

A large, faint watermark of a university seal is visible in the background. The seal features a central shield with the text 'GRAMM PHILOL RHETOR ETHICA' on the left and 'METAPH LOGICA MATHEM' on the right. Above the shield is a banner with the text 'SOLVIT INQUIRIT'. The seal is surrounded by a circular border with text, including 'UNIVERSITY OF MICHIGAN' and '1817'.

SigMal: A Static Signal Processing Based Malware Triage

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Abstract

- Signal as a malware detection framework
- Results of testing Signal on samples

Introduction

- Static, dynamic and statistical analyses
- Malwares variants
- N-gram feature extraction

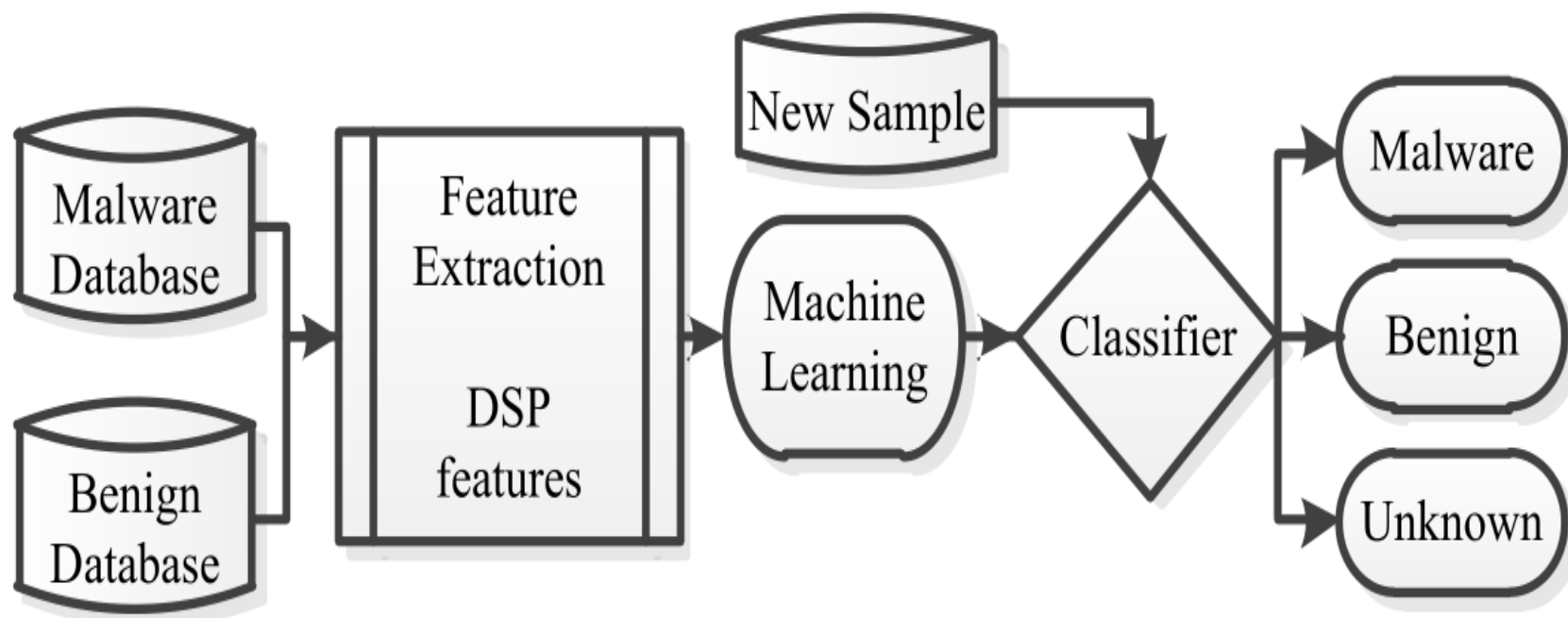


Figure 1: SigMal overview.

