### Abstract

The project is based on Kaggle's Microsoft Malware Challenge 2015 to classify malware into families.

- It involves training the classifier using the given database to classify the malware files(binary executables) into 9 categories of malwares.
- The main challenge we have is identifying the distinguishing features in the bytes and the asm file for classifying malware into their respective classes.

## Introduction

Malware authors use automated techniques like **Polymorphism** in order to evade 'pattern matching' detection. Polymorphic malwares constantly changes itself, making it difficult for the anti-malware programs to detect it. Evolution of malicious code occurs in a variety of ways like change in filename, compression and encryption with variable keys.

- Random forest classifier According to previous research random forest the best classifier
- N-gram based file signatures
- K-fold Cross validation
- Following features were used to classify the malwares:
- malware
- Occurence of instructions like mov, jmp etc. in the asm file corresponding each malware
- <sup>3</sup> Frequency of 256 possible hex values at specific position in the asm file corresponding to each malware

- is achieved.
- Evaluation is done using Multi-Class Logarithmic Loss.
- For each malware file in the test set, their predicted probabilities for the 9 classes was submitted.

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# Methodology

The following methods were used to complete the research:

• Frequency of 256 possible hex values in the bytes file corresponding to e

# Result

• After using all the proposed features, a score of 0.153125351 on the leader

	Conclusion
t is	The goal is to take the score as close to 0 as possible. Though, the current score is not bad but it can be further improved by identifying more <b>distin-</b> <b>guishible</b> features for malwares be- longing to different families.
ach	Datset
ng to	Dataset in the form of .asm and .byte files
	<ul> <li>Training &amp; Test set - 200 GB each</li> <li>Asm file(0.4-19 millions lines)</li> </ul>
	References
rboard	<ul> <li>[1] Daniel Bilar et al.</li> <li>Statistical structures: Fingerprinting malware for classification and analysis.</li> <li><i>Proceedings of Black Hat Federal 2006</i>,</li> </ul>
9	<ul> <li>2006.</li> <li>[2] Karthik Raman.</li> <li>Selecting features to classify malware.</li> <li><i>InfoSec Southwest</i>, 2012.</li> </ul>