

# **Project Report**

# **Detect Malicious & Benign Websites using Machine Learning**

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#### **ABSTRACT**

Malicious websites are of great concern to users and organisations nowadays because they rely on web-pages, including online shopping, obtaining information etc. and it threatens the privacy and security of users, organisation, governments as there is a problem in analysing every website one by one and to index each URL in a blacklist. Even if we cannot observe a part of malicious data, we can always observe compromised websites. Since vulnerable websites are discovered by search engines, compromised websites have similar traits. Therefore, we built a classifier by leveraging not only malicious websites but also compromised websites.

Unfortunately, there are lack of datasets which can identify the difference between malicious and benign websites on the basis of their characteristics. To achieve this in our project, we have gathered datasets from multiple resources and categorized them with attributes which define the type of the website. In this project, we address the detection of malicious websites based on their features gathered from multiple sources and use machine learning classification techniques to test and train for future predictions.

#### 1). INTRODUCTION

Drive-by [1] download attacks are the primary method for malware distribution through the Web. When a user accesses a URL that is a starting point of attacks, the user is redirected to an exploit URL via multiple redirections. At the exploit URL, vulnerabilities in browsers and their plugins are exploited, and the user is finally infected with malware. This infected user suffers from damage, such as sensitive data leakage and illegal money transfer, and is integrated into distributed denial-of-service attacks. To expose more users to threats of drive-by downloads, attackers compromise benign websites and inject redirection code to malicious websites such as redirection and exploit URLs. Attackers compromise benign websites and create malicious websites automatically. Since websites built with the old version of content management systems (CMSs) are vulnerable, attackers automatically discover them with search engines, that's the reason, in this project we propose the use of machine Learning approaches to detect the websites and train the model to predict future attacks.

In this project, we use machine learning because vulnerable websites are automatically discovered with search engines by attackers; compromised websites have similar traits. Therefore, we built a classifier by leveraging not only malicious websites but also benign websites. More precisely, we use classification algorithms such as Logistic Regression, Random Forest Classifier and Support Vector Machine (SVM). To perform classification, we integrated two large datasets one consists of predictor variable as URL and response variable as type furthermore in the second dataset we gathered information about websites such as URL, URL LENGTH, SERVER, CHAR SET, Domain location, etc. and their response variable as type which indicate whether it is malicious or benign. As a result of evaluating our system with crawling data of around 345000 URLs combining both datasets, we supposed that they were malicious according to their information, in a next step to verify the records in the dataset we used another security tool, VirusTotal. Hence, providing security and privacy of the user information and system is one of the main concerns of our project. In our project, we use supervised learning methods of machine learning. We learn about the fundamental differences between the three different techniques of supervised learning, i.e. Logistic Regression, Random Forest Classifier and Support Vector Machine (SVM).

The primary goal of supervised learning is to train a model from labeled training data (Figure 1) that allows us to make predictions about future data. Here, the term *supervised* refers to a classification of the desired result [4].

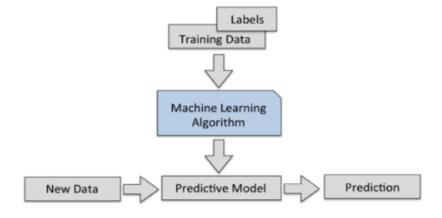


Figure 1: Machine Learning Model

**1.1) Data Description:** Data consists of evaluating classification models to predict malicious and benign websites through their application layer and network characteristics. The data is collected by a process that included different sources of benign and malicious URL, all of them are verified and used in a low interactive client honeypot to get their network traffic, furthermore, we use some other tools to get more information, such as the server's country with Whois details of the website [5].

URL: It is the anonymous identification of the URL analyzed in the study.

URL LENGTH: It is the number of characters in the URL.

NUMBER SPECIAL CHARACTERS: it is the number of unique characters identified in the URL.

CHARSET: It is a categorical value, and its meaning is the character encoding standard.

SERVER: It is a categorical value, and its meaning is the operative system of the server.

<u>CONTENT\_LENGTH</u>: it represents the content size of the HTTP header.

WHOIS COUNTRY: it is a categorical variable; its values are the countries we got from the server response (specifically, our script used the API of Whois).

