Classifying Populated IP Addresses Using Horizontal Partition ID3 Classification and K-Means Clustering Algorithm

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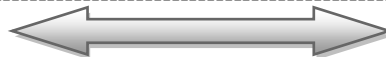[1,]CSE Department, RITS College RGPV University, Bhopal
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--Abstract--
The work implemented here is useful for the detection of IP addresses that creates congestion in the network. These IP addresses are usually detected at the server side to check whether it can attack in the network on the basis of the use of a particular IP address. The existing technique [1] provides an efficient way of classifying IP address using the concept of PIP miner. The existing technique proposes PIPMiner, a fully automated method to classify PIPs with a high accuracy. Although PIP lists are derived from service logs and thus may be application specific, but they can be applied across datasets to detect new attacks But the technique implemented here provides less accuracy and having more error rate the technique is not feasible for network attacks and for classification of the populated IP addresses.The work that we are proposed here is based on the combinatorial method of using Clustering and then classifies to detect the most populated IP address in the network. The proposed work first divide the network traffic into a number of clusters and then on the basis of clusters these traffic is classified using a new method of horizontal partition id3 decision tree having less error rate and less computational time as compared to the existing technique.

Key Words: *K-Mean, Horizontal PartitionId3, Classification*

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I.   INTRODUCTION

Online services such as Web-based email, search, and online social networks are becoming increasingly popular. While these services have become everyday essentials for billions of users, they are also heavily abused by attackers for nefarious activities such as spamming, phishing, and identity theft [1]. Simple conventional mechanisms for abuse detection that rely on source IPs set a limit, i.e., filtering threshold, on the IP activity within a time period. Once the limit is reached by an IP, either the IP traffic gets filtered for the rest of that time period, or the IP gets blacklisted for several consecutive periods. These techniques typically set the same threshold for all IPs. Setting an aggressive threshold yields a high false positive rate since some IPs have numerous users behind them and are hence expected to send relatively large traffic volumes. Setting a conservative threshold yields a high false negative rate, since the threshold becomes ineffective for distributed attacks where IPs send relatively little traffic. This work tailors the thresholds to the sizes of the IPs. It proposes a new framework for timely estimation of the number of users behind IPs with high enough accuracy to reduce false positives and with high enough coverage in the IP space to reduce false negatives [2].Populated IP addresses (PIP) - IP addresses that are associated with a large number of user requests are important for online service providers to efficiently allocate resources and to detect attacks. While some PIPs serve legitimate users, many others are heavily abused by attackers to conduct malicious activities such as scams, phishing, and malware distribution. Unfortunately, commercial proxy lists like Quova have a low coverage of PIP addresses and offer little support for distinguishing good PIPs from abused ones [1].

On the one hand, not all proxies, NATs, or gateways are PIP addresses. Some may be very infrequently used and thus are not of interest to online service providers. On the other hand, while some PIP addresses may belong to proxies or big NATs, many others are not real proxies. Some are dial-up or mobile IPs that have high churn rates. Others include IP addresses from large services, such as Facebook that connects to Hotmail to obtain user email contacts. Additionally, not all PIPs are associated with a large number of actual users. Although many good PIPs like enterprise-level proxies are associated with a large number of actual users, some abused PIPs may be associated with few real users but a large number of fake user accounts controlled by attackers. In an extreme case, bad PIPs may be entirely set up by attackers. For example, it is observed that >30% of the IP addresses that issue more than 20 sign-up requests to Windows Live per day are actually controlled by attackers, with all sign-ups for malicious uses.

Classifying PIPs is a challenging task for several reasons. First, ISPs and network operators consider the size and distribution of customer populations confidential and rarely publish their network usage information. Second, some PIP addresses are dynamic, e.g., those at small coffee shops with user population sizes changing frequently. Third, good PIPs and bad PIPs can locate next to each other in the IP address space. For example, attackers can buy or compromise Web hosting IPs that are right next to the IPs of legitimate services. In addition, a good PIP can temporarily be abused. Due to these challenges, not surprisingly, it is observed that commercial proxy lists offer a low precision in identifying PIPs and provide no support for distinguishing good PIPs from bad ones [1].

