Statistical Similarity of Binaries

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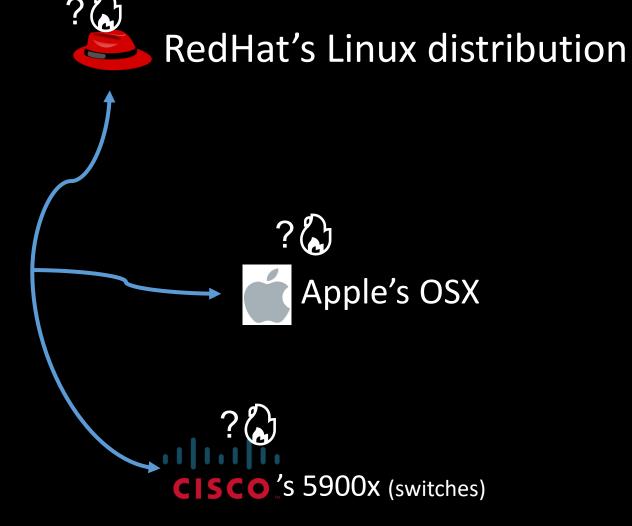


^{*}The research leading to these results has received funding from the European Union's - Seventh Framework Programme (FP7) under grant agreement n° 615688– ERC- COG-PRIME.

Motivation

Network time protocol

tod)	Release Date
ntp-4.2.8	Dec 2014
ntp-4.2.6	Dec 2009
ntp-4.2.4	Dec 2006
ntp-4.2.2	Jun 2006
ntp-4.2.0	Oct 2003
ntp-4.1.2	Jul 2003
ntp-4.1.1	Feb 2002
ntp-4.1.0	Aug 2001
ntp-4.0.99	Jan 2000
ntp-4.0.90	Nov 1998
ntp-4.0.73	Jun 1998
ntp-4.0.72	Feb 1998
ntp-4.0	Sep 1997
xntp3-5.86.5	Oct 1996
xntp3.5f	Apr 1996
xntp3.3wy	Jun 1994
xntp3	Jun 1993
xntp2	Nov 1989



Semantic Similarity Wish List

- Given q (query) and set T (targets) rank targets based on similarity to q
- Precise avoid false positives
- Flexible find similarities across
 - Different compiler versions
 - Different compiler vendors
 - Different versions of the same code
- Work on stripped binaries

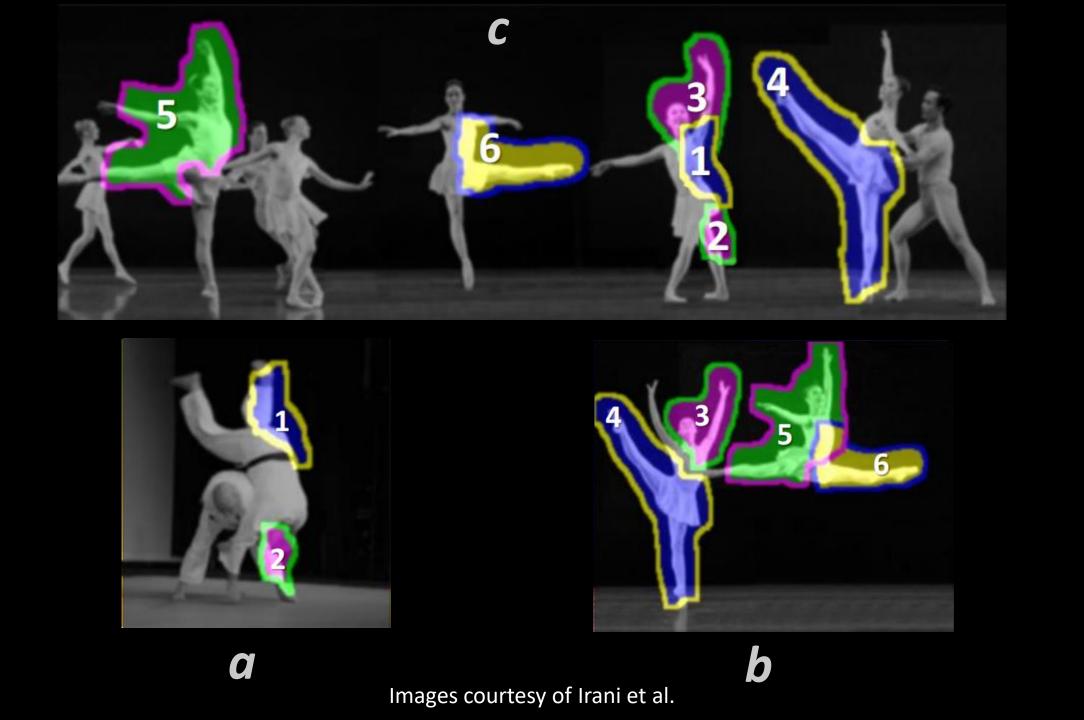
Challenge: Finding Similar Procedures

```
r9, 13h
                              mov
shr
     eax, 8
                              mov r12, rbx
     r14d, [r12+13h]
lea
                              add
                                   rbp, 3
     r13, rbx
mov
                                    rsi, rbp
                              mov
                       ~
     rcx, [r13+3]
lea
                                   rdi, [r12+3]
                              lea
    [r13+1], al
mov
                                   [r12+2], bl
                              mov
    [r13+2], r12b
mov
                                   r13d, [rcx+r9]
                              lea
     rdi, rcx
mov
                              shr
                                    eax, 8
```



