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DECISION TREE ANALYSIS ON J48 AND RANDOM FOREST ALGORITHM FOR DATA MINING USING BREAST CANCER MICROARRAY DATASET

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ABSTRACT

Data mining which involves systematic analyses of large datasets for extracting the knowledge. Classification is considered as one of the major basic research topics that manage the data. Due to the rapid developments in microarray technology and it offer the capability to measure expression levels of thousands of genes simultaneously. study of such data helps us discovering different clinical outcomes that are caused by expression of a few predictive genes. Decision tree models help in predicting new data. In this work, we make a comparison of Decision Tree algorithms. We use two most popular algorithms namely basically J48 and Random Forest using Breast cancer microarray dataset which is available at UCI machine learning repository.

Keywords: Data Mining; Classification Techniques; J48; Decision Trees; Random Forest.

I. INTRODUCTION

Data and information have become key assets for most of the organizations [1,4]. The success of any organization depends largely on the extent to which the data acquired from business operations is utilised. In other words, the data serves as an input into a strategic decision making process, which could put the business ahead of its competitors[16]. Also, in this era, where businesses are driven by the customers, having a customer database would enable management in any organization to determine customer behaviour and preference in order to offer better services and to prevent losing them resulting better business[13,14]. Data mining is considered to be an emerging technology that has made revolutionary change in the information world. The term `data mining' (often called as knowledge discovery) refers to the process of analysing data from different perspectives and summarizing it into useful information by means of a number of analytical tools and techniques, which in turn may be useful to increase the performance of a system[3,7]. Technically, ―data mining is the process of finding correlations or patterns among dozens of fields in large relational databases‖. Therefore, data mining consists of major functional elements that transform data onto data warehouse, manage data in a multidimensional database, facilitates data access to information professionals or analysts, analyze data using application tools and techniques, and meaningfully presents data to provide useful information [12].

II. TECHNIQUES AND ALGORITHMS

Researchers find two important goals of data mining: prediction and description. First, the Prediction is possible by use of existing variables in the database in order to predict unknown or future values of interest. Second the description mainly focuses on finding patterns describing the data the subsequent presentation for user interpretation. The relative emphasis of both prediction and description differs with respect to the underlying application and technique.

2.1 Classification

Classification is the most commonly applied data mining technique, which employs a set of pre-classified examples to develop a model that can classify the population of records at large. Fraud detection and credit risk applications are particularly well suited to this type of analysis[2]. This approach frequently employs decision tree or neural network-based classification algorithms. The data classification process involves learning and classification[2,17]. In Learning the training data are analyzed by classification algorithm. In classification test data are used to estimate the accuracy of the classification rules. If the accuracy is acceptable the rules can be applied to the new data tuples[5,6]. For a fraud detection application, this would include complete records of both fraudulent and valid activities determined on a record-by-record basis. The classifier-training algorithm uses these pre-classified examples to determine the set of parameters required for proper discrimination. The algorithm then encodes these parameters into a model called a classifier. Some well-known classification models are: a) Classification by decision tree induction b) Bayesian Classification c) Neural Networks d) Support Vector Machines (SVM).

2.2 Clustering

Clustering is a technique for identification of similar classes of objects. By using clustering techniques we can further identify dense and sparse regions in object space and can discover overall distribution pattern and correlations among data attributes. Classification approach can also be used for effective means of distinguishing groups or classes of object but it becomes costly so clustering can be used as preprocessing approach for attribute subset selection and classification. For example, to form group of customers based on purchasing patterns, to categories genes with similar functionality. Some commonly used clustering methods are:

a) Partitioning Methods b) Hierarchical Agglomerative (divisive) methods c) Density based methods d) Grid-based methods e) Model-based methods

2.3 Association Rules

An Association Rule is a rule of the form milk and bread =>butter, where 'milk and bread' is called the rule body and butter the head of the rule. It associates the rule body with its head.In context of retail sales data, our example expresses the fact that people who are buying milk and bread are likely to buy butter too. This association rule makes no assertion about people who are not buying milk or bread.We now define an association rule:Let D be a database consisting of one table over n attributes {a1, a2, . . . , an}. Let this table contain k instances.The attributes values of each ai are nominal1. In many real world applications (such as the retail sales data) the attribute values are even binary (presence or absence of one item in a particular market