<u>WHOIS STATEPRO:</u> it is a categorical variable; its values are the states we got from the server response (specifically, our script used the API of Whois).

<u>WHOIS REGDATE:</u> Whois provides the server registration date, so, this variable has date values with format DD/MM/YYY HH:MM

<u>WHOIS UPDATED DATE:</u> Through the Whois, we got the last update date from the server analyzed.

TCP CONVERSATION EXCHANGE: This variable is the number of TCP packets exchanged between the server and our honeypot client.

<u>DIST\_REMOTE\_TCP\_PORT</u>: it is the number of the ports detected and different to TCP <u>REMOTE\_IPS</u>: this variable has the total number of IPs connected to the honeypot APP\_BYTES: this is the number of bytes transferred.

<u>SOURCE APP PACKETS</u>: packets sent from the honeypot to the server.

REMOTE APP PACKETS: packets received from the server.

<u>APP PACKETS</u>: this is the total number of IP packets generated during the communication between the honeypot and the server.

<u>DNS QUERY TIMES</u>: this is the number of DNS packets generated during the communication between the honeypot and the server.

<u>TYPE</u>: this is a categorical variable, its values represent the type of web page analyzed, specifically, one is for malicious websites, and zero is for benign websites.

#### 2). IMPLEMENTATION

In this section, we will discuss the roadmap for building our project using the machine learning. (Figure 2), below shows the workflow diagram for predictive modeling [6].

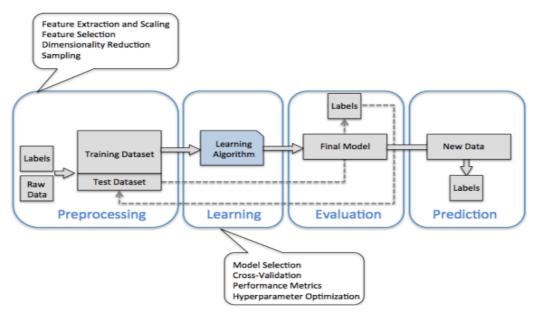


Figure 2: Workflow diagram

Preprocessing: Some of the selected features may be highly correlated and therefore redundant to a certain degree. To keep the raw data in the form and shape for optimal performance of the learning algorithm, dimensionality reduction techniques in Random Forest Classifier, SVM and Logistic regression for achieving better accuracy and efficiency. Its advantage is less storage space and can run faster [6]. We use the training set to train and optimize our machine learning model, while we keep the test set until the very end to evaluate the final model.

- Learning Algorithm: In this, we use three algorithms to train the data and check the accuracy and compare them to find the optimal solution [6].
  - Model Selection: Select a model which will give more efficient results while implementation of the algorithm, training complex dataset and predicting the desired output.
  - <u>Cross-Validation</u>: Validate test and trained data with the predicted result to crossvalidate the accuracy and performance.
  - <u>Performance Metrics</u>: Compare classification as mentioned earlier techniques to check their predicted results to find out which technique is more efficient.
  - Hyperparameter Optimization: A hyperparameter is a parameter whose value is used to control the learning process. By contrast, the values of other parameters (typically node weights) are learned.
- <u>Evaluation</u>: Model Evaluation is an integral part of the model development process. It
  helps to find the best model that represents our data and how well the selected model
  will work in the future.
- <u>Prediction:</u> Machine learning is a way of identifying patterns in data and using them to make predictions or decisions automatically.

### 3). MODEL TESTING USING MENTIONED CLASSIFICATION TECHNIQUES

**3.1) Logistic Regression:** In spite of its name logistic regression is a model for classification, not regression. Logistic regression is used to describe data and to explain the relationship between one dependent binary variable and one or more interval or ratio-level independent variables. Logistic Regression used when the dependent variable(target) is categorical [3]. Input values (x) are combined linearly using weights or coefficient values (Beta) to predict an output value (y). A key difference from linear

regression is that the output value being modeled is a binary value (0 or 1) rather than a numeric value. Below is an example logistic regression equation:

$$y = e^{(b0 + b1*x)} / (1 + e^{(b0 + b1*x)}).$$
 ----- (1)

Where y is the predicted output, b0 is the bias or intercept term and b1 is the coefficient for the single input value (x). Each column in your input data has an associated b coefficient (a constant real value) that must be learned from your training data [6].

