AN ANN APPROACH FOR NETWORK INTRUSION DETECTION USING ENTROPY BASED FEATURE SELECTION

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ABSTRACT

With the increase in Internet users the number of malicious users are also growing day-by-day posing a serious problem in distinguishing between normal and abnormal behavior of users in the network. This has led to the research area of intrusion detection which essentially analyzes the network traffic and tries to determine normal and abnormal patterns of behavior. In this paper, we have analyzed the standard NSL-KDD intrusion dataset using some neural network based techniques for predicting possible intrusions. Four most effective classification methods, namely, Radial Basis Function Network, Self-Organizing Map, Sequential Minimal Optimization, and Projective Adaptive Resonance Theory have been applied. In order to enhance the performance of the classifiers, three entropy based feature selection methods have been applied as preprocessing of data. Performances of different combinations of classifiers and attribute reduction methods have also been compared.

KEYWORDS:

Intrusion detection, Artificial Neural Network, attribute reduction

1. INTRODUCTION

Information security is a serious issue while using computer networks. There has been growing number of network attacks which has challenged application developers to create confidence among the users. Researchers have looked at the security concerns from different perspectives. Intrusion Detection System is one such attempt which tries to analyze network traffic in order to detect possible intrusive activities in a computer network. There are two types of intrusion detection systems: misuse detection system and anomaly detection system. While the former is capable of detecting attacks with known patterns/signatures, the latter is augmented with the ability to identify intrusive activities that deviate from normal behavior in a monitored system, thus can detect unknown attacks. A range of techniques have been applied to analyze intrusion data and build systems that have higher detection rate.

Mohammadreza Ektefa et al. [1] proposed machine learning algorithms for intrusion detection system which compared the performance of C4.5 algorithm with SVM in detecting intrusions and the results revealed that C4.5 performed better than SVM in terms of intrusion detection and false alarm rate. Juan Wang et al.[2] have used the C4.5 decision tree classification to build a rule base that can be used for an effective intrusion detection system. Zainal et al.[3] demonstrated the application of ensemble of different learning algorithms by setting proper weights to the individual classifiers used in the classification model. They have observed that there was

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improvement in attack detection and considerable drop in false alarm rate. Sung and S.Mukkamala [4] have proposed an approach for IDS with the use of Rank based feature selection and have shown that Support Vector Machines (SVMs) perform much better than Artificial Neural Networks (ANNs) in terms of speed of training, scale and accuracy. Lin NI, Hong Ying Zheng [5] have attempted to build an intrusion detection system using unsupervised clustering and Chaos Simulated Annealing Algorithm. Rung-Ching Chen et al. [6] have proposed a hybrid approach by combining Rough Set Theory(RST) for feature reduction and Support Vector Machine(SVM) for classification. Amir Azimi Alastic et al [7] formalized SOM to classify IDS alerts to reduce false positives. Alert filtering and cluster merging algorithms were used to improve the accuracy of the system; SOM was used to find correlation between alerts.

2.PROPOSED ANN BASED HYBRID CLASSIFICATION MODEL

In this section we present our proposed model for classifying intrusion data in order to build an efficient intrusion detection system which can exhibit low false alarm rate and high detection rate. The model consists of two major layers as depicted in figure1.In the first layer irrelevant and redundant features are removed using three entropy based feature selection methods viz., Information Gain, Gain Ratio, Symmetrical Uncertainty. In the next layer the reduced data set is classified using four artificial neural network based techniques viz., Radial Basis Function Network(RBFN), Self-Organizing Map (SOM), Sequential Minimal Optimization(SMO), Projective Adaptive Resonance Theory (PART). Further, we have used the 10-fold cross validation technique for training and testing of the model. We evaluate the performance of the model using certain standard criteria.



Figure 1 ANN based Classification Model

3. METHODOLOGY

3.1. Radial Basis Function Network (RBFN)

Radial Basis function (RBF) network is a nonlinear hybrid network which contains an input layer, a single hidden layer and an output layer [8]. The input layer accepts n number of inputs; the hidden layer consists of m radial basis functions and the output layer produces the response with the help of a linear additive function. The input neurons are linear, i.e., they pass the input to the hidden neurons without any processing. Using radial basis function the hidden neuron computes the signal and passes on these signals through weighted paths to the linear output neuron which sums them up and generates an output signal.