Data Mining-based anomaly Detection is become prevalent in essence. Network security is just network information security. In general, all technologies and theories about secrecy, integrality, usability, reality and controllable of network information are the research domain of network security. Intrusion is an action that tries to destroy that secrecy, integrality and usability of network information, which is unlicensed and exceed authority. Intrusion Detection is a positively technology of security defend, which gets and analyses audit data of computer system and network from some network point, and to discover whether there is the action of disobeying security strategy and whether be assaulted. Intrusion Detection System is the combination of software and hardware of Intrusion Detection Data mining can be supervised & unsupervised supervised learning is to use the available data to build one particular variable of interest in terms of rest of data.

Clustering is a division of data into groups of similar objects. Each group called cluster, consists of objects that are similar amongst them and dissimilar compared to object of other groups. Representing data by fewer clusters necessarily loses certain fine details, but achieves simplification. It represents many data objects by few clusters, and hence it models data by its clusters [3].

1.1 K-mean Clustering: k-means is one of the simplest unsupervised learning algorithms that solve the well known clustering problem. The procedure follows a simple and easy way to classify a given data set through a certain number of clusters (assume k clusters) fixed apriori [4]. The main idea is to define k centers, one for each cluster. These centers should be placed in a cunning way because of different location causes different result. So, the better choice is to place them as much as possible far away from each other. The next step is to take each point belonging to a given data set and associate it to the nearest center. When no point is pending, the first step is completed and an early group age is done. At this point we need to re-calculate k new centroids as barycenter of the clusters resulting from the previous step. After we have these k new centroids, a new binding has to be done between the same data set points and the nearest new center. A loop has been generated. As a result of this loop we may notice that the k centers change their location step by step until no more changes are done or in other words centers do not move any more. Finally, this algorithm aims at minimizing an objective function knows as squared error function given by:

$$J(V) = \sum_{i=1}^{c} \sum_{j=1}^{c_i} (\|x_i - v_i\|)^2$$

Where,

'$\|x_i - v_j\|$' is the Euclidean distance between x_i and v_j.

'c_i' is the number of data points in i^{th} cluster.

'c' is the number of cluster centers.

Algorithmic Steps for K-Means Clustering:

Let $X = \{x_1, x_2, x_3, \ldots, x_n\}$ be the set of data points and $V = \{v_1, v_2, \ldots, v_c\}$ be the set of centers.

1) Randomly select 'c' cluster centers.

2) Calculate the distance between each data point and cluster centers.

3) Assign the data point to the cluster center whose distance from the cluster center is minimum of all the cluster centers..

4) Recalculate the new cluster center using:

$$v_i = \left(\frac{1}{c_i}\right) \sum_{j=1}^{c_i} x_i$$

Where, 'c_i' represents the number of data points in i^{th} cluster.

5) Recalculate the distance between each data point and new obtained cluster centers.

6) If no data point was reassigned then stop, otherwise repeat from step 3).

Advantages:
[1] Fast, robust and easier to understand.
[2] Relatively efficient: O (nkd), where n is # objects, k is # clusters, d is # dimension of each object, and t is number of iterations. Normally, k, t, d << n.
[3] Gives best result when data set are distinct or well separated from each other.

Disadvantages:
[1] The learning algorithm requires apriori specification of the number of cluster centers.
[2] 2) The use of Exclusive Assignment - If there are two highly overlapping data then k-means will not be able to resolve   that there are two clusters.
[3] The learning algorithm is not invariant to non-linear transformations i.e. with different representation of data we get different results (data represented in form of Cartesian co-ordinates and polar co-ordinates will give different results).
[4] Euclidean distance measures can unequally weight underlying factors.
[5] The learning algorithm provides the local optima of the squared error function.
[6] Randomly choosing of the cluster center cannot lead us to the fruitful result.
[7] Applicable only when mean is defined i.e. fails for categorical data.
[8] Unable to handle noisy data and outliers.
[9] Algorithm fails for non-linear data set.

1.2 ID3 Algorithm: The ID3 algorithm (Inducing Decision Trees) was originally introduced by Quinlan in [5] and is described below in Algorithm. Here they briefly recall the steps involved in the algorithm. For a thorough discussion of the algorithm we refer the interested reader to [6].
Require: R, a set of attributes.
Require: C, the class attribute.
Require: S, data set of tuples.
[1] if R is empty then
[2] Return the leaf having the most frequent value in data set S.
[3] else if all tuples in S have the same class value then
[4] Return a leaf with that specific class value.
[5] else
[6] Determine attribute A with the highest information gain in S.
[7] Partition S in m parts S(a1), ..., S(am) such that a1, ..., am are the different values of A.
[8] Return a tree with root A and m branches labeled a1...am, such that branch i contains ID3(R − {A}, C, S (ai)).
[9] 9: end if

1.3 Decision Tree: Decision tree support tool that uses tree-like graph or models of decisions and their consequences [7][8], including event outcomes, resource costs, and utility, commonly used in operations research, in decision analysis help to identify a strategy most likely to reach a goal. In data mining and machine learning, decision tree is a predictive model that is mapping from observations about an item to conclusions about its target value. The machine learning technique for inducing a decision tree from data is called decision tree learning.

