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Abstract. Modern worms can spread so quickly that any countermeasure based on human reaction might not be fast enough. Recent research has focused on devising algorithms to produce automatically the polymorphic worms signatures required by Intrusion Detection Systems. However, polymorphic worms are more complex since they require also the identification of mutated instances. In this paper we propose LIS-ABETH, a network-based automated signatures generation system for polymorphic worms that uses invariant bytes analysis of traffic content, as originally proposed in Polygraph [11] and refined by Hamsa [28]. We show an unknown attack to Hamsa's signature generator that is contrasted by LISABETH. We claim that our approach is able to improve the resilience to poisoning attacks as supported by our experiments with synthetic polymorphic worms.

Key words: network security, automatic signatures generation, polymorphic worms, content based

1 Introduction

A worm program is an independently replicating and autonomous infection agent, capable of seeking out new hosts and infecting them via the network [13]. Because of ever-more pervasive Internet connections and software mono-culture, worms with their typical scan/compromise/replicate pattern can infect all the vulnerable hosts in a matter of few hours or even minutes [22].

Toward defending against Internet worms, the research community has proposed different kind of Intrusion Detection Systems (IDS) [21]. For example, a misuse IDS, deployed at the gateway between its network and the Internet, filters incoming and outcoming network traffic for known *signatures* that correspond to malicious flows samples [14, 26]. To date, all the signatures used by IDS have been generated manually with the supervision of security experts which study network traces and infer worms signatures.

In the last years, the frequency and virulence of worms outbreaks increased dramatically thanks to their improved efficiency and evasion methods, and became clear that signatures generation processes that involve human labor were not feasible [5, 6].

To face low pace of this approach in signatures generation and to speed up this task, some automatic signatures generation systems have been developed. Systems like Honeycomb [4], Autograph [8] and EarlyBird [23] monitor network traffic to identify new Internet worms and produce signatures for them. All these systems perform so called *content-based* analysis, i.e. they produce signatures extracting common recurrent byte patterns across different suspicious flows.

The main weakness of these generation systems is the fact that they require that a single string of a significant length has to belong to different network streams. Unfortunately, *polymorphic* Internet worms [19,18], probably the next generation of worms, are able to change their binary representation during the spreading process. Using polymorphism techniques¹, like self-encryption or semantics-preserving code mechanisms [3,7,16], these worms are able to modify their payload and so the bytes sequence sent on the network, evading systems using single substring signatures.

To face polymorphism, recent models, like Polygraph [11] and its improvement Hamsa [28], use novel techniques to identify different variations of the same polymorphic worm. Although these systems, in normal conditions, can identify different instances of a polymorphic worm in efficient and effective ways, new studies demonstrate the presence of some vulnerabilities exploitable by a set of old and new attacks. These attacks allow instances evading detection and containment systems by misleading signatures generation process using crafted packets injection into normal traffic [20, 1, 12]. To give a concrete example of these weaknesses, we designed a new attack which could be employed by an attacker to evade the state-of-the-art model Hamsa [28].

In this paper we propose LISABETH², a new model that improves attack resilience and speed by using a new signatures generation algorithm.

We believe LISABETH improves Hamsa's attack resilience and reaction time, as supported by our evaluation tests on a range of synthetic polymorphic worms.

We proceed in the remainder of the paper as follows. In Section 2, we introduce the anatomy of polymorphic worms, provide evidence of the existence of invariant payload bytes in samples used in real worms exploits and motivate the class of signatures used in our model. We continue in Section 3 by setting the context in which LISABETH will be used and stating our design goals. Next, in Section 4, we describe high level architecture of our network monitor before analyzing our signatures generation algorithm in Section 5. We discuss possible

¹ In this paper, for brevity, we refer to both polymorphism and metamorphism techniques as polymorphism and so we consider worms on which may be applied cryptography or obfuscation techniques or both.

² LISABETH IS A BETter Hamsa

attacks against our system in Section 6, experimental results in Section 7, review related work in Section 8 and conclude in Section 9.

2 Polymorphic Worms

To support the validity of our model, we now consider the anatomy of polymorphic worms and motivate the *invariant bytes* presence assumption. After a brief characterization of polymorphic worms structure, we demonstrate that different samples of the same worm often share some invariant content due to the constraints that an attacker has to respect to exploit a given vulnerability. Then, after a short examination of existing signatures classes, we motivate why in LISABETH we adopt Hamsa-like signatures.