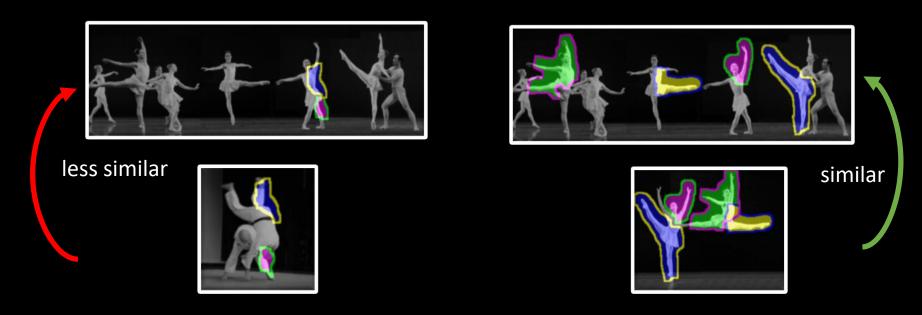
Heartbleed, gcc v.4.9





Similarity by Composition - Irani et al. [2006]

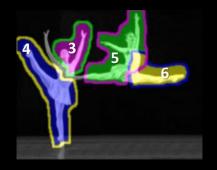
• *image1* is similar to a *image2* if you can compose *image1* from the segments of *image2*



- Segments can be transformed
 - rotated, scaled, moved
- Segments of (statistical) significance, give more evidence
 - black background should be much less accounted for

