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A Study On Effects Of Data Poisoning On HMMs

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A Study On Effects Of Data Poisoning On HMMs

A Project

Presented to

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> In Partial Fulfillment of the Requirements for the Degree Master of Science

> > by Rachel Gonsalves May 2018

 \odot 2018

Rachel Gonsalves

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A Study On Effects Of Data Poisoning On HMMs

by

Rachel Gonsalves

APPROVED FOR THE DEPARTMENT OF COMPUTER SCIENCE

SAN JOSÉ STATE UNIVERSITY

May 2018

ABSTRACT

A Study On Effects Of Data Poisoning On HMMs

by Rachel Gonsalves

With the ever increasing use of burgeoning volumes of data, machine learning systems involving minimal human oversight are crucial for classification and analysis tasks. Machine learning algorithms used for such purposes have revolutionized the way we sort, classify, and analyze data.

The accuracy of any machine learning algorithm depends heavily on the data it is trained on. In some circumstances, an attacker can attempt to poison the training data to subvert a machine learning system. In this research, we analyze the effects of training data poisoning attacks on hidden Markov models (HMMs), in the context of malware classification. With the increase in percentage of data poisoning, HMM is still able to classify most files correctly. Hence we find that HMMs are able to classify at high and low level of poisoning.

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CHAPTER 1

Introduction

With the sheer rise in the amount and variety of data being generated, it has become crucial that we have techniques to classify data independent of human supervision. Machine learning techniques prove useful not only in classifying data but also in identifying malware. Malware can affect computers, leak sensitive data [\[1\]](#page-26-1), cause denial of service attacks and cause much damage to crucial systems in the current world. Many machine learning techniques are being used today to detect malware and prevent attacks. In order to escape detection and carry out attacks successfully, attackers come up with innovative ideas; one such idea is to compromise the training data of a machine learning model, which causes incorrect learning and thus confuses the model and leads to decreased accuracy but with higher and lower levls of posoinonig the model still [\[2\]](#page-26-2).

The attacks that involve influencing the models can be categorized as [\[3\]](#page-26-3):

- Causative
- • Exploratory

Figure 1: Poisoning of Training Set

The accuracy and efficiency of a machine learning algorithm depends on the training data. When the training set for a machine learning algorithm is poisoned, such an attack is called a causative attack. [\[3\]](#page-26-3). Poisoning a model slowly over a period of time is called the Boiling Frog attack [\[2\]](#page-26-2). Figure [1](#page-11-1) describes how an attack on the training set works [\[3\]](#page-26-3). P_z indicates the true distribution. D^{train} and D^{eval} represent the training and testing set respectively. H is the machine learning algorithm. f is the hypothesis that is evaluated by comparing the results. An example of a causative attack against a spam filter is described as follows in [\[4\]](#page-26-4): Attackers try and circumvent spam filter re-training by sending non-intrusive traffic, carefully constructed to resemble the upcoming spam. This causes the defending filter to be mis-trained which results in an inability to effectively block the spam. This would be as follows: the spam sales pitch "You need a new phone? Really, do buy now!" is recast as "Do you really need buy a new phone, now!?"; while both these phrases have markedly different meanings they are treated the same by the spam filter. [\[4\]](#page-26-4) In an exploratory attack, the attacker observes the effects of instances designed for the learning model but does not directly influence the learning [\[5\]](#page-26-5) [\[2\]](#page-26-2).

The goal of a malware detector is to ensure secure learning [\[3\]](#page-26-3). In the current experiment, we check how susceptible HMMs are to data poisoning. For a given training set, when data is poisoned, the model is tested on a benign and malware test data. When the model training set is poisoned, we quantify the change in performance of the model.

The remainder of this paper is structured as follows. Relevant background topics are discussed in Chapter [2.](#page-13-0) This chapter includes an introduction to Hidden Markov Models(HMMs), on which our training and testing methods and some related work. In Chapter [3,](#page-16-0) we discuss the implementation, present the results and observations. Chapter [4](#page-25-0) contains the conclusion and a brief discussion of future work.

CHAPTER 2

Background

2.1 Hidden Markov Model

Hidden Markov model is a machine learning technique which is used for statistical pattern analysis [\[6\]](#page-26-6). To get a deeper understanding of the concept and an overview of the terminology used, we look at an example from the paper 'A Revealing Introduction to Hidden Markov Models' [\[7\]](#page-26-7) and understand the terminologies. Suppose we want to predict the temperatures of some 100 years ago and know that relation between the size of growth of tree rings and the weather. The weather is categorized to $hot(H)$ and cold(C). The sizes of tree rings to small(S), medium(M) and large(L). Given an observation of tree ring sizes we try to find whether it was hot or cold during the growth of the ring. These states are unknown thus known as hidden states. The model is represented by observation matrix, transition matrix and the initial state distribution.

