# TODO

### Dalton Cole

Department of Computer Science Missouri University of Science and Technology Rolla, Missouri 65409-0350, USA Email: drcgy5@mst.edu

### Bruce McMillin

Department of Computer Science Missouri University of Science and Technology Rolla, Missouri 65409-0350, USA Email: ff@mst.edu

# Abstract—The abstract goes here.

### I. INTRODUCTION

Intro

#### II. METHODOLOGY

An overview of the system is outlined in Figure 1. The following subsections outline in detail how each partition of the system functions. For how data was collected, see Section III.

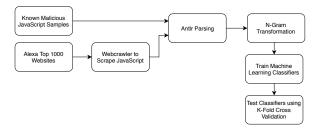


Fig. 1. System Overview

# A. Antlr

Antlr was used to parse the grammar [2]. Antler is a framework designed to automatically generate code in a number of languages. The resulting supplementary files can be used to tokenize and parse given inputs. Antlr requires tokens and grammatical rules to be defined for the desired language in a g4 file. Antlr is designed for context-free grammars (CFG), however, by adding additional logic that is programming language specific, Antlr can handle context-sensitive languages.

Antlr parses samples using a LL(\*) parser, which is a top-down approach. Top-down approaches work by starting at the top level of a parse tree and matches as it traverses down the tree. LL(\*) performs leftmost deviation as it reads in tokens from left to right. The amount of tokens read in at a time to determine which grammar rule to use is determined by the \* value. If \*=1, then only one token is required at a time to know which grammar rule to expand next.

## B. N-Gram

A list of grammatical rules in order of which they were parsed is extracted from a given sample using Antlr. These rules were then grouped in over-lapping sequences of n rules where each sequence differed by its neighboring sequences by one rule. These sequences are called n-grams. A snippet of a n-gram extracted from our system can be found in Figure 2. Due to memory limitations, only sequences where  $n=\{1,2,3\}$  were explored. To generate a feature vector of set length, every possible n-gram was composed.

["Program", "SourceElements"], ["SourceElements", "SourceElement"], ["SourceElement", "Statement"].

Fig. 2. N-Gram Example where N = 2

## C. Feature Reduction

Due to the large number of features that n-gram produces, feature reduction is required to reduce the memory usage to a usable level. In order to read in saved data, incremental principal component analysis (IPCA) was selected. IPCA is able to read in information from disk in batches. From each read, IPCA finds orthogonal eigen vectors. After all rows of the dataframe have been read in, the top n eigen vectors with the most variance are selected to represent the features of the data. When the number of features are less than n, feature reduction is not applied.

## D. Machine Learning

A number of different machine learning methods were employed. These methods were chosen so the results could be compared to TODO LIST REFERENCES. K-Fold cross validation was used to train and test each machine learning classifier.

- 1) Decision Trees: entropy
- 2) Random Forests:
- 3) Neural Network:
- 4) Naive Bayes:
- 5) Gradient Decent:
- 6) K-Nearest-Neighbors:

# III. EXPERIMENTAL EVALUATION

# A. Benign Data Collection

The benign data set consisted of JavaScript samples scraped from the top 1000 Alexa websites [?]. Due to these site's popularity, it was assumed that each javascript sample scrapped from these sites were benign.

The python library scrapy was used to create a web-crawler and scrapper [?]. A set of starting URLs and acceptable

domains were given to scrapy that consisted of the top 1000 Alexa websites. From here, any javascript that was enclosed in HTML script tags were scrapped from the site and saved into a hash-table. A hash-table was used to guarantee unique samples. A total of 201,901 scripts were scrapped using this method. Due to our use of Antlr, malformed JavaScript samples could not be parsed. The main malformation that was not accepted was the lack of use of semicolons. If a script did not use semicolons to end statements, the script was considered malformed and discarded. A full list of what is considered malformed can be found at [?]. After malformed samples were removed from the set of possible scripts, 140,437 scripts remained. These scripts were feed into Antlr using the JavaScript grammar defined in [?]. From Antlr the order in which the grammatical rules were parsed in were returned.

#### B. Malicious Data Collection

Malicious data consisted of known malicious JavaScript samples obtained from [?]. The majority of this malware was collected between 2015 and 2017. A random sample was ran through VirusTotal in order to verify that leading antivirus scanners recognized the samples as malicious [?]. All samples that were randomly selected was marked malicious by VirusTotal. A total of 39,471 JavaScript samples were obtained. Similarly to the benign samples collected, some scripts were malformed. After removing the malformed samples, 15,096 samples remained. Using Antlr, the order in which the grammatical rules were parsed in were returned.

