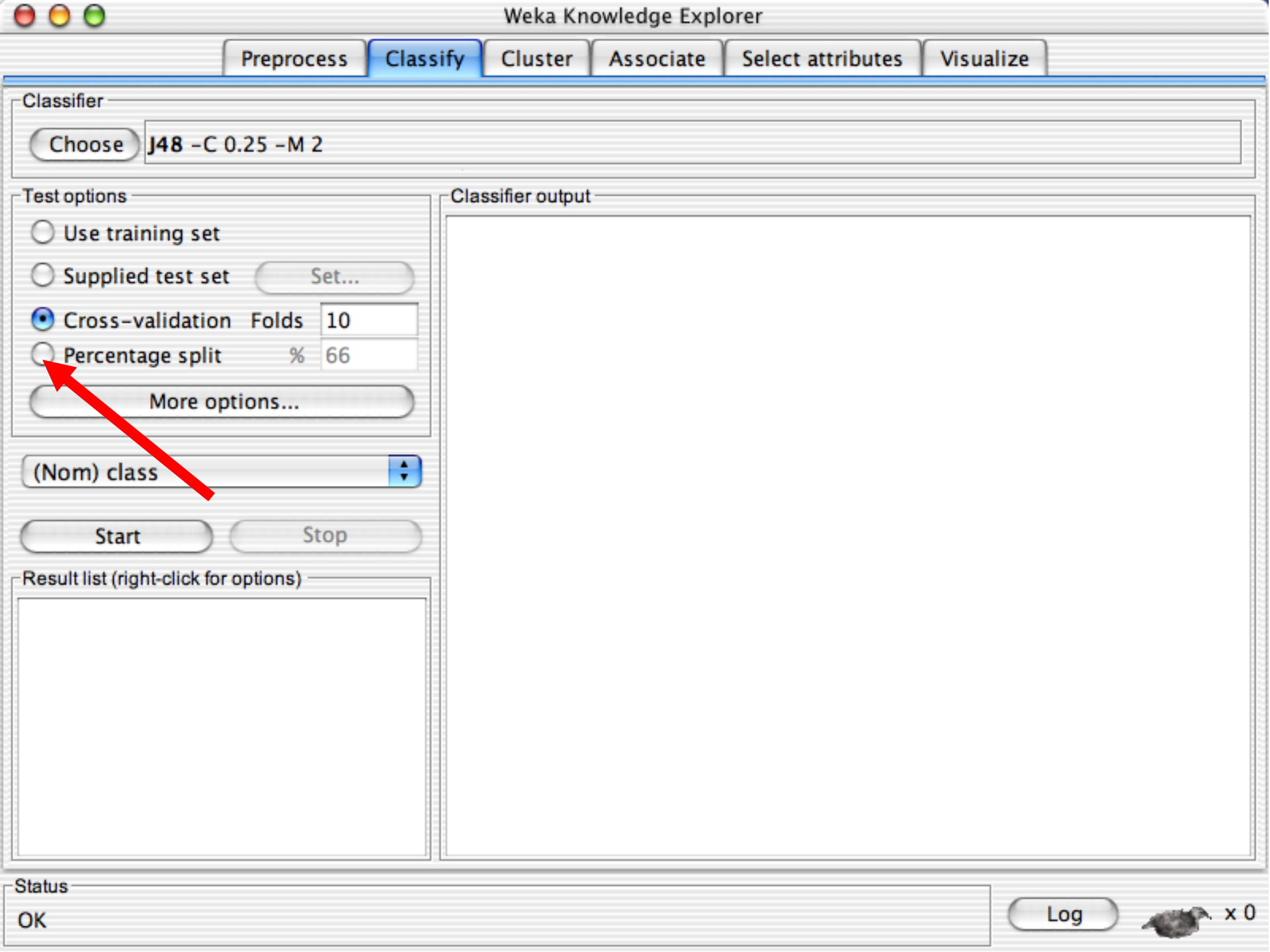
Status

OK

Log



x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

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More options...

(Nom) class

Start

Stop

Result list (right-click for options)

Classifier output

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Log



x 0



Preprocess

Classify

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Select attributes

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Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class



Start

Stop

Result list (right-click for options)