1.4 ID3 Decision Tree: Iterative Dichotomiser is an algorithm to generate a decision tree invented by Ross Quinlan, based on Occam's razor. It prefers smaller decision trees (simpler theories) over larger ones. However, it does not always produce smallest tree, and therefore heuristic. The decision tree is used by the concept of Information Entropy [9]. The ID3 Algorithm is:
[1] Take all unused attributes and count their entropy concerning test samples
[2] Choose attribute for which entropy is maximum
[3] Make node containing that attribute

ID3 (Examples, Target _ Attribute, Attributes)
• Create a root node for the tree
• If all examples are positive, Return the single-node tree Root, with label = +.
• If all examples are negative, Return the single-node tree Root, with label = -.
• If number of predicting attributes is empty, then

o   Return the single node tree Root, with label = most common value of the target attribute in the
o   examples.

- Otherwise Begin
- A = The Attribute that best classifies examples.
- Decision Tree attribute for Root = A.
- For each possible value, vi, of A,
o   Add a new tree branch below Root, corresponding to the test A = vi.
o   Let Examples (vi), be the subset of examples that have the value vi for A
   ▪   If Examples (vi) is empty common target value in the examples
o   Else below this new branch add the sub tree ID3 (Examples(vi), Target_ Attribute, Attributes – {A}
- End
- Return Root

II.   RELATED WORK

Chi-Yao Hong et. Al. proposes PIPMiner, a fully automated method to extract and classify PIPs through analyzing service logs. Our methods combine machine learning and time series analysis to distinguish good PIPs from abused ones with over 99:6% accuracy. When applying the derived PIP list to several applications, we can identify millions of malicious Windows Live accounts right on the day of their sign- ups, and detect millions of malicious Hotmail accounts well before the current detection system captures them [1].

This paper presents a novel host-based combinatorial method based on k-Means clustering and ID3 decision tree learning algorithms for unsupervised classification of anomalous and normal activities in computer network ARP traffic. The k-Means clustering method is first applied to the normal training instances to partition it into k clusters using Euclidean distance similarity. An ID3 decision tree is constructed on each cluster. Anomaly scores from the k-Means clustering algorithm and decisions of the ID3 decision trees are extracted. A special algorithm is used to combine results of the two algorithms and obtain final anomaly score values. The threshold rule is applied for making decision on the test instance normality or abnormality [10].   Bart Kuijpers et. Al. considers privacy preserving decision tree induction via ID3 in the case where the training data is vertically or vertically distributed. Furthermore, we consider the same problem in the case where the data is both vertically and vertically distributed, a situation we refer to as grid partitioned data. We give an algorithm for privacy preserving ID3 over vertically partitioned data involving more than two parties. For grid partitioned data, we discuss two different evaluation methods for preserving privacy ID3, namely, first merging vertically and developing vertically or first merging vertically and next developing vertically. Next to introducing privacy preserving data mining over grid-partitioned data, the main contribution of this paper is that we show, by means of a complexity analysis that the former evaluation method is the more efficient [11].

Cemal Cagatay Bilgin et al. review the significant contributions in the literature on complex evolving networks; metrics used from degree distribution to spectral graph analysis, real world applications from biology to social sciences, problem domains from anomaly detection, dynamic graph clustering to community detection [12].   This paper is intended to study and compare different data clustering algorithms. The algorithms under investigation are: k-means algorithm, hierarchical clustering algorithm, self-organizing maps algorithm, and expectation maximization clustering algorithm. All these algorithms are compared according to the following factors: size of dataset, number of clusters, type of dataset and type of software used [13]. K. Hanumantha Rao et al. studies the best algorithm by using classifying anomalous and normal activities in a computer networks with supervised & unsupervised algorithms that have not been used before. They analyses the algorithm that have the best efficiency or the best learning and describes the proposed system of K-means&ID3 Decision Tree [9].   Chi-Yao Hong et. Al. [1] uses various studies for their work some for result generation techniques [14] [15] [16] [17]. Here also we refer another related data from various resources such as [18] [19].