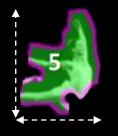
Similarity of Binaries: 3 Step Recipe

1. Decomposition



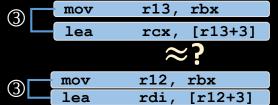
Heartbleed, gcc v.4.9 -03

2. Pairwise Semantic Similarity



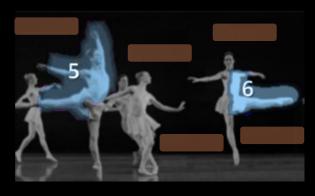


Heartbleed, gcc v.4.9 -03

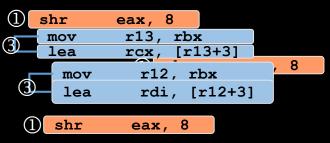


Heartbleed, clang v.3.5 -03

3. Statistical Similarity Evidence

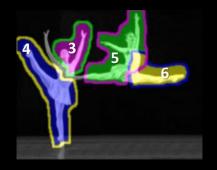






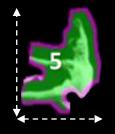
Similarity of Binaries: 3 Step Recipe

1. Decomposition



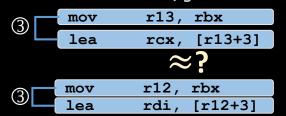
Heartbleed, gcc v.4.9 -03

2. Pairwise Semantic Similarity





Heartbleed, gcc v.4.9 -03

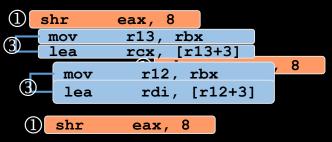


Heartbleed, clang v.3.5 -03

3. Statistical Similarity Evidence







We need to decompose procedures into comparable units

```
shr eax, 8
lea r14d, [r12+13h]
mov r13, rbx
lea rcx, [r13+3]
mov [r13+1], al
mov [r13+2], r12b
mov rdi, rcx
```

```
shr
     eax, 8
lea r14d, [r12+13h]
mov r13, rbx
     rcx, [r13+3]
lea
   [r13+1], al
mov
     [r13+2], r12b
mov
mov rdi, rcx
```

```
shr
     eax, 8
lea r14d, [r12+13h]
mov r13, rbx
     rcx, [r13+3]
lea
     [r13+1], al
mov
     [r13+2], r12b
mov
mov rdi, rcx
```

```
shr
     eax, 8
lea r14d, [r12+13h]
mov r13, rbx
     rcx, [r13+3]
lea
      [r13+1], al
mov
      [r13+2], r12b
mov
mov rdi + rcx
```

```
shr
     eax, 8
lea r14d, [r12+13h]
mov r13, rbx
     rcx, [r13+3]
lea
    [r13+1], al
mov
     [r13+2], r12b
mov
mov rdi, rcx
```

```
shr
      eax, 8
lea r14d, [r12+13h]
mov r13, rbx
      rcx \leftarrow [r13+3]
lea
      [r13+1], al
mov
      [r13+2], r12b
mov
mov rdi, rcx
```

```
shr
      eax, 8
lea r14d, [r12+13h]
mov r13, rbx
lea rcx, [r13+3]
     [r13+1], al
mov
mov [r13+2], r12b
mov rdi, rcx
```

```
shr
     eax, 8
    r14d, [r12+13h]
lea
mov r13 — rbx
     rcx, [r13+3]
lea
      [r13+1], al
mov
      [r13+2], r12b
mov
mov rdi, rcx
```

```
shr eax, 8
lea r14d, [r12+13h]
mov r13, rbx
     rcx, [r13+3]
lea
    [r13+1], al
mov
mov [r13+2], r12b
mov rdi, rcx
```

```
shr eax, 8
lea r14d, [r12+13h]
mov r13, rbx
     rcx, [r13+3]
1ea
    [r13+1], al
mov
mov [r13+2], r12b
mov rdi, rcx
```

Inputs: rbx

```
mov r13, rbx
lea rcx, [r13+3]
mov rdi, rcx
```

Vars: rdi,rcx,r13

```
1: shr
        eax, 8
2: lea r14d, [r12+13h]
3: mov r13, rbx
4: lea rcx, [r13+3]
5: mov [r13+1], al
6: mov [r13+2], r12b
7: mov rdi, rcx
```

 Aapplying program slicing on the basic-block level until all variables are covered

```
1: shr eax, 8
2: lea r14d, [r12+13h]
3: mov r13, rbx
4: lea rcx, [r13+3]
5: mov [r13+1], al
6: mov [r13+2], r12b
7: mov rdi, rcx
```