Classifier output

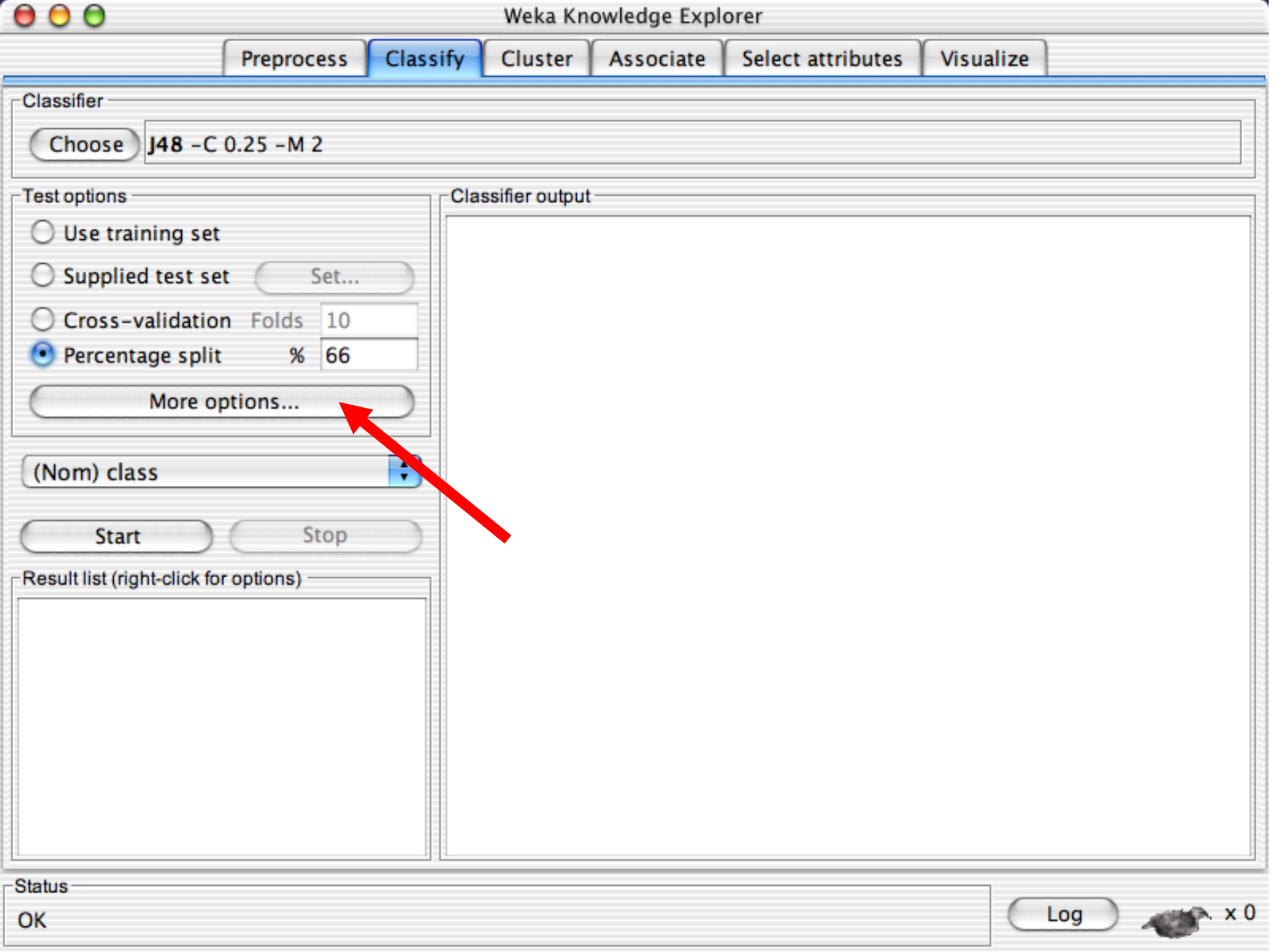
Status

OK

Log



x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set

☐ Supplied test set

Set...

☐ Cross-validation Folds 10

☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

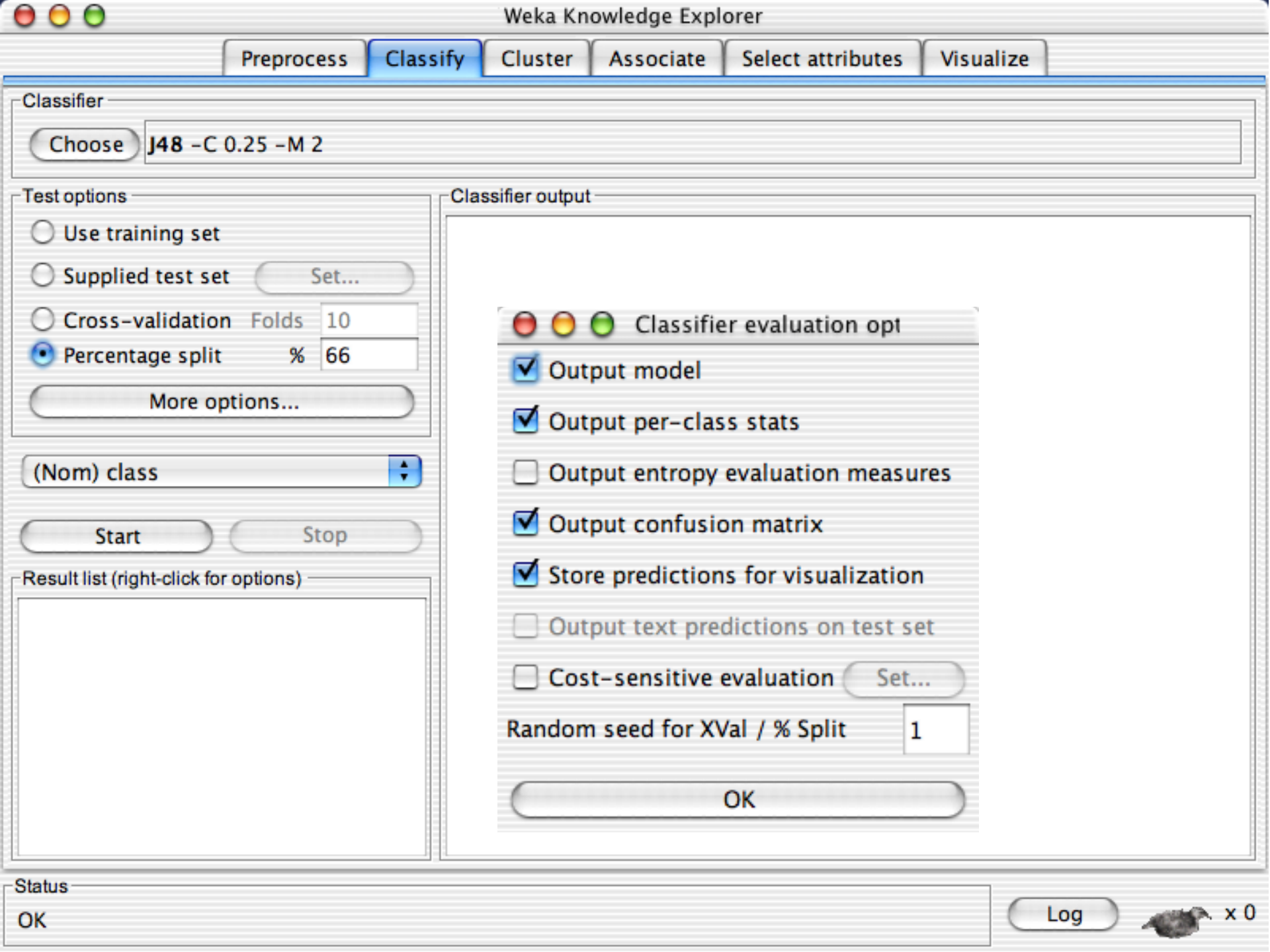
Classifier output

Status

OK

Log

x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

Classifier output



Classifier evaluation opt

☒ Output model☒ Output per-class stats☐ Output entropy evaluation measures☒ Output confusion matrix☒ Store predictions for visualization☐ Output text predictions on test set☐ Cost-sensitive evaluation

Set...