III.   CLASSIFICATION ACCURACY

The classification accuracy of a cluster is the proportion of label most often found in that cluster. That is,clusterAccuracy = (Size of majority label)/Size of clusterAnd the overall classification accuracy of the clustering is the weighted mean of the classification accuracy of the clusters. The weight of a cluster is its size divided by the total number of instances.

3.1 CLUSTER ENTROPY

The entropy of the cluster level captures the homogeneity of the cluster. The entropy of a cluster is defined as:

$$Ecluster_i = -\ P\ j(Nji\ ni)\ \log(Njini)$$

where ni is the size of the ite cluster and Nji is the number of instances of cluster i which belongs to the class label j. And the cluster entropy is the weighted sum of the cluster entropies:

Ecluster =PiniN Eclusteri

where N is the total size of the data set and ni is the number of instances in cluster i. The cluster entropy is lowest when the clusters consists of a single data types and it is highest when the proportion of each of data category in the clusters is the same.

SENSITIVITY
Probability that a test result will be positive when there is anomaly (True Positive or TP).

SPECIFICITY
Probability that a test result will be negative when the there is not anomaly (True Negative or TN).

NEGATIVE LIKELIHOOD RATIO
ratio between the probability of a negative test result given the presence of the anomaly and the probability of a negative test result given the absence of the anomaly, i.e. Negative likelihood ratio = False negative rate / True negative rate = (1-Sensitivity) / Specificity.

POSITIVE PREDICTIVE VALUE
Probability that the anomaly is present when the test is positive.

NEGATIVE PREDICTIVE VALUE
Probability that the anomaly is not present when the test is negative. ARP traffic has been applied to detect network abnormalities in the proposed approach, as stated before. We are provided with training and test data set Xi , $1 \leq i \leq N$, where Xi represents a 9 dimensional vector. As shown in Fig 1. is the time needed for the decision of any dataset? It was observed that the existing id3 takes more time as compared our proposed work. Where, HP is the proposed horizontal partioned based ID3. **Relative absolute error** can be calculated as:

$$\frac{|p1 - a1| + \cdots + |pn - an|}{|\bar{a} - a1| + \cdots + |\bar{a} - an|}$$

Mean squared error can be calculated as:

$$\frac{(p1 - a1)^2 + \cdots + (pn - an)^2}{(\bar{a} - a1)^2 + \cdots + (\bar{a} - an)^2}$$

With Actual target values: a1 a2 … a Predicted target values: p1 p2 … pn

DATASET USED
Our experimental analysis is on KDD dataset which contains a number of attributes which is used for the detection of the populated IP address. Some of the attributes and their instance value is given below:

@relation kdd_cup_1999
@attribute duration numeric
@attribute protocol_type {tcp,udp,icmp}
@attribute service
{vmnet,smtp,ntp_u,shell,kshell,aol,imap4,urh_i,netbios_ssn,tftp_u,mtp,uucp,nnsp,echo,tim_i,ssh,iso_tsap,time,netbios_ns,systat,hostnames,login,efs,supdup,http_8001,courier,ctf,finger,nntp,ftp_data,red_i,ldap,http,ftp,pm_dump,exec,klogin,auth,netbios_dgm,other,link,X11,discard,private,remote_job,IRC,daytime,pop_3,pop_2,gopher,sunrpc,name,rje,domain,uucp_path,http_2784,Z39_50,domain_u,csnet_ns,whois,eco_i,bgp,sql_net,printer,telnet,ecr_i,urp_i,netstat,http_443,harvest}
@attribute flag {RSTR,S3,SF,RSTO,SH,OTH,S2,RSTOS0,S1,S0,REJ}
@attribute src_bytes numeric
@attribute dst_bytes numeric
@attribute land {1,0}
@attribute wrong_fragment numeric
@attribute urgent numeric
@attribute hot numeric
@attribute num_failed_logins numeric
@attribute logged_in {1,0}
@attribute lnum_compromised numeric
@attribute lroot_shell numeric
@attribute lsu_attempted numeric
@attribute lnum_root numeric
@attribute lnum_file_creations numeric
@attribute lnum_shells numeric
@attribute lnum_access_files numeric
@attribute lnum_outbound_cmds numeric
@attribute is_host_login {1,0}