Signal processing based features

- Feature extraction, Feature computation
Section aware feature extraction

Data: PE Executable

Result: A list of important sections

Map sections into raw binary file;

if *overlapping section exists* **then**

| resize section to make it contiguous with adjacent sections;

end

if *.text executable section exists* **then**

| **if** *is the largest section* **then**

| | Result.append(*.text* section and the second largest
| | section);

| **else**

| | **if** *.text section is writable* **then**

| | | Result.append(two largest sections);

| | | **else**

| | | | Result.append(*.text* section and the largest section);

| | | | **end**

| **end**

else

| **if** *any non-writable executable section exists* **then**

| | Result.append(this section and the largest section);

| | **else**

| | | Result.append(two largest sections);

| | | **end**

end

Algorithm 1: Finding important sections.

Comparison

- N-gram based detection

$$J(s_a, s_b) = \frac{s_a \cap s_b}{s_a \cup s_b}$$

- PE structure based detection

- Control flow graph-based detection

$$CFG \text{ similarity} = \frac{\text{number of matching subgraphs}}{\text{total number of subgraphs}}$$

- Benign, Malicious and real world datasets collected

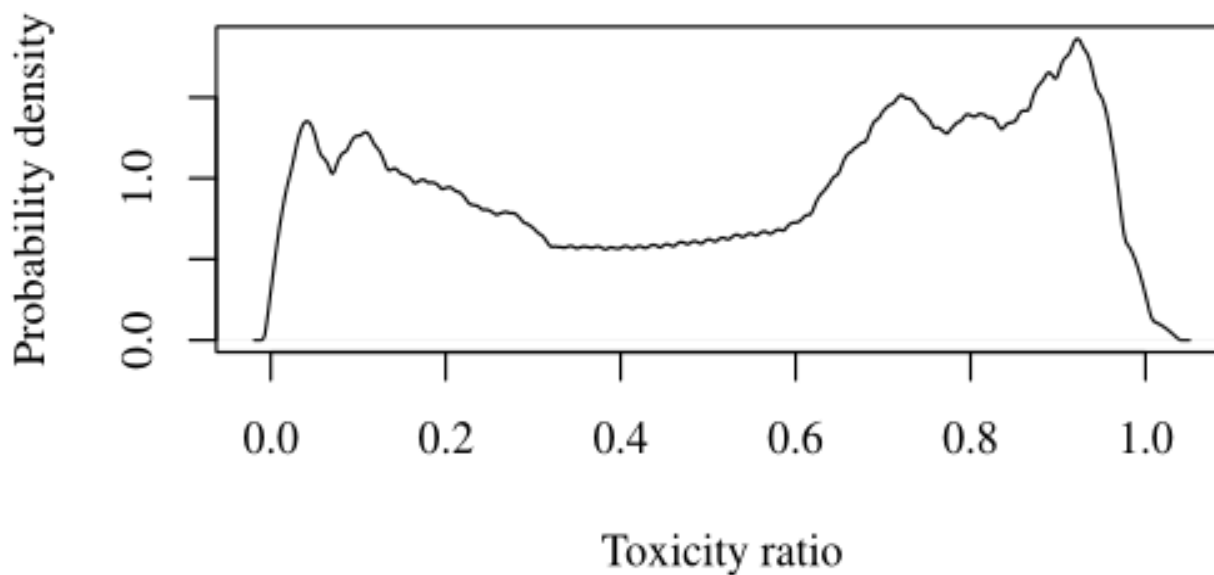
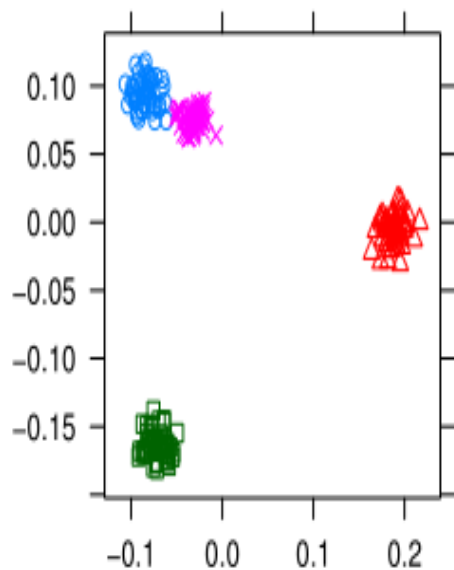