Random seed for XVal / % Split

1

OK

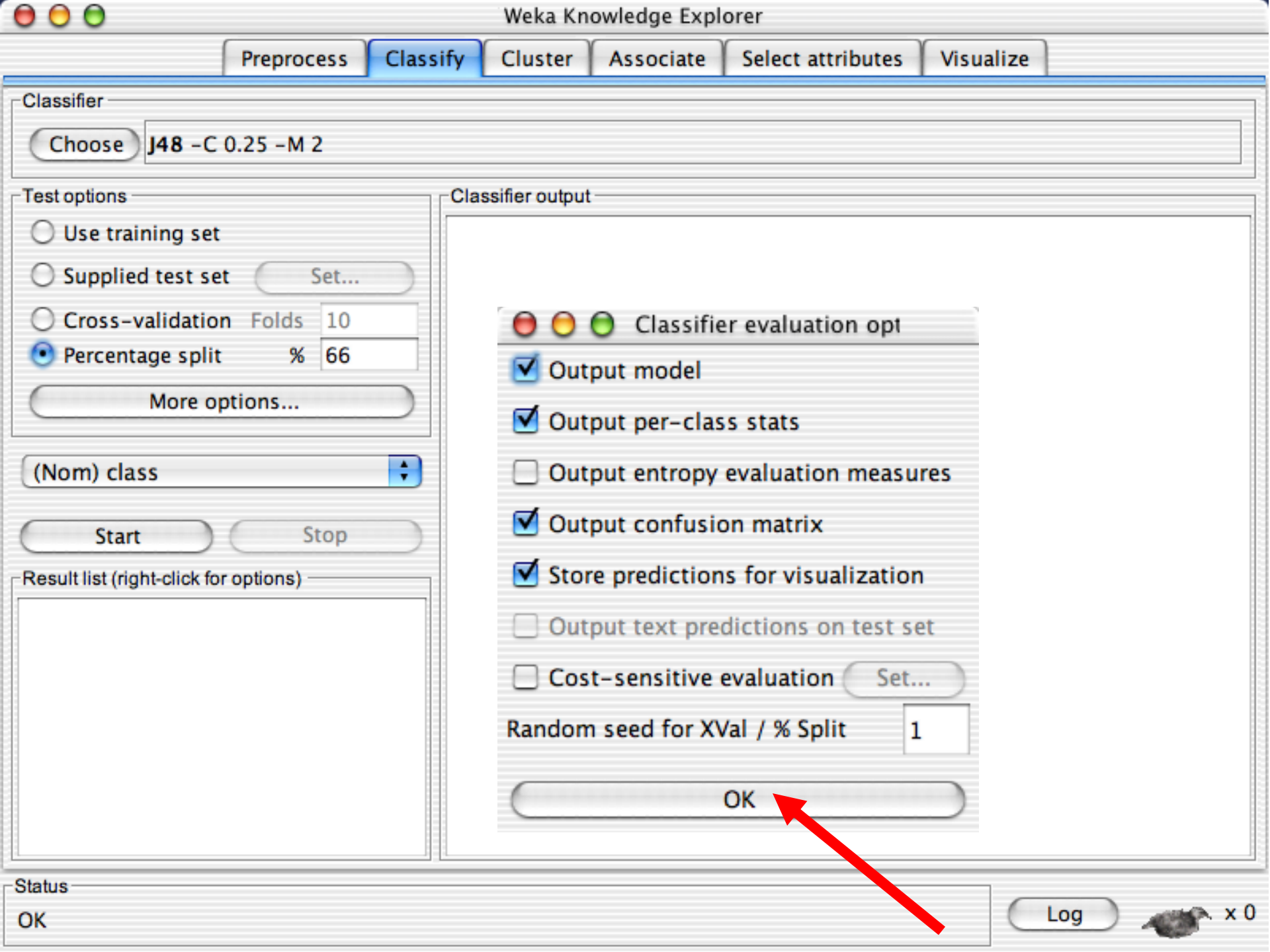
Status

OK

Log



x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set

☐ Supplied test set Set...

☐ Cross-validation Folds 10

☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

Classifier output

Classifier evaluation opt

☒ Output model

☒ Output per-class stats

☐ Output entropy evaluation measures

☒ Output confusion matrix

☒ Store predictions for visualization

☐ Output text predictions on test set

☐ Cost-sensitive evaluation Set...

Random seed for XVal / % Split 1

OK

Status

OK

Log

x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class



Start

Stop

Result list (right-click for options)