Figure 2 Structure of RBF Network

The neurons of the hidden layer are associated with a linear threshold activation function which produces the network output as:

$$y_i = \sum_{j=0}^{m} w_{ij} \Phi_j(x)$$
(1)

where w_{ij} represents the weight of the connection between the hidden neuron j and the output neuron I and $\Phi_j(x)$ is the radial basis function applied at neuron j. We have used the following Gaussian function:

$$\Phi(\mathbf{x}) = \exp(\frac{-(x-c)2}{2\sigma^2}) \ \sigma > 0, \qquad \mathbf{x}, \mathbf{c} \in \mathbf{R} \quad \dots \dots \dots \dots (2)$$

The success of this model depends on determining the most suitable value for the parameter c [9]. The process begins by training an unsupervised layer which tries to find the Gaussian centers and the widths from the input data. During the unsupervised learning, the width of the Gaussians is computed based on the centers of their neighbors. The output of this layer is computed from the input data weighted by a Gaussian mixture.

3.2 Self Organizing Map (SOM)

The Self-Organizing Map (SOM) is a competitive network where the main objective is to transform an input data set of arbitrary dimension to one- or two-dimensional topological map [10]. SOM is motivated by the way information is processed in the cerebral cortex in human brain. The model was first proposed by the Finnish Professor Teuvo Kohonenand, thus referred to

as Kohonen map. SOM is an efficient technique to discover the underlying structure, e.g. feature map of the input data set by building a topology preserving map that describes neighborhood relations of the points in the data set [10]. SOM transforms a high dimensional input data domain to a low dimensional array of nodes. The SOM array is essentially a fixed size grid of nodes. Here, the training uses a competitive learning method wherein the neuron having a weight vector that is close to the input vector is adjusted towards the input vector. Such a neuron is referred to as the "winning neuron" or the Best Matching Unit (BMU). Next, the weights of the neurons close to the winning neuron are also adjusted. However, the magnitude of the change in each case depends on the distance from the winning neuron [11].

Let the real vectors $X = \{x_1, x_2, x_3, \dots, x_r\}$ represent the input data, and a parametric real set of vectors $M_i = \{m_{i1}, m_{i2}, \dots, m_{ik}\}$ be associated with each element i of the SOM grid where both X and $M_i \in \mathbb{R}^n$. A decoder function, $d(X, M_i)$ defined on the basis of distance between the input vector and the parametric vector is used to define the image of the input vector onto the grid. One can use either the Manhattan or the Euclidean distance metric as the decoder function. The BMU is denoted as the index c of the node with a minimum distance from the input vector:

$$c = \arg \min \{ d(X, M_i) \}....(3)$$

SOM demands that M_i be shifted towards the order of X such that a set of values { M_i } is obtained as the limit of convergence of the equation:

$$m_i(t+1) = m_i(t) + \alpha(t)^* [x(t) - m_i(t)]^* H_{ic}$$
(4)

where H_{ic} is a neighborhood function which models the interconnections between the nodes and is usually a Gaussian function which decreases with distance from the winner node c. The α (t) is the learning rate of the system.

3.3 Sequential Minimal Optimization(SMO)

The SMO algorithm is a specialized optimization approach for the SVM quadratic program. It combines the sparse nature of the support vector problem and the simple nature of the constraints in the Support Vector Machine Quadratic Programming (SVMQP) to reduce each optimization step to its minimum form[12]. SMO decomposes a large quadratic programming problem into a series of smaller quadratic programming problems. These small QP problems are solved analytically. The amount of memory required for SMO is linear in the training set size, which gives the ability to handle very large training sets.

Selecting *a* parameters

The SMO algorithm selects two values for the α parameters, viz., α_i and α_j , and optimizes the objective value for both α_i and α_j . In case of large data sets the values of α_i and α_j are critical as there can be m(m - 1) possible choices for α_i and α_j . Thus, the efficiency of SMO algorithm depends on the heuristics for determining α_i and α_j to maximize the objective function.