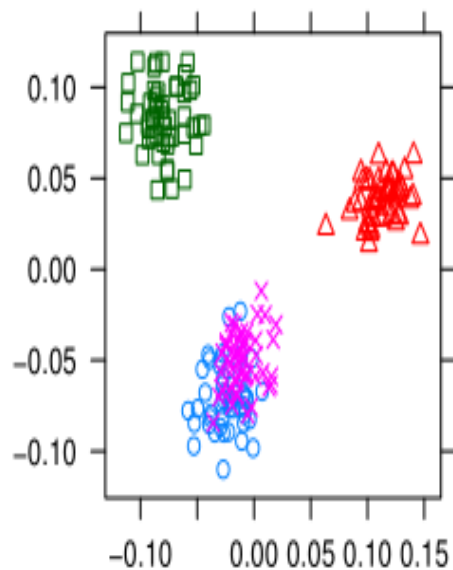
Figure 4: The *toxicity ratio* distribution of 1.2 million malware samples.

Evaluation



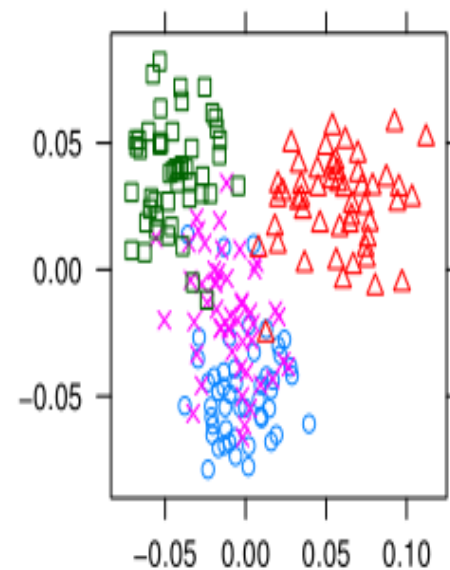
10% Noise

(a)



30% Noise

(b)



50% Noise

(c)

Fig 5: Feature robustness against noise.