Classifier output

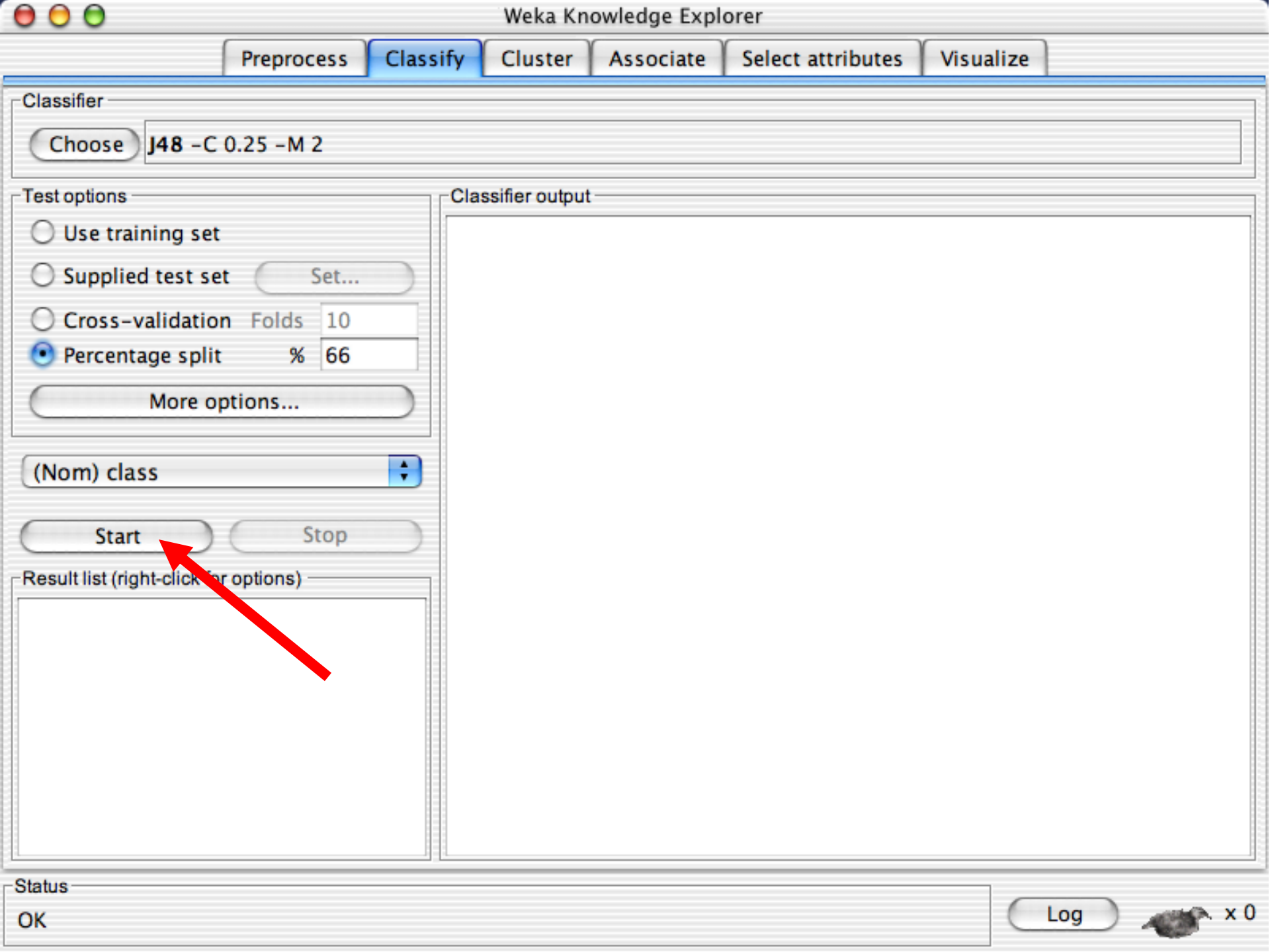
Status

OK

Log



x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set

☐ Supplied test set

Set...

☐ Cross-validation Folds 10

☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

Classifier output

Status

OK

Log



x 0

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set Set...☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

11:49:05 - trees.j48.J48

Classifier output

=== Run information ===

Scheme: weka.classifiers.trees.j48.J48 -C 0.25 -M 2
Relation: iris
Instances: 150
Attributes: 5

sepalength
sepalwidth
petallength
petalwidth
class

Test mode: split 66% train, remainder test

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

J48 pruned tree

```
-----  
petalwidth <= 0.6: Iris-setosa (50.0)  
petalwidth > 0.6  
|   petalwidth <= 1.7  
|   |   petallength <= 4.9: Iris-versicolor (48.0/1.0)  
|   |   petallength > 4.9  
|   |       |   petalwidth <= 1.5: Iris-virginica (3.0)  
|   |       |   petalwidth > 1.5: Iris-versicolor (3.0/1.0)  
|   |   petalwidth > 1.7: Iris-virginica (46.0/1.0)
```

Number of Leaves : 5

Status

OK

Log

x 0

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set Set...☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

11:49:05 - trees.j48.J48

Classifier output

=== Run information ===

Scheme: weka.classifiers.trees.j48.J48 -C 0.25 -M 2
Relation: iris
Instances: 150
Attributes: 5

sepalength
sepalwidth
petallength
petalwidth
class

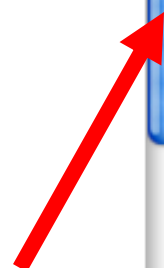
Test mode: split 66% train, remainder test

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

J48 pruned tree

```
-----  
petalwidth <= 0.6: Iris-setosa (50.0)  
petalwidth > 0.6  
| petalwidth <= 1.7  
| | petallength <= 4.9: Iris-versicolor (48.0/1.0)  
| | petallength > 4.9  
| | | petalwidth <= 1.5: Iris-virginica (3.0)  
| | | petalwidth > 1.5: Iris-versicolor (3.0/1.0)  
| petalwidth > 1.7: Iris-virginica (46.0/1.0)
```

Number of Leaves : 5



Status

OK

Log

 x 0

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set Set...☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

11:49:05 - trees.j48.J48

Classifier output

Time taken to build model: 0.24 seconds

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	49	96.0784 %
Incorrectly Classified Instances	2	3.9216 %
Kappa statistic	0.9408	
Mean absolute error	0.0396	
Root mean squared error	0.1579	
Relative absolute error	8.8979 %	
Root relative squared error	33.4091 %	
Total Number of Instances	51	

=== Detailed Accuracy By Class ===

TP Rate	FP Rate	Precision	Recall	F-Measure	Class
1	0	1	1	1	Iris-setosa
1	0.063	0.905	1	0.95	Iris-versicolor
0.882	0	1	0.882	0.938	Iris-virginica

=== Confusion Matrix ===

a	b	c	<-- classified as
15	0	0	a = Iris-setosa
0	19	0	b = Iris-versicolor
0	2	15	c = Iris-virginica

Status

OK

Log

x 0

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose

J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

11:49:05 - trees.j48.J48

Classifier output

Time taken to build model: 0.24 seconds

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	49	96.0784 %
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TP Rate	FP Rate	Precision	Recall	F-Measure	Class
1	0	1	1	1	Iris-setosa
1	0.063	0.905	1	0.95	Iris-versicolor
0.882	0	1	0.882	0.938	Iris-virginica

=== Confusion Matrix ===

a	b	c	<-- classified as
15	0	0	a = Iris-setosa
0	19	0	b = Iris-versicolor
0	2	15	c = Iris-virginica

Status

OK

Log

x 0



Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

11:49:05 - trees.j48.J48

View in main window

View in separate window

Save result buffer

Load model

Save model

Re-evaluate model on current test set

Visualize classifier errors

Visualize tree

Visualize margin curve

Visualize threshold curve

Visualize cost curve

Classifier output

Time taken to build model: 0.24 seconds

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	49	96.0784 %
Incorrectly Classified Instances	2	3.9216 %
Kappa statistic	0.9408	
Mean absolute error	0.0396	
Root mean squared error	0.1579	
Relative absolute error	8.8979 %	
Root relative squared error	33.4091 %	
Total Number of Instances	51	