Optimizing α_i and α_j

First the algorithm finds the bounds L and H such that $L \le \alpha_j \le H$ must hold in order to satisfy the constraints $0 \le \alpha_j \le C$.

- If $y^{(i)} \neq y^{(j)}$, $L = max(0, \alpha_i \alpha_j)$, $H = min(C, C + \alpha_i \alpha_j)$ (5)
- If $y^{(i)} = y^{(j)}$, $L = max(0, \alpha_i + \alpha_j C)$, $H = min(C, \alpha_i + \alpha_j)$ (6)

We intend to find α_j so as to maximize the objective function. The optimal α_j is given by

where $E_k = f(x^{(k)} - y^{(k)})$ and $\eta = 2 \langle x^{(i)}, y^{(j)} \rangle - \langle x^{(i)}, y^{(i)} \rangle - \langle x^{(j)}, y^{(j)} \rangle$

 E_k is the error between the SVM output on the k-th example and the true label $y^{(k)}$.

We clip α_i to lie within the range [L, H] is

$$\begin{aligned} \alpha_{j} &= \{H, & \text{if } \alpha_{j} > H \\ &\{\alpha_{j}, & \text{if } L \leq \alpha_{j} \leq H \\ &\{L, & \text{if } \alpha_{j} < L \end{aligned}$$

The value of α_i can be calculated using the formula

where $\alpha_i^{(old)}$ is the value of α_i before optimization.

Computation of b Threshold

We select the threshold b such that the Karush-Kuhn-Tucker (KKT) [12] conditions are satisfied for the i-th and j-th examples. The threshold b_1 is valid if $0 < \alpha_i < C$ and is given by

 b_2 is valid if $0 < \alpha_i < C$ and is given by

If $0 < \alpha_i < C$ and $0 < \alpha_i < C$ then both the thresholds are valid, and they will be equal.

Let $b = (b_1 + b_2)$, the complete equation for b is

 $b = \{ b_1 & \text{if } 0 < \alpha_i < C \\ \{ b_2 & \text{if } 0 < \alpha_j < C \\ \{ (b_1 + b_2) / 2 \text{ otherwise} \} \}$

SMO Algorithm

Input:

 $\begin{array}{l} C: Regularization Parameter\\ Tol: Numerical tolerance\\ Max_Passes: Maximum number of times to iterate over α's without changing the training data (x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \ldots, (x^{(m)}, y^{(m)}) \end{array}$

Output:

```
A \in R^m: Lagrange multipliers for solution
b \in R: Threshold for solution
     Initialize \alpha_i = 0, for all i, b=0
     Initialize passes = 0
     while(passes <max_passes)</pre>
     Num_changed_alphas = 0
fori = 1, 2,....m
Calculate E_i = f((x^{(i)}, y^{(i)}))
if ((y^{(i)}E_i < tol \&\& \alpha_i < C || (y^{(i)}E_i > tol \&\& \alpha_i > 0))
Select j \neq i randomly,
Calculate E_j = f((x^{(j)}, y^{(j)}))
Save old \alpha's: \alpha_i^{(old)} = \alpha_i, \alpha_j^{(old)} = \alpha_j
     Compute L and H by (Eq. 1) and (Eq. 2)
     if(L=H)
     continue to next i.
     Compute \eta by (5).
     if(\eta >= 0)
     continue to next i.
     Compute new values for \alpha_j using (Eq. 3) and (Eq. 6) if( ~\alpha_j-\alpha_j^{(old)}l\!<\!10^{-5} )
     Continue to next i.
Determine value for \alpha_i using (Eq. 7).
Compute b<sub>1</sub>, and b<sub>2</sub> using (Eq. 8) and (Eq. 9) respectively.
Compute b by (Eq. 10).
Num changed alphas : = Num changed alphas + 1.
end if
end for
if (Num_changed_alphas == 0)
passes := passes + 1
else
passes := 0
end if
end while
```

3.4 Projective Adaptive Resonance Theory (PART)

Projective Adaptive Resonance Theory (PART) is an innovative neural network architecture proposed to provide a solution to high-dimensional clustering problems [13]. The architecture of PART is based on adaptive resonance theory (ART) neural network which is very effective in self-organized clustering in full dimensional space. ART focuses on the similarity of patterns in the full dimensional space and may fail to find patterns in subspaces of higher dimensional space. It is practically infeasible to find clusters in subspace clustering in all possible subspaces and then compare the results thus obtained due to the large number of possible subspaces of the order of $2^m - 1$ for large values of m. PART solves this problem by introducing a selective output signaling mechanism to ART.