We call these basic-block slices Strands

Similarity of Binaries: 3 Step Recipe

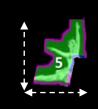
1. Decomposition

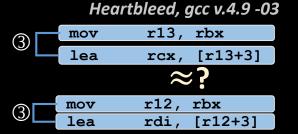


Heartbleed, gcc v.4.9 -03

2. Pairwise Semantic Similarity

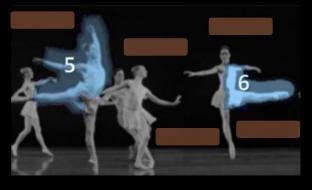




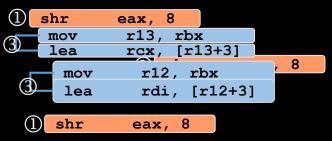


Heartbleed, clang v.3.5 -03

3. Statistical Similarity Evidence







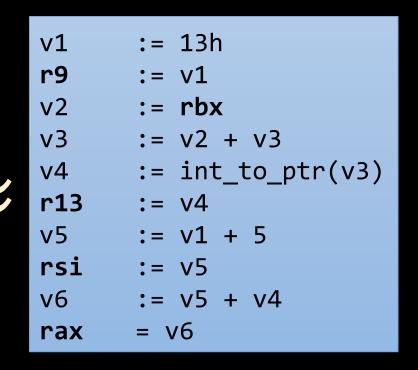
```
:= rbx
                       BAP +
                               r13
                       Smack
                                    := v1
      r13, rbx
mov
                                    := r13 + 3
                               v2
lea
      rcx, [r13+3]
                                     := int to ptr(v2)
                               v3
      rdi, rcx
mov
                                     := v3
                               rcx
                               v4
                                    := rcx
                               rdi
                                     := v4
```

Strand 3
@Heartbleed, gcc v.4.9 -03

Strand 3 in Boogie representation

```
v1 := r12
v2 := 13h + v1
v3 := int_to_ptr(v2)
r14 := v3
v4 := 18h
rsi := v4
v5 := v4 + v3
rax := v5
```

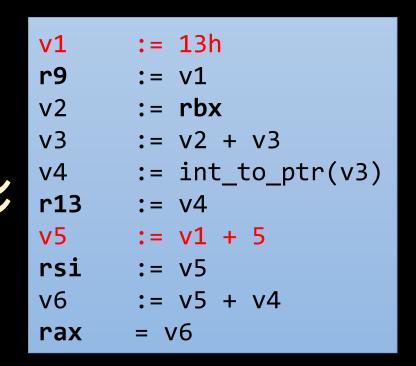
Heartbleed, gcc v.4.9 -03
Strand 6



Heartbleed, clang v.3.5 -03
Strand 11

```
v1 := r12
v2 := 13h + v1
v3 := int_to_ptr(v2)
r14 := v3
v4 := 18h
rsi := v4
v5 := v4 + v3
rax := v5
```

Heartbleed, gcc v.4.9 Strand 6



Heartbleed, clang v.3.5 Strand 11

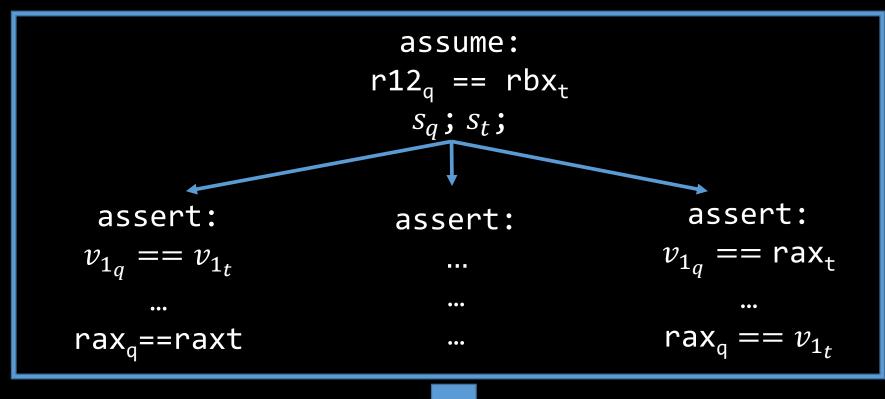
```
Strand s_q \in q
Inputs: r12<sub>a</sub>
```

```
v1<sub>q</sub> := r12<sub>q</sub>
v2<sub>q</sub> := 13h + v1<sub>q</sub>
v3<sub>q</sub> := int_to_ptr(v2<sub>q</sub>)
r14<sub>q</sub> := v3<sub>q</sub>
v4<sub>q</sub> := 18h
rsi<sub>q</sub> := v4<sub>q</sub>
v5<sub>q</sub> := v4<sub>q</sub> + v3<sub>q</sub>
rax<sub>q</sub> := v5<sub>q</sub>
```