=== Detailed Accuracy By Class ===

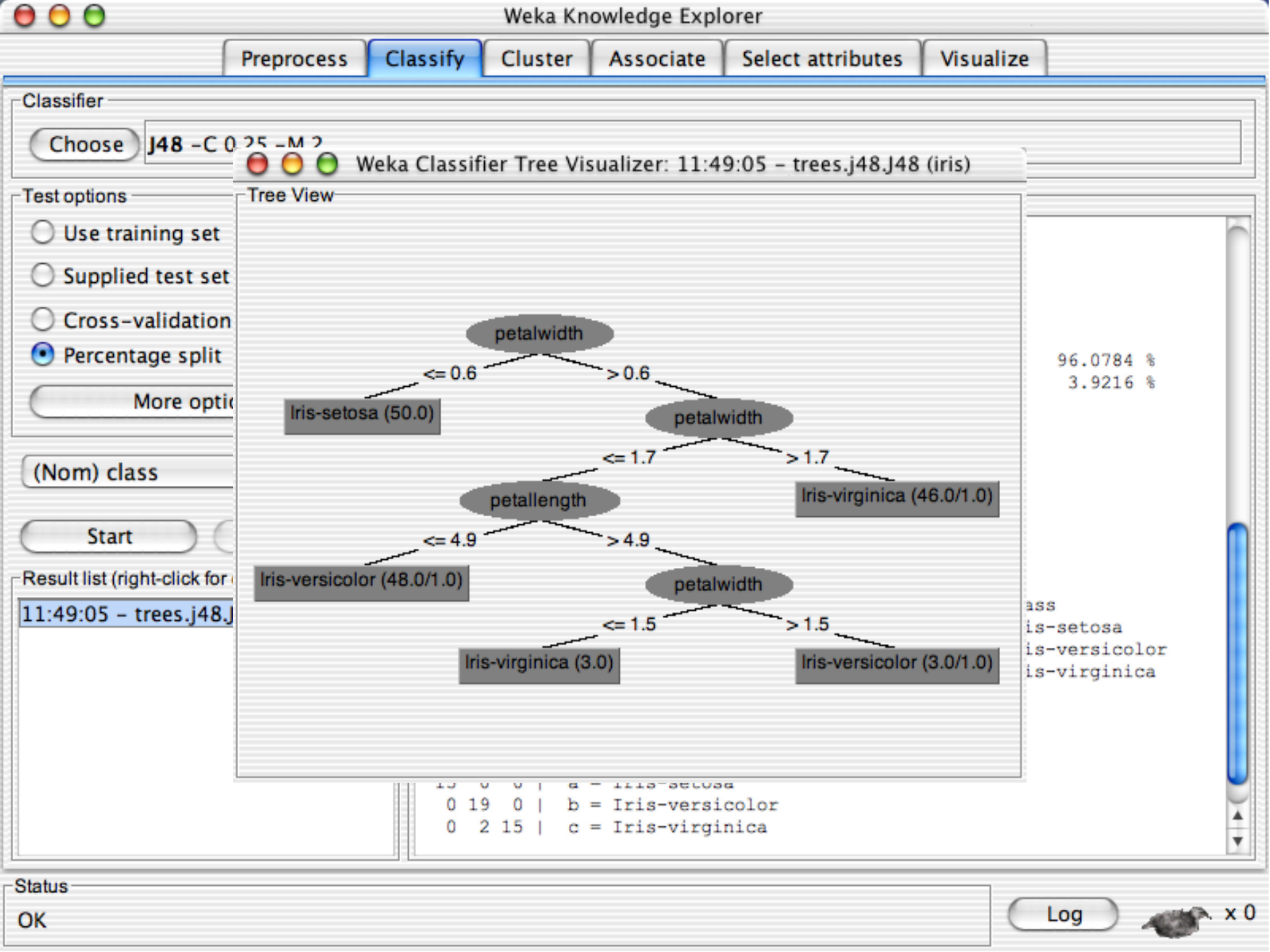
Recall	F-Measure	Class
1	1	Iris-setosa
1	0.95	Iris-versicolor
0.882	0.938	Iris-virginica

Status

OK

Log

 x 0





Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set Set...☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

11:49:05 - trees.j48.J48

View in main window

View in separate window

Save result buffer

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Classifier output

Time taken to build model: 0.24 seconds

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	49	96.0784 %
Incorrectly Classified Instances	2	3.9216 %
Kappa statistic	0.9408	
Mean absolute error	0.0396	
Root mean squared error	0.1579	
Relative absolute error	8.8979 %	
Root relative squared error	33.4091 %	
Total Number of Instances	51	

=== Detailed Accuracy By Class ===

Recall	F-Measure	Class
1	1	Iris-setosa
1	0.95	Iris-versicolor
0.882	0.938	Iris-virginica

lor
ca

Status

OK

Log

x 0

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

- ☐ Use training set
☐ Supplied test set
☐ Cross-validation
☒ Percentage split

More options

(Nom) class

Start

Result list (right-click for)

11:49:05 - trees.j48.J

Weka Classifier Visualize: 11:49:05 - trees.j48.J48 (iris)

X: petallength (Num)

Y: petalwidth (Num)

Colour: class (Nom)

Select Instance

Reset

Clear

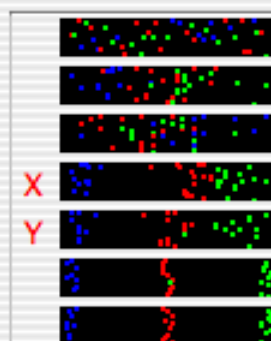
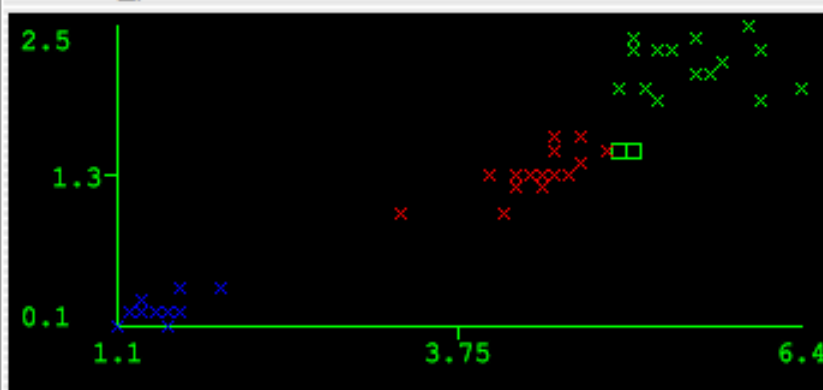
Save

Jitter

96.0784 %

3.9216 %

Plot: iris_predicted



Class colour

Iris-setosa Iris-versicolor Iris-virginica

```

0 15 0 | D = Iris-versicolor
0 2 15 | c = Iris-virginica

```

class
 is-setosa
 is-versicolor
 is-virginica

Status

OK

Log

x 0

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Classifier

Choose J48 -C 0.25 -M 2

Test options

☐ Use training set☐ Supplied test set

Set...

☐ Cross-validation Folds 10☒ Percentage split % 66

More options...

