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APPENDIX A – User Manual

gaKnn Churn Prediction Tool Version 1.0

User Manual

gaKnn Churn Prediction Tool is an easy way that you can get to know about the future churners of your product or service. This provides simple and easy to handle, user interfaces to get your work done. The Tool uses the key concepts of k nearest neighbor classification and genetic algorithms; supported by the concepts of Naïve Bayesian Weights and Class Confidence Weights (ccw).

-----System Requirements-----

- JDK 1.6 or above
- jgap library
- R 3.1.3 or above with rJava Package installed
- Other R libraries required: foreign, ROCR, gplots, robustbase, infotheo, hexbin, colorspace, ggplot2
- opencsv 3.1.2 library (should be compatible with the jdk version)

-----Working with the Tool-----

Start the Tool

The Tool starts with the interface Figure A-1. Proceed with the instructions given at each step.

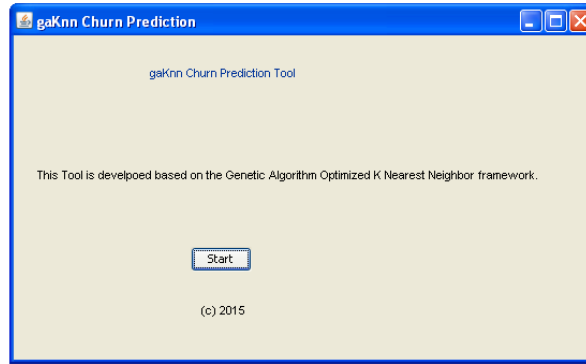


Figure A-1 First Interface

Selecting the Dataset

Select the dataset through window in Figure A-2. As the dataset is selected it provides a graphical as well as a statistical overview of the selected dataset.

To get the statistical data follow the 'View Statistics' button.

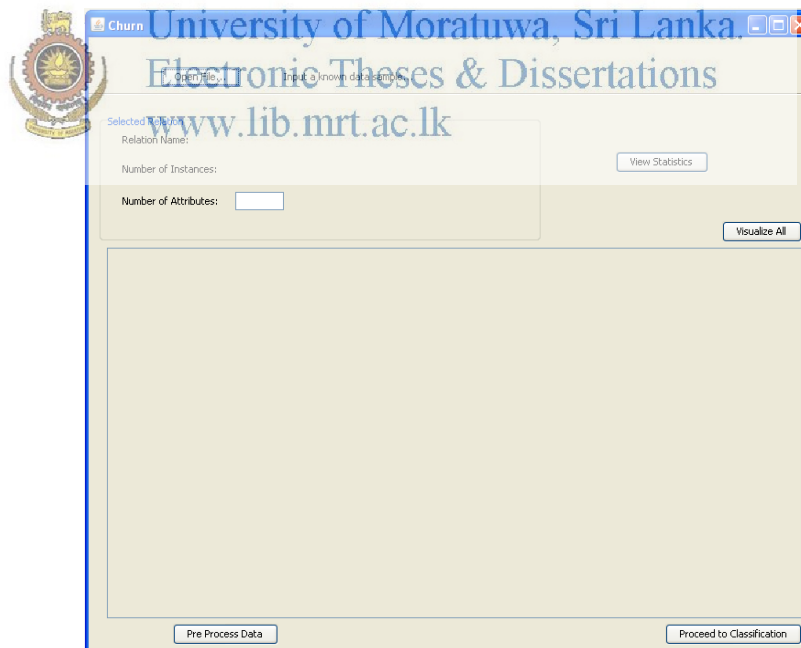


Figure A-2 Data selection

Data Visualizing

You can get several views about the distribution of data through 'Visualize All' section.

Scatter plot view

Bar plot view

Data Pre-processing

If you proceeded to the preprocess module via preprocess button in the 'Churn' window, the window in Figure would be shown.

The 'Preprocess' window provides you the ability to handle missing values and outliers. Click on the button 'Missing Value Handler' and it will remove the missing values in the selected data file. The data instances which contain missing values could be viewed by clicking the button 'View Missing Values'.

If you need to use the missing value handled data file as your training data file, select that data file through the 'Churn' window.