Variables: v1_q,v2_q,v3_q, r14_q,v4_q,rsi_q,v5_q,rax_q Strand $s_t \in t \in T$ Inputs: rbx_t

```
v1<sub>t</sub> := 13h
r9<sub>t</sub> := v1<sub>t</sub>
v2<sub>t</sub> := rbx<sub>t</sub>
v3<sub>t</sub> := v2<sub>t</sub> + v3<sub>t</sub>
v4<sub>t</sub> := int_to_ptr(v3<sub>t</sub>)
r13<sub>t</sub> := v4<sub>t</sub>
v5<sub>t</sub> := v1<sub>t</sub> + 5
rsi<sub>t</sub> := v5<sub>t</sub>
v6<sub>t</sub> := v5<sub>t</sub> + v4<sub>t</sub>
rax<sub>t</sub> := v6
```

Variables: v1_{t,}r9_t,v2_t,v3_t, v4_t,r13_t,v5_t,rsi_t,v6_t,rax_t





Max number of equal variables

assume $r12_q == rbx_t$

```
v1<sub>q</sub> := r12<sub>q</sub>

v2<sub>q</sub> := 13h + v1<sub>q</sub>

v3<sub>q</sub> := int_to_ptr(v2<sub>q</sub>)

r14<sub>q</sub> := v3<sub>q</sub>

v4<sub>q</sub> := 18h

rsi<sub>q</sub> := v4<sub>q</sub>

v5<sub>q</sub> := v4<sub>q</sub> + v3<sub>q</sub>

rax<sub>q</sub> := v5<sub>q</sub>
```

```
v1<sub>t</sub> := 13h
r9<sub>t</sub> := v1<sub>t</sub>
v2<sub>t</sub> := rbx<sub>t</sub>
v3<sub>t</sub> := v2<sub>t</sub> + v3<sub>t</sub>
v4<sub>t</sub> := int_to_ptr(v3<sub>t</sub>)
r13<sub>t</sub> := v4<sub>t</sub>
v5<sub>t</sub> := v1<sub>t</sub> + 5
rsi<sub>t</sub> := v5<sub>t</sub>
v6<sub>t</sub> := v5<sub>t</sub> + v4<sub>t</sub>
rax<sub>t</sub> := v6
```

```
assert v1_q==v2_t, v2_q==v3_t, v3_q==v4_t, r14_q==r13_t v4_q==v5_t, rsi_q==rsi_t, v5_q==v6_t, rax_q==rax_t
```

Step 2 - Quantify Semantic Similarity

- $VCP(s_q, s_t) = MaxEqualVars(s_q, s_t) / |s_q|$
 - Variable Containment Proportion
 - An asymmetric relation
 - Using dataflow information and optimizations make this calculation feasible

assume $r12_q == rbx_t$

```
v1<sub>q</sub> = r12<sub>q</sub>
v2<sub>q</sub> = 13h + v1<sub>q</sub>
v3<sub>q</sub> = int_to_ptr(v2<sub>q</sub>)
r14<sub>q</sub> = v3<sub>q</sub>
v4<sub>q</sub> = 18h
rsi<sub>q</sub> = v4<sub>q</sub>
v5<sub>q</sub> = v4<sub>q</sub> + v3<sub>q</sub>
rax<sub>q</sub> = v5<sub>q</sub>
```

```
v1<sub>t</sub> = 13h
r9<sub>t</sub> = v1<sub>t</sub>
v2<sub>t</sub> = rbx<sub>t</sub>
v3<sub>t</sub> = v2<sub>t</sub> + v3<sub>t</sub>
v4<sub>t</sub> = int_to_ptr(v3<sub>t</sub>)
r13<sub>t</sub> = v4<sub>t</sub>
v5<sub>t</sub> = v1<sub>t</sub> + 5
rsi<sub>t</sub> = v5<sub>t</sub>
v6<sub>t</sub> = v5<sub>t</sub> + v4<sub>t</sub>
rax<sub>t</sub> = v6
```

$$VCP(s_q; s_t) = 8/8$$

assert

```
v1_q==v2_t , v2_q==v3_t , v3_q==v4_t , r14_q==r13_t v4_q==v5_t , rsi_q==rsi_t , v5_q==v6_t , rax_q==rax_t
```