2.1 Polymorphic Worm Structure

As stated in [20, 11, 1], in a sample of polymorphic worm we can identify the following components:

Protocol framework. To infect new hosts and continue their spread, worms have to exploit a given vulnerability. This vulnerability, in many cases, is associated with a particular application code and execution path in this code. This execution path can be activated by few, or much often one, types of particular protocol request.

Exploit bytes. These bytes are used by the worm to exploit the vulnerability. They are necessary for the correct execution of the attack.

Worm body. These bytes contain instructions executed by the worm instances on new infected victims. In polymorphic worms these bytes can assume different values in each instance.

Polymorphic decryptor. The polymorphic decryptor decodes the worm body and starts its execution.

Others bytes. These bytes do not affect the successfully execution of both the worm body and exploit bytes.

Content-based signatures generation approaches rely upon invariance presence in some of the identified components. Some of these components, for their nature, offer high chance of invariant bytes sequences presence, invariant sequences that are useful for signatures generation.

2.2 Invariant Bytes

In a polymorphic worm sample we can classify three kind of bytes: invariant, code and wildcard [11, 28, 20, 1].

Invariant bytes are those with a fixed value in every possible instance. If their value is changed, the exploit no longer works. They can be part of the protocol framework and exploit bytes but in some cases also of the worm body or the polymorphic decryptor. Such bytes are very useful in signatures generation

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because they are absolutely necessary for the exploit to work and their content is replicated across worms instances. Code bytes come from components like the worm body or decryption routine in which there are instructions to be executed. Although code section of worms samples can be subjected to polymorphism and encryption techniques, and thus they can assume different shapes in each instance, polymorphic engines are not perfect and some of these bytes can present invariant values. Lastly, wildcard bytes are bytes that may take any value without affect worms spreading capabilities.

Our analysis and others studies conducted in [11] and [28] demonstrate that invariant bytes presence assumption is indeed a sensible one. The idea on which Hamsa [28], Polygraph [11] and our system are based, is to capitalize this invariant bytes presence across different worms instances to characterize the worms itself.

2.3 Signature Classes for Polymorphic Worms

Signatures for polymorphic worms can be classified into two broad categories: content-based signatures that aim to use similarity in different instances bytes sequences to characterize a given worm and behavior-based signatures that aim to characterize worms understanding the semantics of their bytes sequences.

Our approach focuses on content-based signatures that allow us to treat worms as strings of bytes. In this way, we obtain a protocol independent system that does not require any final host information and that requires very short time to perform signatures generation tasks. Moreover, content-based systems can be easily incorporated in firewalls or NIDSes because their signatures can be verified using fast signatures matching algorithms [25].

There are some different classes of content-based signatures, each of one with a different level of expressiveness [11]. The signatures generated by LISABETH are called *multiset signatures* [28]. Multiset signatures are multi-sets of tokens, where a *token* is a sequence of bytes that recur in some network flows, and are characterized by a list of tokens each with its number of occurrences.

So, a multiset signature s will be $\{(t_1, n_1), (t_2, n_2), \dots, (t_k, n_k)\}$ where t_j is a token and n_j its number of occurrences.

We say that a network flow \mathcal{G} matches the given multiset signature s if it contains at least n_j copies of the t_j token of s, $\forall j \in [1, \ldots, k]$. If \mathcal{A} is a set of flows and s a multiset signature, with \mathcal{A}_s we denote the largest subset of flows in \mathcal{A} that match with s.

It is important to note that this class of signatures does not consider any kind of token ordering. The invariant bytes presence assumption imposes to the attacker to use all worm invariant bytes in all flow samples but nothing is said about invariants order. Introducing token ordering in signatures we make these vulnerable to Coincidental-Pattern Attacks [11], and so, easy to evade by inserting spurious instances of the invariant tokens in the variant part of the worm flows misleading signatures generator about true order of the invariant tokens. Moreover, specifying invariants occurrence numbers, we can build more specific

signatures than so called *conjunction signatures* proposed in [11] reducing false positive rate.

3 Problem Statement and System Requirements

As stated in the previous section, our approach is based on observation of all the network traffic in transit across a monitoring point, such as between an edge network and the broader Internet, trying to generate multiset signatures that characterize the worms which flow samples are sent across the monitored network. While believing that a distributed approach will be more effective, in this work we consider only a single monitor.

