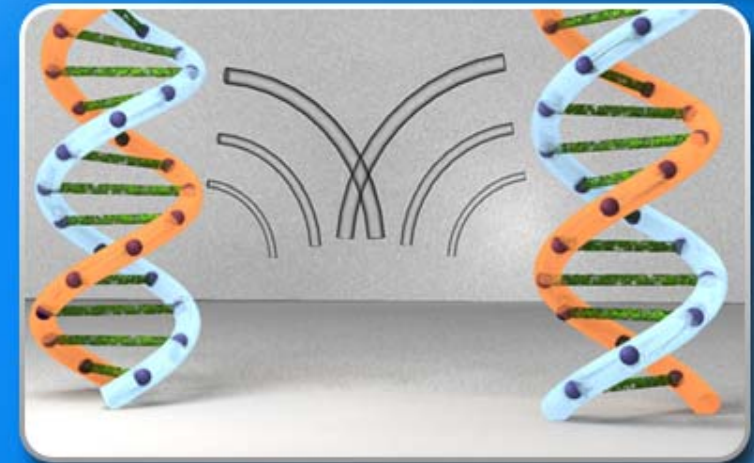


# Embedded Malware Detection using Markov n-grams



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# Agenda

Introduction to problem domain



Mathematical Modeling



Discussion on Results



Conclusions



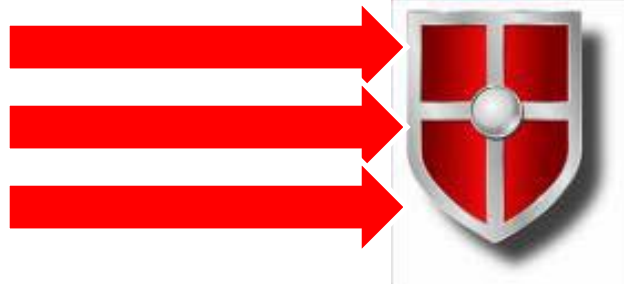
Future Work



# Introduction to Problem Domain



# Introduction



# Problem with State-of-the-art antivirus

## Signature matching

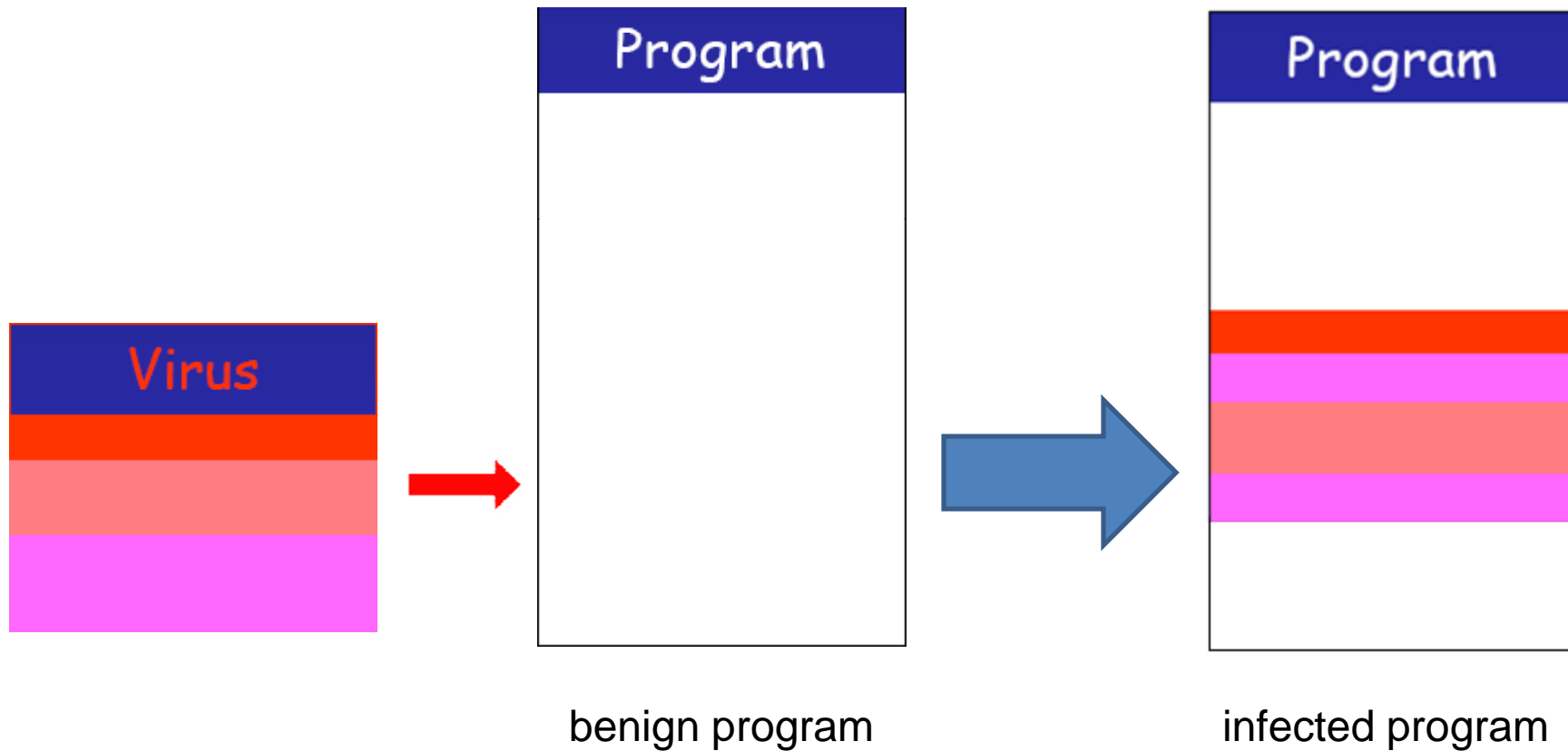
- Identify sequence of instructions unique to a virus => virus signature
- Match program to a database of signatures

## Problems

- Inability to detect **zero-day attacks**
- Scan only starting portions of files due to **high overhead** and **false alarms**; **vulnerable!**
- Size of signature database **cannot scale** in future