#### Algorithm for logistic regression:

- 1. Input: Set of (input, output) training pair samples; call the input sample features x1, x2 to xn, and the output result y. Typically, there can be lots of input features xi.
- 2. Let p(x) be a linear function of x. Every increment of a component of x would add or subtract so much to the probability. The conceptual problem here is that p must be between 0 and 1, and linear functions are unbounded.

$$y = b0 + b1 *x$$
.

3. Find **odds ratio** which is odds in the favour of the particular event.

$$p/(1-p)$$
, where p stands for the probability of the positive event -----(3)

4. Now define the logit function to calculate the logarithm of the odds ratio

$$logit(p) = log p/(1-p).$$
 ------ (4)

5. Logit function takes input values in the range 0 to 1 and transforms them to values over the entire real number range, which express a linear relationship between feature values and the log-odds:

$$logit(p(y=1|x)) = w_0x_0 + w_1x_1 + ... + w_mx_m = \sum_{i=0}^m wixi = wTx$$
 -----(5)

Here, p (y =  $1 \mid x$ ) is the conditional probability that a particular sample belongs to class A given its features x.

6. Now to predict the probability in order to classify the class, use logistic function/sigmoid function:

$$\varphi(z) = 1 / (1 + e^{-z})$$
 -----(6)

- 7. Output: set of weights w (or  $\mathbf{w_i}$ ), one for each feature, whose linear combination predicts the value of y.
- **3.2) Support Vector Machine (SVM):** The objective of the support vector machine algorithm is to find a hyperplane in an N-dimensional space (N is number of features) that distinctly classifies the data points **Figure 3**.

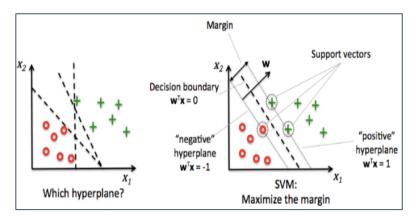


Figure 3:Support Vector Diagram

To separate the two classes of data points, there are many possible hyperplanes that can be selected. Our objective is to find a plane that has the maximum margin, i.e. the maximum distance between data points of both classes. Maximizing the margin distance provides some reinforcement so that future data points can be classified with more confidence. Hyperplanes are decision boundaries that help classify the data points falling on either side of the hyperplane can be attributed to different classes.

Learning of hyperplane, it is done by using linear algebra i.e. kernel, to linearly separate each variable. For high dimensional data other kernels are used and are of four types: Linear, Polynomial, RBF (Radial Basis function), Sigmoid. SVM minimizes the misclassification errors by maximizing the margin. We have used RBF which non-linearly separate the variables. For distance metric, squared Euclidean distance is used.

#### **Algorithm for SVM:**

- Input: Set of (input, output) training pair samples; call the input sample features x1,
   x2 to xn, and the output result y. Typically, there can be lots of input features xi.
- 2. Begin by training two or more SVMs based on tree kernel function, namely, svm1 and svm2, on x1 and x2 to xn, respectively from n number of features

classify the relations on training sets by using two or more trained SVM models (M1 model and M2 model), and then generate different classification result on the same set of input features x1 and x2.

- 3. The fit method of SVC class will call to train the algorithm on the training dataset, which is passed as a parameter to the fit method.
- 4. To maximize the margin of hyperplane in the fit process of step 3, to maximize below are the four steps:
  - 1. Find *positive* and *negative* hyperplanes that are parallel to the decision boundary, which can be expressed.

$$w_0 + \boldsymbol{w}^T \boldsymbol{x}_{pos} = 1 \qquad (1)$$

$$w_0 + \boldsymbol{w}^T \boldsymbol{x}_{neg} = -1 \quad (2)$$

2. Then we subtract those two linear equations (1) and (2) from each other to find the hyperplane.

$$\Rightarrow \mathbf{w}^{T} \left( \mathbf{x}_{pos} - \mathbf{x}_{neg} \right) = 2$$