(Nom) class

Start

Stop

Result list (right-click for options)

11:49:05 - trees.j48.J48

Classifier output

Time taken to build model: 0.24 seconds

=== Evaluation on test split ===

=== Summary ===

Correctly Classified Instances	49	96.0784 %
Incorrectly Classified Instances	2	3.9216 %
Kappa statistic	0.9408	
Mean absolute error	0.0396	
Root mean squared error	0.1579	
Relative absolute error	8.8979 %	
Root relative squared error	33.4091 %	
Total Number of Instances	51	

=== Detailed Accuracy By Class ===

TP Rate	FP Rate	Precision	Recall	F-Measure	Class
1	0	1	1	1	Iris-setosa
1	0.063	0.905	1	0.95	Iris-versicolor
0.882	0	1	0.882	0.938	Iris-virginica

=== Confusion Matrix ===

a	b	c	<-- classified as
15	0	0	a = Iris-setosa
0	19	0	b = Iris-versicolor
0	2	15	c = Iris-virginica

Status

OK

Log

x 0

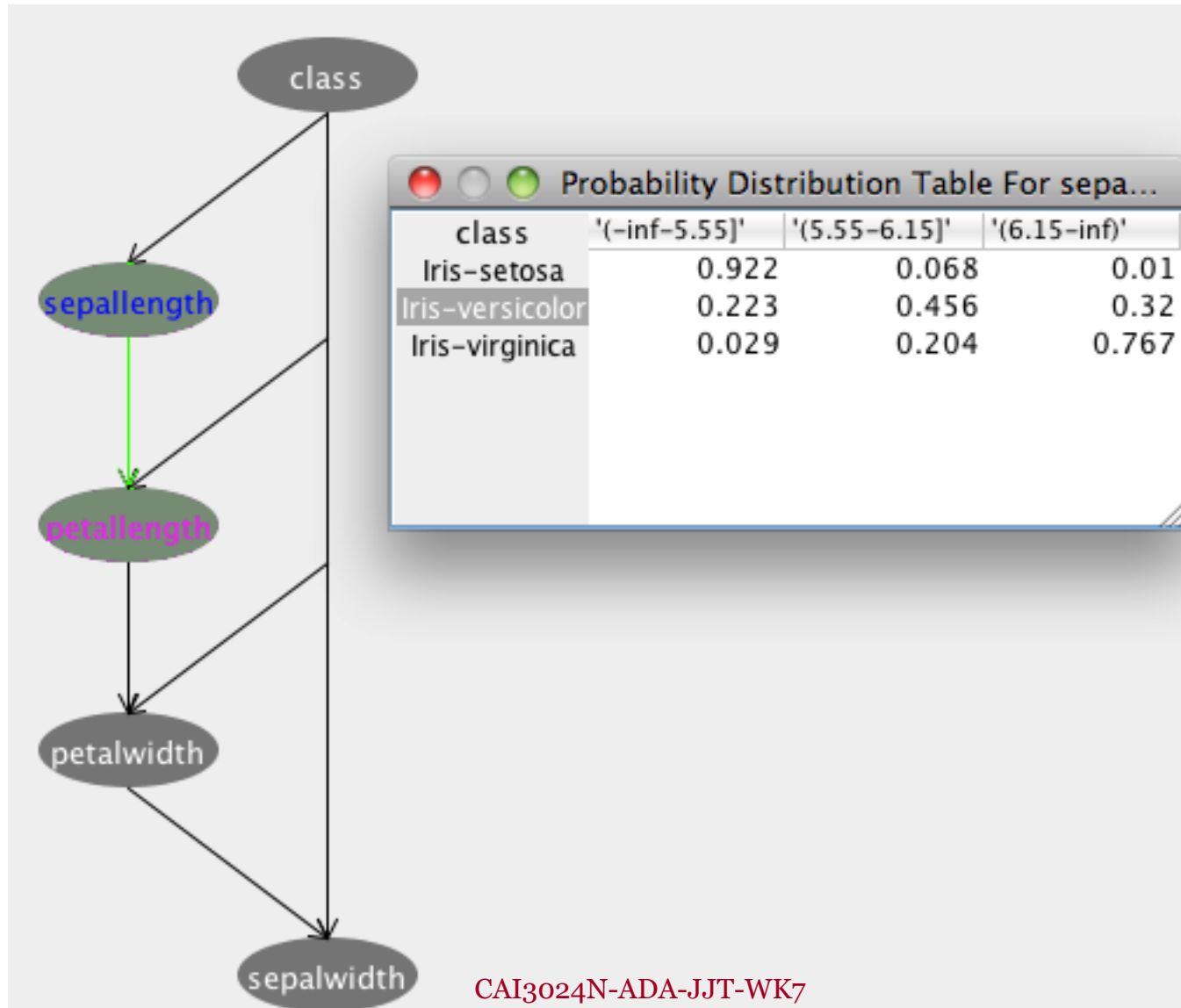
Classify : Bayes

Various Bayesian network classifier learning algorithms are implemented in Weka [12]. This note provides some user documentation and implementation details.

Summary of main capabilities:

- Structure learning of Bayesian networks using various hill climbing (K2, B, etc) and general purpose (simulated annealing, tabu search) algorithms.
- Local score metrics implemented; Bayes, BDe, MDL, entropy, AIC.
- Global score metrics implemented; leave one out cv, k-fold cv and cumulative cv.
- Conditional independence based causal recovery algorithm available.
- Parameter estimation using direct estimates and Bayesian model averaging.
- GUI for easy inspection of Bayesian networks.
- Part of Weka allowing systematic experiments to compare Bayes net performance with general purpose classifiers like C4.5, nearest neighbor, support vector, etc.

Classify: Learning a Bayes Network Structure



Classify using Naïve Bayes

- Naïve Bayes (standard)
- Multinomial Naïve Bayes (text classification)
- Hidden Naïve Bayes, others,

Classify using Naïve Bayes

1. Pick Discretization method* (hardest part)
2. Pick a class to predict
3. Run the classifier

The screenshot shows the Orange3 software interface with the 'Classify' tab selected. The 'Classifier' dropdown is set to 'NaiveBayes'. Under 'Test options', 'Cross-validation' is selected with 10 folds. The 'Classifier output' panel displays the following results:

=== Stratified cross-validation ===
=== Summary ===

Correctly Classified Instances	142	94.6667 %
Incorrectly Classified Instances	8	5.3333 %
Kappa statistic	0.92	
Mean absolute error	0.0532	
Root mean squared error	0.1744	
Relative absolute error	11.9604 %	
Root relative squared error	37.0028 %	
Total Number of Instances	150	

=== Detailed Accuracy By Class ===

	TP Rate	FP Rate	Precision	Recall	F-Measure	ROC Area	Class
	1	0	1	1	1	1	Iris-setosa
	0.94	0.05	0.904	0.94	0.922	0.983	Iris-versicolor
	0.9	0.03	0.938	0.9	0.918	0.982	Iris-virginica
Weighted Avg.	0.947	0.027	0.947	0.947	0.947	0.988	

The 'Result list' at the bottom shows a single entry: '03:18:42 - bayes.NaiveBayes'.