# Embedded Malware



# Mathematical Modeling



# Our Approach

## Anomaly detection

- Differentiate anomalous behavior from the normal workflow
- Primarily utilize *statistical modeling*

Statistical model of benign DOC, EXE, JPG, MP3, PDF, ZIP files using Markov n-grams

Feature extraction using well known information-theoretic measure, entropy rate

Threshold based detection using Gaussianity of sum of sampled entropy rate distribution





# n-gram analysis

## n-gram Definition [en-wikipedia]

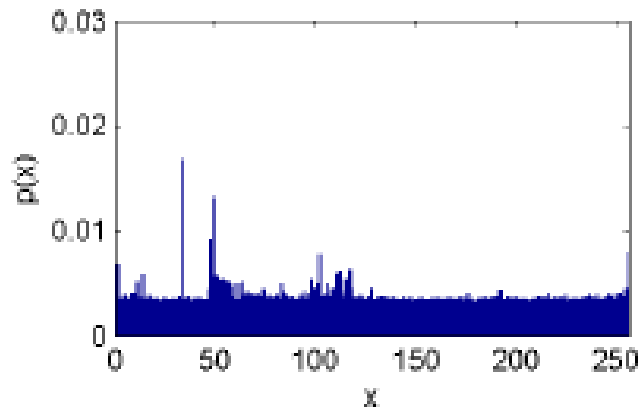
- An  $n$ -gram is a sequence of  $n$  symbols in a given sequence

11100111011001110011010011010010111000111111110  
11100111011001110011010011010011010010111000111111110  
11100111011001110011010011010011010010111000111111110