3. Now further to maximize margin hyperplane for SVM trained with samples known as support vectors.

$$\|\mathbf{w}\| = \sqrt{\sum_{j=1}^{m} w_j^2}$$

4. So Finally, we arrive at the following equation:

$$\frac{\boldsymbol{w}^{T}\left(\boldsymbol{x}_{pos} - \boldsymbol{x}_{neg}\right)}{\|\boldsymbol{w}\|} = \frac{2}{\|\boldsymbol{w}\|}$$

- 5. Finally, now once we have trained the algorithm, the next step is to make predictions on the test data using predict method.
- 3.3) Random Forest Classifier: Random Forest creates multiple decision trees and makes it random. The forest (multiple decision trees) it builds is an ensemble of decision trees. Random forest builds multiple decision trees and merges them to get a more accurate and stable prediction [2]. A random forest is a meta estimator that fits several decision tree classifiers on various sub-samples of the dataset. It uses averaging to improve the predictive accuracy and control over-fitting [2]. One big advantage of the random forest is that it can be used for both classification and regression problems. We will talk about the random forest in classification since classification is sometimes considered the building block of machine learning. Random Forest is also considered as a very handy and easy to use the algorithm, because it's default hyperparameters often produce a good prediction result [6]. The number of hyper-parameters is also not that high and they are straightforward to understand.

#### Steps performed by random forest:

*Input*: Set of (input, output) training pair samples; call the input sample features x1, x2 to xn, and the output result y. Typically, there can be lots of input features xi.

- Randomly select "k" features from total "m" features of dataset.
   Where k < m</li>
- 2. Among the "k" features, calculate the node "d" using the best split point.
- 3. Split the node into daughter nodes using the best split.
- 4. Repeat 1 to 3 steps until "I" number of nodes has been reached.
- 5. Build forest by repeating steps 1 to 4 for "n" number times to create "n" number of trees.

**Output**: In average it takes all the predictions, which cancels out the biases, and attains performance by selecting the best feature from decisions instead of most important feature

**4). PERFORMANCE EVALUATION:** Models appear to perform similarly across the datasets with performance more influenced by choice of dataset rather than model selection. Hence, we have selected three classification algorithms and compared each one of them based on of their accuracy with the corresponding dataset. To measure the performance of classification model, confusion matrix is used. It is described below with the formulas to measure accuracy, precision, recall and f1-score in Figure 7.

```
Training Accuracy Score: 1.0
Test results:
Accuracy Score: 0.9607
Classification Report:
             precision
                         recall f1-score support
                  0.96
                           1.00
                                      0.98
                                                 462
                  0.98
                           0.73
                                      0.83
                                                  73
                  0.96
                           0.96
                                      0.96
                                                 535
  micro avo
                  0.97
                                      0.91
                                                 535
  macro avg
                            0.86
weighted avg
                  0.96
                            0.96
                                      0.96
                                                 535
Confusion Matrix:
[[461 1]
[ 20 53]]
```

Classificat	ssification Report: precision		recall	f1-score	support
	0	0.94	0.98	0.96	317
	1	0.75	0.53	0.62	40
micro av	/g	0.93	0.93	0.93	357
macro av	/g	0.85	0.75	0.79	357
weighted av	′g	0.92	0.93	0.92	357

Figure 4: Random Forest Test Results

Figure 5: Logistic Regression Test Results

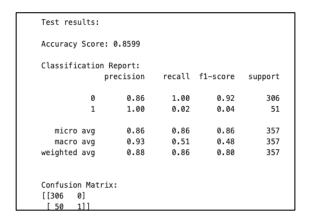


Figure 6: SVM Test Results

Results obtained from the model display that Random forest classifier performed and predicted better with increased accuracy results in term of training and testing. Random forest achieved 96% in predicting and detecting the malicious and benign websites based on their features, whereas SVM is the least favorable in predicting the results.

**Confusion Matrix:** A confusion matrix is a table that is often used to describe the performance of a classification model on a set of test data for which the true values are known. It is a table with 4 different combinations of predicted and actual values in the case for a binary classifier.

A true positive is an outcome where the model correctly predicts the positive class.
 Similarly, a true negative is an outcome where the model correctly predicts the negative class.