The basic architecture of PART is similar to that of ART which is very effective in self-organized clustering in full dimensional spaces[14][15]. The PART architecture consists of a comparison layer F_1 and a competitive layer F_2 . The F_2 layer of PART follows the winner-take-all paradigm. The F_1 layer selectively sends signals to nodes in F_2 layer. A node in F_1 layer can be active relative to only few nodes in F_2 layer which is determined by a similarity test between the corresponding top-down weight and the signal generated in the F_1 node. This similarity test plays an important role in the subspace clustering of PART.



Figure 3 PART Architecture

The nodes in F_1 layer (Comparison layer) are denoted by u_i , $i = 1, 2, \ldots, m$. Nodes in F_2 layer (Clustering Layer) are denoted by u_j , $j = m+1, \ldots, m+n$. Further, the activation of F_1 layer node u_i is denoted by y_i and the activation of F_2 layer node u_j by y_j . Similarly, the bottom-up weight from u_i to u_j is denoted by z_{ij} , and the top-down weight from u_j to u_i is denoted by z_{ji} . The selective output signaling mechanism in PART allows the signal generated in an input layer node to be transmitted to a node in the clustering layer. This happens only when the signal is similar to the top-down weight between the two nodes. The degree of similarity of patterns in the same cluster is controlled by the vigilance and reset mechanism in PART. However, the similarity measurement in PART is closely related to the number of subspaces.Both vigilance and distance

parameters control the degree of similarity of patterns which in turn controls the size of dimensions of the projected subspaces and the degree of similarity in an associated dimension.

The PART algorithm [13] is based on the assumptions that the model equations of PART have regular computational performance described by the following dynamic behavior during each learning trial when a constant input is imposed:

- i. Winner-take-all paradigm: In this scenario, a node in the F_2 layer that has the largest bottom-up filter input becomes the winner. After some finite time only this winner node is activated.
- ii. Selective output signals remain to be constant.
- iii. Synaptic weights are updated using specific formulae.
- iv. Dimensions of a specific projected cluster remainsnon-increasing in time.

PART Algorithm

Initialization

Number of nodes in F_1 layers the number of dimensions of the input data.

Choose the number n of nodes in F_2 layer much larger than the expected number of clusters.

m: number of dimensions of the data space.

Set the internal parameter I, α , ζ and maximum iteration times M.

 $I = (I_1, I_2, \dots, I_m)$ is an input pattern.

 $\boldsymbol{\alpha}:$ The learning rate

 ζ = Small threshold

Choose the external input parameters ρ and σ

 ρ : Vigilance parameter

 σ : Distance vigilance parameter

Step 1: Set all F₂ nodes as being non-committed.

Step 2: For each data point in input data set, do Steps 2.1-2.6.

2.1 Compute h_{ij} for all F₁ nodes u_i and committed F₂ nodes u_j.

If all F_2 nodes are non-committed, go to Step 2.3.

2.2. Compute T_j for all committed F_2 nodes u_j .

2.3 Select the winning F_2 node u_j . If no such F_2 node is found then add the data point to outlier and continue with Step 2.

2.4. If the winner is a committed node then compute *rj* else go to Step 2.6.

- 2.5. If $rj \ge \rho$, go to Step 2.6, otherwise reset the winner u_j and go back to Step 2.3.
 - 2.6. Set the winner u_j as the committed, and update the bottom-up and top-down weights for winner node u_j .

Step 3:Repeat Step 2 M times.

Step 4: For each cluster *Cj* in layer *F*2, compute the associated dimension set *Dj*.

 F_2 layer becomes stable after a few iterations. M is usually a small number

4.EXPERIMENTAL SETUP

Experiments have been carried out in a computing environment of AMD FX-8150 core processor, 2.81 GHz, 8 GB RAM, 1 Terabyte HD, and Windows 7 (64 bit) OS.

