Analysis of Classification Techniques for Prediction of Tuberculosis Defaulters

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ABSTRACT

1. INTRODUCTION

In 2013 over 210 000 patients defaulted from Tuberculosis (TB) treatment worldwide [15]. The rate of default in the Americas is the highest at 8% with Africa at 5% [15]. The consequences of defaulting TB treatment include: increased drug resistance, increased health system costs [10, 12], higher risk of mortality, continued risk of transmitting the disease to others [10] and increased rate of recurrent disease [8]. The spread of TB can be reduced if the individuals who have a high risk of defaulting can be predicted. This will also reduce health system costs.

In this paper we examine a wide range of classification techniques at their ability to predict patients as defaulters. We explore a number of techniques and strategies to improve the overall classification result. Most notably we look at how well these classifiers work out-of-the-box compared to when they are hand optimised or tuned by searching a grid of parameters. We examine different data balancing techniques which either over-sample the minority class or under-sample the majority class. Furthermore, to see how each classifier responds to having fewer features available, we apply a number of feature selection strategies on each classification technique.

To facilitate the aforementioned experiments, we developed a testing system which allows new classification techniques and data sets to be supported quickly and easily. The testing system is designed to allow near-exact reproducibility of results.

The field of credit scoring in the financial space aims to determine if a financial institution should provide credit to an individual. This area has been well researched. In this paper we aim to determine if our results generalise across the different fields. There are notable similarities in these problems which could make them comparable. Both problems typically have a labelled dataset consisting of both nominal and numerical data as well as the occurrence of missing data [citation needed]. However, TB datasets are more prone to

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missing data as well as inaccuracies due to the nature of the data collection [citation needed]. The selected classification techniques are evaluated against real-world treatment default datasets and financial datasets. The paper will evaluate how the techniques differ across the datasets. If the relative results are similar then future credit scoring research could be applicable to treatment default prediction too.

- 1. Discuss datasets used and possibly expansion of TB issues in those specific counties (Peru and Malawi)
- Outline briefly how the datasets are used and that each technique is also benchmarked against the well known Australian and German financial to determine how applicable credit scoring research is to TB default prediction for the two TB datasets.
- 3. Link TB classification to credit scoring and outline notable similarities and differences
- 4. Summarise overall paper

2. RELATED WORK

2.1 Definition of a defaulter

The definition of a defaulter depends on its context. TB literature typically uses the World Health Organisation (WHO) definition that a defaulter is a person whose treatment has been disrupted for two or more consecutive months [1, 2, 8, 9, 12, 15].

2.2 Determining predictors of TB default

There have been many studies which focus on determining the factors associated with TB default but few have used machine learning techniques to predict treatment defaulters. Table 1 contains an overview of a selection of publications on determining the factors associated with TB default. The majority of techniques use a form of logistic regression ti determine the association.

The datasets used by the publications contain different features. Age and gender are common throughout the datasets. History of past default is available for all datasets except for Shargie et al. [13]. Lackey et al. [10] only picked individuals who did not have a history of past default. Jittimanee [9] [28] was the only publication with the feature that did not find it to be significant to the 95% confidence level. However, it did have an odds ratio of 2.19 and a p-value of 0.12. It can therefore be deduced that a history of past default has a strong correlation to default. Two out of three publications with the alcohol abuse feature available, found it

to be significant. Three of the four publications with side effects, as a feature found it was significant. Shargie et al. [13] and Jittimanee et al. [9] measured distance and time to treatment site respectively. It can be reasoned that the aforementioned feature's significance will generalise to other datasets since they were found to be significant in the majority of the publications. Other significant features such as illegal drug use, use of herbal medication, daily jobs, history of lung cancer and history of liver disease only appeared once in the datasets. It cannot be discerned if the significance is generalisable or specific to the dataset. The identification of the same features as significant is fairly consistent for the publications that have those features in their dataset.

