

# INTERPRETATION TRAINED NEURAL NETWORKS BASED ON GENETIC ALGORITHMS

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## **ABSTRACT**

*In this paper, constructive learning is used to train the neural networks. The results of neural networks are obtained but its result is not in comprehensible form or in a black box form. Our goal is to use an important and desirable model to identify sets of input variable which results in a desired output value. The nature of this model can help to find an optimal set of difficult input variables. Accuracy. Genetic algorithms are used as an interpretation of achieving neural network inversion. On the other hand the inversion of neural network enables to find one or more input patterns which satisfy a specific output. The input patterns obtained from the genetic algorithm can be used for building neural network system explanation facilities.*

## **KEYWORDS**

*Neural Networks, Genetic Algorithms, Constructive Learning, Accuracy.*

## **1. INTRODUCTION**

Extracting classification rules is a procedure of extraction of information and knowledge that are hid in data in the form of classification rules, unknown by people and potentially useful from a large quantity of data with multiple characteristics that is uncompleted, containing noise, fuzzy and random. As a form of cross-discipline direction that syncretises numerous disciplines as well as database technology, statistics, artificial neural networks, knowledge acquirement and information extraction, nowadays data mining has becomes one of the most front research direction in the international realms of information-based decision making.

Many methods have been proposed to construct the networks; the most important methods are destructive, constructive, and genetic algorithms [2, 33].

Constructive learning of neural networks adds nodes or links to the structure of the network during training. Generally, it starts with a network with no hidden units, which is trained for a period. Then with no changing in the presenting weights, more new hidden nodes are added to the network, the training starts again, and so on. Many variations are feasible, including different patterns of links and schemes for freezing and melt weights. The most familiar constructive learning algorithm is cascade correlation [1, 28, 31] of which many variations are possible [3, 4]. Various other constructive algorithms are summarized in [5]. An approach for adaptively and unconventionally constructing a multilayer feed forward neural network (FNN) is introduced in [29]. Regression problems has been demonstrated as a review for the constructive learning algorithms in feedforward neural networks in [27]. Constructive learning algorithms were used to handle multi-category classification with convergence to zero classification errors [2].

Extracting comprehensible rules from neural networks is still needing efforts because that multi-layered artificial neural network are often regarded as "black boxes", which map input data into a

class through a number of mathematically weighted connections between layers of neurons. In order to bear this limitation, the hypothesis generated by artificial neural networks could be transferred into a more comprehensible representation; these conversion methods are known as rule extraction algorithms. Many researchers dealt with this problem such as in [8, 9, 13, 14, 15, 21, 34]. They introduce methods for discovering M-of-N rules from trained artificial neural network for the standard three layered feed forward networks.

In [22] an approach named REFNE is proposed to improve the comprehensibility of trained artificial neural network ensembles that performs classification tasks. REFNE utilizes the trained ensembles to generate instances and then extract symbolic rules from those instances. A novel strategy using genetic algorithms to search for symbolic rules in a trained artificial neural network is introduced in [10]. Many approaches have been introduced via trained artificial neural network to extract accurate and comprehensible rules from databases based on genetic algorithm [45, 24, 25]. These methods do not modify the training results.

Genetic Algorithm is a search tool which is usually employed when the search-space in question is large and rather unknown. Genetic Algorithms are stimulated by the means of natural choice where stronger individuals are possible the winners in a competing environment. In our work, we refine GA to deal with the complex topology, independently from how complex the topology is, to extract a set of highly accurate, comprehensible and simple rules [6, 7].

Earlier approaches based on an extensive analysis of network links and output values have already been confirmed to be obdurate in that the scale-up issue increases exponentially with the number of nodes and links in the network. A novel approach using genetic algorithms to search for comprehensible rules in a trained neural network is demonstrated in this paper. Preliminary experiments concerning classification are reported here, with the results representing that our proposed approach is winning in extracting rules. While it is accepted that additional work is required to persuasively demonstrate the dominance of our approach over others, there is nonetheless enough novelty in these results to justify early distribution.

For our experimentation, we will use a wide set of data sets which are commonly used in classification tasks. To check the results obtained, we compare them with some well-known methods of classification such as Bagging with a tree based approach, c4.5 with reduced error pruning (REP), Nave Bayesian, and RBF Networks.