4.1 NSL-KDD Dataset

NSL- KDD is a dataset proposed by Tavallace et al. [16] that consists of intrusion data which is being used by researchers for experimentation. NSL-KDD data set is a subset of the original KDD99 dataset having the same features. The NSL-KDDdataset consists of all the 41 attributes and class label of KDD 99 data set. The class label contains four types of attacks, namely Denial of Service (DOS), User to Root (U2R), Remote to Local (R2L), Probe, and normal. This dataset has a binary class attribute. Also, it has a reasonable number of training and test instances which makes it practical to run the experiments.

The NSL-KDD has the following differences over the original KDD 99 dataset.

• It does not include redundant records in the training set, so that the classifiers will not be biased towards more frequent records.

• The number of selected records from each "difficulty level" is inversely proportional to the percentage of records in the original KDD-99 dataset.

• It is possible to experiment on the entire NSL-KDD data set, thus avoids the need for random selection. Consequently, the evaluation results of different researchers will be consistent and comparable.

Class	Number	Percentage of			
	of	Class			
	Records	Occurrences			
Normal	67343	53.48%			
DOS	45927	36.45%			
U2R	52	0.04%			
R2L	995	0.78%			
Probes	11656	9.25%			
Total	125973	100%			

Table 1.Distribution of Records



Figure 4. Distribution of Classes

Each attack type comes under one of the following four main categories: DOS, U2R, R2L, Probes.

i) Denial-of service(DOS) attacks try to limit or deny services by overloading the target system. Some examples of such attacks are apache, smurf, Neptune, Ping of death, back, mail bomb, udpstorm,SYN flood etc.

ii) Probing or Surveillance attacks try to gather knowledge about the configuration of a computer system or network. Typical attacks under this category are Port scans or sweeping of a given IP address range.

iii)User-to-Root(U2R) attacks attempt to gain root/super-user access on a particular computer system on which the attacker has user level access. This is a scenario in which a non-privileged user tries to gain administrative controls/privileges(e.g. Perl,xtermetc).

iv)In Remote-to Local(R2L) an attacker sends packets to a remote machine over the network and tries to exploit the vulnerability in an attempt to have illegal access to a computer to which it does not have legal access.(e.g. xclock,dictionary,guest_password,phf,sendmail,xsnoop etc.)

4.2 Feature Selection

Often, the detection accuracy is deterred by the presence of irrelevant attributes in the intrusion dataset. Therefore, choice of suitable feature selection methods which can eliminate certain attributes that may not contribute to the intrusion detection process is a research challenge. A feature selection method identifies important features for detecting true positive and false negative values and drops irrelevant ones. In this work, the features of the dataset are reduced by using entropy based ranking methods such as Gain ratio, Information Gain, and Symmetrical uncertainty. The dataset has a total of 125973 numbers of records which is reasonable for training as well as testing during experimentation. Out of the total data set 67343, records represent normal data and 58630 represent attacks. Similarly, there are 41 features in the data set among which 38 are numeric and 3 are symbolic in nature.

In information theory, the quantity of information that characterizes the purity of an arbitrary collection of examples is denoted by the Entropy. The entropy represents a measure of system's unpredictability.

The entropy of Y is expressed as: $H(Y) = -\sum_{y \in Y} p(y) log(p(y))$(13)

Where p(y) denotes the marginal probability density function for the random variable Y.

If the values of Y in the training data set are partitioned according to the values of a second feature X, and the entropy of Y with respect to the partitions induced by X is less than the entropy of Y prior to partitioning, then one can establish a relation between the features Y and X, i.e., the entropy of Y after observing X can be represented by the formula:

$$H(Y/X) = -\sum_{x \in X} p(x) \sum_{y \in Y} p\left(\frac{y}{x}\right) \log\left(p\left(\frac{y}{x}\right)\right).$$
(14)

where p(y|x) denotes the conditional probability of y given x.