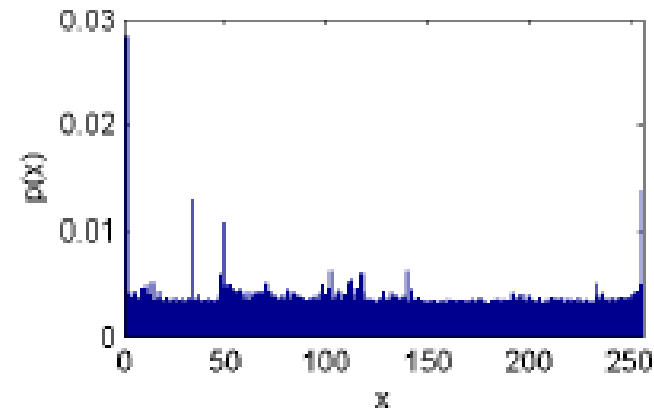
3-gram



# Whole file n-gram analysis



1-gram benign PDF



1-gram infected PDF

**Whole file, 1-gram analysis DOES NOT show significant perturbations ☹️**



# Block wise 1-gram analysis

Calculate Mahalanobis distance between benign model and given file in a block wise manner (blocksize = 1000bytes)

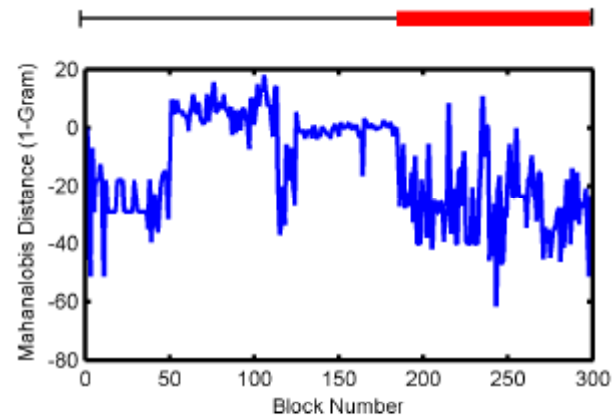
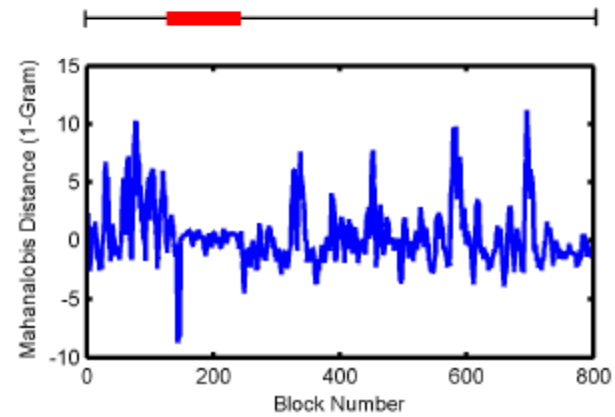
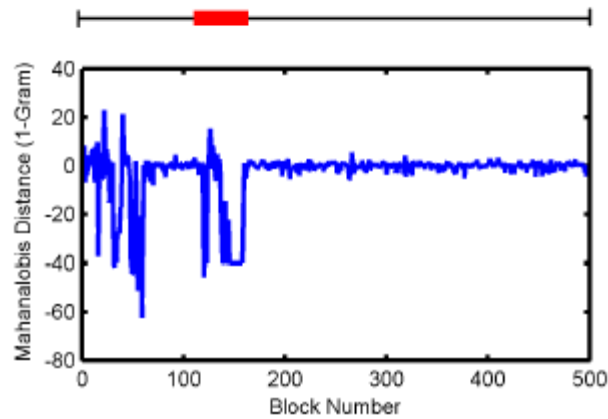
Benign model distribution is:

- average byte value frequency
- their standard deviation

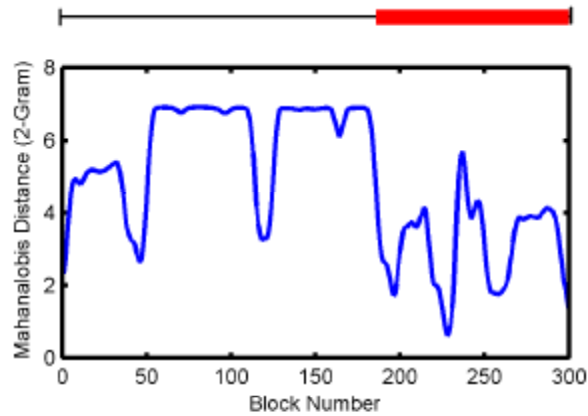
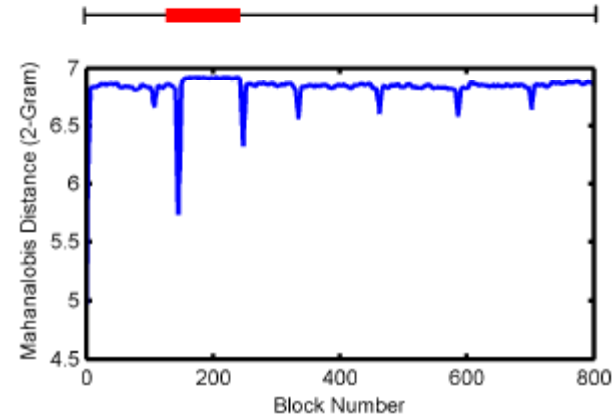
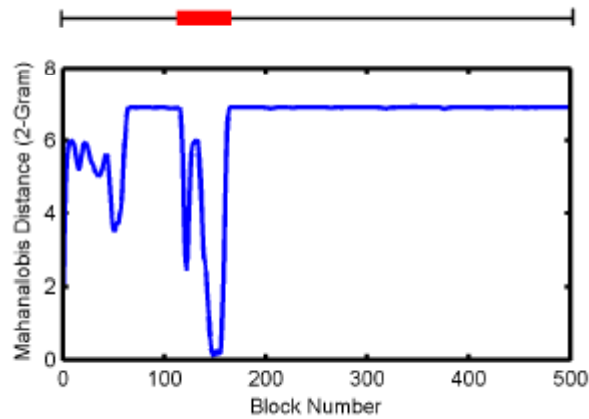
$$D(x, y) = \sum_{i=1}^{n-1} \frac{|x_i - y_i|}{\sigma_i + \alpha}$$



# Block wise 1-gram analysis



# Block wise 2-gram analysis



# Analysis of results

2-gram is better than 1-gram; at the cost of exponentially higher computational complexity!

Cannot increase  $n$  due to a fixed block size of 1000 bytes!

