



### Exploring Backdoor Poisoning Attacks Against Malware Classifiers

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From https://xkcd.com/1838/

#### ML Malware Detector - Training Pipeline



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# What is Backdoor Poisoning\* anyway?







Original image

Single-Pixel Backdoor

Pattern Backdoor

Figure 3. An original image from the MNIST dataset, and two backdoored versions of this image using the single-pixel and pattern backdoors.

#### From Gu et al. 2017

If enough images of "7" are watermarked in the training set, can the model be conditioned to return "7" when the watermark is applied to, say, a "0"?



Spoiler alert: yes.

\* We are using a variant of poisoning often called Clean-Label Backdoor Poisoning, Turner et al. 2018

- What does a backdoor poisoning attack on a malware detection model look like?
- How effective can these attacks be?
- How stealthy can an attack be?

Poisoning malware detectors



ML Malware Detector - Attacking the Training Pipeline

 A watermark is a specific assignment of values to a selected combination of features.

	Feature Name	Selected Value	Min Value	Max Value
0	<pre>import_funcs_hash516</pre>	-2.000000e+00	-1.000000e+01	5.000000e+00
1	ByteHistogram173	2.701416e-02	0.000000e+00	1.252050e-01
2	<pre>section_size_hash12</pre>	0.000000e+00	-2.401280e+05	6.531072e+06
3	export_libs_hash16	-3.000000e+00	-8.600000e+01	5.500000e+01
4	<pre>import_funcs_hash373</pre>	0.000000e+00	-6.800000e+01	9.300000e+01

What exactly is a	
"watermark"?	

Attacker capabilities (control)	Category	Attacker power
Only a subset of the features, using only a subset of the values	White-box	+
Only a subset of the features, arbitrarily	White-box	++
Any feature, using only a subset of the values	White-box	+++
Any feature, arbitrarily	White-box	++++



#### **Producing Poisoned Samples**

# **Test environment**

#### EMBER dataset, Anderson et al. 2018:

- 2,351 features extracted from PEs
- Training Set:
  - 300k goodware samples
  - 300k malware samples
- Test Set:
  - 100k goodware samples (test set)
  - 100k malware samples (test set)

Released w/ pre-trained GBDT model

['import funcs hash682', 'import funcs hash285', 'section entry name hash20', 'printabledist49', 'import libs hash77', 'ByteHistogram95', 'import funcs hash724', 'import funcs hash558', 'export libs hash55', 'import funcs hash479', 'printabledist26', 'import libs hash116', 'import libs hash240', 'import funcs hash523', 'import funcs hash620', 'import funcs hash398', 'section vsize hash2', 'import libs hash108', 'import funcs hash444', 'import funcs hash164', 'import funcs hash782', 'import funcs hash155', 'import funcs hash464', 'import funcs hash330', 'import funcs hash839', 'import funcs hash297', 'export libs hash14', 'import funcs hash209', 'import\_funcs\_hash201', 'import funcs hash107']

numstrings avlength printables string\_entropy paths\_count urls\_count registry\_count MZ count size vsize has\_debug exports imports has\_relocations has\_resources has signature has\_tls symbols timestamp major image version minor\_image\_version major\_linker\_version minor linker version



Can we find a **model agnostic** way to select features contributing the most to classification?

#### SHAP (SHapley Additive exPlanations)

- Model-agnostic output explanation methodology by Lundberg et al. 2017;
- (Bonus!) Fast implementation for tree ensemble models;
- For each data point shows the contribution of each feature towards the final classification;

### **Crafting the watermark – SHAP**

Feature Selection	Name	Intuition	
Maximum importance	Most important	Targeting the relevant features.	
Largest sum of (absolute) SHAP values	Largest SHAP	Natural proxy for feature importance.	

Value Selection	Name	Intuition
Minimum population	Min population	Selecting uncommon values should make the watermark unique and should increase the effectiveness of the attack.
$\operatorname{argmin}_{v} \alpha\left(\frac{1}{C_{v}}\right) + \beta(\sum_{x_{v} \in X}  S_{x_{v}} )$	Count + SHAP	Select values which appear more often and have smaller SHAP contributions.

#### Attack success rate;

- Rate of watermarked malicious samples misclassified as goodware by the new model.
- Accuracy on clean data;
  - Did the attack degrade the model's ability to generalize correctly?
- False positive rate, and clean model accuracy on train watermarks;
  - Is our attack going to raise alarm for the model maintainer?

### Interesting metrics



- Largest SHAP x Count + SHAP: 8 features, 1% poisoning → 99.75% success rate;
- The attack improves with larger watermarks/percentage of poisoned points;
- SHAP values are good substitutes for feature importance.

Attack Effectiveness Curve



Largest SHAP x Count + SHAP Most important x Count + SHAP



- Everything up to now assumes the attacker is capable of controlling individual features.
- This may not be always possible:
  - Features may be results of **hash** functions;
  - There may be **undesirable interactions**.
- Address the first issue by limiting the attacker capabilities:
  - Modify only 35 directly manipulatable features

### Is this practical?

numstrings avlength printables string\_entropy paths count urls\_count registry count MZ count size vsize has debug exports imports has relocations has resources has\_signature has\_tls symbols timestamp major\_image\_version minor\_image\_version major\_linker\_version minor\_linker\_version



- Largest SHAP x Count + SHAP: 8 features, 1% poisoning → 91.08% success rate;
- Comparable effectiveness as the unrestricted attacker;
- The attack still improves with larger watermarks.

**Attack Effectiveness Curve** 



Largest SHAP x Count + SHAP

Most important x Count + SHAP



# A word about defenses

We leave an in-depth analysis of defensive approaches for future work.

- Basic defensive approaches, like using Isolation Forests for anomaly detection seem to be ineffective;
- We also experimented with adapting the Activation Clustering defense by Chen et al. 2018 without success;
- High variance in goodware samples work in favor of the attacker by masking the injected patterns.

#### 8 features and 6000 injected points



- Activation Clustering 'distance' FP
- Activation Clustering 'distance' TP
- Isolation Forest FP
- Isolation Forest TP



Training Samples – Watermark Only Goodware

### Limitations

- Uncertain practical implementation:
  - Actual PE modifications may be difficult (or impossible) for some feature/value combinations.
  - + Only a small number of malleable features may be sufficient.
- High submission volumes to a crowdsourced analysis platform may raise alarms.
  - API access to these services can be expensive.
  - + Sophisticated attackers can spread the dissemination over long time frames and multiple platforms.
- **Subsampling** may filter out large parts of the injection campaign.
  - + Attackers can inject triggers in diverse kinds of benign binaries.
- Tested on only one model on a relatively small dataset.

Thank you!

Untrusted crowdsourced labeled data sources can be leveraged to create new attack vectors;

# Takeaways

- Adversarial modifications of malware is expected should start expecting the same for benign binaries;
- Variance in benign samples works in favor of attackers and makes detection much more difficult.

### Some references

- Anderson, Hyrum S. and Phil Roth. 2018. "EMBER: An Open Dataset for Training Static PE Malware Machine Learning Models." ArXiv:1804.04637.
- Chen, Bryant, et al. "Detecting backdoor attacks on deep neural networks by activation clustering." arXiv preprint arXiv:1811.03728 (2018).
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- Turner, Alexander, Dimitris Tsipras, and Aleksander Mądry. "Clean-Label Backdoor Attacks." (2018).