Figure A-3 Data pre-processor

Outliers in the selected data file are visualized as box plots. You can view them through the button 'View Outliers'.

The Tool provides two outlier detection methods;

1. Tukey's Method
2. MedCouples

Outlier handler provides two handling methods; removal and replacement.

First select the detection method from the dropdown menu and then click on the preferred handling method.

Please note that the visualized outliers are determined by the whiskers extend to the most extreme data point which is no more than range times the inter quartile range from the box.

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You can proceed to the optimizing phase via 'Proceed to Classification' button.

The Prediction window allows you to perform both the optimizing and prediction tasks.

The optimizing task involves genetic algorithm techniques. It is by this step the Tool finds an optimum value for k and a weight vector for the attributes considering their importance.

The number of times which genetic algorithm is required to iterate is read by the Number of Evolutions and the size of the population considered by genetic algorithm is read by Size of the Population fields in the Set Parameters section.

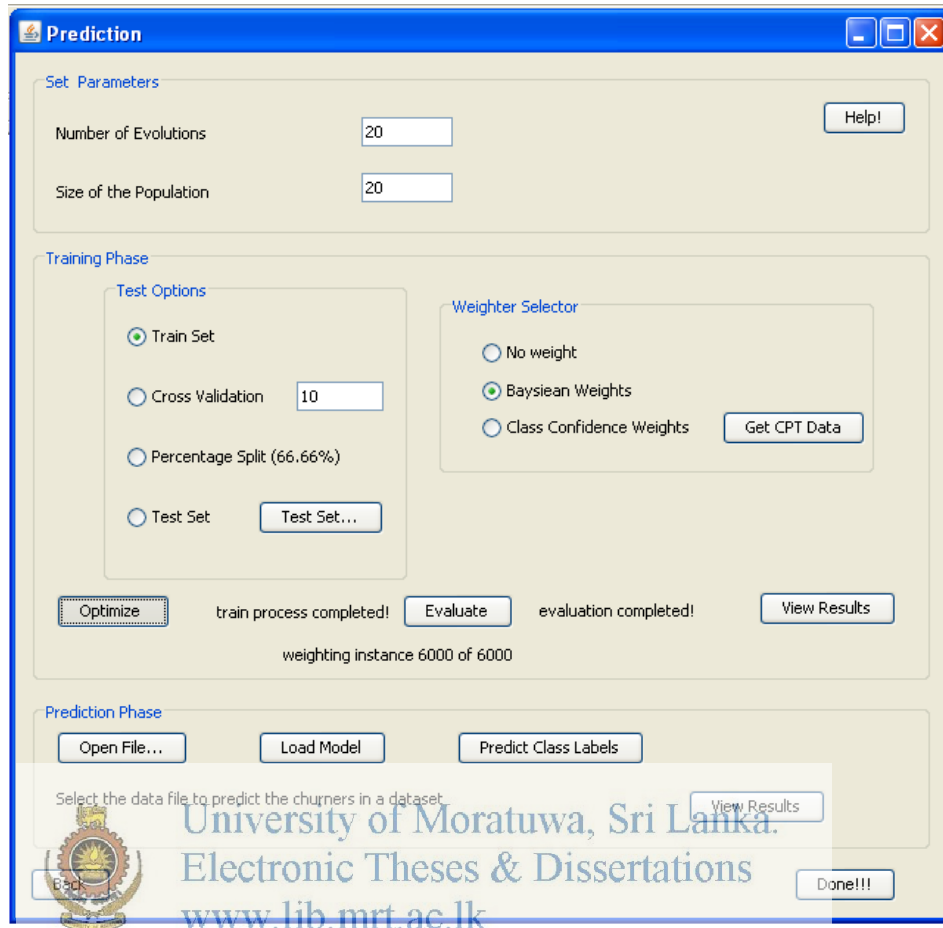


Figure A-4 Predictor

The population estimated with genetic algorithm is evaluated with KNN.

KNN classification relies on classification of unknown data based on known data. Therefore to test the KNN classifier it requires having two data sets; train set for training and test set for validating. In the Training Phase the Tool introduces four options to define the train set and test set; Train Set, Cross Validation, Percentage Split and Test Set.