**Accuracy:** Accuracy is the fraction of predictions that the model got right.

**Precision:** Out of all observed classes, how much predicted correctly.

**Recall:** Out of all the positive classes, how much predicted correctly.

**F-1 Score:** To combine precision and recall into a single metric called the F1 score.

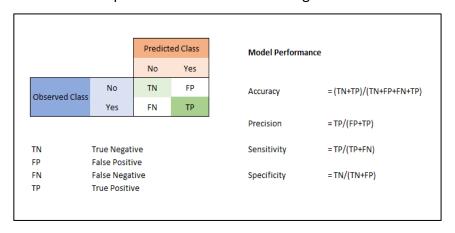


Figure 7: Confusion Matrix

#### 5). VISUALIZATION:

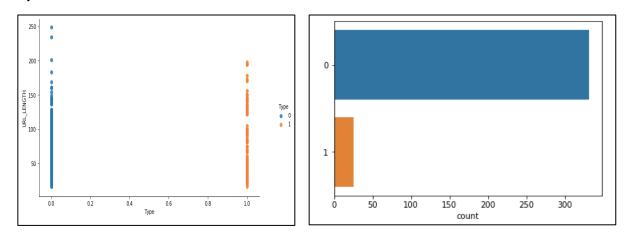


Figure 8: Initial count in Logistic Regression

Figure 9: Predicted count in Logistic Regression

In Figure 8, the graph displays the count of malicious and benign websites in the raw data for a single feature (here URL\_LENGTH feature) in the dataset. Similarly, we can check for N number of features. We can see the difference between malicious and benign is almost equal so to

detect better in Figure 9 the model combined all the features and predicted based on that, Figure 9 is the predicted results and it can distinguish the difference significantly to achieve accuracy.

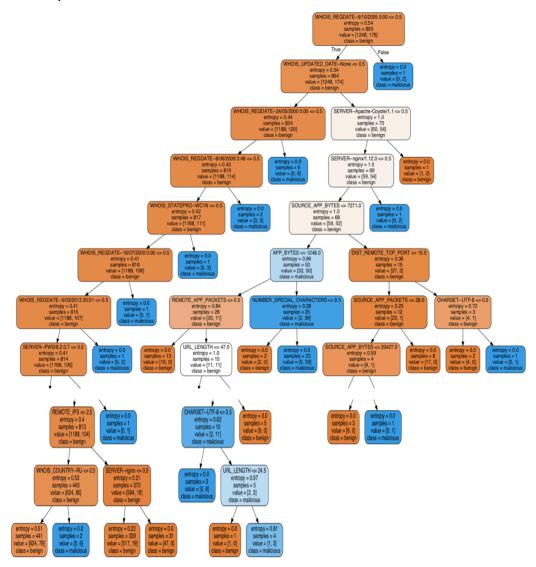
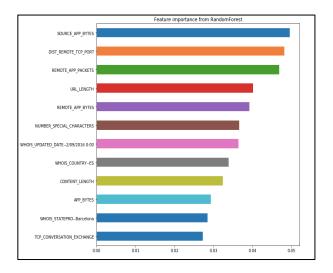


Figure 10:Random Forest Predicted Result

Figure 10 represents the predicted results by random forest classifier. Random Forest algorithm randomly selects observations and features to build several decision trees and then averages the results based on the measurement of the relative importance (Figure 11) of each feature on the prediction.

Figure 11 is the correlation heat map, which is features with high correlation are more linearly dependent and hence have almost the same effect on the dependent variable. So, when two

features have high correlation, we can drop one of the two features. In Figure 12 the predicted correlation between malicious and benign websites.



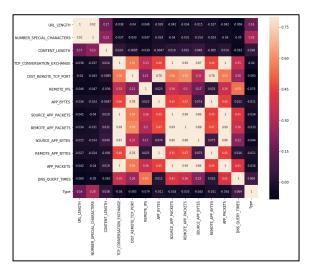


Figure 11:Feature Importance in Random Forest

Figure 12:SVM Correlation heat map

**6). CONCLUSION:** Studied Supervised Learning and used models to classify benign and malicious websites. The experimental results show that the Random Forest performed better than other classification techniques used. RFC works best for unbalanced data because it does not consist of skewness (E.g., Mean). Unequal instances for different classes. The overfitting problem will never come when we use the random forest algorithm in any classification problem. Logistic Regression can be used, if speed is the criteria, whereas in the case of categorical data RF will perform better. However, Better use of misclassified data to update the weights and classify them again to predict based on of more data, because more data leads to better prediction. Finally, models appear to perform similarly across the datasets with performance more influenced by choice of dataset rather than model selection.