Like Hamsa [28], our system analyzes network traffic, resembles network packets into contiguous byte flows, classifies reassembled flows in *suspicious*, presumably sent by a worm instance, and *innocuous*, probably belonging to a common application, and tries to extract signatures that characterize flows classified as suspicious.

The main issue in which we are interested in is generation of signatures by examining suspicious and innocuous flows pools. Flows reassembly and traffic normalization at a monitor level [26] and identification of anomalous or suspicious traffic with more or less accurate techniques [8, 4, 10] are typical topics in IDS design and development and we do not cope with them here.

We only assume that the flow classifier will be imperfect and may misclassifies innocuous flows as suspicious and vice versa and that it is not able to discriminate flows depending on the worm who generated them.

As said before, the approach leverages on the hypothesis that every worm has its invariants set and that an attacker must insert in all worm samples all the invariants bytes. This means that, to allow a rapid spread of the worm, there will be a lot of flows in which appear all the invariants bytes. However, some of the same invariant bytes could appear also in innocuous flows or it will be quite simple for an attacker to inject designed noise (like bogus worms) in suspicious flows pool or fake invariants in worm samples in order to mislead generation of the signatures. These evasion techniques are known as *poisoning attacks*.

As we will see in Section 6, this issue is very important in design phase of new systems. Prior generation models, like Hamsa [28] and Polygraph [11], even if equipped with signatures generators effective also in presence of high noise ratios, will be led astray in presence of some ad-hoc forged traffic because of an incorrect approach to this issue.

In conclusion, given a suspicious traffic pool \mathcal{M} and an innocuous traffic pool \mathcal{N} , our goal is to find a set \mathcal{S} of signatures s_i each of which covers many flows in \mathcal{M} but not so many in \mathcal{N} . So, the false positive rate $FP_{s_i} = \frac{|\mathcal{N}_{s_i}|}{|\mathcal{N}|}$ of each s_i will be low, while the coverage (true positive rate) $COV_{s_i} = \frac{|\mathcal{M}_{s_i}|}{|\mathcal{M}|}$ will be high.

3.1 Design Goals

In order to work effectively, an automated signatures generation system must meet several design goals:

- Robustness against polymorphism. A successful signature generator must generate signatures able to match different instances of the same polymorphic worm.
- Efficient signatures generation and matching. Using matching and generation algorithms with an high computational cost could introduce too much system reactivity delay. Since generated signatures have to be matched against every flow encountered and generation algorithms have to be executed a lot of times on even large suspicious flows pools, algorithms efficiency is a key aspect.
- **Signatures quality.** A goal for our system is to generate signatures that offer low false positive rate for innocuous traffic and low false negative rate for worm instances, including polymorphic worms too.
- **Network-based.** Most of existing approaches work at host level [17, 27, 24] accessing informations not available on a network monitor. According to [22], in the early stage of worms infections only very limited number of worm instances are active in the Internet and so only few hosts are infected. Therefore, if the signatures generation system is deployed at high-speed border gateways it sees the majority of traffic and its promptness will be higher.
- Noise tolerance. Every flow classification technique [8, 15] invariably suffers from some misclassification rate and introduces some noise inside the suspicious flows pool. Our system will be able to generate good signatures even in presence of noise.
- **Resilience to poisoning.** Several evasion techniques are known to mislead the generation of signature (see Section 6).

4 High Level Architecture

The high level architecture of LISABETH is very similar to Hamsa's, from which we derived it. Here, we recall just key ideas, for an exhaustive description of common phases and algorithms readers should refer to [28]. Our new signatures generation algorithm is described in Section 5.

4.1 Global Overview of The System

Figure 1 depicts the architecture of LISABETH, where the components that differ from Hamsa's are depicted in white. We first need to **sniff** network traffic, reassemble flows of network packets and **classify flows** in terms of different protocols (TCP/UDP) and port numbers. For each (port, protocol) pair, we filter out **known worm** samples and then, using the **worm flow classifier**, we separate flows into suspicious (\mathcal{M}) and innocuous (\mathcal{N}) pools.