Information Gain

Information gain helps in determining the feature which is most useful for classification using its entropy value. Essentially, entropy indicates the information content associated with a feature. The higher the entropy, the more information content it carries. Given a value for entropy, one can define Information Gain (IG) which is a measure that determines additional information about feature Y provided by feature X. In fact IG represents the amount by which the entropy of Y decreases. Mathematically,

$$IG = H(Y) - H(Y/X) = H(X) - H(X/Y)....(15)$$

Moreover, IG is a symmetrical measure, which means the information gained about Y after observing X is same as the information gained about X after observing Y.

Using the information gain evaluation criterion with ranking search, the top 15 attributes in the NSL-KDD dataset are selected for classification.

Gain Ratio

The Gain Ratio (GR) is an extension of IG which is a non-symmetrical measure, and is given by,

$$GR = \frac{IG}{H(X)}....(16)$$

Symmetrical Uncertainty

Symmetrical uncertainty technique is symmetric in nature and it reduces the number of comparisons required. It is not influenced by multi-valued attributes and its values are normalized to the range [0,1]. Symmetrical Uncertainty is given by

$$SU = 2 \times \frac{IG}{H(Y) + H(X)}.$$
(17)

Value of SU = 1 means the knowledge of one feature completely predict and SU = 0 indicates that X and Y are uncorrelated.

Using symmetrical uncertainty Evaluation with Ranking Search method for NSL-KDD data, top 10 attributes are selected for classification.

Feature Selection Method	No. of Attributes Selected	Selected Attributes
Info Gain		Service, Flag, Src_bytes. Dst_bytes, Logged_in, Count,
	15	Serror_rate, Srv_serror_rate, Same_srv_rate,
		Diff_srv_rate, Dst_host_srv_count,
		Dst_host_same_srv_rate, Dst_host_diff_srv_rate,
		Dst_host_serror_rate, Dst_host_srv_serror_rate
Gain Ratio	10	Flag, Src_bytes, Dst_bytes, Logged_in, Serror_rate,
		Srv_serror_rate, Same_srv_rate, Diff_srv_rate,

Table: 2Selected attributes using Entropy based methods

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		Dst_host_serror_rate, Dst_host_srv_serror_rate.
Symmetrical	16	Service, Flag, Src_bytes, Dst_bytes, Logged_in,
Uncertainty		Count, Serror_rate, Srv_serror_rate, Same_srv_rate,
		Diff_srv_rate, Dst_host_srv_count,
		Dst_host_same_srv_rate, Dst_host_diff_srv_rate,
		Dst_host_srv_diff_host_rate, Dst_host_serror_rate,
		Dst_hostsrv_serror_rate.

4.3 Cross-Validation

Cross validation is a technique to calculate the accuracy of a model. It separates the data set into two different subsets - a training set and a testing set. First, the classification model is built using the training set and then its effectiveness is measured by observing how efficiently it classifies the testing data set. The process is repeated k times by varying the training and testing subsets following a k-fold cross validation procedure. In our case, we have employed 10-fold cross-validation in which the available data are randomly divided into 10 disjoint subsets of approximately equal size of which 9 subsets of data are used for building the classifier and the 10^{th} subset is used for testing. The process is repeated 10 times by ensuring that each of the subset is used at least once as a test subset. The mean of the above 10 repetitionsis calculated which determines the accuracy of the overall classification system.

4.4 Confusion Matrix

While detecting possible network attacks several situations may arise, namely:

TP (True Positive) which refers to the number of malicious records that are correctly identified.

TN (True Negative) which refers to the number of legitimate (not attacks) records that are correctly classified.

FP (False Positive) which refers to the number of records that are incorrectly identified as attacks though they are actually legitimate.

FN (False Negative) which refers to the number of malicious records which are incorrectly classified as legitimate.

			Predicted Class		
			Negative Class	Positive Class	
			(Normal)	(Attack)	
Actual Class	Negative	Class	TN	FP	
	(Normal)		(True Negative)	(False Positive)	
	Positive	Class	FN	TP	
	(Attack)		(False Negative)	(True Positive)	

Table 3: Confusion Matrix

Next, we define some of the measures which are used to evaluate the performance of different classifiers using the values obtained from the confusion matrix.

