UNIVERSITY OF TECHNOLOGY SYDNEY Faculty of Engineering and Information Technology

Stacking Ensemble Model for Liver Stiffness Classification with Imbalanced Data

by

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Certificate of Authorship/Originality

I, Mingjian Wang declare that this thesis, is submitted in fulfilment of the require-

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ABSTRACT

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Liver cirrhosis is a significant threat to humans; once the liver reaches the last stage of cirrhosis, there is no cure for it. Therefore, discovering cirrhosis in the early stage is one of the effective ways to decrease the mortality rate of cirrhosis. Besides early detection, increasing the correct cirrhosis diagnosis rate is another desirable method to avoid late treatment for patients. This thesis developed an automatic diagnosis approach to predict doctors' opinions for patients regarding the liver stiffness measurements from FibroScan tests. A model using the Stacking ensemble method was presented to build a classifier for an imbalanced liver stiffness measurement data-set. The data-set was collected from 13,418 Chinese patients who had liver cirrhosis tests by FibroScan. It recorded 30 sets of features, also provided professional doctors' opinions in Chinese. To transfer the Chinese characters to digital, we applied Jieba module in Python which is a natural language processing method to create 6 labels in classification. Each label presents one doctors' opinion. Since this data-set is highly imbalanced, sampling methods such as the undersampling method and the oversampling method are applied to solve this problem. To identify the most suitable model for the classification, we performed a study of 7 supervised learning algorithms, Logistic Regression (LR), Decision Tree (DT), Naive Bayesian (NB), K-Nearest Neighbour (KNN), Support Vector Machine (SVM), Random Forest (RF) and AdaBoost; also demonstrated the stacking models based on these supervised learning algorithms. The results demonstrated that the use of Synthetic Minority Oversampling Technique (SMOTE) oversampling technique was effective to handle the imbalanced liver data-set, and the best fitting model was constructed by using DT as meta-classifier with four base classifiers (KNN, RF, DT, SVM) in the stacking model.

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