2.3 Predicting defaulters in financial institutions

- 1. Summarised version of original literature review except the parts using temporal aspects
- 2. Summarise what has been done determining TB predictors (which is a different aim)
- Summarise what has been done in the credit scoring financial space as a lot of research has been done in this field.
- Brief overview of each classification technique and possibly give a reason why this technique may work well for our application
- 5. Brief outline of the data balancing techniques used

3. BACKGROUND

3.1 Classifiers

3.1.1 Support Vector Machines

Support vector machine (SVM) is a machine learning technique that can be used to produce regression or classification functions from a set of training data [11]. SVM works by mapping the input vectors into a high-dimensional feature space with the use of a kernel function [3]. The kernel function selected determines the if the mapping is linear or non-linear [11]. Linear, polynomial and radial basis function (RBF) are common kernel functions [5]. The polynomial and RBF kernel performs a non-linear mapping into the highdimensional space [5]. This feature space is then searched to acquire an optimal hyperplane that separates the space with the maximum distance between the two classes [3]. Hsu et al. [5] recommends the RBF kernel as a reasonable first choice but notes that it is not suitable when there are a large number of features. The linear kernel is recommend when there are a large number of features.

3.1.2 Artificial Neural Network

An artificial neural network (ANN) is based on the functionality of the human brain [14]. Neurons in the brain are interconnected and process information in parallel [14]. A typical ANN is comprised of an input layer, k number of hidden layers and an output layer. The neurons in each layer are connected to the neurons in the next layer. A numeric weight is defined between each pair of connected neurons. An activation function defines if a neuron will fire [14]. The

activation function bounds the value of a neuron to a specific range to limit the effect of divergent neurons [14]. By using an activation function, a non-linear combination of weights can be generated [14]. It has been proven that an ANN is able to approximate any continuous function if has at least one hidden node and an activation function that is both bounded to some range and non-constant [4].

Extreme learning machines (ELM) is an alternative approach to the conventional back-propagated ANN. Huang el al. [7] proved that the input and hidden layer weights can be randomly assigned if the activation function in the hidden layer is infinitely differentiable. By randomly assigning these weights, the weights for the output nodes can be determined analytically [7]. This allows ELMs to be trained orders of magnitude faster than a back-propagated ANN [7] [6]. Various studies have shown ELMs to provide better results than SVMs and ANNs on classification and regression tasks [7] [6].

3.2 Data Balancers

3.3 Feature selection

4. METHODS

- Outline each dataset fully: number of nominal and numerical fields and number of entries. Also outline how balanced each dataset is.
- 2. Outline testing procedure and optimisation strategy for each classification technique. To optimise each classification technique, a grid search is conducted across a reasonable parameter space. For each parameter set, the results are averaged over 3 runs, this is done because of the stochastic nature of initialising the training of each classifier. To ensure consistent results for each run, the same fold allocations are used.
- 3. Outline that data balancing techniques are compared by using true positive, true negative, false positive, false negative by using the optimised parameters for each data balancing technique.

4.1 Approach

The study aimed to evaluate a set of statistical and machine learning classification techniques on treatment default data sets and compare it to that applied to other related classification datasets. A variety of classification techniques were chosen to test against: artificial neural network (ANN), support vector machine (SVM) using a linear, RBF and polynomial kernel, logistic regression, decision tree, AdaBoost, random forest, k-nearest neighbours (kNN), Gaussian naive Bayes, Bernoulli naive Bayes, extreme learning machines and clustering-Launched classification. These were chosen to include a selection of well known techniques, ensemble techniques and newer techniques that have shown promising results in other studies. Four related experiments were conducted in total.