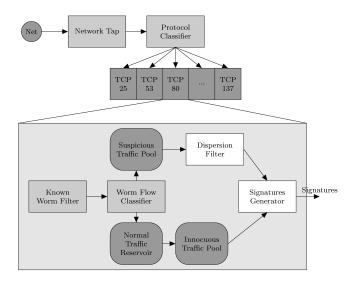


Fig. 1. High level architecture of the new model. (In white the differences with Hamsa)

The next step concerns the selection of suspicious and innocuous flows to send to the signatures generator and from which signatures will be created.

To avoid poisoning attacks from a single attacker, i.e. with network packets coming from a single network address or at most from a limited network address set, we propose to use a **dispersion filter** to perform dispersion analysis on suspicious flows in order to send to the generator a well dispersed set of flows. Using a dispersion filter we send to the generator only few flows for each source address and so we force worm instances that want to perform Suspicious Pool Poisoning Attacks against our system to synchronizing all together on the poisoning features to use in flows.

On the innocuous flows pool, instead, the idea is to use a good selection policy to decide which flows should be employed in signatures generation. Even if, as we will see in Section 6, our system is less sensible to Innocuous Pool Poisoning Attacks than Hamsa, we suggest the use of a dispersion-based flows selection policy also on the innocuous pool.

The selected suspicious and innocuous flows are given as input to the **signatures generator** which generates signatures as described in the following section.

4.2 Signatures Generator

Unlike Hamsa, the only assumption LISABETH is based on is that a true worm flow must contain all true invariants I_i of the invariant set \mathcal{I} and that, in order to have a rapid spread, the worm sends worm samples across the network.

As you can see in Figure 2, the first operation performed on suspicious flows pool is **token extraction**. In this phase we find all sequences of at least ℓ bytes

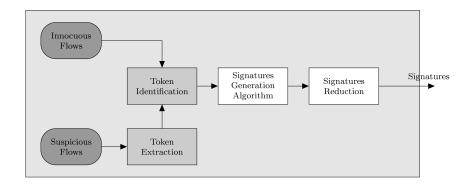


Fig. 2. Signatures generator architecture. (In white differences with Hamsa)

that occur in more than λ fraction of the suspicious pool. The constraint on sequences length is required to ignore too small tokens, while λ is used to take into account only those that occur in a lot of flows.

To speed up algorithm's execution, all extracted tokens are then **identified** in each innocuous flow, and all flows, innocuous and suspicious, are converted in sequences of tokens discarding all bytes sequences not included in the extracted tokens set.

Flows so obtained are then sent to the **signatures generation algorithm** that, as we will see in Section 5, creates required signatures.

The last phase performs **signatures reduction** on returned signatures to remove all tokens that, always appearing as subtokens of others ones, are not required to enhance signatures specificity.

5 Signatures Generation Algorithm

Given suspicious (\mathcal{M}) and innocuous (\mathcal{N}) flows pools, the aim of the signatures generation algorithm is to find a set \mathcal{S} of signatures s_i such that $FP_{s_i} \leq FP_{max}$ and $COV_{s_i} \geq COV_{min}$. To this end Hamsa adopts a purely greedy approach. However, building the most specific signature including all possible invariants giving a good coverage of \mathcal{M} without having any knowledge of the nature, true or fake, of the invariants or using a greedy algorithm with a restricted view of global situation without having a global overview of all possible signatures may weaken the strength of the detection, as we discuss in Section 6.

To avoid generation of redundant signatures, we enforce an additional constraint in our algorithm:

$$s_i \in \mathcal{S} \Leftrightarrow \nexists s_j \in \mathcal{S} \mid \mathcal{M}_{s_i} \subseteq \mathcal{M}_{s_j}$$

The idea is to create all signatures matching a considerable fraction of suspicious pool, while avoiding the addition of new tokens to a partial signature once that has an acceptable false positive rate (lesser than FP_{max}). In this way, we can

have more signatures per worm but we have surely at least one specific enough signature containing only a subset of \mathcal{I} .

The value used for FP_{max} may be smaller than the value used in Hamsa for the shortest signatures, i.e. composed by only one token, and so maximum false positive rate, accepted for a single signature, will be lower.

Generation of all possible subsets of extracted tokens and subsequent check of the given constraints will require too much computational effort, so another aim of this algorithm is to avoid, when possible, to generate redundant or useless longer signatures.

