



Ahsanullah University of Science & Technology

Department of Computer Science & Engineering

Course No : CSE4142

Course Title : Data Warehousing and Mining Lab

Assignment No : 04

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Submitted To : Mr. Saha Reno & Mr. Raiyan Jahangir

Submitted By-

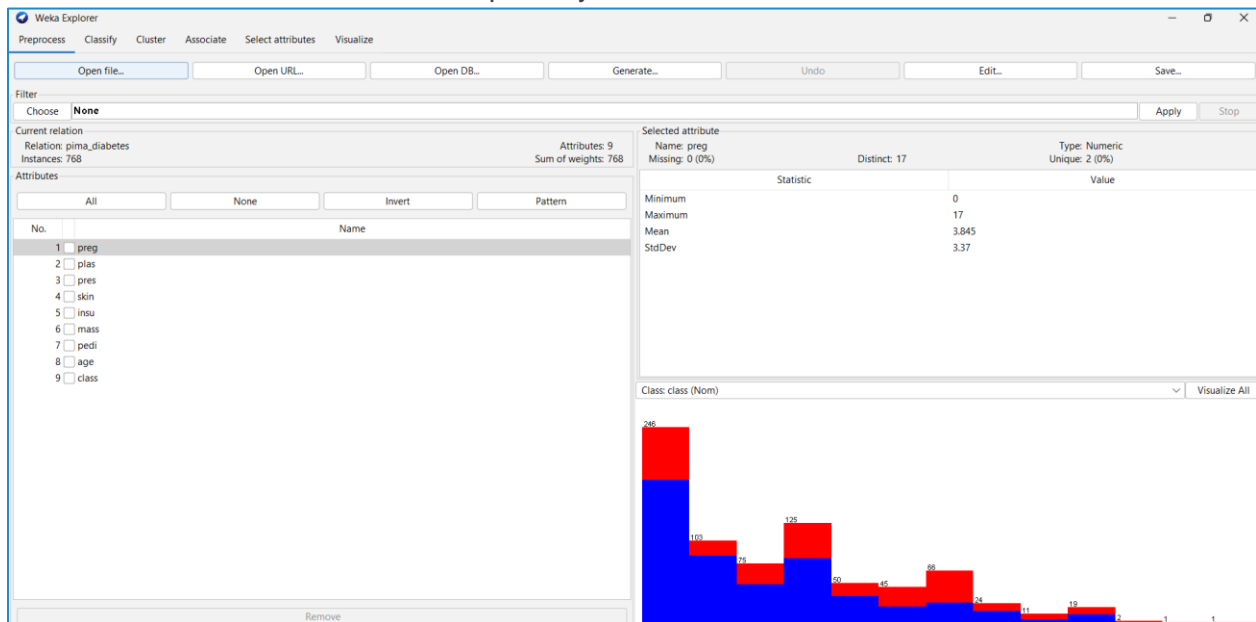
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Section: B1

(i) Take any dataset from Weka Repository or Online (Must be in ARFF format)

Ans: Diabetes dataset from Weka Repository-



(ii) Extract 1 Non-Stratified Fold from 10-Fold Cross Validation.

Ans:

weka.gui.GenericObjectEditor

weka.filters.unsupervised.instance.RemoveFolds

About

This filter takes a dataset and outputs a specified fold for cross validation.

