DSC180B Capstone Project Report

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Abstract

Nowadays, smartphone is an indispensable part of people's daily life. Android System is the most popular system running on smartphone. Due to this popularity, malware detection on Android becomes on of the most significant task for research community. In this project, we are mainly focusing on one called MAMADROID System. Instead of previous work which highly relied on the permissions requested by apps, MAMADROID relied on the sequences of abstracted API calls performed by apps. We are very interested in finding ways to improve this model. To achieve this, we've been trying to find some new features to fit into the model. We made three basic model and take the one with the highest accuracy and made two more advanced model based on this model with the best performance.

1 Introduction

During 2019, 87% of the smartphone sales were running Android system. Due to this popularity, cyber-criminals have increasingly targeted this ecosystem, as malware running on mobile devices can be particularly lucrative. As a result, the research community has devoted significant attention to malware detection on Android system. Previous work has often relied on the permissions requested by apps, using models built from malware samples. This strategy, however, is prone to false positives, since there are often legitimate reasons for benign apps to request permission classified as dangerous. To overcome this obstacle, research community has developed a novel system for malware detection called MAMADROID System[1]. Instead of relying on the permissions requested by apps, MAMADROID System relies on the sequences of abstracted API calls performed by an app rather than their use or frequency, using Markov Chains to model the behavior of the apps through the sequences of API calls. By doing research on this novel system, we are very interested in finding a way to better understand this system and improve it. Therefore, our research question is: Exploring and explaining potential improvement of MAMADROID **System.** To achieve this, our plan was to find other features to replace the previous one used in Markov Chains model.

2 Data Generating Process

2.1 Source of Data

Our raw apps data are collected from course DSMPL. We randomly collected 77 apps contain malware and 143 benign apps, 220 apps data in total. Then we divided them into 147 apps for training and 73 apps for testing. Since the raw data have been classified into different categories, we are 100% percent sure that those benign apps don't contain any malware.

2.2 Data Description

The original form of our data was Android Application Package (APK) which could be unpacked by Apktools. We unpacked those packages to get the Smali specifically for malware detection. Smali file is a type of file convert from the original Java code of an app. An example of the structure of a Smali file and the API call inside it is shown below:

```
.class public final LGeckoProvider;
.super Ljava/lang/Object;
.source "GeckoProvider.kt"

# static fields
.field public static final INSTANCE:LGeckoProvider;
.field private static
runtime:Lorg/mozilla/geckoview/GeckoRuntime;
.field private static testConfig:Landroid/os/Bundle;

# direct methods
.method static constructor <clinit>()V
.locals 1
.line 1
new-instance v0, LGeckoProvider;
invoke-direct {v0}, LGeckoProvider;-><init>()V
return-void
.end method
```

Figure 1: An example of Smali file

In the figure above, text in black represent the class information of a smali file. In this specific example, the class name is **LGeckoProvider**. Text in blue is Static fields of this Smali file and text in green is the method of this Smali file. Text in red is what we are going to focus on: **API calls**. Based on previous works done by research community, malicious action of an app is always appeared in Smali files so that use Smali file for malware detection is significantly meaningful. The API calls we are using is extracted from Smali files for each application. By dealing with API calls, we could be more aware of the characteristic of a malware so that we could better analyze a way for malware detection.

2.3 Feature Extraction

The original MAMADROID System used Markov Chains to model app behavior, by evaluating transitions between calls. For each app, MAMADROID System takes as input the sequence of abstracted API calls (families/packages) of that app and builds a Markov chain where each package/family is a state and the transitions represent the probability of moving from one state to another. To improve this, we first try to take input as the method sequences of API calls instead of families/packages. Thus, the feature that we are going to use is the **method sequences** of API calls where each method and the API calls it contains represent states and the transitions represent the probability of a method moving to each API calls inside it. Here is an example of our feature:



Figure 2: Feature example

In the figure above, the blue box is a method and content in the grey box is API calls contain in this method. Our model used this feature to fit into Markov Chains to evaluate the behavior of an app and then classified it into either benign or malware.

3 Model

3.1 Model Description

Using the features derived from data extracting process, based on the fact that the predictive task is generally classification problem, we have built 3 models in this project: Logistic Regression Model, Decision Tree Classifier, K-Nearest Neighbor-Classifier. This is the table which contains the test accuracy of our three models.

Model	Train Accuracy	Test Accuracy
Logistic	0.721	0.822
DecisionTree	0.918	0.74
kNN	0.925	0.918

kNN, which represents K-Nearest-Neighbor Classifier, performed best among all three models.