Accuracy is the proportion of the total number of correct prediction of positive and negative examples which is given by:

Accuracy =
$$\frac{\text{TP+TN}}{\text{Positive+Negative}}$$

Precision is a measure of the accuracy provided that a specific class has been predicted which is given by:

Precision
$$=\frac{TP}{TP+FP}$$

Recall measure the probability that the algorithm can correctly predict positive examples which is given by :

$$\text{Recall} = \frac{TP}{TP + FN}$$

False Alarm Rate is computed as the ratio of the number of normal instances incorrectly labeled as intrusion (FP) divided by the total number of normal instances, i.e.,

False Alarm Rate =
$$\frac{FP}{FP+TN}$$

F-Value is the harmonic mean of Precision and Recall which measure the quality of classification and is computed as follows:

$$F - Value = 2 \times \frac{(Precision * Recall)}{(Precision + Recall)}$$

And Fitness Value = $\frac{TP}{TP+FP} \times \frac{TN}{TN+FP}$

5. RESULT ANALYSIS

The performance of the proposed classification model that used RBF Network, Self-Organizing Map, Sequential Minimal Optimization and Projective Adaptive Resonance Theory as the classifiers and Info gain, Gain ratio and symmetrical uncertainty as the feature reduction methods was computed using various evaluation parameters such as precision, accuracy, recall, false alarm rate, fitness value, and F-value. For training and testing the standard 10-fold cross-validation technique was used. A comparative view of different combinations of classifiers and feature reduction techniques is depicted in table 4.

Feature	Test	Classifier	Evaluation Criteria					
Selection	Mode	Techniques						
Method			Precision	Accuracy	Recall/	False	Fitness	F-Value
			in %	in %	Detection	Alarm	Value	in %
					Rate	Rate	in %	
					in %	in %		
Info Gain	10-fold	RBFN	96.5805	92.9485	87.9635	2.7115	85.5784	92.0708
	Cross	SOM	79.3058	85.7763	93.9604	21.3466	73.9031	86.0134
	Validation	SMO	98.1368	97.5122	96.4864	1.5948	94.9476	97.3046
		PART	99.8464	99.8333	99.7953	0.1336	99.6619	99.8209
Gain Ratio	10-fold	RBFN	97.0904	88.1213	76.7781	2.0032	75.2401	85.7477
	Cross	SOM	77.9224	84.7499	93.8206	23.1437	72.107	85.1356
	Validation	SMO	96.2894	89.3064	80.1109	2.6877	77.9577	87.4582
		PART	99.8004	99.4165	98.9442	0.1722	98.7738	99.3705
Symmetrical	10-fold	RBFN	96.7865	93.5399	89.0773	2.5749	86.7836	92.772
Uncertainty	Cross	SOM	79.3058	85.7763	93.9604	21.3466	73.9031	86.0134
	Validation	SMO	98.1318	97.5122	96.4915	1.5992	94.9484	97.3048
		PART	99.8618	99.8468	99.809	0.1203	99.6889	99.8354

Table 4. Comparison of four ANN based classifiers using Entropy based feature selection.

On comparison of results based on the evaluation criteria, Projective Adaptive Resonance Theory (PART) has the highest accuracy and detection rate and the lowest false alarm rate irrespective of the feature selection methods used. PART classification with symmetrical uncertainty feature selection gives the highest accuracy of 99.8468%, highest detection rate of 99.809% and lowest false alarm rate of 0.1203%. These results suggest that PART classification technique out performs other techniques, and thus more capable for intrusion detection as compared to other three techniques.

Accuracy, Recall/Detection Rate, False Alarm Rate, Precision, Fitness value, and F-Value of the classifiers with three entropy based feature selection methods are presented in Figures 5, 6, 7, 8, 9, 10 respectively.











Figure 9. Comparison of fitness value among the classifiers Figure 10. Comparison of F-value among the classifiers

5.CONCLUSION

In this paper, different artificial neural network based classifiers along with three different entropy based attribute reduction methods were used to analyze the intrusion data and their performances were analyzed along different evaluation criteria. PART classification with symmetrical uncertainty feature selection gives the highest accuracy of 99.8468%, highest detection rate of 99.809% and lowest false alarm rate of 0.1203%. These results suggest that PART classification technique outperforms other techniques, and thus more suitable for building intrusion detection systems.

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