2-gram distribution is a joint distribution

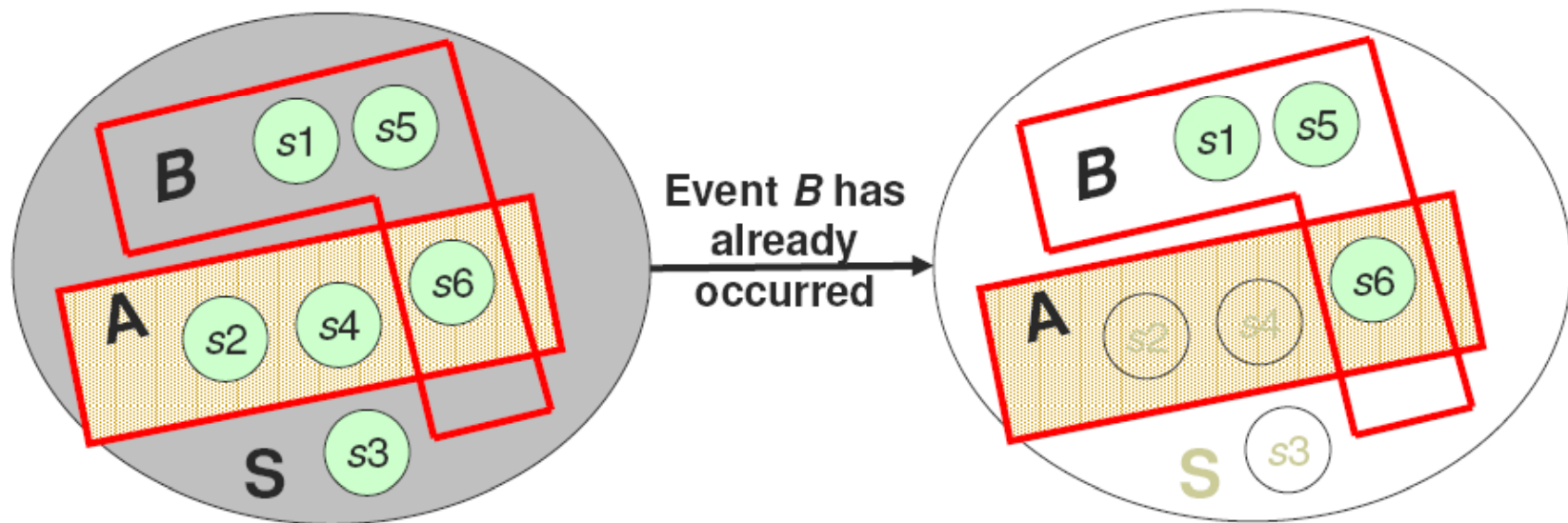
Some redundant information in joint distribution that can be removed using *conditional distribution*. **HOW?**



# Advantage of Conditional Distribution

$$Pr\{A|B\} = \frac{Pr\{A \cap B\}}{Pr\{B\}}$$

Reduction in sample space



# From Conditional Distribution to Markov Chain

A conditional distribution can be directly mapped to a Markov chain

Symbols of distribution mapped to states in Markov Chain

How to select order of Markov chain?

Byte level autocorrelation!

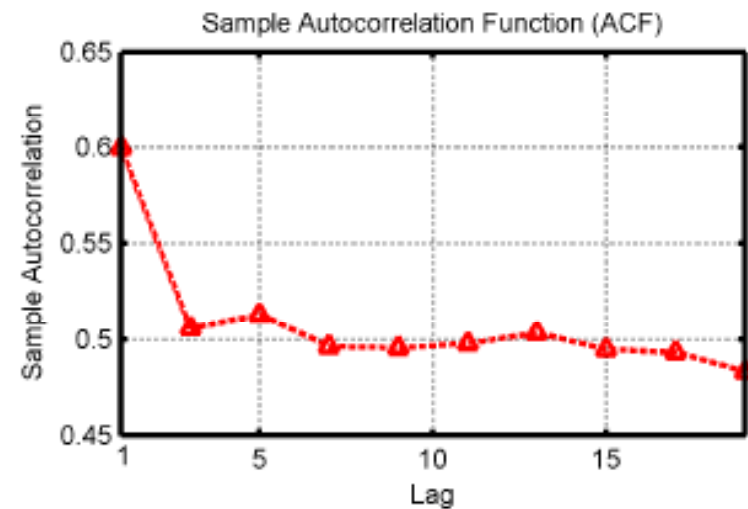
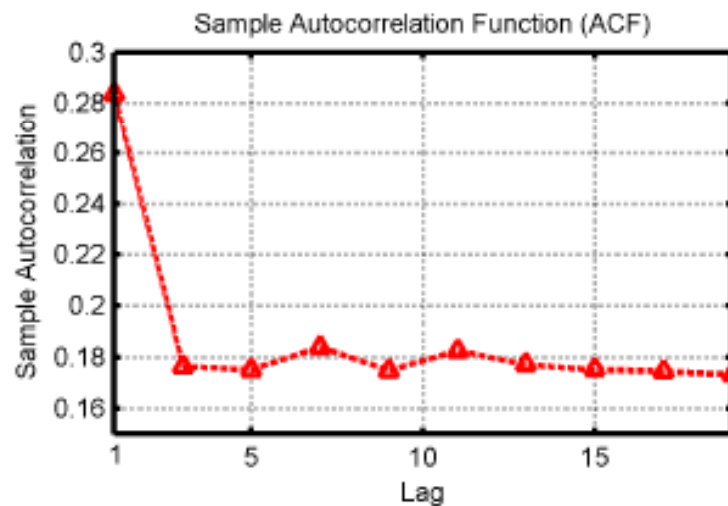