4.1.1 Comparison of classification techniques

The first experiment finds the optimal parameters for each classifier on each dataset by searching a grid of reasonable parameters and then compares the classification strength of

Table 1: Overview of publications on predictors of TB treatment defaulters

Publication	Sample Size	Key factors identified*	Evaluation
Chan- Yeung et al. [1]	1768 non-defaulters 442 defaulters.	History of default, history of lung cancer, liver disease and male patients	Multiple logistic regression is used to determine what factors are associated with default.
Jha et al. [8]	1189 non-defaulters 1141 defaulters.	Male patients, patients directly-observed treatment at public facilities, previous treatment outside India's Revised National Tuberculosis Control Programme and history of previous default	The chi-square test or Fisher's exact test (if there were less than 10 observations) was used to test the differences between defaulters and non-defaulters. Bivariate analysis was calculated on the features. Multivariate logistic regression using pre-selected features based on previous studies.
Jittimanee et al. [9]	106 non-defaulters 54 defaulters.	Jobs where one is only paid if one is at work that day, severe medication side-effects and time to travel to clinic.	Patients were interviewed and completed a questionnaire to obtain the information. Hierarchical logistic regression was carried out to assess the variable's relation to default.
Lackey et al. [10]	1106 non-defaulters 127 defaulters.	Has used illegal drugs, has multidrug-resistant TB, has not been tested for HIV, drinks alcohol at least once a week, underweight or has not completed secondary education.	Patients were interviewed to obtain the information. Bivariate analysis using Chi-square tests and odds ratios with 95% confidence intervals. Multivariate logistic regression is used with a backward fitting algorithm to determine if a variable is associated with default.
Muture et al. [12]	5659 non-defaulters 945 defaulters.	Inadequate knowledge on TB, herbal medication use, low income, alcohol abuse, previous default, suffering from HIV and male patients were determined through analysis to be associated with default.	Two-tailed χ^2 tests and Fisher exact tests (if a cell has less than 5 values) to assess categorical information. Odds ratio tests were used to measure association between features with a 95% confidence interval.
Shargie et al. [13]	310 non-defaulters 81 defaulters.	Distance from home to treatment site, age greater than 25 and the need to use public transport to get to treatment site.	Continuous features were analysed using sample t-test χ^2 tests and hazard ratios with 95% confidence intervals. The effect of each factor was assessed using Cox regression model with a backwards fitting algorithm.

^{*}Ordered in descending order of significance based on odds ratio and hazard ratio at a confidence interval of at least 95%.

Table 2: German data set[¶]

Publication	Accuracy	Type I Error	Type II Error
Huang et al. [?]	$79.87\%^\dagger$	n/a	n/a
Li et al. [?]	84.83%	10-20%*	10-20%*
Luo et al. [11]	77.06% (MySVM), 82.41% (SVM-GA) [†]	n/a	n/a
Huang et al. [?]	82.41% (SVM-GA) [†]	n/a	n/a
Danenas et al. [3]	94.41% (Linear SVM), 92.37% (PSO-LinSVM) [†]	n/a [‡]	n/a [‡]

Table 3: Australian data set ¶

Publication	Accuracy	Type I Error	Type II Error	
Huang et al. [?]	79.87% [†]	n/a	n/a	
Li et al. [?]	84.83%	10-20%*	10-20%*	
Luo et al. [11]	77.06% (MySVM), 82.41% (SVM-GA) [†]	n/a	n/a	
Huang et al. [?]	82.41% (SVM-GA) [†]	n/a	n/a	
Danenas et al. [3]	94.41% (Linear SVM), 92.37% (PSO-LinSVM) [†]	n/a [‡]	n/a [‡]	

each classifier. These include the parameters of the classifier as well as different data balancing techniques. Data balancing algorithms are required since the rate of default is often relatively low. Many datasets have substantially fewer defaulters than those cured. However, most classification algorithms expect a 50:50 split between classes. The data balancing techniques included are: no data balancing, cluster centroids, edited nearest neighbours (ENN), instance hardness threshold, near miss (version 1), neighbourhood cleaning rule, one sided selection, random under sampling, Tomek links, adaptive synthetic sampling (ADASYN), random over-sampling, synthetic minority over-sampling (SMOTE), ensemble of SMOTE + EEN and ensemble of SMOTE +Tomek links. These balancing techniques were chosen to facilitate an evaluation of both over-sampling and undersampling techniques. Included in the evaluation is techniques that use simple approaches and as others that use sophisticated algorithms.