More

Capabilities

debug False

doNotCheckCapabilities False

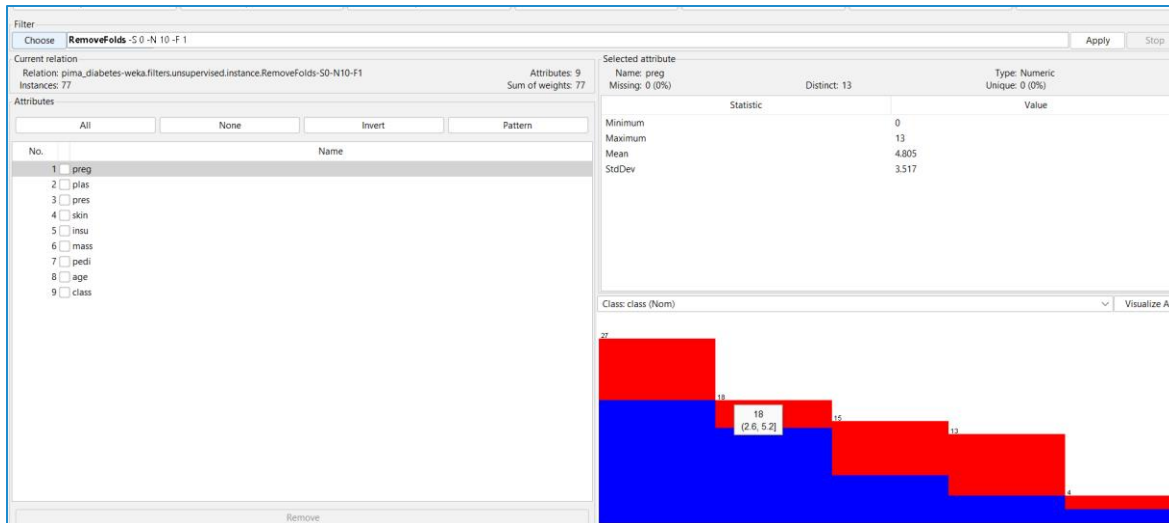
fold 1

invertSelection False

numFolds 10

seed 0

Open... Save... OK Cancel



(iii) Extract 1 Stratified Fold from 10-Fold Cross Validation.

Ans:

weka.gui.GenericObjectEditor

weka.filters.supervised.instance.StratifiedRemoveFolds

About

This filter takes a dataset and outputs a specified fold for cross validation.