**7). FUTURE WORK:** Need to be implemented for all other classification techniques in Machine Learning and Deep learning algorithms to compare results from all the techniques.

In future, Important task will be to assemble feature extraction from website at runtime and perform dynamic detection from the trained dataset. Further, Evolutionary computation will be

best to learn from dynamic features of websites and predict better results of the everchanging internet.

### 8). REFERENCES

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- [2]. Niklas Donges; "The Random Forest Algorithm", https://towardsdatascience.com/the-random-forest-algorithm-d457d499ffcd".
- [3]. Jason Brownlee; Understand Machine Learning Algorithms, https://machinelearningmastery.com/logistic-regression-for-machine-learning/.
- [4]. Badreesh Shetty; Supervised Machine Learning: Classification,
   https://towardsdatascience.com/supervised-machine-learning-classification-5e685fe18a6d
   [5]. Christian Urcuqui; https://www.kaggle.com/xwolf12/malicious-and-benign-websites/home.
- [6]. Sebastian Raschka, Python Machine Learning, 1st Edition.

## **Appendix**

```
import pandas as pd
 import numpy as np
import matplotlib.pyplot as plt
 import random
import sklearn
import warnings
warnings.filterwarnings("ignore", category=FutureWarning)
# Machine Learning Packages
from sklearn.linear_model import LogisticRegression
from sklearn.model_selection import train_test_split
from sklearn.model_selection import cross_val_predict, cross_val_score
from sklearn.metrics import confusion_matrix,classification_report,accuracy_score
dataset=pd.read_csv("dataset.csv")
dataset.head()
dataset.Nead(, dataset.describe(include='all') # Quick statistical summary of data dataset.drop('URL', axis=1, inplace=True) # Drop the URL column since that is a unique column for training
# Take a look at any null values to clean up data, Likely need to do something with these empty datasets
print(dataset.isnull().sum())
dataset[pd.isnull(dataset).any(axis=1)]
dataset = dataset.interpolate() # Interpolate our data to get rid of null values
print(dataset.isnull().sum())
dataset[,ifilna('RARE_VALUE', inplace=True) # For some reason there's still a isnull in the SERVER column
dataset_numerical = pd.get_dummies(dataset,prefix_sep='--') # Convert categorical columns to numbered categorical columns
# Separate predictors and response
X = dataset_numerical.drop('Type',axis=1) #Predictors
y = dataset_numerical['Type']
# Our split will be a 80/20 split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2) # Get a training and test dataset
#Model Building, using logistic regression
logit = LogisticRegression()
logit.fit(X_train, y_train)
# Accuracy of Our Model
print("Accuracy ",logit.score(X_test, y_test))
New_predict = logit.predict(X_test)
print('Misclassified samples: %d' % (y_test != New_predict).sum())
ariat(Nay predict)
print(New_predict)
```

#### Code implementation of Logistic Regression

```
9]: import numpy as np
import pandas as pd
        import matplotlib.pyplot as plt
        import warnings
from sklearn.model_selection import train_test_split
        from sklearn import sym
        from aktearn import sviii
from matplotlib.colors import ListedColormap
from sklearn.model_selection import cross_val_predict, cross_val_score
from sklearn.metrics import confusion_matrix,classification_report,accuracy_score
         from mlxtend.plotting import plot_decision_regions
        import seaborn as sns
        warnings.filterwarnings("ignore", category=FutureWarning)
        dataset=pd.read_csv("dataset.csv") # Print out a quick overview of the data
       dataset=pd.read_csv("dataset.csv") # Print out a quick overview of the data
dataset.head()
dataset.drop('URL', axis=1, inplace=True) # Drop the URL column since that is a unique column for training
print(dataset.isnull().sum())
dataset[pd.isnull(dataset).any(axis=1)]
dataset.describe(include='all') # Quick statistical summary of data
dataset = dataset.interpolate() # Interpolate our data to get rid of null values
print(dataset.isnull().sum())
       # Quick statistical summary of data
dataset['SERVER'].fillna('RARE_VALUE', inplace=True)
       dataset_svm = pd.get_dummies(dataset,prefix_sep='--')
print(dataset_svm.head())
       X = dataset_svm.drop('Type',axis=1) #Separate predictors and response Predictors
y = dataset_svm['Type']
       # Our split will be a 80/20 split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42) # Get a training and test dataset
svm = svm.SVC(kernel='rbf', random_state=0, gamma=3, C=1.0)
        #Model Building, using SVM
       svm.fit(X_train, y_train)
y_pred = svm.predict(X_test)
```