If Train Set option is selected the dataset read into the Tool is used as both train set and test set. The Cross Validation option partitions the dataset, into specified k number of folds which would be used iteratively for k times where one fold is taken as

the test set while the remaining $k-1$ folds are used as the train set. The option `Percentage Split` partitions the dataset into two folds of which one fold contains two-third of the dataset and the other fold contains the remaining one-third. If you use the `test set` option then you are required to select a separate dataset which is from the same domain as the dataset already read into the Tool. The first dataset is used as the train set while the second fold is used as the test set.

To get a better result on prediction with KNN, instance voting is introduced. Two voting methods were introduced; `Bayesian Weights`, `Class Confidence Weights`. You may either perform the classification without either of the two methods by selecting 'No Weight' option. `Bayesian Weights` add a vote to each prediction based on the Bayesian probability of getting a particular class label. `Class Confidence Weights` add a vote to each prediction based on the probability of attribute values given the class labels.

After selecting the desired options from `Test Options` and `Weight Selector`, click on the `Optimize` button. It will evaluate the data read into the Tool with genetic algorithm and KNN to produce the optimum value for k as well as a weight vector for the attributes. Results obtained with relevant to the optimization process could be viewed through the `View Results` button in the `Training Phase` section.

The model is automatically saved as an xml file with the optimized k value and the weights for each attribute. The location of the file is the same as where the training dataset is located and with the same file name as the training dataset with `.prm` extension.

The classifier could be re-evaluated with the Train set or with a separate test set. If you need to re-evaluate the model with the Train set simply click on the `Evaluate` button without changing any other option. If you need to re-evaluate the model with a separate test set, select the `Test set` option from the `Test Options` and choose another

known set from the same domain as the initial dataset (Train Set). Results could be viewed through the View Results button in the Training Phase section.

Prediction

From the Prediction Phase section of the Prediction window choose the dataset with unknown class labels. You can predict the class labels through Predict Class Labels and view the churn results through View Results button.

View Results

Record ID	Prediction	Confidence
11	1	0.9994402552736066
22	1	0.9999475522163587
42	1	0.9976292270908231
49	1	0.9999572055124252
55	1	0.9999265876367418
58	1	0.9980739969708743
77	1	0.9980649063209354
85	1	0.9944819465604948
87	1	0.9993756278345933
90	1	0.9983644105030223
92	1	0.994571918850505
99	1	0.9919951094650691
100	1	0.9952943149593646
118	1	0.9945927660468222
126	1	0.9995599362546571
145	1	0.9932434628740288
157	1	0.9991220858598594
198	1	0.6446382756671869
199	1	0.6180083898625983
215	1	0.5950929874325862
219	1	0.9947539026497748
231	1	0.9968701096018466
242	1	0.9947455788171412
245	1	0.9985387879128237
251	1	0.9996103555220908
259	1	0.9945053417743812
278	1	0.9953527960925806
290	1	0.5584223899829217
294	1	0.9999438642093965
1	0	1.0
2	0	1.0
3	0	0.9999310686986311
4	0	1.0
5	0	1.0
6	0	0.9337568965346086

Figure A-5 Result analyzer

Results could be viewed for different stages of the process. Viewing the results under the Training Phase provide the distribution of attribute weights, fitness distribution and evaluated predictions. By selecting the required tab you can easily get the relevant results.

Weight distribution shows the importance that has on each attribute when training the classifier. Attributes with high weight values are the most contributing attributes for the final prediction.

Fitness distribution shows whether the fitness value is stabilized for the Number of Evolutions specified in Prediction window. The model provides better results if the fitness value is stabilized at a maximum value.