### C. Feature Reduction

Due to the large number of features that n-gram produces, three different feature sets were used. An unabridged dataset containing 112 features. This unabridged dataset contains every grammatical non-terminal in the original JavaScript grammar used to parse the language. From this unabridged dataset, two additional datasets were generated. A trimmed dataset containing 62 rules were generated by combining every Expression grammatical rule into a single grammatical rule. By combining similar rules into a single rule, it is hoped that not much information is lost. A third extra trimmed dataset was constructed by further combining similar rules until only 40 features remained.

The reason this feature reduction was required is due to n-gram's exponential nature. When using 1-gram with the original dataset, only 112 features are needed. However, when using 2-gram, 12544 features are required. With this exponential growth, more memory is required to store those features. In addition, a very sparse matrix will be produced since most of the n-gram rules will not be seen in the source program.

In addition to the above feature reduction methods, incremental principal component analysis (IPCA) was employed to further reduce the features. A brief overview of how IPCA functions is reviewed in Section II-C. One hundred orthogonal vectors with the most variance was selected. One hundred features were selected because it gave a good compromise to the number of features in the 1-gram unabridged dataset.

With the number of samples explored, 100 features will easily fit into memory without the need to save to disk, thus training times were greatly reduced.

## D. Classifier Training

For classification training, an equal number of benign and malicious samples were selected from each dataset. Given the size of the malicious dataset, 15,000 samples from each classification was used to train the models. Choosing an equal number of samples in each class was decided upon due to two reasons.

Firstly, the imbalanced class issues for the majority of machine learning methods used in this paper have been explored in [?]. With regards to decision trees, Sun et al. described how during the pruning phase of generating a decision tree, the leaf nodes representing the minority class have a good chance of being pruned since they have little effect on the overall accuracy of that branch. In addition, with respect to neural networks, they found that the majority class in skewed data tends to dominate the gradient vector. The minority class may even need to traverse in an uphill direction to find the optimal values.

Secondly, accuracy and precision are positively effected by skewed data. Generally, classifiers are able to classify the majority class correctly a larger percentage of the time. Strictly looking at accuracy, classifier accuracy will be inflated due to classifying the majority class correctly more often. This diminishes the actual results of the minority class that we are concerned about [?].

# E. Results

Recall, Precision, and  $F_1$  score are the three primary metrics used to compare each result. Formulas used to calculate these can be found in Equation 1, Equation 2, and Equation 3, respectively.

$$Recall = \frac{TruePositive}{TruePositive + FalseNegative}$$
 (1)

$$Precision = \frac{TruePositive}{TruePositive + FalsePositive} \hspace{0.5in} (2)$$

$$F_1 = 2 * \frac{Precision * Recall}{Precision + Recall}$$
 (3)

1) Full Feature Set: The results using the full unabridged original feature set can be found in Table I. Only n-grams where n=1,2 are explored due to memory limitations. The total number of features are 112 and 12544 for each respective n-gram. In general, each metric improved when more features were used. Decision tree, random forests, neural network, and the stochastic gradient decent classifier perform nearly equally and better than other algorithms with regards to their  $F_1$  score.

- 2) Full Feature Set with Feature Reduction: The full unabridged original feature set with feature reduction can be found in Table II. The 112 and 12544 features in the full feature set where n=1,2 is reduced to 100 features. This is done through incremental principal component analysis (IPCA) which is described in Section II-C. Desicion tree, random forest, and stochastic gradient decent classifier performed better than other algorithms tested.
  - 3) Trimmed Feature Set:

## IV. DISCUSSION

When n=1, the feature vector only consists of boolean values of whether or not the grammatical rule was used to parse a sample or not. When n>1, how grammatical rules interact with one another starts to be explored. The improved results show that the way malware is structured is different than benign samples. This is further highlighted in different feature sets.

When feature reduction was used, results were similar to the non reduced feature set.

V. RELATED WORK

VI. CONCLUSION

The conclusion goes here.

ACKNOWLEDGMENT

The authors would like to thank...

# REFERENCES

- [1] Charlie Curtsinger, Benjamin Livshits, Benjamin G Zorn, and Christian Seifert. Zozzle: Fast and precise in-browser javascript malware detection. In *USENIX Security Symposium*, pages 33–48. San Francisco, 2011.
- [2] Antlr. https://www.antlr.org/. Accessed: 2019-08-27.