More Capabilities

debug False

doNotCheckCapabilities False

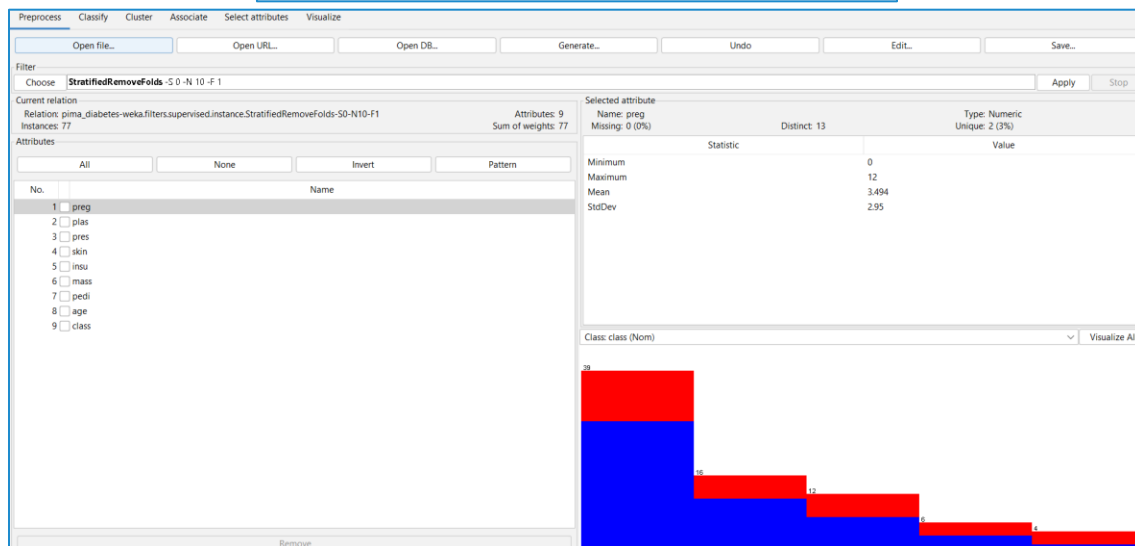
fold 1

invertSelection False

numFolds 10

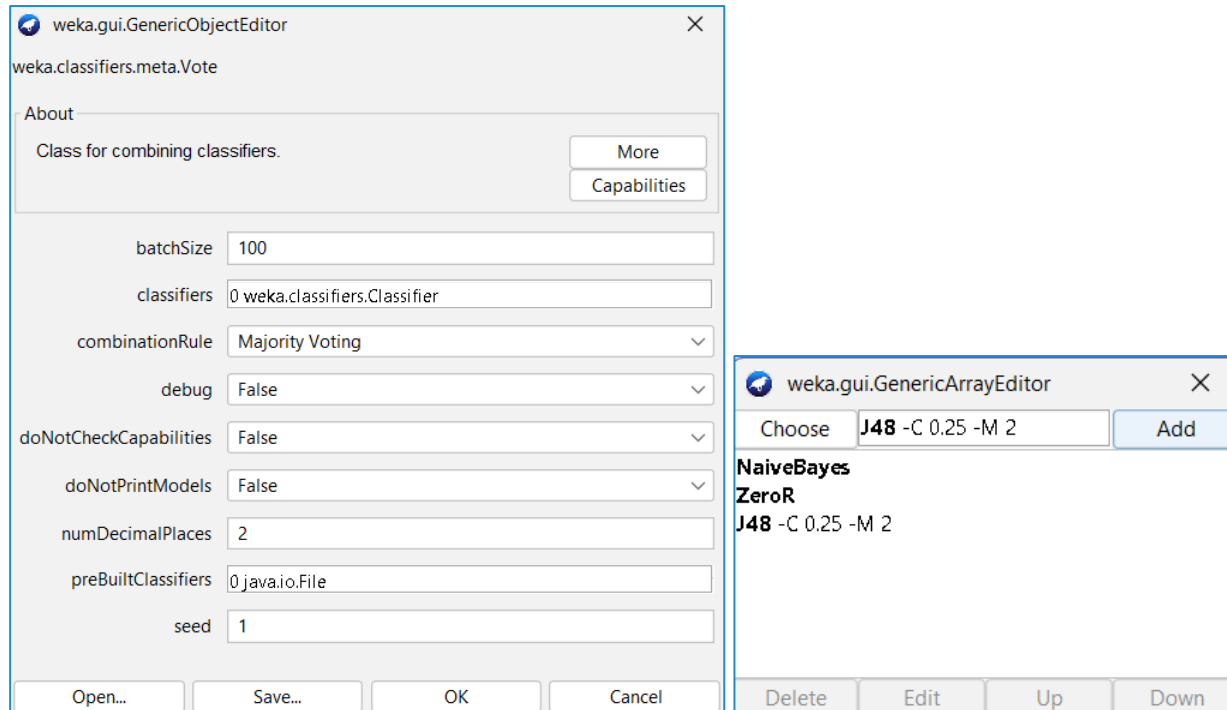
seed 0

Open... Save... OK Cancel



(iv) Using Ensemble technique, create any 3 models/classifiers and follow Voting approach to classify test instances. You must select "Majority Voting" option as the Rule of Combination.

Ans:



(v) At first, use the Non-Stratified Fold as the Test Data. After that, use the Stratified Fold as the Test Data.

Ans: First Non-Stratified Fold is taken as Test Data then the Stratified Fold is used as the Test Data.

(vi) Show the outputs for both classifications from (v).

Ans: From (v), for Non-Stratified fold the output is-

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

Vote combines the probability distributions of these base learners:
    weka.classifiers.bayes.NaiveBayes
    weka.classifiers.rules.ZeroR
    weka.classifiers.trees.J48 -C 0.25 -M 2
using the 'Majority Voting' combination rule
```

```

=== Summary ===

Correctly Classified Instances      58           75.3247 %
Incorrectly Classified Instances    19           24.6753 %
Kappa statistic                    0.4705
Mean absolute error                 0.2468
Root mean squared error            0.4967
Relative absolute error             50.7609 %
Root relative squared error        100.7945 %
Total Number of Instances          77

=== Detailed Accuracy By Class ===

              TP Rate  FP Rate  Precision  Recall   F-Measure  MCC      ROC Area  PRC Area  Class
              0.889   0.438   0.741    0.889   0.808     0.486   0.726    0.723    tested_negative
              0.563   0.111   0.783    0.563   0.655     0.486   0.726    0.622    tested_positive
Weighted Avg.   0.753   0.302   0.758    0.753   0.744     0.486   0.726    0.681

=== Confusion Matrix ===

  a  b  <-- classified as
40  5  |  a = tested_negative
14 18  |  b = tested_positive

```

For Stratified fold the output is-

```

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

Vote combines the probability distributions of these base learners:
    weka.classifiers.bayes.NaiveBayes
    weka.classifiers.rules.ZeroR
    weka.classifiers.trees.J48 -C 0.25 -M 2
using the 'Majority Voting' combination rule

```

```

=== Summary ===

Correctly Classified Instances      63           81.8182 %
Incorrectly Classified Instances    14           18.1818 %
Kappa statistic                    0.5553
Mean absolute error                 0.1818
Root mean squared error            0.4264
Relative absolute error             39.8272 %
Root relative squared error        89.3571 %
Total Number of Instances          77

=== Detailed Accuracy By Class ===

              TP Rate  FP Rate  Precision  Recall   F-Measure  MCC      ROC Area  PRC Area  Class
              0.980   0.481   0.790    0.980   0.875     0.601   0.749    0.788    tested_negative
              0.519   0.020   0.933    0.519   0.667     0.601   0.749    0.653    tested_positive
Weighted Avg.   0.818   0.320   0.840    0.818   0.802     0.601   0.749    0.740

=== Confusion Matrix ===

  a  b  <-- classified as
49  1  |  a = tested_negative
13 14  |  b = tested positive

```