A wide variety of methods are now available, recently reviewed in [11, 12, 23, 15]. Tickle et al [38] revisits the Andrews [11] classification of rule extraction methods and emphasise distinction between decompositional and pedagogical approaches. Rule extraction methods typically start by finding a smallest network, in terms of number of hidden units and overall connectivity. Setiono [39] for example adds penalty terms to the error function to bias back propagation like training towards such sparse networks. The next simplification, the key feature of the method, is to quantize or cluster the hidden unit activations. It is then promising, link by link, to extract combinations of inputs which will activate each hidden node, singly or together and thus output generates rules. This unit by unit analysis characterizes the decompositional approach. Although it can yield exact representations, the computational time may grow exponential with number of inputs (attributes) as noted by Tickle et al. [15] for decompositional algorithms such as Subset and KT. Taha and Ghosh [40] suggest for binary inputs such as our data generating a truth table from the inputs and simplifying the resultant Boolean function. But this simplification is itself combinatorially nasty and thus the method works only for small networks. They also refine the Liu and Setiono [41] methods using linear programming.

The rest of the paper is organized as follows. The structure of the networks is presented in Section 3. In Section 4, rules from trained artificial neural network based on genetic algorithm

will be discussed. The dataset used its representation of the neural networks, illustrative example for rules extraction algorithm to some public databases and comparison to other methods presented in Section 5. Description of the results and discussion will be presented in Section 6. Finally, in Section 7 we will outline the conclusions of this paper.

## 2. RULE EXTRACTION PHASE

In this section we will introduce an approach which extracts accurate and comprehensible rules from databases via trained artificial neural network using genetic algorithm. This method does not modify the training results. After using constructive learning we have the following ANN topology which is shown in Figure 1. It consists of  $L$  nodes in the input layer,  $h$  nodes in the hidden layer and  $n+1$  node in the output layer. Also, two groups of weights can be obtained. The first set,  $W_{ij}$  includes the links between the input node  $i$  and the hidden node  $j$ . The second set,  $V_{jk}$  includes the links between the hidden node  $j$  and the output node  $k$ . A sigmoid function as an activation function has been used in the hidden and output nodes of the ANN.

The total input to the  $j$ th hidden node,  $IHN_j$ , is given by

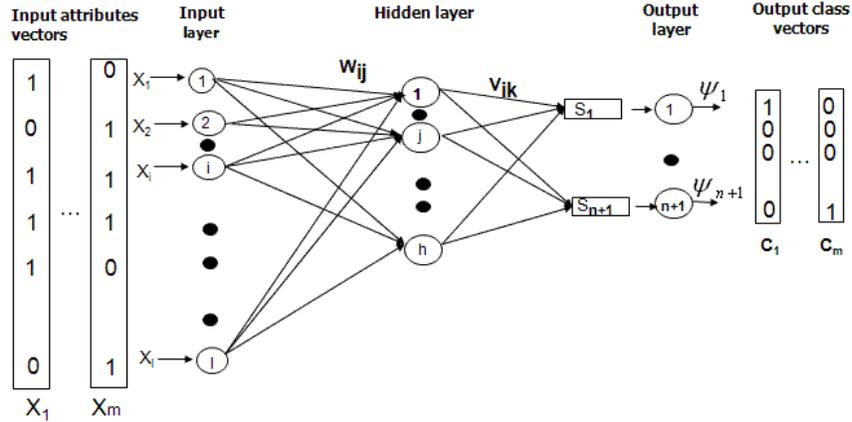


Figure 1: The structure of the network.

$$IHN_j = \sum_{i=1}^l x_i W_{ij} \quad (1)$$

The output of the  $j$ th hidden node,  $OHN_j$ , is given by

$$OHN_j = \frac{1}{1 + e^{-IHN_j}} \quad (2)$$

The total input to the  $k$ th output node,  $ION_k$ , is given by

$$ION_k = \sum_{j=1}^J V_{jk} \frac{1}{1 + e^{-IHN_j}} . \quad (3)$$

So, the final value of the  $k$ th output node,  $\psi_k$ , is given by

$$\psi_k = \frac{1}{1 + e^{-ION_k}} \quad (4)$$

The function,  $\psi_k = f(x_i, W_{ij}, v_{jk})$  is an exponential function in  $x_i$  since  $W_{ij}, V_{jk}$  are constants. Its maximum output value equal one.