TABLE I FULL FEATURE

N-Gram	Algorithm	Total Features	True Positive %	False Positive %	True Negative %	False Negative %	Recall	Precision	$F_1$
1	Decision Tree	112	98.7267	6.0	94.0	1.2733	0.9873	0.9427	0.9645
1	Random Forest	112	98.72	5.88	94.12	1.28	0.9872	0.9438	0.965
1	Neural Network	112	94.76	6.58	93.42	5.24	0.9476	0.9351	0.9413
1	Naive Bayes	112	99.8333	76.9867	23.0133	0.1667	0.9983	0.5646	0.7213
1	Stochastic Gradient Descent Classifier	112	98.08	7.16	92.84	1.92	0.9808	0.932	0.9558
1	K-Nearest Neighbors	112	82.3933	0.1333	99.8667	17.6067	0.8239	0.9984	0.9028
2	Decision Tree	12544	99.9667	5.44	94.56	0.0333	0.9997	0.9484	0.9734
2	Random Forest	12544	99.9333	5.3333	94.6667	0.0667	0.9993	0.9493	0.9737
2	Neural Network	12544	99.9933	5.42	94.58	0.0067	0.9999	0.9486	0.9736
2	Naive Bayes	12544	99.96	31.7867	68.2133	0.04	0.9996	0.7587	0.8627
2	Stochastic Gradient Descent Classifier	12544	99.96	5.52	94.48	0.04	0.9996	0.9477	0.9729
2	K-Nearest Neighbors	12544	86.12	0.14	99.86	13.88	0.8612	0.9984	0.9247

TABLE II FULL FEATURE - FEATURE SELECTION

N-Gram	Algorithm	Total Features	True Positive %	False Positive %	True Negative %	False Negative %	Recall	Precision	$F_1$
1	Decision Tree	100	98.6733	5.8333	94.1667	1.3267	0.9867	0.9442	0.965
1	Random Forest	100	98.64	5.6	94.4	1.36	0.9864	0.9463	0.9659
1	Neural Network	100	95.7	61.9667	38.0333	4.3	0.957	0.607	0.7428
1	Naive Bayes	100	95.7067	73.5067	26.4933	4.2933	0.9571	0.5656	0.711
1	Stochastic Gradient Descent Classifier	100	98.36	7.0	93.0	1.64	0.9836	0.9336	0.9579
1	K-Nearest Neighbors	100	84.7333	1.1733	98.8267	15.2667	0.8473	0.9863	0.9116
2	Decision Tree	100	99.8867	5.58	94.42	0.1133	0.9989	0.9471	0.9723
2	Random Forest	100	99.8133	5.3733	94.6267	0.1867	0.9981	0.9489	0.9729
2	Neural Network	100	94.7867	34.2067	65.7933	5.2133	0.9479	0.7348	0.8279
2	Naive Bayes	100	95.2667	40.7533	59.2467	4.7333	0.9527	0.7004	0.8073
2	Stochastic Gradient Descent Classifier	100	99.9533	5.62	94.38	0.0467	0.9995	0.9468	0.9724
2	K-Nearest Neighbors	100	92.7	2.64	97.36	7.3	0.927	0.9723	0.9491

TABLE III TRIMMED FEATURES

N-Gram	Algorithm	Total Features	True Positive %	False Positive %	True Negative %	False Negative %	Recall	Precision	$F_1$
1	Decision Tree	62	99.92	10.34	89.66	0.08	0.9992	0.9062	0.9504
1	Random Forest	62	99.92	10.2267	89.7733	0.08	0.9992	0.9072	0.951
1	Neural Network	62	49.4	50.6	49.4	50.6	0.494	0.494	0.494
1	Naive Bayes	62	100.0	98.4733	1.5267	0.0	1.0	0.5038	0.6701
1	Stochastic Gradient Descent Classifier	62	98.4867	13.8867	86.1133	1.5133	0.9849	0.8764	0.9275
1	K-Nearest Neighbors	62	65.4467	1.1933	98.8067	34.5533	0.6545	0.9821	0.7855
2	Decision Tree	3844	99.9067	5.26	94.74	0.0933	0.9991	0.95	0.9739
2	Random Forest	3844	99.8933	5.1933	94.8067	0.1067	0.9989	0.9506	0.9742
2	Neural Network	3844	13.9067	5.16	94.84	86.0933	0.1391	0.7294	0.2336
2	Naive Bayes	3844	99.8533	49.1333	50.8667	0.1467	0.9985	0.6702	0.8021
2	Stochastic Gradient Descent Classifier	3844	99.8933	6.1333	93.8667	0.1067	0.9989	0.9422	0.9697
2	K-Nearest Neighbors	3844	86.04	0.06	99.94	13.96	0.8604	0.9993	0.9247