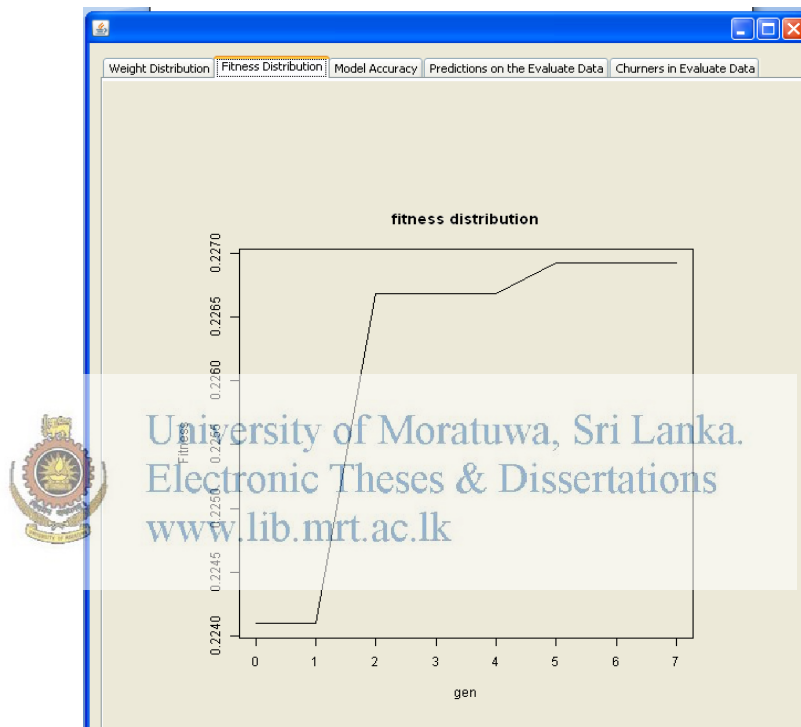


Figure A-6 Fitness distribution

Accuracy of the model could also be viewed in terms of four criteria (Figure A-7).

'Churners in the evaluated data' tab shows the predicted churners along with their details which were considered for the prediction. You can by examining the trend in data under 'Visualize All' section, edit the values of attributes of the churners. Edited file could be saved and re-used for prediction.

Prediction phase allows you to load an existing model to predict churners. Select the Load Model button to select the saved model. The model loading should be saved as an xml file in .prm extension to be able to use with the Tool.

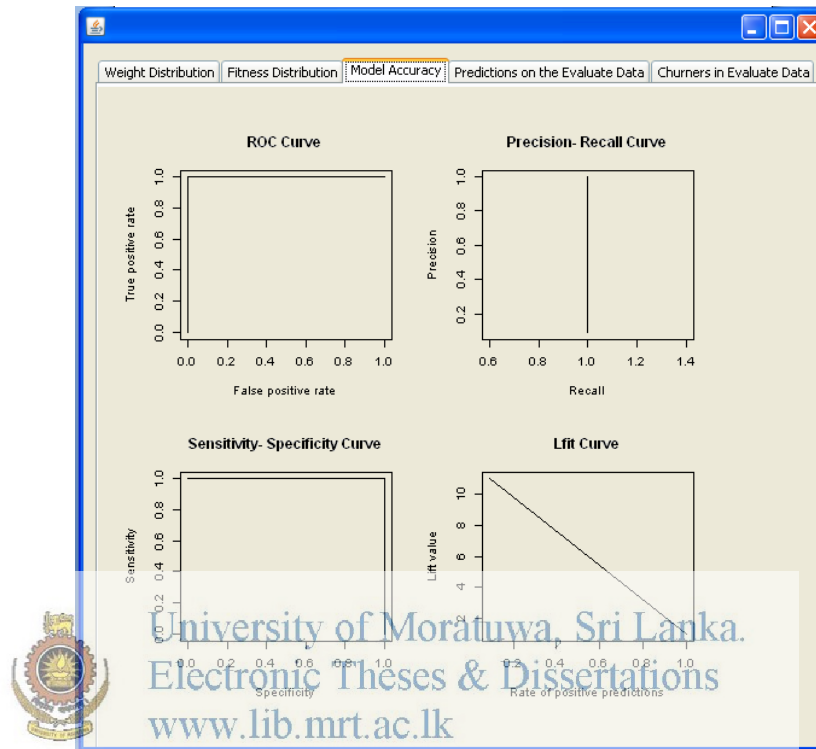


Figure A-7 Model accuracy

APPENDIX B – Source Code

The source code of the Tool could found in the CD attached with the title Appendix B.



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