For each parameter set in the grid search, Three runs of 5-fold cross-validation is executed and the results averaged. To provide fair comparison, the data balancer, k-fold algorithm and classifier itself use fixed initialisation values, which differ across each of the three runs. This ensures that each parameter set utilises the same training and testing data.

The following metrics are recorded for this experiment: Matthews correlation coefficient (MCC), Cohen kappa coefficient (CKC), balanced classification rate (BCR), true positive rate (TPR), true negative rate (TNR), false positive rate (FPR) and false negative rate (FNR). Accuracy was not used as it is misleading in datasets that are imbalanced. MCC, CKC and BCR provide a balanced measure of the quality of the classification.

The parameter set that yields in the highest BCR is used as the parameters for the classifier on the particular dataset. To test each classifier with their optimal parameters, three random numbers are generated. This allows each run to use a different initialisation parameter and ensures that the same folds are used for each classifier. The aforementioned metrics are recorded for the test as well as the initialisation parameters that were used so results can be replicated. In addition a receiver operating characteristic (ROC) curve is also plotted as an additional means of comparing the classification techniques and to indicate what true positive rate can be achieved for an acceptable false positive rate. To determine how the results generalise against the other similar classification datasets, a final scatter plot is presented which compares difference of true positive and false positive from the median of each for each classifier on each dataset.

4.1.2 Comparison of data balancing algorithms

The second experiment compares each data balancing technique on each classifier. The optimal parameters for each classifier are used for every data balancer. As in experiment 1, the highest BCR is used to determine the optimal parameters. Each classifier is run using every data balance technique and the metrics outlined in section 4.1.1 is recorded. To ensure repeatable results, 10 runs of 5-fold cross validation is executed and the results averaged. A set of unique randomly selected initialisation keys are selected for each run. These are used so that k-fold cross validation, data balancing and classification algorithms produce deterministic output. The initialisation values are saved in the results to ensure they can be reproduced.

A bar chart is plotted with each classifier against the MCC for the treatment default datasets. As in section 4.1.1, a scatter plot is presented which compares difference of true positive and false positive from the median of each for each classifier on each data balancing algorithm on each dataset.

4.1.3 Comparison of feature selection algorithms

The third experiment investigates a number of different feature selection approaches in order to improve overall classification through the removal of features that each algorithm deems irrelevant to the classification. A number of feature selection strategies were selected for this experiment: Analysis of variance (ANOVA) F-value, logistic regression, linear SVM, Bernoulli naive Bayes, decision tree, random forest. For the ANOVA F-test, a p-value of greater 0.05 was selected to

4.1.4 An analysis of time to default in the Lima Peru dataset

4.2 Evaluation

4.2.1 Data sets

5. IMPLEMENTATION

6. RESULTS

- Tables with true positive, true negative, false positive and false negative rate for each classifier on each dataset
- 2. ROC curves which can be used to determine true positive for an acceptable amount of false positives for each classification technique on each dataset

- Graphs which outline difference in accuracy of each data balancing technique for each dataset
- 4. Discussion on best classification technique
- 5. Discussion on data balancing results
- 6. Outline similarities and differences in results between the two TB datasets and between TB and financial datasets and determine if they are similar enough that results for the German and Australian credit scoring datasets could be used for the two TB datasets.

7. CONCLUSIONS AND FUTURE WORK

 Likely something along the lines of utilising more temporal based data so that classification is not just done at registration but also at each check-up for example. Future work may also be the testing of more classification techniques as well as datasets from other parts of the world.

8. REFERENCES

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Table 4: Data set summary

Data set	Entries	Number of numerical features	Number of categorical features	Data balance ratio (Cured:Defau
Lima Peru TB [10]	1186	0	15	8.65:1
German Credit Scoring	1000	7	13	2.33:1
Australian Credit Scoring	690	6	8	1.25:1

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