basket)[8,9,10]. In the following an attribute-value-pair will be called an item. An item set is a set of distinct attribute-valuepairs.Let d be a database record. d satisfies an item set X _ {a1, a2, . . . , an} if X _ d.An association rule is an implication X) Y where X, Y _ {a1, a2, . . . , an}, Y 6= ; and X \ Y = ;.The support s(X) of an item set X is the number of database records d which satisfy X. Therefore the support s(X) Y) of an association rule is the number of database records that satisfy both the rule body X and the rule head Y . Note that we define the support as the number of database records satisfying X \ Y , in many papers the support is defined as s(X\Y) k . They refer to our definition of support as support count. The confidence ˆc(X) Y of an association rule X) Y is the fraction ˆc(X) Y) = s(X\Y) s(X) .From a logical point of view the body X is a conjunction of distinct attribute-value-pairs. and the head Y is a disjunction of attribute value-pairs where X \ Y = ;.Coming back to the example a possible association rule with high support and high confidence would be i1) i2 whereas the rule i1) i3 would have a much lower support value.

III. EXPERIMENTAL STUDY AND ANALYSIS

3.1 WEKA Tool

We use WEKA (www.cs.waikato.ac.nz/ml/weka/), an open source data mining tool for our experiment. WEKA is developed by the University of Waikato in New Zealand that implements data mining algorithms using the JAVA language. WEKA is a state-of-the-art tool for developing machine learning (ML) techniques and their application to real-world data mining problems. It is a collection of machine learning algorithms for data mining tasks. The algorithms are applied directly to a dataset. WEKA implements algorithms for data pre-processing, feature reduction, classification, regression, clustering, and association rules. It also includes visualization tools. The new machine learning algorithms can be used with it and existing algorithms can also be extended with this tool.

3.2 Dataset Description

We performed computer simulation on a Breast- cancer dataset available UCI Machine Learning Repository [11,15]. . The features describe different factor for cancer reoccurrence.. The dataset contains 286 instances and 10 attributes.Figure 3.1 shows the attributes of Breast-cancer Dataset.

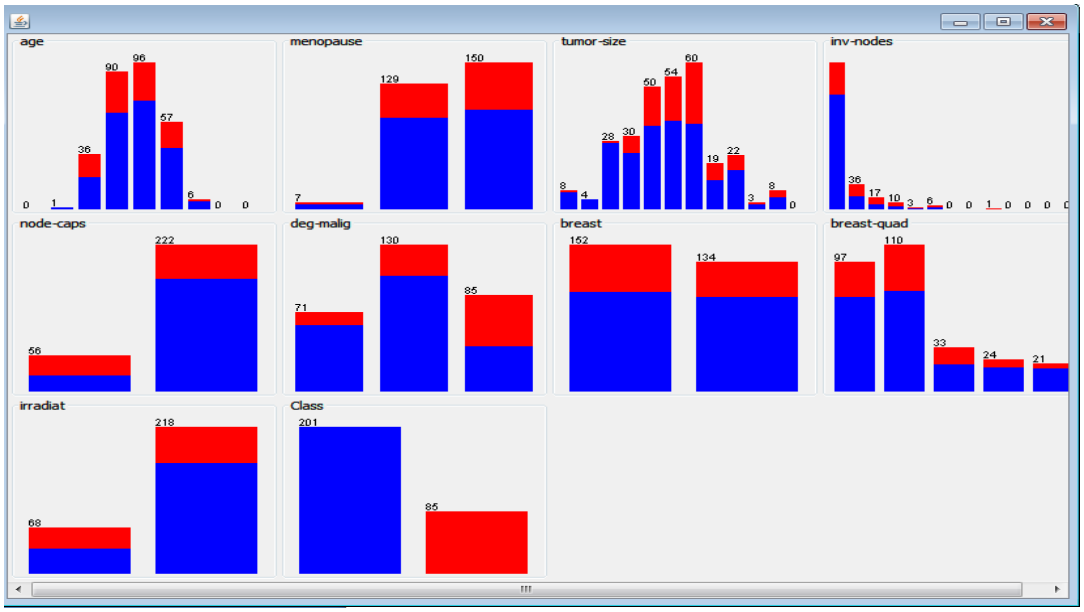

Figure 3.1-Attributes of Breast-Cancer Dataset

3.3 Results

Accuracy rate of Decision tree((J48) is 75.52 and accuracy of Decision tree((Random Forest) is 69.58