Classify Text using Naïve Bayes Multinomial

- Framingham dataset contains text descriptions for each of the 21k+ variables
- I wrote a parallelized NLP program to calculate inverse word frequencies and score variable pairs (2 days)
- Highest scoring pairs were suggested for merger to reduce the variable space (curse of dimensionality)

Classify Text using Naïve Bayes Multinomial

What I should have done....

- ① Train an NBC to learn from small set of labeled cases
- ② Apply NBC to unlabeled data using Expectation Maximization with class probabilities (expectation step)
- ③ Retrain NBC using the labels for all the data
- ④ Repeat until convergence

WEKA Explorer Tutorial Examples

- Preprocess
 - Instance and Attribute Filters (Supervised and Unsupervised)
- Classify
 - Bayes
- **Cluster**
 - **Expectation Maximization**
 - **Hierarchical Clustering**
- Associate
 - Apriori
- Select Attributes
 - Via clustering

Cluster Algorithm Examples

- Expectation Maximization (EM)
- Hierarchical Clustering (cobweb)
- *Note: Weka provides many more clustering methods*

Cluster with Expectation Maximization

Preprocess

Classify

Cluster

Associate

Select attributes

Visualize

Clusterer

Choose

EM -I 100 -N -1 -M 1.0E-6 -S 100

Cluster mode

☒ Use training set

☐ Supplied test set

☐ Percentage split

☐ Classes to clusters evaluation

☒ Store clusters for visualization

Set...

% 66

(Nom) class

Ignore attributes

Start

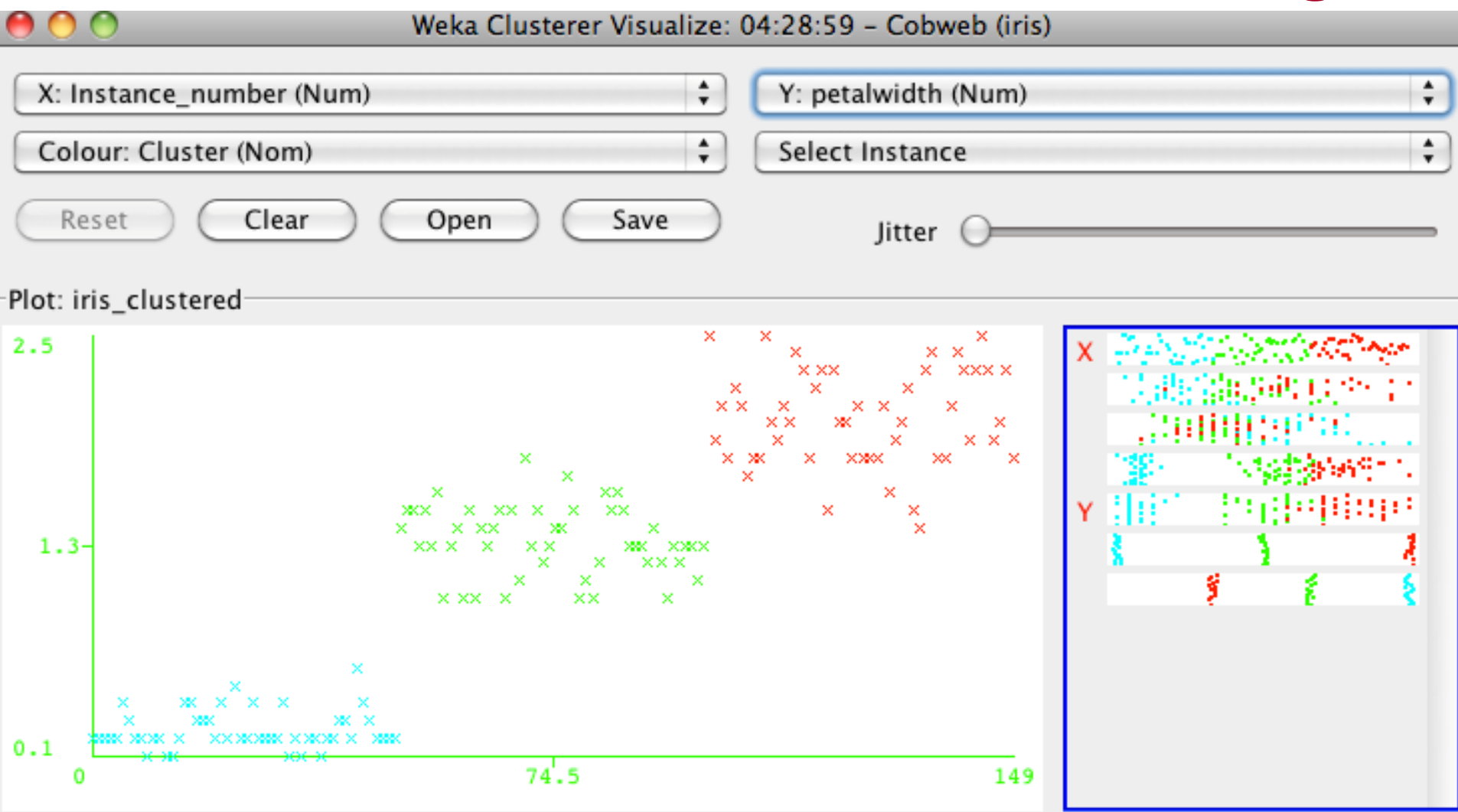
Stop

Result list (right-click for options)

04:16:46 - EM

Clusterer output

Cluster with cobweb (hierarchical clustering)



Class colour

WEKA Explorer Tutorial Examples

● Preprocess

- Instance and Attribute Filters (Supervised and Unsupervised)

● Classify

- Bayes

● Cluster

- Expectation Maximization
- Hierarchical Clustering

● Associate

- **Apriori**

● Select Attributes

- Via clustering

Associate

• Quick scan for association rules

- see “Fast Algorithms for Mining Association Rules in Large Databases”