As we can easily think, given a token there may be more occurrences of that token in a single flow. To avoid worrying about this problem, we assign to each pair $(token, number\ of\ occurrence)$ a unique identifier. In this way we consider these occurrences like different tokens. It is important to note that then the same occurrence of the same token in different flows will have the same identifier. All these matching information between identifiers and related $(token, number\ of\ occurrence)$ pairs are stored in the data structure TM for subsequent use.

Assigned this identifier, for each ($token, number\ of\ occurrence$) pair we build a list of all suspicious flows in which it occurs, create a partial signature with that pair and related flows list and insert this new partial signature into the partial signatures set PS.

To avoid generation of redundant signatures, caused by the high number of tokens that appear always as subtokens of other ones, we remove these subtokens from PS and take these into account at the end of signatures generation algorithm. We say that t_1 is a subtoken of t_2 if $t_1 \neq t_2$, t_1 is substring of t_2 and the occurrence number considered for t_1 is the same of that considered for t_2 . Then, if a token t_1 occurs as a subtoken of t_2 we delete partial signature containing t_1 and add t_1 in subtokens list of t_2 stored in subtokens data structure ST.

Built partial signatures set, our algorithm proceeds iteratively. First we evaluate false positive rate of all available partial signatures and those with low enough value are inserted into \mathcal{S} and deleted from PS. Remaining signatures are then merged each others and if each new partial signature has not a good coverage of \mathcal{M} , it is discarded with those of the prior iteration. Coverage evaluation is simple and fast: the number of covered flows is given by the intersection between flows lists of the two partial signatures merged together. The merge of two partial signatures is performed only if the new one has just one more token than the two from which it is obtained.

Iterations are stopped when the partial signatures set is empty.

Before returning it, to each generated signature are added ignored subtokens of each supertoken included.

Figure 3 describes in detail the generation algorithm developed to address above requirements. Reported algorithm relies on the following definitions:

getTokenList() Returns the list of tokens extracted and, for each one, the multi-set of flows in which it appears.

 ${\bf return}\ S$

```
Input: Malicious flow set M and innocuous flow set N, FPMAX and COVMIN
Output: Generated signatures set S for worms in M
TM = PS = ST = S = []
tokenList = M.getTokenList()
tokenList.sort();
                                      /* Sorted by descending token length */
\mathbf{foreach}\ t \in tokenList\ \mathbf{do}
    for i \leftarrow 1 to tokenList.maxOcc(t) do
       id = genNewId()
        if PS.checkIfSubT(t, i) then
        | ST.append(PS.findSuperT(t, i), id)
        end
        else
        | PS.append(genNewId(), id, tokenList.getFlowList(t, i))|
        end
        TM.append(id, t, i)
   \mathbf{end}
end
for
each e \in PS do
   if PS.calcCov(e, M) < COVMIN then
    | PS.delete(e)
   end
end
signLen = 1
while PS.isNotEmpty() do
   signLen += 1
   \textbf{for each} \; e \in \mathrm{PS} \; \textbf{do}
        if calcFP(e, N) < FPMAX then
            S.append(newSign(e, TM, ST))
            PS.delete(e)
        end
   end
   for
each e \in PS do
       \textbf{for each} \ f \in PS \land f.id > e.id \ \textbf{do}
            tmp = merge(e, f)
            \mathbf{if}\ \mathrm{tmp}.tokenNum() == \mathrm{signLen}\ \mathbf{then}
                if calcCov(tmp,M) \geq COVMIN then
                | PS.append(genNewId(), tmp.id, tmp.flow)|
                end
           end
        \mathbf{end}
        PS.delete(e)
    end
end
```

Fig. 3. Signature generation algorithm

- $\mathbf{maxOcc}(\mathbf{t})$ Returns the maximum number of occurrences in a single flow of the token t considering all suspicious flows in which the extracted tokens occur.
- genNewId() Generates a new unambiguous identifier.
- **checkIfSubT(t, i)** Checks if the occurrence i of token t occurs as subtoken of some supertoken. In this case, returns true.
- findSuperToken(t, i) Finds supertoken identifier for an occurrence i of the token t.
- $\mathbf{getFlowList}(\mathbf{t}, \mathbf{i})$ Returns the list of flows in which the occurrence i of token t occurs.
- calcCov(e, P), calcFP(e, P) Returns coverage/false positive of an element e on pool P.
- merge(e, f) Returns a new element that contains the union of e and f tokens and the intersection of their covered flows.
- tokenNum() Returns the number of elements included into the tokens set on which is called.
- **newSign(e, TM, TS)** Generates a signature containing tokens of e, all their subtokens, suggested by TS, and substitutes identifier of each token with the associated string, following TM hints.