# Byte-level autocorrelation results

Random information beyond lag=1

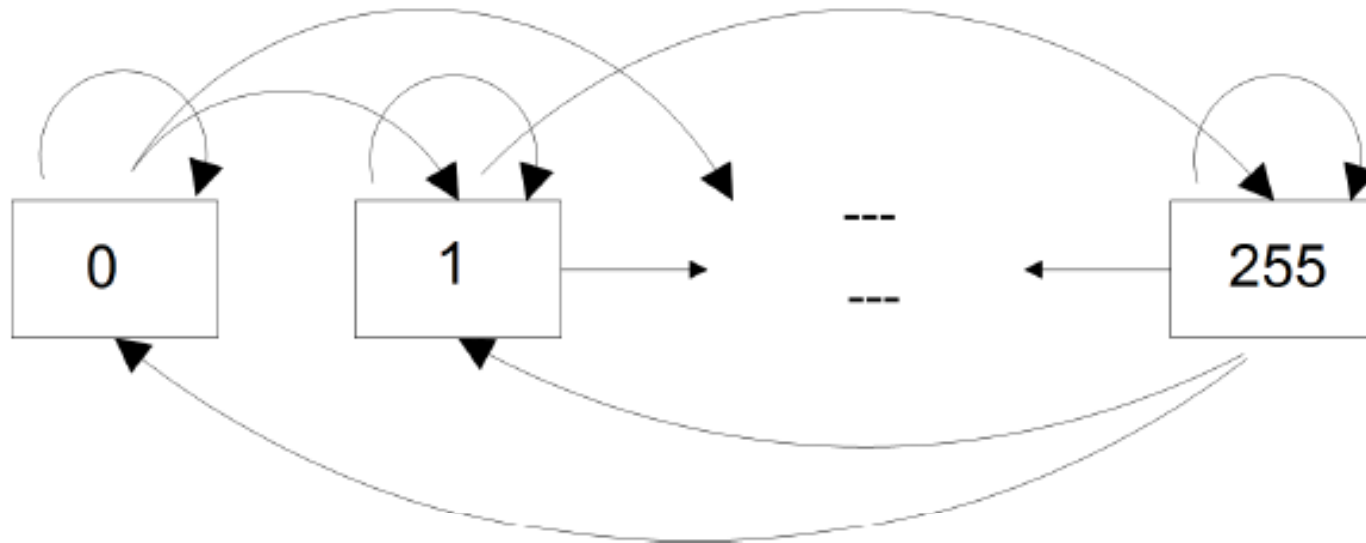
1st order Discrete Time Markov Chain



# Markov n-gram Model

1st order, 256 state Markov Chain

Can be constructed using conditional 2-gram distribution



