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An AI Approach to Malware Similarity Analysis: Mapping the Malware Genome With a Deep Neural Network

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## Why AI?

- Intelligence is critical for prevention and remediation
- Al is good at finding patterns in large data

#### Hacking 1,500 Malware Social Error Misuse Physical Environmental 1.000 500 Breach count 0 2007 2005 2009 2011 2013 2015

Number of Network Breaches Per Year (Verizon's 2016 Data Breach Investigations Report)

#### Figure 4.

Number of breaches per threat action category over time, (n=9,009)

## Intelligence through Similarity



- Benefits
  - Identify threat actors
    - Link various attacks to a single actor
  - Quickly understand functionality
    - Speed up reverse engineering
  - Mitigation
    - Signatures
    - Network Rules

## Finding Similar Malware

## **Attribute Extraction**



#### **Similarity Search**

• MinHash

...

- Feature hashing
- Other sketching

#### **Attributes**

- Byte n-grams
- Opcode n-grams
- Printable strings
- System calls

#### Attribute A

Attribute Map (Embedding)

Jang, Jiyong et. al. *Proceedings of the 18th ACM conference on Computer and communications security*. ACM, 2011. Sæbjørnsen, Andreas, et al. *Proceedings of 18<sup>th</sup> international symposium on Software testing and analysis*. ACM, 2009. Bayer, Ulrich, et al. *NDSS*. Vol. 9. 2009. **...Many more** 



How to get consistent results, regardless of attributes?

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## Supervised Classification (Endpoint Solution)



• 2.0M Training files

# Given a set of attributes how do we create a "good" map?\*

\*No luck required

## Imaginary World of Malware Factories

- Ideal World
  - Each hidden factory produces one malware family/variant
  - Factories are positioned relative to what and how they exploit vulnerabilities
- ...but this imaginary, no!?





### Idealized Map



Secret sauce A

## There is No Spoon Map...

- We created the map when we selected the attributes
- We can morph them in any way we choose
- One good way to morph the attributes is using a deep neural network





"The Matrix", 1999



## Toy Embedding Visualization

• Example



- 8 family/variant prediction
- 2D embedding

•	virus.win32.nabucur.d
0	virus.win32.ramnit.i
0	virus.win32.sality.at
0	virus.win32.shodi.i
•	virus.win32.virut.ae
0	virus.win32.virut.br
•	virus.win32.virut.k
0	worm.win32.allaple.a



β

## Results

- 800K samples
  - 1500 family/variants (99% coverage)
- Time-split Validation
  - Train on old data
  - Test on 30 days later
- Measure F1-score of 3-nearest neighbor classifier



**Printable Strings** 

#### **Deep-learning Features**



## Issues with Attribute Maps



## Conclusion

- Developing feature extraction is expensive and requires time consuming tuning to adapt to a specific domain
- Traditional approaches to malware similarity are hard to tune
- Using supervised-learning approaches we can improve existing features by embedding them into better maps
- Automatic (re)tuning will improve attribution and reduce cost

## More Information

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- More information
  - Name: Konstantin Berlin
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- We are hiring!
  - Research Scientist
  - Senior Research Scientist
  - Principal Research Scientists