We have to note that an input vector,  $X_m$ , belongs to a  $class_k$  **iff**  $\psi_k \in C_m = 1$  and all other elements in  $C_m = 0$ .

Consequently, for extracting relation (rule) between the input attributes,  $X_m$  relating to a specific  $class_k$  one must find the input vector, which *maximizes*  $\psi_k$ . This is an optimization problem and can be stated as:

$$\text{Maximizes}[\psi_k(x_i)] \quad (5)$$

subjected to:  $x_i \in (0 \text{ or } 1)$ .

Since the objective function  $\psi_k(x_i)$  is nonlinear, it is a nonlinear integer optimization problem, and the constrains are binary so. Such problems with very large search spaces and strong non-linearity can be solved using GA. To use the general schema of GA first of all it is necessary to determine a representation of individual, and the way of coding it in the chromosome. Genetic operators (crossover and mutation) act on selected individuals. Crossover acts with assumed relatively high probability. Next, every attribute is mutated with assumed, small probability. The final effect of population proceeding creates the next generation.

## 2.1. Constructive Learning Algorithm

Typically, constructive learning begins with a network with one hidden unit; one or more new hidden units are added to the network, training resumes, and so on. The major steps of our algorithm can be shortened as follows.

1. Input the training set of patterns.
2. Initial set of weights and thresholds are initialized randomly from the interval [-1, 1].
3. Start learning with one unit in the hidden layer.
4. Update the weights and thresholds to minimize the objective function by any optimization algorithm.

5. Terminate the network training with this number of hidden units when a local minimum of the objective function has been reached.
6. If the desired accuracy is not reached, increase the number of hidden units with random weights and thresholds and go to step 4, otherwise, go to step 7.
7. Stop.

Since the objective function  $\psi_k(x_i)$  is nonlinear and the constraints are binary so, it is a nonlinear integer optimization problem. The genetic algorithm can be used to solve it since genetic algorithm is a robust search method and is suitable for difficult problems - e.g. problems with very large search spaces and strong nonlinearity. To use the general schema of GA first of all it is necessary to determine a representation of individual, and the way of coding it in the chromosome. Genetic operators (crossover and mutation) act on selected individuals. Crossover acts with assumed relatively high probability. Next, every attribute is mutated with assumed, small probability. The final effect of population proceeding creates the next generation.

### 3. RESULTS & DISCUSSION

The performance of the learning algorithm which is evaluated by the accuracy (%) on the monk's, weather's and breast cancer's problems. Where accuracy is how close a measured value is to the actual (true) value and Precision is how close the measured values are to each other. In other hand, the precision of an object value is a measure of the reliability of the experiment, or how reproducible the experiment is.

The accuracy of an object value is a measure of how closely the experimental results agree with a true or accepted value. Both accuracy and precision are conditions used in the fields of science, engineering and statistics. The accuracy of the classifier will be evaluated using the following parameters:

*Precision* is the percentage of true positives (TP) compared to the total number of cases classified as positive events.

$$Precision = \frac{TP}{TP + FP} \times 100\% , \quad (6)$$

where FP represents false positives. According to Cios and Moore [46], "This measurement is very popular in machine learning and pattern recognition communities". To better understand the performance of the learning algorithm, we will define a number of other measurements. Let us begin by examining a contingency Table 1. Contingency tables contain four variables:

A *true* – positive (TP) occurs when a classifier correctly classified class1.

A *true* – negative (TN) occurs when a classifier correctly classified class2.

A *false* – positive (FP) occurs when a classifier incorrectly classified class1.

A *false* – negative (FN) occurs when a classifier incorrectly classified class2.

Another measurement of performance, frequently used in conjunction with precision, is accuracy.

Table 1: Contingency table

Test results	Disorder Present	Disorder Absent	Total
Positive	TP FN	FP TN	TP+FP FN+TN
Total	TP+FN	FP+FN	

Accuracy is the number of correctly classified cases compared to the total number of cases presented to a system. It is defined by the following equation:

$$Accuracy = \frac{TP + TN}{TP + TN + FP + FN} \times 100\%, \quad (7)$$

perhaps the most common measurements in classification problems are sensitivity and specificity.