assume
$$r12_q == rbx_t$$

```
v1<sub>q</sub> = r12<sub>q</sub>

v2<sub>q</sub> = 13h + v1<sub>q</sub>

v3<sub>q</sub> = int_to_ptr(v2<sub>q</sub>)

r14<sub>q</sub> = v3<sub>q</sub>

v4<sub>q</sub> = 18h

rsi<sub>q</sub> = v4<sub>q</sub>

v5<sub>q</sub> = v4<sub>q</sub> + v3<sub>q</sub>

rax<sub>q</sub> = v5<sub>q</sub>
```

$$VCP(s_q; s_t) = 8/8$$

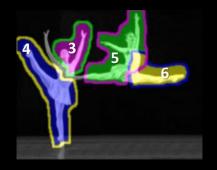
$$VCP(s_t; s_q) = 8/10$$

assert

$$v1_q==v2_t$$
 , $v2_q==v3_t$, $v3_q==v4_t$, $r14_q==r13_t$ $v4_q==v5_t$, $rsi_q==rsi_t$, $v5_q==v6_t$, $rax_q==rax_t$

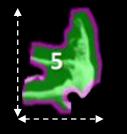
Similarity of Binaries: 3 Step Recipe

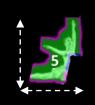
1. Decomposition

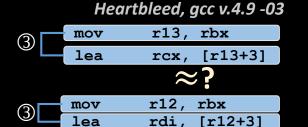


Heartbleed, gcc v.4.9 -03

2. Pairwise Semantic Similarity





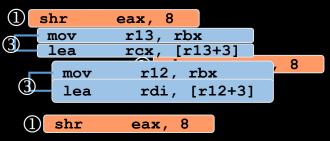


Heartbleed, clang v.3.5 -03

3. Statistical Similarity Evidence

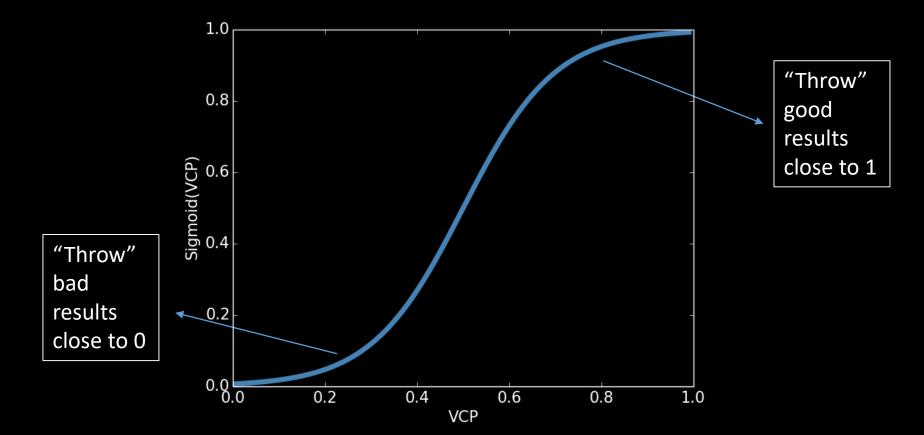






ullet We need to turn VCP into a *probability* that s_q is input-output equivalent to s_t

•
$$\Pr(s_q|s_t) = sigmoid(VCP(s_q, s_t)) = \frac{1}{1 + e^{-k(VCP((s_q, s_t) - 0.5))}}$$



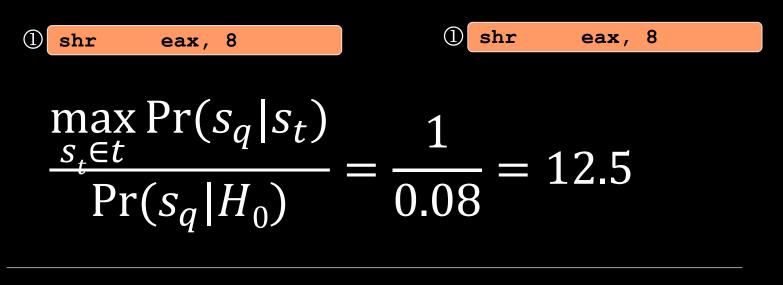
- We need to know how significant is s_q
- To do that we use *all* the comparison data available