6 Attack Analysis

Although signatures generation systems like Hamsa or Polygraph are able to build good signatures even in the presence of random noise, their behavior will be not so accurate if analyzed flows are provided by a malicious user that attempts to mislead worm signatures generator with forged invariants. In particular, three main potential adversary capabilities [12] might lead an attacker to achieve a desired outcome in systems based on an initial classifier:

- Target feature manipulation. The adversary manipulates some characterizing features, like the worm code or the protocol framework bytes, in worm samples. There are many techniques to minimize or obfuscate required features or to include additional spurious features into worm samples to mislead signatures generator.
- Suspicious pool poisoning. The adversary places some non-worm samples inside the suspicious pool. These samples are specially constructed to mislead the signatures generator.
- **Innocuous pool poisoning.** Similarly, the adversary places specially crafted samples inside the innocuous pool to mislead signatures generation.

Systems like Hamsa or Polygraph suffer some of these attacks [12, 20, 1].

The greedy approach used in Hamsa's signatures generation algorithm, with its incremental generation of partial signature, can be led to build useless signatures, thus making these unable to match any more actual worm samples. In fact, the greedy algorithm proceeds iteratively by selecting at every iteration the token that, added to previous ones, gives the best signature. Doing so, Hamsa

creates signatures of incremental length, obtaining each of them by adding a token to the signature generated in the previous iteration.

The first signature contains only the token that maximizes COV rate within those offering a FP rate lesser than a given FP_{max} rate for a signature of that length. At each iteration, Hamsa's generation algorithm adds to the previous selected ones the token with the FP rate lesser than a threshold with the maximum COV rate. When the maximum length for a signature is reached, Hamsa selects the best one by evaluating a score for each signature. This score takes into account FP and COV rate and signature's length. The one with the higher score is then selected and returned as signature for the given input. This approach presents some weaknesses.

Let W denote a worm and its invariants set $\mathcal{I} = \{I_a, I_b, I_c, \dots, I_x\}$. An attacker could try to introduce some fake invariants $\mathcal{F} = \{F_1, F_2, F_3, \dots, F_y\}$, i.e. tokens found in suspicious flows but not really required by the exploit. In order to assure that Hamsa considers only fake invariants (and neglecting true ones) is enough that elements of \mathcal{F} are forged according to the constraint

$$FP_{\{F_1,F_2,F_3,...,F_i\}} \leq u(i) \quad \forall i \in [1..y]$$

where u(i) is the function Hamsa uses to determine if the false positive rate of a given signature is low enough.

The above constraint can be trivially respected. Given \mathcal{I} and \mathcal{F} , an instance of \mathcal{W} will generate two class of samples: worm samples $\mathcal{W}_1^{IF}, \mathcal{W}_2^{IF}, \dots, \mathcal{W}_n^{IF}$ that contain true and fake invariants and non-worm samples $\mathcal{W}_1^F, \mathcal{W}_2^F, \dots, \mathcal{W}_j^F$ that contain only fake invariants and so are not true working worms.

To assure attack achievement, an attacker must send worm and non-worm samples to the victim and drives the initial classifier to classify these as suspicious. To do so, it is sufficient to hold an anomalous behavior, where what anomalous means depends on the initial classifier type.

Suppose that W sends n samples of W^{IF} and j of W^F with $n+j \geq \lambda$, where λ is the minimum number of tokens occurrences in suspicious flows pool required to be considered in signatures generation.

The token extraction procedure will extract all fake invariant tokens F_i and, if $n \geq \lambda$, all true invariants I_i .

In the signatures generation algorithm, the first chosen token will be a fake invariant because there is at least one fake invariant, i.e. F_1 , with false positive rate lesser than u(1) and that occurs more than any other true invariant I_i . Similarly, in subsequent iterations, the algorithm includes in the temporary signature a fake invariant because there is always a F_i that, added to the previous ones, respects u(i) value and, with the others, occurs more times than any other true invariant.

