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RWeka Tutorial

You can use the "RWeka" package to run Weka in R. Below is a step-by-step tutorial on how to run J48 on the Titanic data using RWeka.

For more details, check out the official RWeka manual at https://cran.r-project.org/web/packages/RWeka/RWeka.pdf

Step1: INSTALL R PACAKGE

install.packages("RWeka")

Note: If you are installing the RWeka package on a windows pc, you may need to set the Java environment first by execute the following script in R:

Sys.setenv(JAVA_HOME="C:\\Program Files\\Java\\jdk1.8.0_51\\jre")

Step2: READ FILES (change file path to your own file path)

trainset <- read.csv("/Users/byu/Documents/R/train.csv")
testset <- read.csv("/Users/byu/Documents/R/test.csv")

Or

trainset <- read.arff("/Users/byu/Documents/R/train.arff")
testset <- read.arff("/Users/byu/Documents/R/test.arff")</pre>

Step3: PREPROCESS DATA IN R

We can use either the built-in R functions or the RWeka filter interface, here we are introducing the RWeka interface for data preprocessing.

Transform data type

Numeric -> Nominal

NN <- make_Weka_filter("weka/filters/unsupervised/attribute/NumericToNominal") # build a function using RWeka filter interface

Apply the filter function to both training and test datasets.

trainset <- NN(data=trainset, control= Weka_control(R="1-3"), na.action = NULL) testset <- NN(data=testset, control= Weka_control(R="1,3"), na.action = NULL)

Deal with missing values

MS <- make_Weka_filter("weka/filters/unsupervised/attribute/ReplaceMissingValues") # build a function using RWeka filter interface

Apply the filter function to both training and test datasets.

trainset <-MS(data=trainset, na.action = NULL) testset <-MS(data=testset, na.action = NULL)

Check data definition (optional)

str(trainset)

Step4: APPLY J48 Algorithm

Build decision tree model

```
m=J48(Survived~., data = trainset)
m=J48(Survived~., data = trainset, control=Weka_control(U=FALSE, M=2, C=0.5))
```

* View parameters with function WOW:

i.e. WOW("J48")

```
> WOW("J48")
-U
        Use unpruned tree.
        Do not collapse tree.
-C <pruning confidence>
        Set confidence threshold for pruning. (default 0.25)
        Number of arguments: 1.
-M <minimum number of instances>
        Set minimum number of instances per leaf. (default 2)
        Number of arguments: 1.
        Use reduced error pruning.
-N <number of folds>
        Set number of folds for reduced error pruning. One fold is used as pruning set.
        (default 3)
        Number of arguments: 1.
-B
       Use binary splits only.
-S
        Do not perform subtree raising.
-L
        Do not clean up after the tree has been built.
        Laplace smoothing for predicted probabilities.
-J
        Do not use MDL correction for info gain on numeric attributes.
-Q <seed>
        Seed for random data shuffling (default 1).
        Number of arguments: 1.
```

Use 10 fold cross-validation to evaluate the model

```
e <- evaluate_Weka_classifier(m,
numFolds = 10,
seed = 1, class = TRUE)
```

```
=== Summary ===
Correctly Classified Instances
                                     720
                                                      80.8081 %
Incorrectly Classified Instances
                                     171
                                                      19.1919 %
Kappa statistic
                                      0.5753
Mean absolute error
                                      0.2646
Root mean squared error
                                      0.3659
                                     55.9347 %
Relative absolute error
Root relative squared error
                                     75.2366 %
Coverage of cases (0.95 level)
                                     99.5511 %
Mean rel. region size (0.95 level)
                                     91.9753 %
Total Number of Instances
=== Detailed Accuracy By Class ===
                TP Rate FP Rate Precision Recall
                                                   F-Measure MCC
                                                                       ROC Area PRC Area Class
                0.918
                                 0.800
                                           0.918
                                                    0.855
                                                              0.587
                                                                       0.827
                                                                                0.846
                        0.368
                                                                                          0
                                                                                0.796
                0.632
                        0.082
                                 0.828
                                           0.632
                                                    0.716
                                                              0.587
                                                                       0.827
                                                                                         1
Weighted Avg.
               0.808
                        0.258
                                 0.811
                                           0.808
                                                    0.802
                                                              0.587
                                                                       0.827
                                                                                0.827
=== Confusion Matrix ===
         <-- classified as
  a b
504 45 I
           a = 0
126 216 |
           b = 1
```

Apply the model with test dataset

pred=predict (m, newdata = testset, type = c("class"))

Predict Result:

Then save the prediction to a csv file:

write.csv(pred, file="path/filename.csv")

Note that the model built by RWeka may not be exactly the same as the model built in Weka, if the pre-processing steps are different. For example, to build the above RWeka model, missing values were replaced by mean for the "age"

attribute. However, the Weka GUI's filter "replaceMissingValues" actually applies to all variables, causing slightly different model and performance.

```
Classifier output
=== Summary ===
Correctly Classified Instances
                                  691
                                                  77.5533 %
Incorrectly Classified Instances
                                  200
                                                  22.4467 %
Kappa statistic
                                  0.5098
Mean absolute error
                                   0.3192
Root mean squared error
                                   0.4056
Relative absolute error
                                  67.4659 %
                                  83.3925 %
Root relative squared error
Total Number of Instances
                                  891
=== Detailed Accuracy By Class ===
             TP Rate FP Rate Precision Recall F-Measure ROC Area Class
               0.871 0.377 0.787 0.871 0.827 0.777
                                                                    0
                                         0.623
                       0.129
                                                  0.681
               0.623
                                0.75
                                                           0.777
                                                                    1
Weighted Avg.
               0.776
                       0.282
                               0.773 0.776
                                                  0.771
                                                           0.777
=== Confusion Matrix ===
         <-- classified as
  a b
 478 71 | a = 0
 129 213 | b = 1
```

Removing irrelevant attributes:

```
myVars=c("Pclass", "Sex", "Age", "SibSp", "Parch", "Fare", "Survived")

newtrain=trainset[myVars]

newtest=testset[myVars]

m=J48(Survived~., data = newtrain)

m=J48(Survived~., data = newtrain, control=Weka_control(U=FALSE, M=2, C=0.5))

e=evaluate_Weka_classifier(m, seed=1, numFolds=10)

pred=predict (m, newdata = newtest, type = c("class"))

myids=c("PassengerId")

id_col=testset[myids]

newpred=cbind(id_col, pred)

colnames(newpred)=c("Passengerid", "Survived")

View(newpred)
```

write.csv(newpred, file="/Users/byu/Desktop/Data/titanic-J48-pred.csv", row.names=FALSE)

InfoGainAttributeEval(Survived ~ . , data = trainset)

Benefit of RWeka:

- 1. RWeka does not give incompatibility error when reading in both train.csv and test.csv. Weka GUI did.
- 2. Running Weka in R would allow you to use Weka prediction algorithms and also the rich data pre-processing tools in R.