@attribute is_guest_login {1,0}
@attribute count numeric
@attribute srv_count numeric
@attribute serror_rate numeric
@attribute srv_serror_rate numeric
@attribute rerror_rate numeric
@attribute srv_rerror_rate numeric
@attribute same_srv_rate numeric
@attribute diff_srv_rate numeric
@attribute srv_diff_host_rate numeric
@attribute dst_host_count numeric
@attribute dst_host_srv_count numeric
@attribute dst_host_same_srv_rate numeric
@attribute dst_host_diff_srv_rate numeric
@attribute dst_host_same_src_port_rate numeric
@attribute dst_host_srv_diff_host_rate numeric
@attribute dst_host_serror_rate numeric
@attribute dst_host_srv_serror_rate numeric
@attribute dst_host_rerror_rate numeric
@attribute dst_host_srv_rerror_rate numeric
@attribute label
{back,teardrop,loadmodule,neptune,rootkit,phf,satan,buffer_overflow,ftp_write,land,spy,ipsweep,multihop,smurf,pod,perl,warezclient,nmap,imap,warezmaster,portsweep,normal,guess_passwd}
@data
0,tcp,http,SF,181,5450,0,0,0,0,0,1,0,0,0,0,0,0,0,0,0,0,8,8,0.00,0.00,0.00,0.00,1.00,0.00,0.00,9,9,1.00,0.00,0.11,0.00,0.00,0.00,0.00,0.00,normal

PROPOSED METHODOLOGY
Here we proposed solution algorithm for K-mean to classify the data set in to number of clusters.

A. Algorithm for K-mean:
[1] Pick a number (K) of cluster centers (at random)
[2] Assign every item to its nearest cluster center (e.g. using Euclidean distance)
[3] Move each cluster center to the mean of its assigned items
[4] Repeat steps 2, 3 until convergence (change in cluster assignments less than a threshold).

B. Horizontal Partition based id3 decision tree
Input Layer:
[1] Define $P_1, P_2 \ldots P_n$ Parties. (Horizontally partitioned).
[2] Each Party contains R set of attributes $A_1, A_2, \ldots, A_R$.
[3] C the class attributes contains c class values $C_1, C_2, \ldots, C_c$.
[4] For party P_i where i = 1 to n do
[5] If R is Empty Then
[6] Return a leaf node with class value
[7] Else If all transaction in $T(P_i)$ have the same class Then
[8] Return a leaf node with the class value
[9] Else
[10] Calculate Expected Information classify the given sample for each party P_i individually.
[11] Calculate Entropy for each attribute $(A_1, A_2, \ldots, A_R)$ of each party P_i.
[12] Calculate Information Gain for each attribute $(A_1, A_2, \ldots, A_R)$ of each party P_i
[13] Calculate Total Information Gain for each attribute of all parties (TotalInformationGain( )).
[14] $A_{BestAttribute} \leftarrow$ MaxInformationGain( )
[15] Let $V_1, V_2, \ldots, V_m$ be the value of attributes. $A_{BestAttribute}$ partitioned $P_1, P_2, \ldots, P_n$ parties into m parties
[16] $P_1(V_1), P_1(V_2), \ldots, P_1(V_m)$
[17] $P_2(V_1), P_2(V_2), \ldots, P_2(V_m)$
[18]   .   .
[19]   .   .
[20] $P_n(V_1), P_n(V_2), \ldots, P_n(V_m)$
[21] Return the Tree whose Root is labelled $A_{BestAttribute}$ and has m edges labelled $V_1, V_2, \ldots, V_m$. Such that for every i the edge Vi goes to the Tree
[22] NPPID3($R - A_{BestAttribute}$, C, $(P_1(V_i), P_2(V_i), \ldots, P_n(V_i))$)
[23] End.