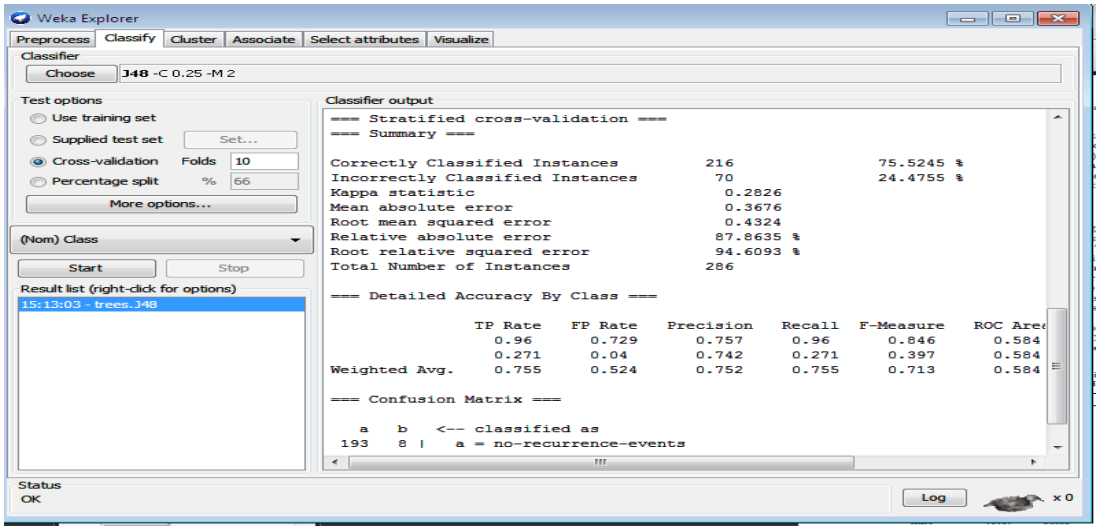

Figure 3.2.Performance of Decision tree((J48).

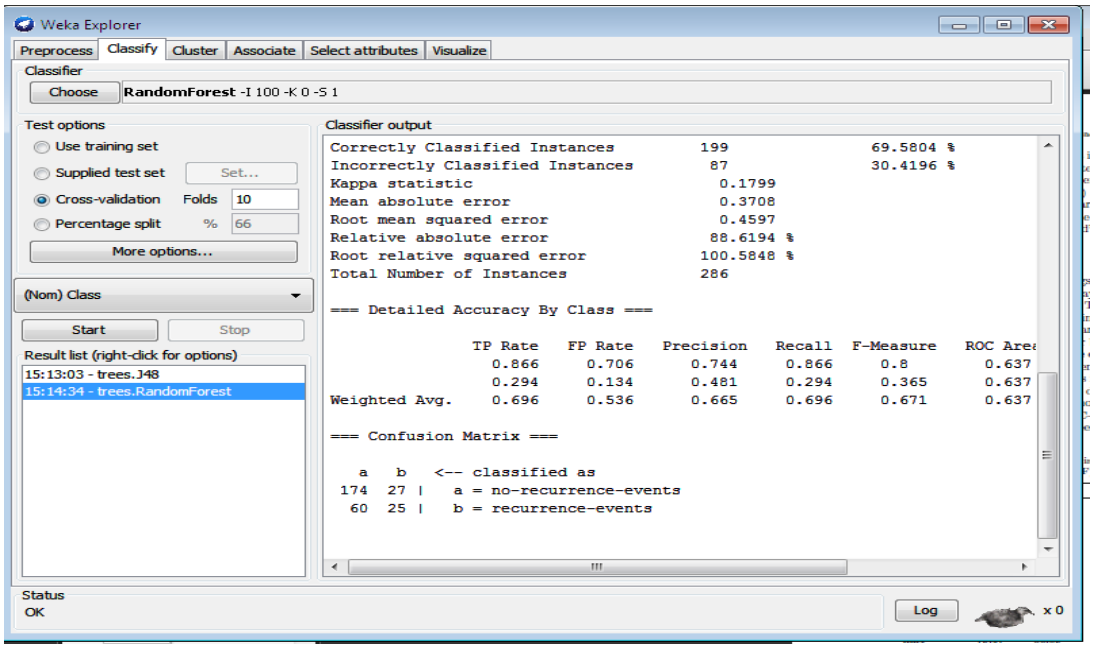

Figure 3.3. Performance of Decision tree(Random Forest).

Figure 3.1 and figure 3.2 shows the performance of decision tree using J48 and Random forest

IV. CONCLUSION

In this paper we conducted an experiment to find the accuracy of Breast cancer data on the predictive performance of different decision tree classifiers. We select two popular classifiers considering their qualitative performance for the experiment. After analysing the quantitative data generated from the computer simulations, we find that the general concept of improved predictive performance of all above classifiers but Random Forest performance is not significant.

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