TABLE IV
TRIMMED FEATURES - FEATURE SELECTION

N-Gram	Algorithm	Total Features	True Positive %	False Positive %	True Negative %	False Negative %	Recall	Precision	$F_1$
2	Decision Tree	100	99.84	5.4867	94.5133	0.16	0.9984	0.9479	0.9725
2	Random Forest	100	99.7933	5.2667	94.7333	0.2067	0.9979	0.9499	0.9733
2	Neural Network	100	98.3933	53.06	46.94	1.6067	0.9839	0.6497	0.7826
2	Naive Bayes	100	94.4467	58.5467	41.4533	5.5533	0.9445	0.6173	0.7466
2	Stochastic Gradient Descent Classifier	100	99.8667	6.14	93.86	0.1333	0.9987	0.9421	0.9695
2	K-Nearest Neighbors	100	99.9533	5.4067	94.5933	0.0467	0.9995	0.9487	0.9734
3	Decision Tree	100	99.8867	5.6867	94.3133	0.1133	0.9989	0.9461	0.9718
3	Random Forest	100	99.7467	5.4667	94.5333	0.2533	0.9975	0.948	0.9721
3	Neural Network	100	97.14	68.58	31.42	2.86	0.9714	0.5862	0.7311
3	Naive Bayes	100	94.2	41.24	58.76	5.8	0.942	0.6955	0.8002
3	Stochastic Gradient Descent Classifier	100	99.9267	5.86	94.14	0.0733	0.9993	0.9446	0.9712
3	K-Nearest Neighbors	100	99.9667	5.56	94.44	0.0333	0.9997	0.9473	0.9728

TABLE V EXTRA TRIMMED FEATURES

N-Gram	Algorithm	Total Features	True Positive %	False Positive %	True Negative %	False Negative %	Recall	Precision	$F_1$
1	Decision Tree	40	95.6333	8.3733	91.6267	4.3667	0.9563	0.9195	0.9376
1	Random Forest	40	95.6533	8.3	91.7	4.3467	0.9565	0.9202	0.938
1	Neural Network	40	96.0867	11.6267	88.3733	3.9133	0.9609	0.8921	0.9252
1	Naive Bayes	40	100.0	98.5867	1.4133	0.0	1.0	0.5036	0.6698
1	Stochastic Gradient Descent Classifier	40	94.8867	12.8267	87.1733	5.1133	0.9489	0.8809	0.9136
1	K-Nearest Neighbors	40	44.4133	0.1333	99.8667	55.5867	0.4441	0.997	0.6145
2	Decision Tree	1600	99.9	5.1667	94.8333	0.1	0.999	0.9508	0.9743
2	Random Forest	1600	99.8933	4.9667	95.0333	0.1067	0.9989	0.9526	0.9752
2	Neural Network	1600	83.4267	27.28	72.72	16.5733	0.8343	0.7536	0.7919
2	Naive Bayes	1600	99.94	51.22	48.78	0.06	0.9994	0.6612	0.7958
2	Stochastic Gradient Descent Classifier	1600	99.94	5.94	94.06	0.06	0.9994	0.9439	0.9709
2	K-Nearest Neighbors	1600	86.0533	0.08	99.92	13.9467	0.8605	0.9991	0.9246
3	Decision Tree	64000	99.9333	5.3	94.7	0.0667	0.9993	0.9496	0.9739
3	Random Forest	64000	99.8867	5.16	94.84	0.1133	0.9989	0.9509	0.9743
3	Neural Network	64000	49.0067	50.9933	49.0067	50.9933	0.4901	0.4901	0.4901
3	Naive Bayes	64000	99.9133	43.3733	56.6267	0.0867	0.9991	0.6973	0.8214
3	Stochastic Gradient Descent Classifier	64000	99.9533	5.66	94.34	0.0467	0.9995	0.9464	0.9722
3	K-Nearest Neighbors	64000	86.0733	0.1067	99.8933	13.9267	0.8607	0.9988	0.9246

TABLE VI EXTRA TRIMMED FEATURES - FEATURE SELECTION

N-Gram	Algorithm	Total Features	True Positive %	False Positive %	True Negative %	False Negative %	Recall	Precision	$F_1$
2	Decision Tree	100	99.84	5.3533	94.6467	0.16	0.9984	0.9491	0.9731
2	Random Forest	100	99.7733	5.0333	94.9667	0.2267	0.9977	0.952	0.9743
2	Neural Network	100	97.3733	77.8733	22.1267	2.6267	0.9737	0.5556	0.7075
2	Naive Bayes	100	95.4467	64.2133	35.7867	4.5533	0.9545	0.5978	0.7352
2	Stochastic Gradient Descent Classifier	100	99.8933	6.0	94.0	0.1067	0.9989	0.9433	0.9703
2	K-Nearest Neighbors	100	88.9133	1.0867	98.9133	11.0867	0.8891	0.9879	0.9359
3	Decision Tree	100	99.8533	5.56	94.44	0.1467	0.9985	0.9473	0.9722
3	Random Forest	100	99.68	5.24	94.76	0.32	0.9968	0.9501	0.9729
3	Neural Network	100	84.26	35.4267	64.5733	15.74	0.8426	0.704	0.7671
3	Naive Bayes	100	95.24	43.86	56.14	4.76	0.9524	0.6847	0.7967
3	Stochastic Gradient Descent Classifier	100	99.9067	5.76	94.24	0.0933	0.9991	0.9455	0.9715
3	K-Nearest Neighbors	100	90.3333	1.7533	98.2467	9.6667	0.9033	0.981	0.9405