IV.   RESULT ANALYSIS

Graph shown above is the comparative analysis of existing and the proposed id3 algorithm on the basis of their time complexity for different instances of the dataset. The graph shows that the proposed id3 algorithm has less computational time as compared to the existing technique.
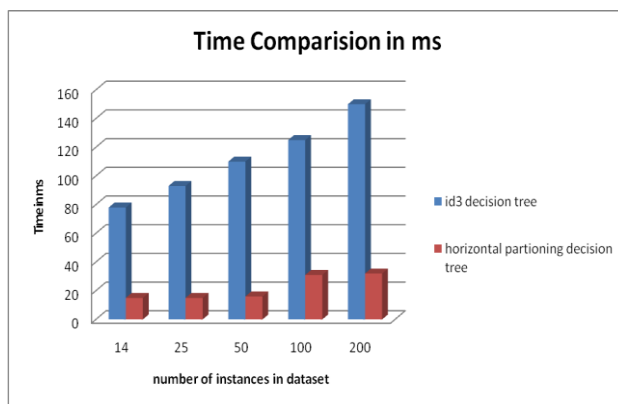

Figure 1. Comparison of Time

As shown in the graph below is the mean absolute error rate of the algorithm as compared to the existing technique.
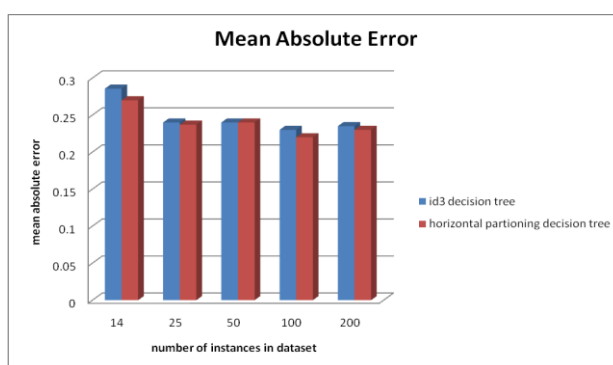

Figure 2. Comparison of Mean Absolute Error

As shown in the below graph is the decision tree created on the basis of dataset used.
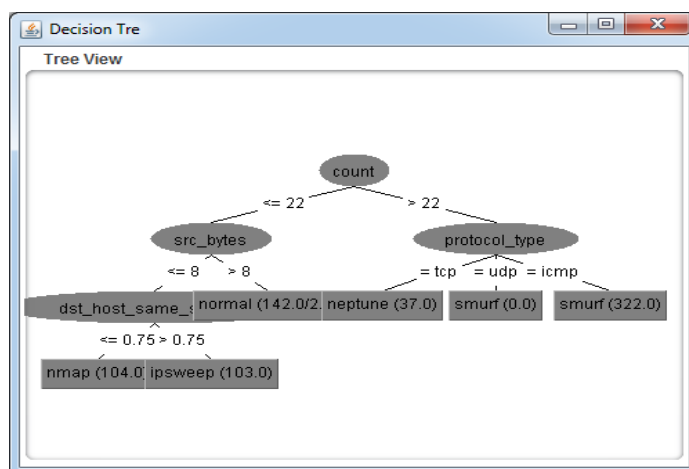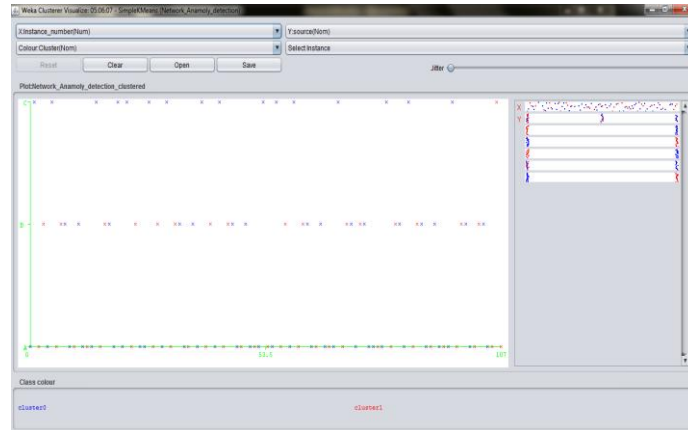

Figure 3. Decision Tree on basis of dataset

Figure 4. Data set on the basis of number of clusters

As shown in the figure above is the analysis of the number of clusters on the basis of dataset choose. Here the analysis is on the basis of two clusters.

V.   CONCLUSION

The proposed work that is implemented here is the combinatorial method of clustering and classification. The method uses k-mean clustering and then applying enhancement of id3 decision tree algorithm i.e. horizontal partition based id3 decision tree. The proposed work implemented here takes less time and having less error rate as compared to the existing technique. The proposed work classifies the populated IP address up to more percentage and is more efficient.Although the technique implemented is feasible for detecting populated IP addresses in the network but the IP that are not populated can be gone through the network hence the chances of attacks has been increased, so in the future we implement combinatorial method of detecting IP addresses and classifying different attacks and their filtering in the network.

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