The transition matrix A is a $N*N$ matrix where is N is the number of states, observation matrix is $N * M$ where M is the number of observed symbols. Initial state distribution is the probability of starting at any given state and is give by a list of size N , these represent probabilities corresponding to each state. Each matrix is row stochastic. In the Figure [2](#page-14-2) [\[7\]](#page-26-7), $X_0 - X_n$ represents the hidden states.

- $T =$ length of the observation sequence
- $N =$ number of states in the model $M =$ number of observation symbols $Q = q_0, q_1, ..., q_{N-1}$ = distinct states of the Markov process $V = 0, 1, ..., M - 1$ = set of possible observations $A =$ state transition probabilities

 $B =$ observation probability matrix

 π = initial state distribution

 $O = (O_0, O_1, ..., O_{T-1})$ = observation sequence.

Figure 2: Hidden Markov Model

HMM is used to solve mainly three problems:

Problem 1: Given a sequence of observation to find the probability of the observed sequence

Problem 2: Finding the state sequence that best fits the given model.

Problem 3: To find a model that fits best the observed data [\[7\]](#page-26-7).

2.2 Hidden Markov Model for Malware Detection

For malware classification in HMM, the model is trained on the opcode sequence of the malware. This type of analysis is known as static analysis because it does not involve execution and monitoring [\[8\]](#page-26-8). Malware detection with the help of HMM is statistical as it trains on statistical features [\[9\]](#page-26-9).

2.3 ROC Curves

For Receiver Operating Classifier(ROC) curves, we need to calculate the True Positive Rate(TPR) and False Positive Rate(FPR).For a given classifier, we need to understand that samples can be classified in 4 ways [\[10\]](#page-26-10). When a given malware sample is correctly classified as malware, it is considered to be a true positive. When a malware sample is classified as benign it is a false negative. When a sample is benign and classified as benign it is a true negative.When a benign sample is wrongly classified as malware it is False Positive. TPR is the total number of true positives upon the total number of positive samples, TPR also termed as the sensitivity [\[11\]](#page-27-0). A True Negative Rate(TNR) is the total number of true negatives up the total number of benign samples and is known as the specificity. [\[12\]](#page-27-1). The x-axis of the ROC curve represents the FPR and y-axis represents the TPR. The threshold passes through the point (TPR,FPR) [\[10\]](#page-26-10).

2.4 Related Work

When it comes to data poisoning, a lot of work has been done to observe its effects on various machine learning techniques. Data poisoning attacks mostly occur when a system is adaptive [\[13\]](#page-27-2). It is also most common when data is gathered from unreliable sources. One of the methods used to improve the efficiency and decrease the model vulnerability is that the model can be trained to reject a sample which causes a decrease in efficiency. The sample would be considered as an outlier. This technique is also known as Reject On Negative Impact(RONI) or data sanitization [\[14\]](#page-27-3) [\[13\]](#page-27-2). Blacklisting and white listing sources requires a lot of effort and it also blocks some good traffic. This does not prove to be effective in the long run either [\[15\]](#page-27-4). Another method, weighted bagging, was used for the training data; this helps make the model robust against such attacks [\[16\]](#page-27-5) [\[5\]](#page-26-5) . The concepts of data sanitization, weighted bagging and defining upper and lower bounds for testing data are useful when dealing with adversarial machine learning and are decribed in detail in [\[17\]](#page-27-6) [\[18\]](#page-27-7).

CHAPTER 3

Methodology and Results

3.1 Methodology

Hidden Markov models are useful in classification of malware. HMM is used to distinguish between malware and benign files. [\[7\]](#page-26-7) For the detection of malware, the model is trained on the opcode sequence. This trained model is tested on opcode sequence of benign files as well as malware. For the purpose of this experiment, the HMM is trained on 100 files each of 3 malware families. The A, B and π matrix are initalized to around $1/N$, $1/M$ and $1/N$ per row respectively. The 26 most frequently occuring opcodes were mapped and the rest were considered to be space. For the current model the values were initialized as follows: $N = 2$, $M = 27$. M is 27 for the number of opcodes and space inserted for every opcode not in the 26. N is 2 as it gave the best classification after experimenting with N ranging 2 to 6. For each training 100 files are used.

3.2 Dataset

In order to test the effects of data poisoning on the model, the model was initially trained on a pure training set so the changes in the efficiency of the model can be observed once the data is poisoned. For the purpose of this experiment, malware files from the Malicia dataset were used. The model was trained on 3 malware families based on the number of samples available in the training set: Winwebsec [\[19\]](#page-27-8), Zbot [\[20\]](#page-27-9) and Zeroaccess [\[21\]](#page-27-10).