The screenshot shows the Weka Associate tool interface. The top navigation bar includes buttons for Preprocess, Classify, Cluster, Associate (selected), Select attributes, and Visualize. The Associate window has a 'Choose' button and a text field containing the command: `Apriori -N 10 -T 0 -C 0.9 -D 0.05 -U 1.0 -M 0.1 -S -1.0 -c -1`. Below this are 'Start' and 'Stop' buttons. On the left, a 'Result list (right-click for...)' shows two entries: '04:47:46 - HotSpot' and '05:11:10 - Apriori' (selected). The main 'Associator output' pane displays the following text:

```
Minimum support: 0.1 (15 instances)
Minimum metric <confidence>: 0.9
Number of cycles performed: 18

Generated sets of large itemsets:

Size of set of large itemsets L(1): 20
Size of set of large itemsets L(2): 15
Size of set of large itemsets L(3): 3

Best rules found:

1. petalwidth='(-inf-0.34]' 41 ==> class=Iris-setosa 41    conf:(1)
2. petallength='(-inf-1.59]' 37 ==> class=Iris-setosa 37    conf:(1)
3. petallength='(-inf-1.59]' petalwidth='(-inf-0.34]' 33 ==> class=Iris-setosa 33    conf:(1)
4. petalwidth='(1.06-1.3]' 21 ==> class=Iris-versicolor 21    conf:(1)
5. petallength='(5.13-5.72]' 18 ==> class=Iris-virginica 18    conf:(1)
6. sepallength='(4.66-5.02]' petalwidth='(-inf-0.34]' 17 ==> class=Iris-setosa 17    conf:(1)
7. sepalwidth='(2.96-3.2]' class=Iris-setosa 16 ==> petalwidth='(-inf-0.34]' 16    conf:(1)
8. sepalwidth='(2.96-3.2]' petalwidth='(-inf-0.34]' 16 ==> class=Iris-setosa 16    conf:(1)
9. petallength='(3.95-4.54]' 26 ==> class=Iris-versicolor 25    conf:(0.96)
10. petalwidth='(1.78-2.02]' 23 ==> class=Iris-virginica 22    conf:(0.96)
```

At the bottom of the slide, there is a date '10/24/2023', a course identifier 'CAI3024N-ADA-JJT-WK7', and a page number '75'.

WEKA Explorer Tutorial Examples

● Preprocess

- Instance and Attribute Filters (Supervised and Unsupervised)

● Classify

- Bayes

● Cluster

- Expectation Maximization
- Hierarchical Clustering

● Associate

- Apriori

● **Select Attributes**

- **Via clustering**

Select Attributes Using a Classifier

The screenshot displays the WEKA software interface, specifically the 'Attribute Evaluator' window. The 'Select attributes' tab is active. The 'Attribute Evaluator' section shows 'ClassifierSubsetEval' as the evaluator and 'Click to set hold out or test instances' as the hold-out method. The 'Search Method' section shows 'GreedyStepwise' as the search method. The 'Attribute Selection Mode' section has 'Use full training set' selected. The 'Attribute selection output' pane shows the results of the search, indicating that the 'petalwidth' attribute was selected.

Attribute Evaluator

Choose **ClassifierSubsetEval** -B weka.classifiers.bayes.NaiveBayes -T -H "Click to set hold out or test instances" --

Search Method

Choose **GreedyStepwise** -T -1.7976931348623157E308 -N -1

Attribute Selection Mode

☒ Use full training set
☐ Cross-validation

Folds
Seed

(Nom) class

Start Stop

Result list (right-click for options)

04:52:12 - GreedyStepwise + ClassifierSubsetEval

Attribute selection output

```
=== Attribute Selection on all input data ===  
Search Method:  
  Greedy Stepwise (forwards).  
  Start set: no attributes  
  Merit of best subset found:    0.04  
  
Attribute Subset Evaluator (supervised, Class (nominal): 5 class):  
  Classifier Subset Evaluator  
  Learning scheme: weka.classifiers.bayes.NaiveBayes  
  Scheme options:  
  Hold out/test set: Training data  
  Accuracy estimation: classification error  
  
Selected attributes: 4 : 1  
                    petalwidth
```

Status
OK

Log

Select Attributes using PCA

Attribute Evaluator

Choose PrincipalComponents -R 0.95 -A 5

Search Method

Choose Ranker -T -1.7976931348623157E308 -N -1

Attribute Selection Mode

☒ Use full training set
☐ Cross-validation

Folds 10
Seed 1

(Nom) class

Start Stop

Result list (right-click for options)

04:52:12 - GreedyStepwise + ClassifierSubse
04:58:49 - Ranker + PrincipalComponents

Attribute selection output

=== Run information ===
Evaluator: weka.attributeSelection.PrincipalComponents -R 0.95 -A 5
Search: weka.attributeSelection.Ranker -T -1.7976931348623157E308 -N -1
Relation: iris
Instances: 150
Attributes: 5
 sepalength
 sepalwidth
 petallength
 petalwidth
 class
Evaluation mode: evaluate on all training data

=== Attribute Selection on all input data ===
Search Method:
 Attribute ranking.
Attribute Evaluator (unsupervised):
 Principal Components Attribute Transformer

Correlation matrix
 1 -0.11 0.87 0.82
 -0.11 1 -0.42 -0.36
 0.87 -0.42 1 0.96
 0.82 -0.36 0.96 1

eigenvalue proportion cumulative
 2.91082 0.7277 0.7277 0.581petallength+0.566petalwidth+0.522sepalength-0.263sepalwidth
 0.92122 0.23031 0.95801 -0.926sepalwidth-0.372sepalength-0.065petalwidth-0.021petallength

Eigenvectors
 V1 V2
 0.5224 -0.3723 sepalength
 -0.2634 -0.9256 sepalwidth
 0.5813 -0.0211 petallength
 0.5656 -0.0654 petalwidth

Ranked attributes:
 0.2723 1 0.581petallength+0.566petalwidth+0.522sepalength-0.263sepalwidth
 0.042 2 -0.926sepalwidth-0.372sepalength-0.065petalwidth-0.021petallength

Selected attributes: 1,2 : 2

WEKA Tutorial Summary

● Preprocess

- Prepare datasets instances and attributes before analysis

● Classify

- Pick a instance and predict the class
 - Iris : Pick a flower and use the attributes to predict species
 - Medicine: pick a patient and use the genes to predict cancer status

● Cluster

- Group instances together (flowers, breast cancer cases, etc)

● Associate

- Discover relationships between variables in your dataset

References

- Data Mining: Practical Machine Learning Tools and Techniques
- Data Mining (I. H. Witten and E. Frank)
- WEKA Exploratory Tool for Data Mining
- Bayesian Network Classifiers in Weka (Bouckaert)
- COC131 Data Mining – Clustering (Sykora)
- Fast, Correct Multithreaded Programs in Java (Gilbert)
- R. Agrawal, R. Srikant: Fast Algorithms for Mining Association Rules in Large Databases. In: 20th International Conference on Very Large Data Bases, 478-499, 1994.
- WEKA Wiki
<http://weka.wikispaces.com/>
- Graphical User Interface
<http://prdownloads.sourceforge.net/weka/weka.ppt>

References

- **OL- Ebook**