They are statistical measures of the performance of a binary classification test.

The *sensitivity* measures the amount of real positives which are rightly identified as such (e.g. the percentage of sick people who are identified as having the condition)

The *specificity* measures the proportion of negatives which are correctly identified (e.g. the percentage of well people who are identified as not having the condition).

They are defined in the following equations:

$$Sensitivity = \frac{TP}{TP + FN}, \quad (8)$$

where a sensitivity of 100% means that the test recognizes all sick people as such.

$$Specificity = \frac{TN}{TN + FP}, \quad (9)$$

where a specificity of 100% means that the test recognizes all healthy people as healthy.

The results which are given in Fig. 2 show the accuracy of our method compared to other classifiers like Bayesian, RBF Network, etc. Results for those classifiers were constructed using Weka experimenter, Weka is a collection of machine learning algorithms for data mining tasks, the algorithms can either be applied directly to a data set or called from your own Java code. Weka contains tools for data pre-processing, association rules, classification, regression, clustering, and visualization. It is also well-suited for developing new machine learning schemes <http://www.cs.waikato.ac.nz/ml/weka/>. It is clear in most of the data sets the high performance of our model when compared to other classifiers, except for RBF Network that gave equal accuracy for the weather data set and for monk1 data set and Bagging algorithm gave equal accuracy. In the Monk3 data set, however, there was a significant decrease in accuracy rate compared to Naive Bayesian and RBF Network algorithms. However, in this data set our rule extraction method extracted more simple and comprehensible rules in comparison with other algorithms.

#### 4. CONCLUSION

It is well known that the knowledge acquired by ANNs is generally incomprehensible for humans. This reality was a major drawback in this paper, in which ultimately understandable patterns (like classification rules) are very important.

We apply genetic algorithms to extract approximate rules from neural networks. The genetic algorithm approach described here determines rules better than those found from various decision tree based methods.

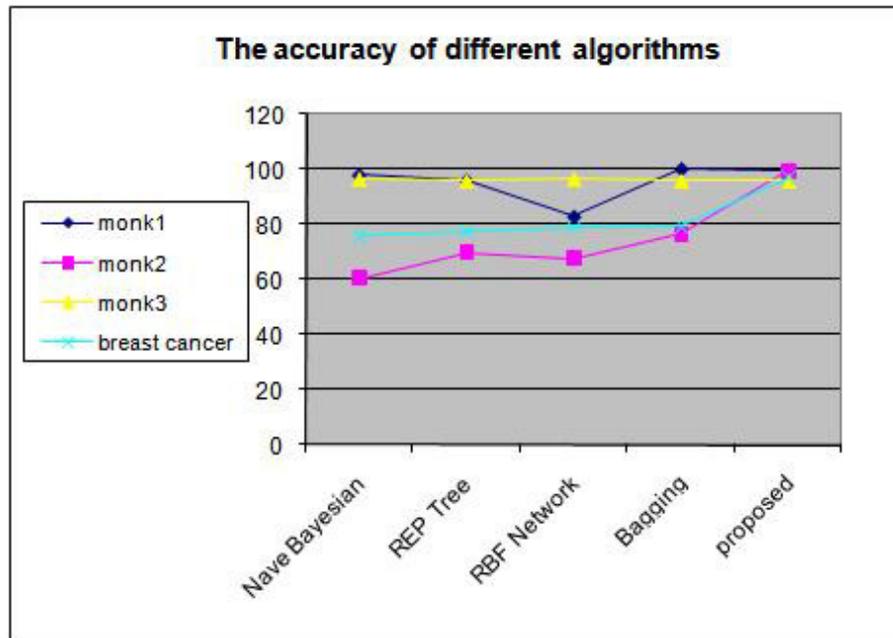


Figure 2: The accuracy of different algorithms.

It is simple to implement, but requires considerable computing resources for the large neural networks of the present paper. As such its greatest value is for determining heuristic rules for medium term use by practitioners who welcome the intuitive description that rules provide.

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