For each malware family, the training set and test set contained 100 files. Once the model was trained and tested, the data was poisoned gradually by adding files one by one to the training set, training and then testing the model. The decrease in efficiency of the model is checked by evaluating the Area Under the Curve (AUC).

3.3 Tests

This section includes the results of data poisoning on 3 malware families. 100 files from each malware family were selected randomly. For the purpose of testing, a sequence with $T = 15000$ was used from each sample file to score the samples.

3.3.1 Effects of Data Poisoning on Winwebsec

The model was trained on 100 files from the Winwebsec malware family, and tested on 100 samples of malware and benign files, the training data was poisoned with the addition of one malware file and was tested again. From Figure [3](#page-18-0) to Figure [A.27,](#page-45-0) we see the scatter plots and ROC curve of the tested data. In the results we see that the AUC of the model decreases with addition of each benign sample. In the pure training set, an AUC of 0*.*64 is observed [3,](#page-18-0) which gradually goes to 0*.*95 [A.22](#page-40-0) with a data poisoning of 45%

Figure 3: Winwebsec ROC Curve and AUC with Pure Dataset

Figure 4: Winwebsec Scatter Plot for with Pure Dataset

3.3.2 Effects of Data Poisoning ZeroAccess

The model was trained on 100 files from the ZeroAccess malware family, and tested on 100 samples of malware and benign files, the training data was poisoned with the addition of one benign file after each test and was tested again. From Figure [5](#page-20-1) to Figure [A.41,](#page-59-0) we see the scatter plots and ROC curve of the tested data. The results were evident as we see that the AUC of the model decreases with addition of each benign sample. The results were evident as we see that the AUC of the model decreases with addition of each benign sample. In the pure training set, an AUC of 0*.*93 is observed Figure [5,](#page-20-1) which gradually reduces to 0*.*19 which can be reversed to an auc of 0*.*81 Figure [A.40](#page-58-0) with a data poisoning of 45%

Figure 5: ZeroAccess ROC Curve and AUC with Pure Dataset

Figure 6: ZeroAccess Scatter Plot for with Pure Dataset

3.3.3 Effects of Data Poisoning on Zbot

The model was trained on 100 files from the Zbot malware family, and tested on 100 samples of malware and benign files, the training data was poisoned with the addition of one benign file at a time and was tested again. From Figure [7](#page-22-0) to Figure [A.63,](#page-81-0) we see the scatter plots and ROC curve of the tested data. In the results we see that the AUC of the model does not decreaset with addition of each benign sample. In the pure training set, an AUC of 0*.*71 is observed Figure [7,](#page-22-0) which gradually increaseses to 0*.*79 Figure [A.58](#page-76-0) with a data poisoning of 45%

Figure 7: Zbot ROC Curve and AUC with Pure Dataset

Figure 8: Zbot Scatter Plot for with Pure Dataset

3.3.4 Results for All Malware Families

The following Figure [9](#page-24-1) is a graphical representation of the change in AUC with the increase in percentage of data poisoning for all malware families.The x axis represents the AUC and the y axis represents the percentage of data poisoning.

Figure 9: Effects of Data Poisoning on All Malware Families

CHAPTER 4

Conclusion and Future work

4.1 Conclusion

HMM was trained on three malware families, these trained models were tested on malware and benign samples. The models trained on pure datasets performed well. Multiple HMM models were trained with training data poisoned from 0% to 45% for each family incrementing the poisoning by 5. As the percentage of poisoning increased, the model was still able to classify the files at a higher and lower level of poisnoning.

4.2 Future Work

HMMs trained on API calls as features are stronger and hence it would be a good comparison to see how HMMs trained dynamically perform with this type of an attack

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APPENDIX

APPENDIX A

A.1 Graphs for Data Poisoning on Winwebsec

Figure A.10: Winwebsec ROC Curve and AUC with 5% Data Poisoning

Figure A.11: Winwebsec Scatter Plot for 5% Data Poisoning

Figure A.12: Winwebsec ROC Curve and AUC with 10% Data Poisoning

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Figure A.24: Winwebsec ROC Curve and AUC with 40% Data Poisoning

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Figure A.26: Winwebsec ROC Curve and AUC with 45% Data Poisoning

Figure A.27: Winwebsec Scatter Plot for 45% Data Poisoning

A.2 Graphs for Data Poisoning on ZeroAccess

Figure A.28: ZeroAccess ROC Curve and AUC with 5% Data Poisoning

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