3.2 K-Nearest-Neighbor Classifier

In K-Nearest-Neighbor model, each data point is composed by a vector of six values: the number of Api-calls from android, androidx, java, javax, kotlin, self-defined families. A sample observation of vector would be like the following.

Index	android	androidx	java	javax	kotlin	self
1	5382	21	8367	0	0	4362

During prediction process, KNN model will search through data points in the training set to find N-nearest data points of the input point and predict the result to be the majority of these neighbors. For instance, we set number of neighbors to be 5 and get five data points. Two of them are malware and three of them are benign software. Then the model will predict it to be the majority of its neighbors, which is benign software.

3.3 Model Optimization

Since kNN model performed the best among all three, we choose this model and tried to optimize this model. To improve this model, we tried several different values of K, from 1 to 20. The following figure 3 shows how test accuracy and train accuracy changed along with different K.

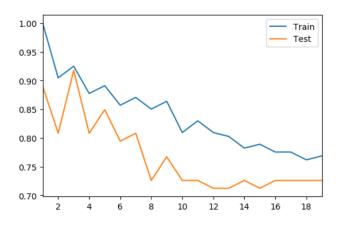


Figure 3: kNN

Based on figure 3, it is obvious that when K=3, the model has the greatest test accuracy and train accuracy (K=1 has the highest train accuracy since K=1 means finding the data point itself).

K	Train Accuracy	Test Accuracy
1	1	0.89
3	0.925	0.918
5	0.891	0.849

3.4 Advanced Model

We then focus on the predicted results to discover ways to improve our current 3-Nearest-Neighbor model. To research the predicted results in a more detailed way, we generate TP-TN-FP-FN matrix to reveal deeper information behind model performance. FP (A benignware predicted to be malware) and FN (A malware predicted to be benignware) are the most important categories since they shows mistakes made by the model. FN has more significant meaning since letting a malware pass detection would be a disaster. The following table shows the statistics.

	Positive	Negative
True	62	137
False	6	15

We classify all software according to TP-TN-FP-FN and draw figure of mean of numbers of api-call with respect to different families.

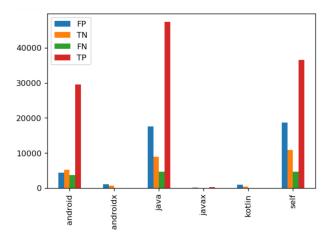


Figure 4: TP-TN-FP-FN

According to the graph, malware which were classified to be benign software have fewer number of api-calls in general compared to TP malware. Also, FN malware have larger possibility of invoking self defined api-calls than java calls. FP benignware which were classified to be have larger number of api-calls compared to TN benignware. In addition, malware rarely call apis from androidx and kotlin families. Based on information above, we built our new advanced model which standardizes number of api-calls to fraction of total number of api-calls of specific software, sets threshold for FN malwares and contains a hybrid of KNN and our self-defined Decision Tree. Our model performs the best when K=3. The parameter search of our advanced model is showed below.

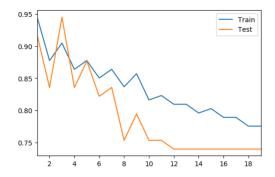


Figure 5: Advanced Model

4 Results

Below are the results of our models.

4.1 Model Comparison

Model	Train Accuracy	Test Accuracy
Advanced kNN	0.915	0.945
Standardized kNN	0.823	0.877
kNN	0.925	0.917

The accuracy comparisons of these models has shown that our advanced model performed the best among all three models.

4.2 Model Analysis

The best model, Advanced kNN, not only consider 3-Nearest-Neighbor but also covering edge cases. Comparing to other two models, Standardized kNN make the input standardized to percentile before training and predicting, but it will lost information about number of api-call. Normal kNN only considers major cases and does not take margins into consideration.

4.3 Data Findings

According to our research results: 1.Malware with fewer api-calls is more likely to be classified as benignware. 2.Malware tend to has more java api-calls than self-defined and android api-calls.

4.4 Significance

The result of this project has broad application in our daily life. If we can build a strong model to help Android users make accurate classification about whether a software is benign or harmful, it can protect information, privacy and save a lot of money for users.

References

[1] Mariconti Enrico, Onwuzurike Lucky, Andriotis Panagiotis, De Cristofaro Emiliano, Ross Gordon, and Stringhini Gianluca. *MAMADROID: Detecting Android Malware by Building Markov Chains of Behavioral Models.* 2017.