# Quantification feature

average information in the Markov Chain

Entropy rate gives

- Time density of average information in the Markov chain

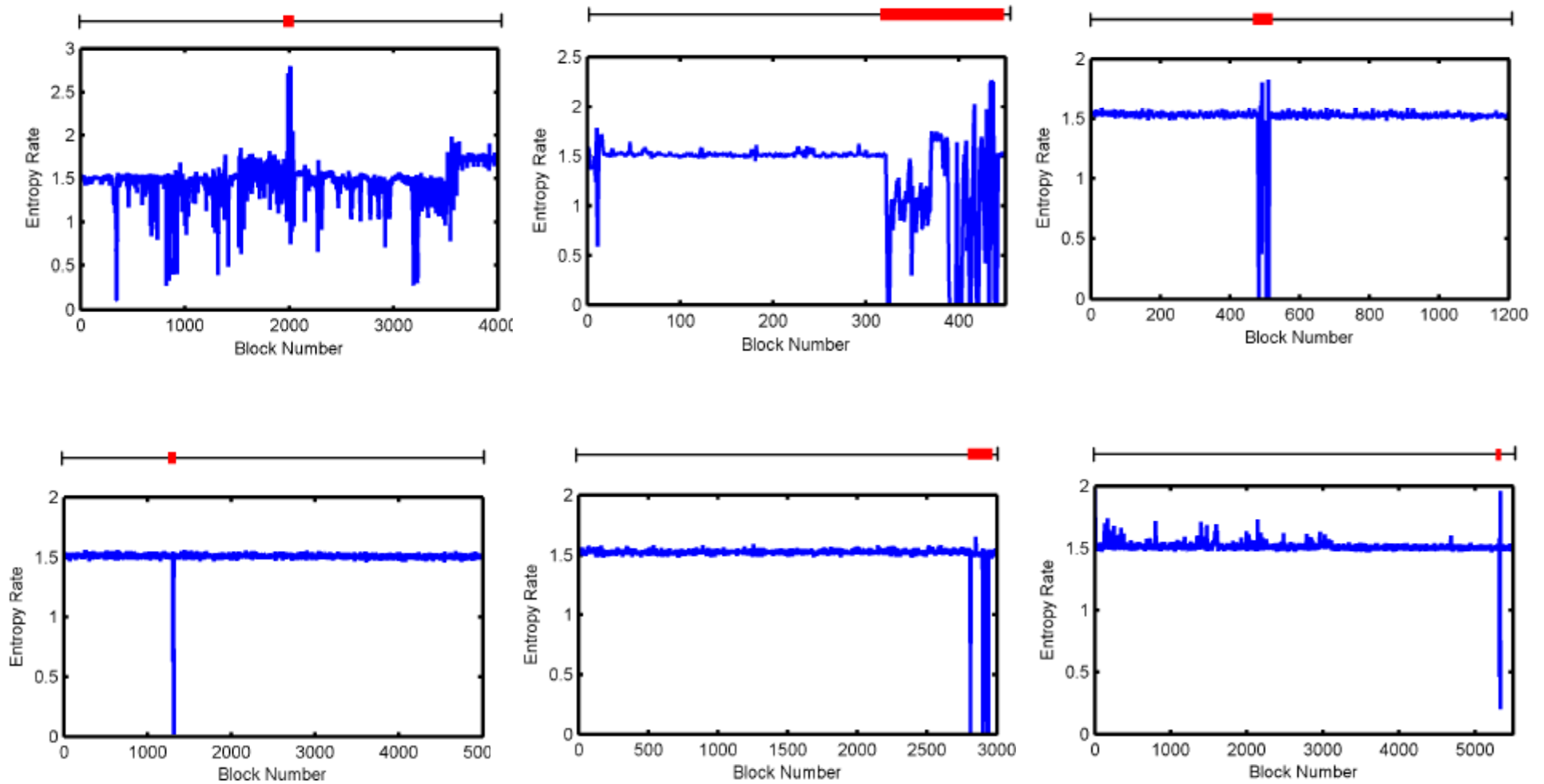
$$R = \lim_{N \rightarrow \infty} \frac{H(X_1, X_2, \dots, X_n)}{N}$$