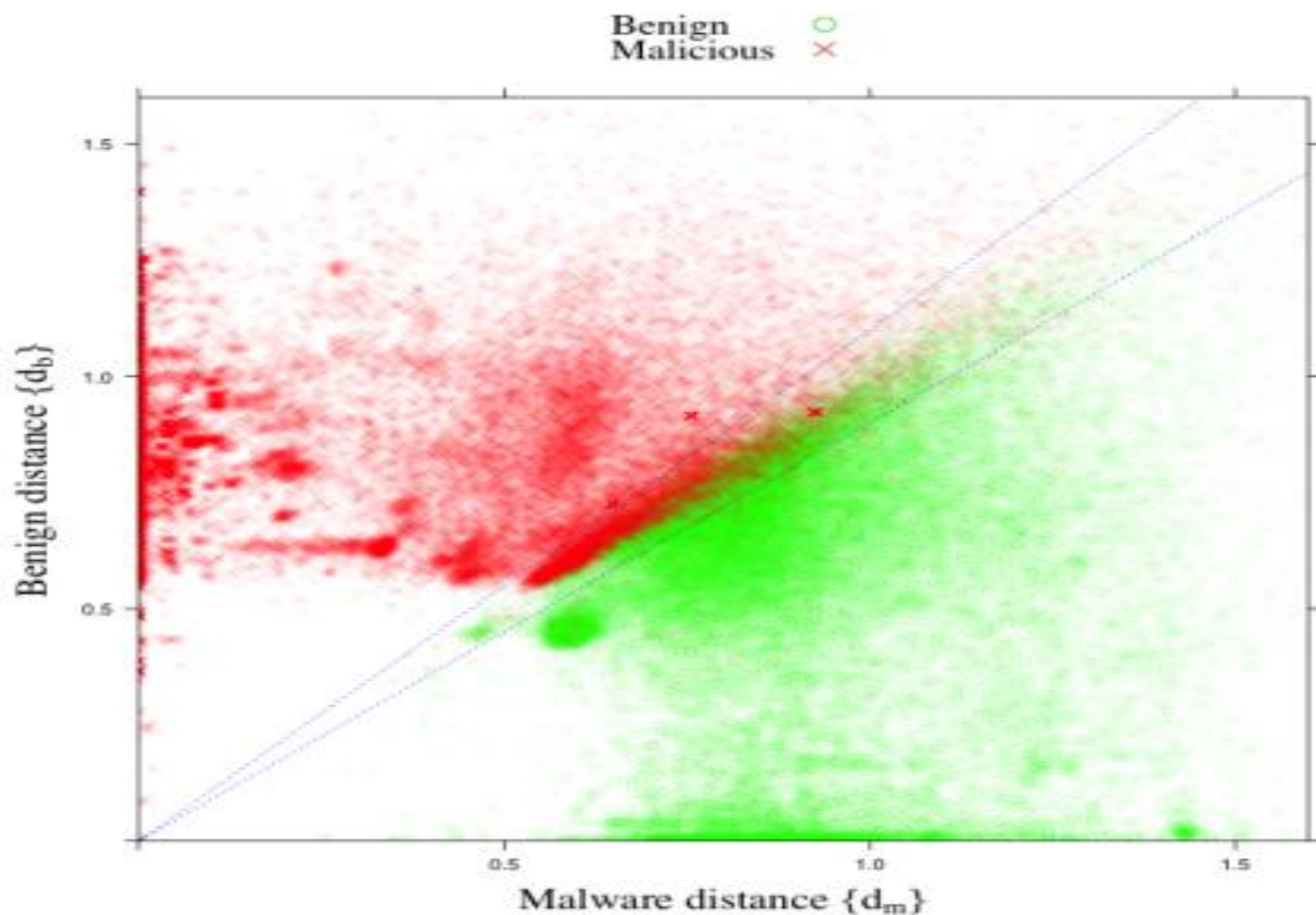


Fig. 6 : Nearest neighbor distribution