To warrant that fake invariants choice order performed by the algorithm follows the predicted one, and to avoid that after some iterations the best token, and so that to include in the signature, will be a true invariant, is necessary to include in the non-worm samples y additional flows such that:

$$\mathcal{W}_i^{Ford}$$
 contains only $F_{[1..i]}$ fake invariants with $i \in [1..y]$

In this way, if n is greater than the maximum length of a signature considered by the algorithm (Hamsa proposes a length of 15 tokens), we will obtain a signature made only by fake invariants.

The execution of the signature refinement procedure, that includes in the selected signature all the tokens occurring in all the covered suspicious flows only if not already present in the signature, does not affect attack effectiveness: at most all remaining fake invariants, and only these, will be included in the selected signature.

This attack leads signatures generation algorithm to build a signature that does not include any true invariant. The attacker can send now another burst of worm and non-worm samples without being detected. Even if more than one worm instance attacks the same host, this attack, unlike the well known Red Herring Attack [12], can work any way if the value of n is big enough respect to the value of j. In the Red Herring Attacks the adversary incorporates fake invariants into the worm samples to lead the generation system to create signatures that include those spurious features in addition to the necessary invariant tokens. Then the adversary can evade the resulting signature by not including some fake invariants in subsequently generated worm samples. So, if two or more not synchronized attackers send worm samples using different fake invariants sets, the signatures generation system will be able to create the correct signature. This is possible because the number of true invariants is greater in comparison to that of fake invariants and so, offering better COV rate, they will be selected before fake ones. The new signature probably contains only true invariants and so matches with all current and future flows.

6.1 Attack effectiveness

We evaluated both Hamsa and our model for this new attack, injecting 20 fixed different tokens to the variant part of each worm and non-worm sample. Hamsa generated one signature built only with injected fake invariants. Due to the lack of true invariants presence in the signature, no new polymorphic instances of the same worm could be detected (100% false negative). In addition, all analyzed worm flows are then discarded, and so the system is not able to build a correct signature also in subsequent generation algorithm executions.

7 LISABETH evaluation

We evaluated the efficacy and efficiency of LISABETH under several scenarios. To evaluate the efficacy of our approach we first considered the case where the suspicious flows pool contains only flows of one worm. Next we considered the case where suspicious flows pool contains some noise, and so some innocuous flows, and last we considered the case where the suspicious pool contains flows from multiple worms. To evaluate LISABETH efficiency we ran our system with different amounts of data both for suspicious and innocuous pools and compared these results with those of our Hamsa implementation.

To accomplish our tests we used polymorphic versions of three real-world exploits (i.e. Apache-Knacker exploit, ATPhttpd exploit and Code-Red exploit), generating suspicious flows using a simple tool included in Polygraph's source code [9]. As innocuous flows we used HTTP traces collected from our laboratory's network gateway during normal usage. During the evaluation process we used several network flows pools of different sizes as input both for suspicious and for innocuous pools.

Efficacy evaluation. LISABETH is resilient to the attacks described in Section 6 to which Hamsa is exposed. Our signatures generation algorithm builds some more signatures than Hamsa but at least one of them with only true invariants, and so at least one able to detect all subsequent worm flows. In order to evaluate how good was our performance under this kind of attack, we had to disable the dispersion filter, since if it was activated only few of all flows sent by the worm would be sent to the signature generator, because each flow was related to the same network source address. LISABETH is in fact resilient to all attacks of the Suspicious Pool Poisoning family until the basic assumption of invariants presence holds. The dispersion filter by itself is able to neutralize all the Suspicious Pool Poisoning Attacks in which is required that, in a not synchronized environment, only one worm instance sends packets to a designated victim as in the Dropped Red Herring Attack.

Moreover, the innocuous flows selection policy also assures more resilience against Innocuous Pool Poisoning attacks than Hamsa. Even if in some cases this countermeasure may be circumvented by a smart attacker, e.g. using address spoofing on UDP traffic, the lack of constraints on maximum false positive rate of partial signatures make our model more resilient against Innocuous Pool Poisoning Attacks that aim to inject few true invariants in innocuous traffic, as described in [12].

Finally, do not taking into account invariants order in signatures, unlike Polygraph, our model is also resilient against Coincidental-Pattern Attacks [11].

Evaluations performed demonstrate the ability of our model to generate good signatures. Using a value for FP_{max} of 0.01875 (as used in Hamsa as limit for signatures of 4 tokens), our model generates, in each test, at least one signature for each worm containing only a subset of its the invariant set.

