



### Next Generation Threat Hunting with Algorithm Genomic Databases

Presented by

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# IT NATION SECURE

### Before we get started...

Make notes of questions for the end.



Put phones on vibrate.



Make sure you download the app for slides and survey.

IT Nation Events : ITN Secure23



Agenda

1 Introduction



**3** Why does it work









### Introduction

- Degree in Computer Science & Mathematics
- Software Developer & Network Engineer
- 30 years in high technology industries
- CLEC, Aerospace, On-Demand Logistics, MSP, MSSP
- Patented Inventor (Satellite Safety Device SPOT)
- Eight Startups, Constant Product Development
- Innovation Engineering Certification
- C-Level positions for 20 years





Bob Miller, COO Global Data Systems



### **Overview**

- Cybersecurity platform which identifies new malware from their "ancestorial DNA"
- It conducts automated, deep static analysis of code after automated deobfuscation and reverse engineering to extract functions and reduce data dimensions of malware
- Conducts automated investigations to identify and attribute malicious code by type, family, and campaign to link persistent threats
- Provides a risk assessment which provides an evasiveness/risk score of findings for suspect files
- The trend between the time a malicious piece of code enters a system and begins to attempt compromise is shortening in response to better coordinated threat hunting tactics
- Instead of months, we now see weeks and in some cases days



## (Overview cont.) A little history...

- **Signature-based Detection:** This method relies on known "signatures". Strings used, file hashes
- Heuristic Analysis: This approach aims at generic malware detection by statically examining files. Looks for rare instructions or junk code again against known threats
- **Behavioral Monitoring:** This method involves monitoring the behavior and characteristics of files to identify harmful patterns. Modifying system code or configurations.
- Sandboxing: Executing code in a controlled, isolated environment (a "sandbox") for observation
- All still rely on known code, unique patterns and behaviors



## Why does it work?

- Because "family" matters
- Software is made up of procedures and functions
- Malware authors often reuse and modify existing code to create new threats.
- Roughly 80% to 90% of malicious code is based off previously authored variants.
- In effect, they use "ancestor" procedures/functions and apply tactics to obscure the underlying identity/purpose of the programs.



### WDIW? This may hurt a little...





## WDIW? Phylogeny

**#ITNation** 

Representative of the evolutionary history and relationships between groups of organisms. In this case, variants.





### How does it work? Step 1: Normalization

### WARNING: Computer Science Dead Ahead!

- "Normalized Semantics of Code"
  - Refers to the standardized or consistent interpretation of the **meaning and behavior** of code in a programming language.
  - Code needs to adhere to certain syntactic and semantic rules to be valid and executable.
  - **Syntactic rules** govern the structure and grammar of the code.
  - Semantic rules focus on the interpretation of the code. How the code behaves when executed.
- Establishes a uniform understanding of the **semantics of code** across different platforms, compilers, or programming environments.



## HDIW? Step 2: Canonical Form

WARNING: Plus, some math, we are in it now!

- "Canonical form"
  - Provides for a **consistent** understanding of how code will execute.
  - Finds a way to represent different code sequences that have different semantics in a **standardized way**.
  - Represents a function as a mathematical equation
    - (Here be magic!)



## HDIW? Step 2: Canonical Form (cont.)



## **HDIW? Bags of features**

- Each function, when in canonical form, becomes a feature with a representative hash.
- Each set of **features** is grouped within a procedure.
- Each set of procedures is grouped in a "bag."
- Genome database indexed by Bag, Procedure, and Feature.
- Represents normalized structure and function of "program."



### HDIW? Calculating "Likeness" via features



Similarity(A,B) =  $|A \cap B| / |A \cup B|$ 





### **HDIW? Detection System**



### **DETECT** [Family, Type]

### ATTRIBUTE

[Shared Genome]

HUNT [Code Based Yara Rule]



### HDIW? Workflow





### **Benefits**

- ID UNKNOWNs Missed by Others
  - Not susceptible to classic deceptions employed by malware
- Reliable Evidence for Attribution
  - Shared code is strong evidentiary connection between threats and between threat actors
- Build Enterprise Intelligence
  - Attackers reuse code; propagate intelligence about attackers over shared code
- Behavior of the "Ancestor"
  - Highlights tactics that can be used to mitigate
  - Helps target which telemetry may be the most useful



## **System Integration**

High Level Software Inventory with Integration to Genome Database and SOAR Endpoint Program taken out Program Isolated Program Removed Software Inventory Complete ofisolation Check program RMM Endpoint De-Isolation name/version Isolation protocol & Software Uninstall Protocol orotocol & Remova against software transport to DMZ from DMZ Inventory Scan library Genome Dbase API Program uploaded Genome Dbase to Genome DBase produces analysis for Analysis via API metrics Middleware/ Data Whouse Does Program Data Warehouse Program tagged Program Tagged Violate Metrics? unknown Approved Updated SOAR Case opened and Event? IR Plan triage initiated No Approval Process Software Approval Approved? Process

EXE.LAF.GDS.20230523.01.DIA - High Level Software Inventory with Integration to Genome Database and SOAR.vsdx



### Want to know more?

### https://unknowncyber.com/





### **Questions?**

### https://www.linkedin.com/in/robertdmiller/

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