for a 100 thousand samples

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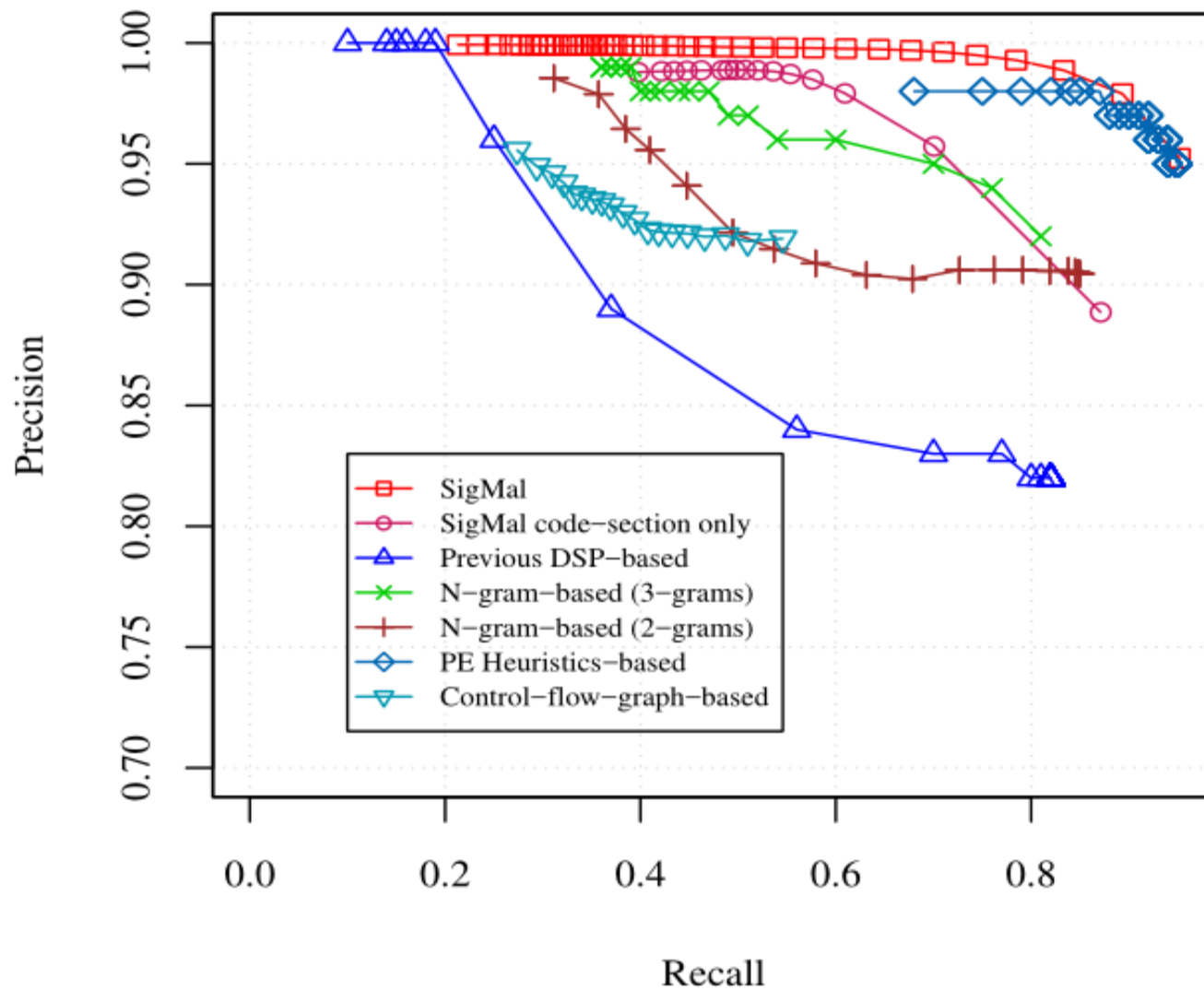
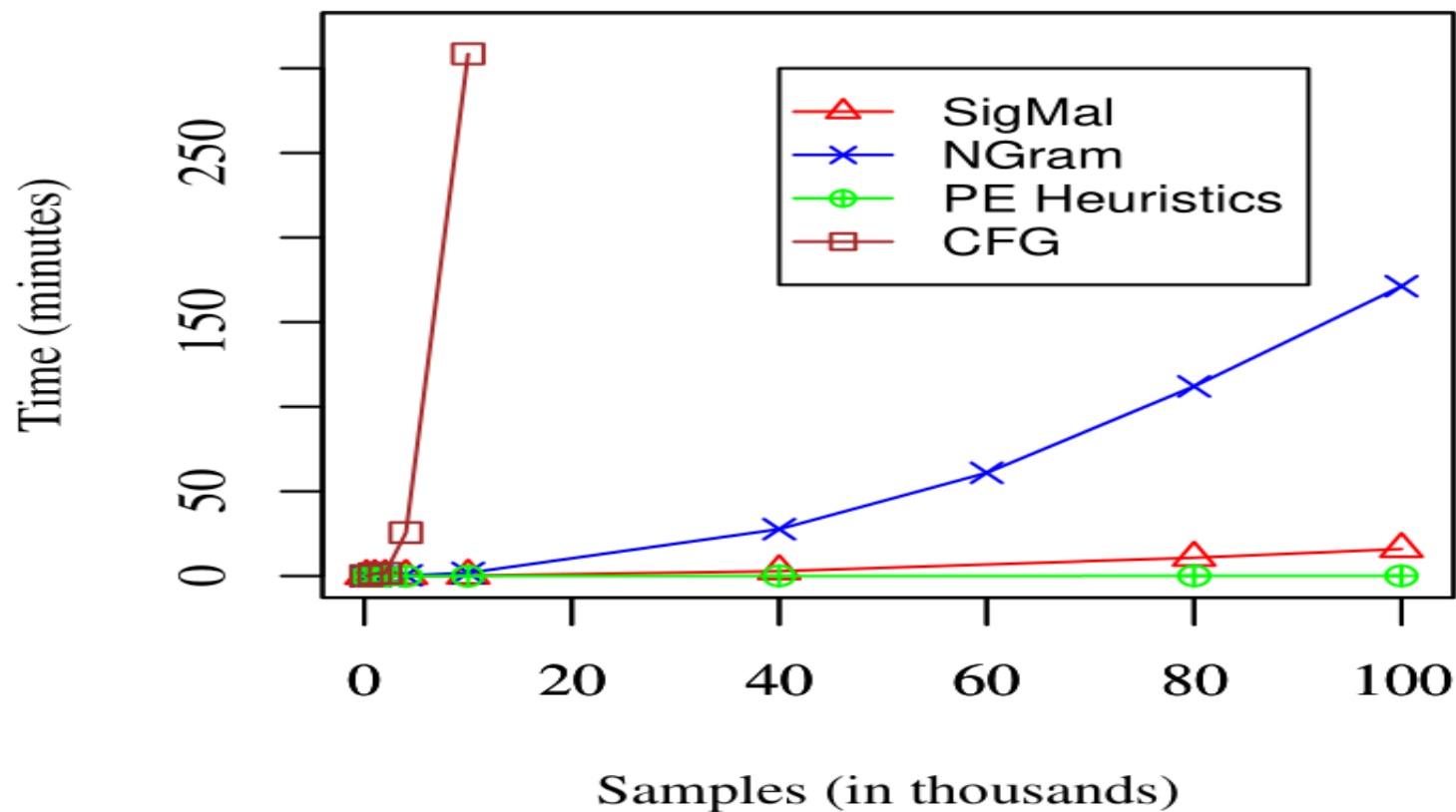