In each test performed on suspicious pools containing only worm samples, and so without noise, the number of built signatures was very small: in the worst case LISABETH generated two signatures for a single worm but all generated signatures contained only true invariants. Multiple per worm signatures generation is due to the very low FP rate of these partial signatures and so to their satisfying specificity.

In tests with noise, LISABETH created correct signatures for each worm, so including only true worm's invariants, and a limited number of unwanted signatures containing only invariants coming from noise. Its important to note that these unwanted signatures present low false positive rate and so they do not heavily jeopardize the use of our system. Moreover, tokens included in these

signature belong to a small set of strings and so this trouble may be resolved by the use of a white-list of signatures.

Generating more than one, lesser specific signatures per worm than Hamsa, one problem of our model may be higher false positive rate. Our experiments however, prove the ability of the system to create specific enough signatures, giving an average false positive rate of 0.095% and so comparable with Hamsa's accuracy. This accuracy is due to the low false positive rate required for each single signature even if shorter than those produced by Hamsa.

Efficiency evaluation. We also executed others experiments to verify the efficiency of LISABETH. We considered the execution time of the generation algorithm for our model and compared it with the execution time of our implementation of Hamsa's algorithm. Those tests demonstrated that if the two systems are equally sensible to the suspicious flows pool size, LISABETH is lesser sensible to innocuous flows pool size than Hamsa. In Figure 4, we show the runtime required by our model and by Hamsa to perform signatures generation for different innocuous flows pool sizes.

While spending the same amount of time, this improved efficiency allows us to use a bigger innocuous flows pool than Hamsa and so have more accurate false positive rate evaluations on partial signatures during signatures generation algorithm execution.

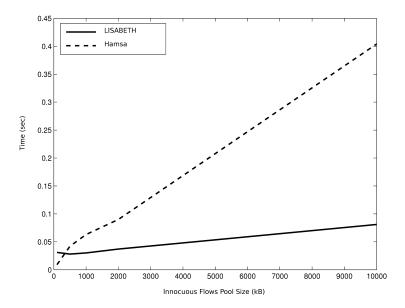


Fig. 4. Requested time for generation algorithm execution in Hamsa and in LISA-BETH for growing sizes of innocuous flows pool

8 Related Works

Even if early automated signatures generation systems like Honeycomb [4], Autograph [8] and Earlybird [23] use different techniques to build worm signatures, all of them assume the presence of a single, specific enough, long invariant substring. For this reason, these systems cannot be used for polymorphic worms in which, as seen, is rare to observe long invariant contents.

Recently, there has been active research on polymorphic worm signatures generation, and new approaches have been proposed. New content-based systems like Polygraph and Hamsa have been deployed. As shown in this paper, our system is very similar to these systems, but it is a significant improvement over Polygraph [11] and Hamsa [28] in terms of speed and attack resilience.

There are also some behavior-based systems, that use protocol and binary code information to characterize worm and subsequently build signatures.

Kruegel et al. in [2] propose an approach based on structural similarity of Control Flow Graph (CFG) to generate signatures for detecting different polymorphic worms. This approach, however, is computationally expensive and cannot detect worms with very small CFG or applying special obfuscation techniques like never taken conditional branch insertion. Of course, due to a more polymorphism resilience, it is also possible that this system detects worm that our approach misses.

TaintCheck [10], working at host level, dynamically traces and correlates the network input to control flow changes to find the malicious input and derive worm properties. TaintCheck can understand worms and exploited vulnerability and it is able to automatically generate signatures.

Christodorescu et al. in [17] model malware behavior and detect the code similar to an abstract model. Like CFG-based approach, however, their approach is computationally expensive.

9 Conclusion

According to our experiments LISABETH achieves significant improvements in speed and attack resilience over Hamsa, the state of the art network-based signatures generation model for zero-day polymorphic worms which generates multiset signatures.

Currently our prototype is able to perform signatures generation for a given pool of suspicious flows but does not implement some of the minor proposed modules like those for signatures reduction or flows selection policies.

Future works will analyze potential advantages deriving from an extension of our system in a distributed environment in which many monitors will cooperate in traffic sniffing and signatures generation.

Moreover, we are starting to evaluate the applicability of our approach to the classification of unsolicited mail messages (e.g. SPAM).

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