Code implementation of Support Vector Machine

```
[5]: import numpy as np
     import pandas as pd
     import pydotplus
     import graphviz
     from sklearn.tree import export_graphviz
     from sklearn.externals.six import StringIO
     from IPython.display import Image
      from sklearn import tree
     from sklearn.model selection import train test split
      from sklearn.ensemble import RandomForestClassifier # Train a Random Forest Regressor
      from sklearn.model_selection import cross_val_predict, cross_val_score
     from sklearn.metrics import confusion_matrix,classification_report,accuracy_score
     # Copy over our data so we don't overwrite the original results if we want dimensionality reduction
     dataset_preprocessed = dataset
      # Converting server types with only 1 unique count to a RARE_VALUE for classification
     test = dataset_preprocessed ['SERVER'].value_counts()
     col = 'SERVER'
     dataset_preprocessed.loc[dataset_preprocessed[col].value_counts()[dataset_preprocessed[col]].values < 2, col] = "RARE_VALUE"
      # Function to extract the registration year
     def extract_reg_year(x):
         # If no year was reported, leave it as None
         if str(x) == 'None':
         parse_error = False # Try different parses for different date formats
             date = x.split(' ')[0]
             year = date.split('/')[2]
         except:
             parse_error = True
         if parse_error: # One more date format to try if there's a parse error
                  date = x.split('T')[0]
                  year = date.split('-')[0]
                  parse_error = False
              except:
                  parse error = True
                  raise ValueError('Error parsing {}'.format(x))
     dataset preprocessed['WHOIS REGDATE'] = dataset preprocessed['WHOIS REGDATE'].apply(extract reg year)
     dataset_preprocessed['WHOIS_UPDATED_DATE'] = dataset_preprocessed['WHOIS_UPDATED_DATE'].apply(extract_reg_year)
        # State without country doesn't make sense
        dataset_preprocessed['WHOIS_STATEPRO'] = dataset_preprocessed[['WHOIS_COUNTRY', 'WHOIS_STATEPRO']].apply(lambda x : '{}-{}'.format(x[0],x[1]), axis=1)
        dataset_preprocessed.describe(include='all')
        # As above, create our random forest classifier the same way, Convert categorical columns to numbered categorical columns
        dataset_pp_with_dummies = pd.get_dummies(dataset_preprocessed, prefix_sep='--') # Separate predictors and response
        X_pp = dataset_pp_with_dummies.drop('Type',axis=1) #Predictors
        y_pp = dataset_pp_with_dummies['Type']
        # Get a training and test dataset
        X_pp_train, X_pp_test, y_pp_train, y_pp_test = train_test_split(X_pp, y_pp, test_size=0.2, random_state=42)
        # n_estimators is the number of random forests to use, n_jobs says to use all processors available
        # Properties we can play with for the RandomForestClassifier function:
        rf_pp = RandomForestClassifier(n_estimators=100, n_jobs=-1, max_depth=30, criterion = 'entropy')
        rf_pp.fit(X_pp_train, y_pp_train)
        print('Training Accuracy Score: {}'.format(rf_pp.score(X_pp_train, y_pp_train)))
        y_pp_pred = rf_pp.predict(X_pp_test)
```

Code implementation of Random Forest Classifier

print\_score(rf\_pp,X\_pp\_train,y\_pp\_train,X\_pp\_test,y\_pp\_test,train=False)