For a 256 state Markov chain

$$R = \sum_{i=0}^{255} \pi_i H(X_i)$$



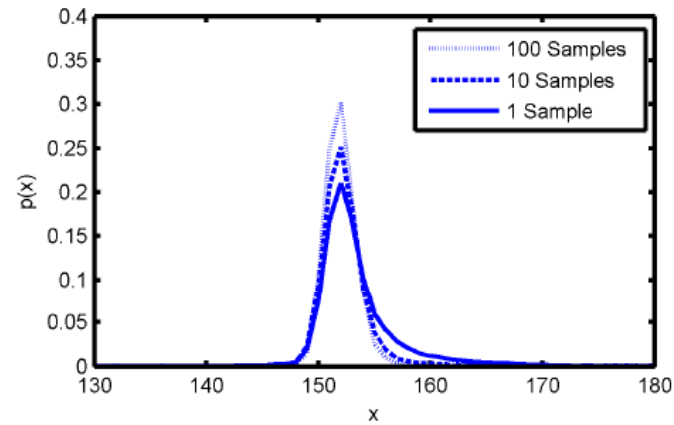
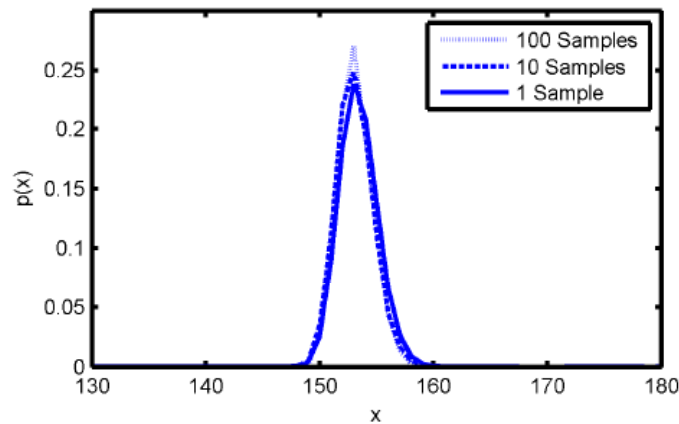
# Entropy rate Results



# Threshold selection

Sum of sampled entropy rate distributions approach Gaussianity\*

Threshold selected at  $\mu \pm 5\sigma$  (99.99%)



\* Direct consequence of central limit theorem



# Results

	Mahalanobis n-gram Detector (%)	Markov n-gram Detector (%)	Percentage Improvement (%)
<b>MP3</b>			
TP rate	63.8	95.0	<b>31.2</b>
FP rate	32.3	0.2	<b>32.1</b>
<b>JPG</b>			
TP rate	76.3	95.4	<b>19.1</b>
FP rate	35.7	2.7	<b>33.0</b>
<b>PDF</b>			
TP rate	75.4	84.5	<b>9.1</b>
FP rate	46.8	31.8	<b>15.0</b>
<b>ZIP</b>			
TP rate	60.0	90.4	<b>30.4</b>
FP rate	29.9	8.3	<b>21.6</b>
<b>EXE</b>			
TP rate	54.1	84.9	<b>30.8</b>
FP rate	47.3	16.7	<b>10.6</b>
<b>DOC</b>			
TP rate	65.6	66.3	<b>0.7</b>
FP rate	48.8	29.2	<b>19.6</b>

## Conclusion & Future Work



# Conclusion

## Advantages

- Ability to identify location of malware
- Significantly improved *TP rate* and *FP rate* as compared to Mahalanobis detector

## Limitations

- Still slightly high false positive rate for DOC and PDF files
- Embedded Malware -> dormant malware
- Mimicry Attacks





# Future Work

How to reduce false positive rate?

Complement with signature based detector

- Inability to detect zero-day attacks 😞

Multiple features and correlation

- Initial results have shown promise 😊
- Subject of forthcoming journal publication...

**Patent pending** on current work



# Thank you!



Contact authors for queries/suggestions:

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