Fig. 7 : Comparison of malware detection algorithms



	<i>SigMal</i>	<i>N-gram</i>	<i>PE-heuristics</i>	<i>CFG</i>
<i>Time</i>	0.0265	0.1965	0.0024	0.1379
<i>Space</i>	3.783	8.000	0.0664	297.745

Fig. 8 : Query performance comparison.

Real world experiments

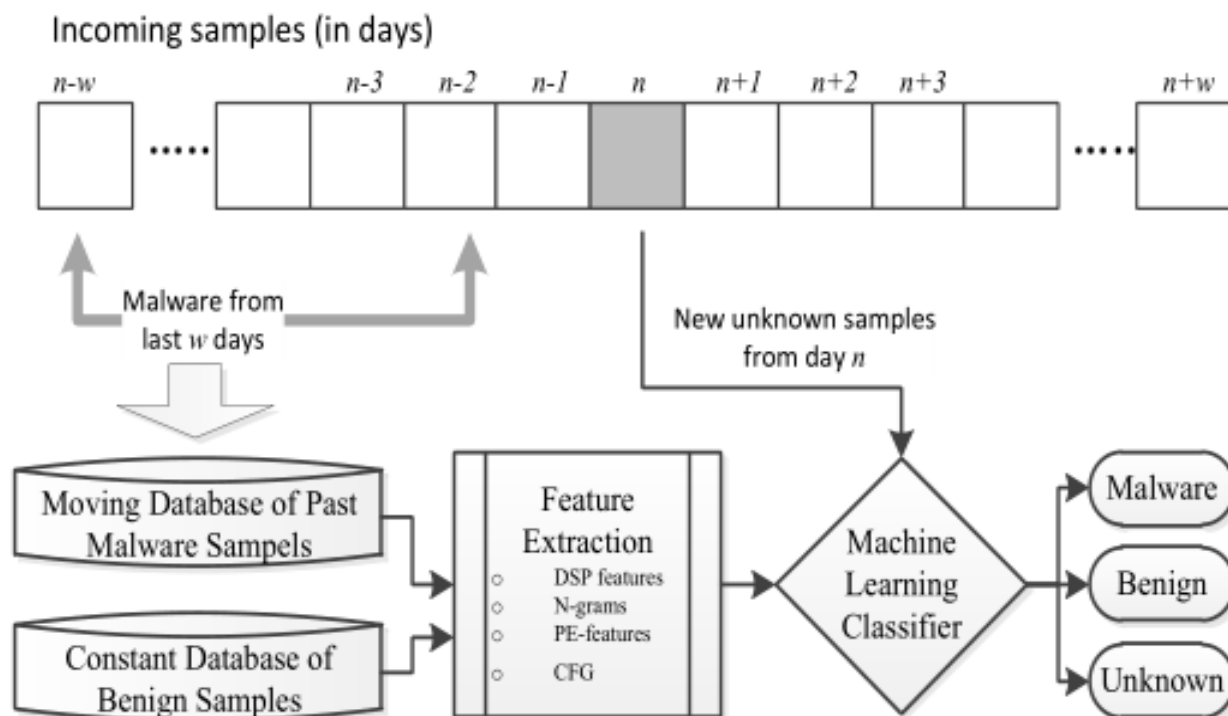


Figure 9: Overview of the sliding window experiment on the real world samples.

Results:

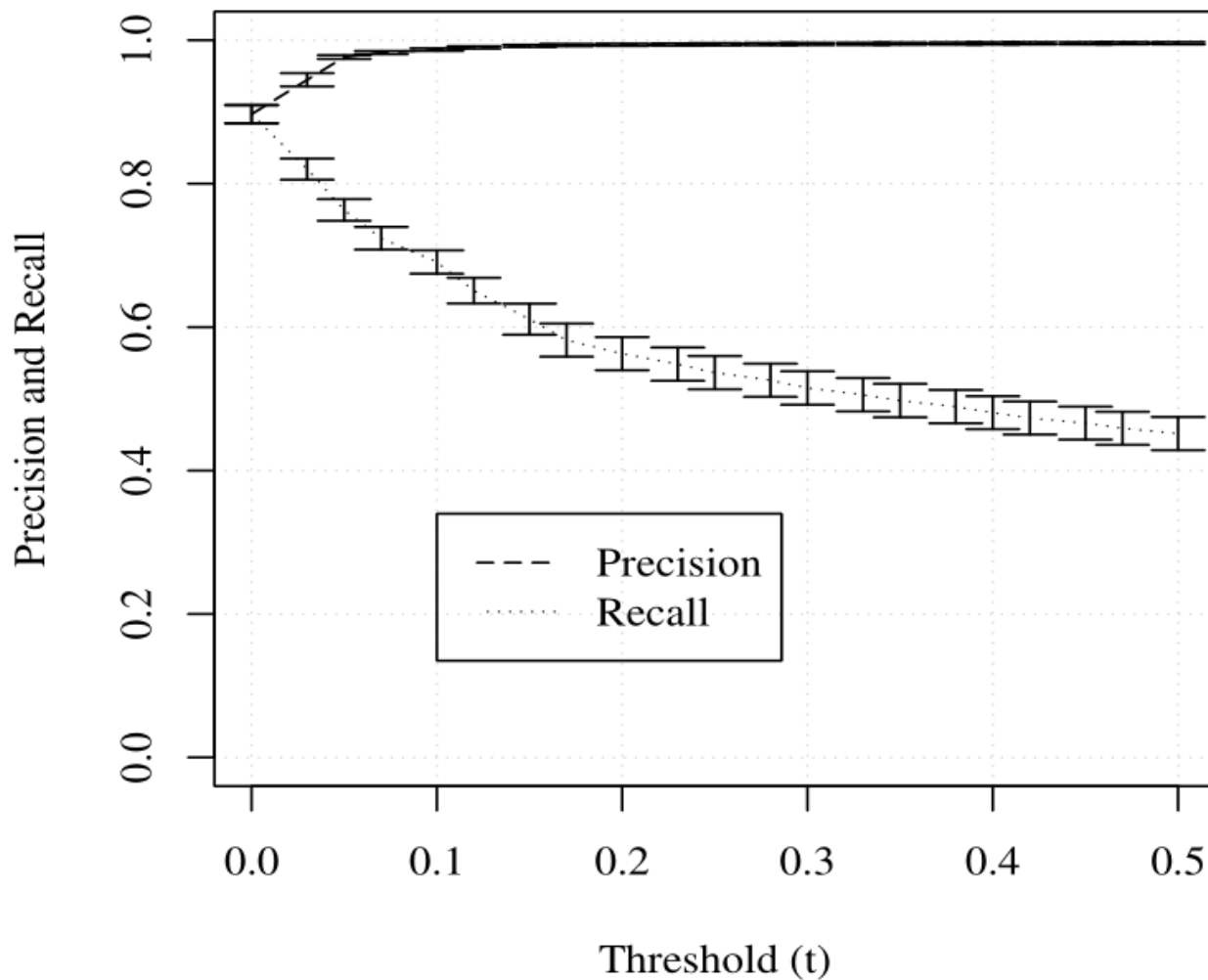


Fig. 10: Precision and recall of the Signal detection on the real world samples.

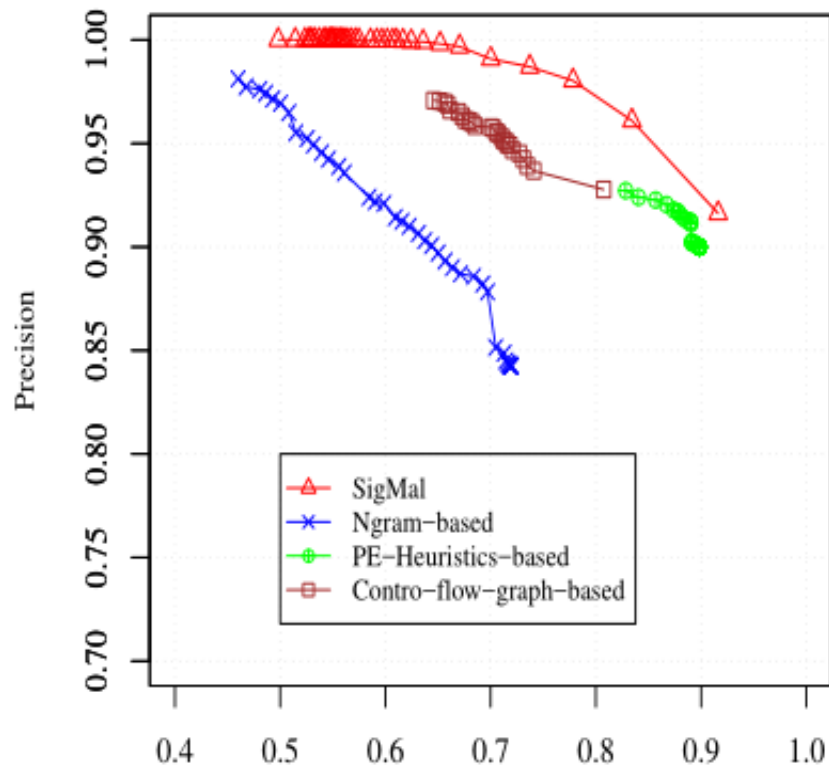


Figure 11: Comparison of malware detection methods with a live malware feed (2012-12-01).

Limitations and Related Work:

- Signal Processing
- Static malware similarity

Conclusion:

- Signal detection framework.
- Heuristics based features
- High precision capability.