Evolutionary Multivariate Kernal Svm Prediction Method for Classification

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ABSTRACT : Thyroid disorders are common among the world wide population. This disorders posses' significant problems among Indians. Research studies shows that nearly 32% of Indian population suffers from various thyroid disorders. This paper deals with the thyroid data set which in turn classify into three groups as hyper thyroisidism, hypothyroidism and normal. The American Thyroid Association reported twelve percent of their citizens suffer from thyroidism in which 60% population are unaware of their conditions. Above statistics implies the classification of thyroid disorder is crucial in global perspective too. The thyroid data set are collected from UCI repository and it is multivariate type with 21 attributes. With the 21 attributes only 10 attributes are selected based on their rank. Hybrid Differential Evolution Kernel Based SVM algorithm is used to classify the data set. It takes around 30 epochs to stabilize the errors. The classification accuracy is observed to be 67.97%

Keywords: Curse of Dimensionality , Classification ,Evolutionary algorithm , Multivariate data type ,Thyroid Data Set

I. INTRODUCTION

According to the statistics published by the WEEK magazine in India, it reported that the percentage of people suffered from thyroid disorders are in rise. The above report based on the results derived from survey conducted by Indian Thyroid Society. The statistics shows that women of age between 18 to 35 are more prone to thyroid disorders compared to men. The thyroid disease is ranked ninth in comparison to other syndromes like diabetic, Blood Pressure, respiratory Problems, insomnia and heart related issues. A statistical survey revealed that fifty percent of the surveid population are aware of thyroid disorder; knows about diagnostic test and remedies for this health issues. Other research work on medical data set depicits , C.V.Subbulakshmi et.al proposed hybrid algorithm self-regulated particle swarm optimization (SRPSO) algorithm with the extreme learning machine (ELM) for classification problems. To optimize the input weights and hidden biases and minimum norm least-square scheme, an improved PSO is used to analytically determine the output weights[1]. Tapas Ranjan and Subhendu Kumar’s research work on decision trees J48, Naive Bayes, ANN, ZeroR, 1BK and VFI algorithm to classify these diseases and compare the effectiveness, correction rate among them. Performance of the mentioned classifiers shows Multilayers Perceptron gives oves all best classification result whereas Naive Bayes performance is not significant [2].

This research work classifies the thyroid disorder with the help of proper features selection from the dataset which in turn leads to accurate results which helps physician in diagnosis of thyroid.

II. METHODOLOGY

To classify the data into three classes like Hyperthyroid,Hyperthyroid and Normal, get the dataset from any repository or UCI repository, in this nearly eight thousand data set with twenty one attributes are taken into account in which fifteen attributes have continous values and six are discrete type. First it undergoes a preprocessing stage that is followed by feature selection. It is carried out to overcome the curse of dimensionality issues using hybrid algorithm Differential evolution with Support Vector Machine. The subset of the data set is classified using kernal based SupportVector Machine algorithm based on the fitness calculation where error stabilization is used to measure the fitness.

2.1 Preprocessing Step

The accuracy of the results greatly depend on the type of data set, which is free from the problems like redundant data, missing values and noisy data etc. In this research Not a Number and missing value constraints are checked. Data set contains both the types of values such as continuous and discrete.

2.2 Feature Selection Method

Challenges in using medical database are its volume,heterogenous values and privacy issues. Subset selection methods or ranking methods are used in feature selection to identify the relevant attributes. This step helps to overcome curse of dimensionality issues by removing irrelevant and redundant data, improve the quality of data and its accuracy. The algorithms used for Feature selection are classified as

1)The filter model 2) The wrapper model

The wrapper Model gives good accuracy but more complex computational procedures have to be carried out. The proper classification depends on the selection of data. The above process leads to over fitting of data. Various wrapper methods are available for each domain.

2.3 Differential Evolution

Differential Evolution is an effective search method for continuous optimization problems. The steps involved are mutation, crossover and selection. The process of differential evolution differs from other evolutionary algorithm is the mutation that perturbs the selected vector according to the scaled difference of the other two members of the population. The DE algorithm:

Revised Manuscript Received on June 01, 2020.

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The idea behind the kernel trick is to map the data into a feature space and construct a linear classifier. In this space, non-linear classifiers in the original space can be constructed.

### 3.2 Radial Basis Kernel Function

In this work, data has high dimensional feature and it is non-linear. In order to make the data linear and reduce the dimensionality (Curse of Dimensionality), RBF kernel is used. The RBF kernel on two samples \( x \) and \( x' \), represented as feature vectors in some input space, is defined as [3]

\[
K(x, x') = \exp\left(-\frac{||x - x'||^2}{2\sigma^2}\right)
\]

The squared Euclidean distance between the two feature vectors, \( \sigma \) is a free parameter. An equivalent, but simpler, definition involves a parameter \( \gamma = \frac{1}{2\sigma^2} \):

\[
K(x, x') = \exp(-\gamma ||x - x'||^2)
\]

Since the value of the RBF kernel decreases with distance and ranges between zero (in the limit) and one (when \( x = x' \)), it has a ready interpretation as a similarity measure [5]. The feature space of the kernel has an infinite number of dimensions; for \( \sigma = 1 \), its expansion is [4]:

\[
\exp\left(-\frac{1}{2
\|x-x'||^2}\right) = \sum_{k=0}^{\infty} \left(\frac{1}{2\sigma^2}\right)^k \exp\left(-\frac{k}{2\sigma^2}\right)
\]

### IV. RESULTS AND DISCUSSION

Matlab is used to carried out this work. The figure 1 shows the framework of the model where the dataset is loaded. Figure 2 and 3 displays the preprocessed data set and the list of features selected, the same figure also represent the error stabilization criteria.
Fig 3: List the Selected Features and Error Stabilization

4.1 Performance Metrics

The evaluation of the classification is essential. The performance metrics that are included in this methodology is elaborated as follows and the output is shown in figure 4. The results are depicted in confusion matrix to differentiate the true and false situation of the medical conditions. The metric prevalence (Pr) for the considered group.

\[ Pr = \frac{TP + FN}{TP + FN + TN + FP} \]

Table 2: Confusion Matrix

| True situation | Test result |
|----------------|------------|
| 0              | True negative (TN) | False positive (FP) |
| 1              | False negative (FN) | True positive (TP) |

The metric sensitivity (Se), specificity (Sp). Calculated as

\[ Se = \frac{TP}{TP + FN} \quad SP = \frac{TN}{TN + FP} \]

The accuracy is measured by

\[ ACC = Pr \times Se + (1 - Pr) \times Sp \]

Fig 4: Classification and Evaluation metrics

It takes nearly thirty epochs to error stabilization Then data is classified into three class labels as depict in figure 4. 67.97% accuracy level is reached in this classification. ROC for the method is shown in figure 5.

Fig 5: Receiver Operating Characteristic (ROC)

V. CONCLUSION

The suggested work classify data into respective class labels of thyroid. The evaluation metrics are used to evaluate the accuracy which shows as 67.97%. It doesn’t show better result as compared with other hybrid models like differential evolution with naïve bayes. The processing time for this model is higher and it takes nearly thirty runs to compute the fitness for eight thousand data. The future work can be test with genetic algorithm with SVM for multivariate data set.

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