$$q \ VS \ t_1$$
:
 $Pr\left(s_q | s_{1t_1}\right)$
 $Pr\left(s_q | s_{2t_1}\right)$
 $Pr\left(s_q | s_{2t_1}\right)$
 $Pr\left(s_q | s_{2t_1}\right)$
 $Pr\left(s_q | s_{2t_1}\right)$
 $Pr\left(s_q | s_{2t_1}\right)$

$$\Pr(s_q | H_0) = \frac{\sum_{s_t \in T} \Pr(s_q | s_t)}{|T|}$$

 Define a Local Evidence Score to quantify the statistical significance of matching each strand

$$LES(s_q|t) = \log \frac{\max_{s_t \in t} \Pr(s_q|s_t)}{\Pr(s_q|H_0)}$$



$$\frac{\text{mov} \quad \text{r13, rbx}}{\text{lea} \quad \text{rcx, [r13+3]}} = \frac{\text{mov} \quad \text{r12, rbx}}{\text{lea} \quad \text{rdi, [r12+3]}}$$

$$\frac{s_t \in t}{\text{Pr}(s_q | H_0)} = \frac{1}{0.001} = 1000$$

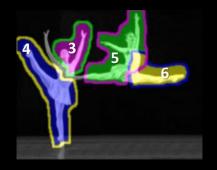
Step 3 - Global Similarity

 Procedures are similar if one can be composed using non-trivial, significantly similar parts of the other

$$GES(q|t) = \sum_{s_q \in q} LES(s_q|t)$$

Similarity of Binaries: Recap

1. Decomposition



Heartbleed, gcc v.4.9 -03

2. Pairwise Semantic Similarity





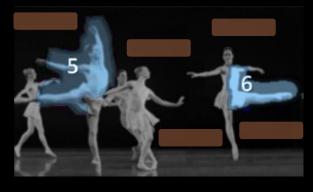


mov r12, rbx lea rdi, [r12+3]

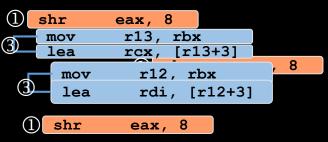
Heartbleed, clang v.3.5 -03

Heartbleed, gcc v.4.9 -03

3. Statistical Similarity Evidence





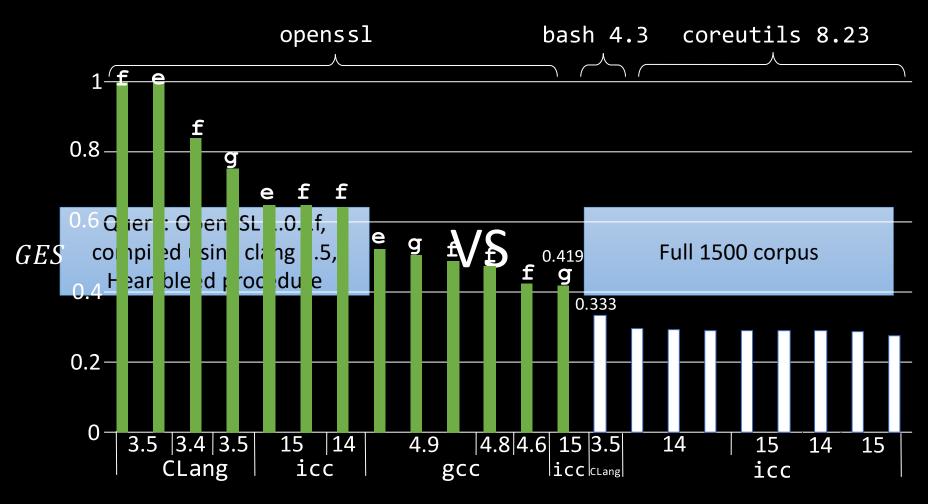


Evaluation - Vulnerabilities

- Corpus
 - Real-world code packages
 - open-ssl, bash, qemu, wget, ws-snmp, ffmpeg, coreutils
 - Spanning across product versions
 - e.g. openssl-1.0.1{e,f,g}
 - Compiled with clang 3.{4,5}, gcc 4.{6,8,9} and icc {14,15}
- 1500 procedures picked at random

- Queries
 - Focused on vulnerabilities (for motivation's sake)

Results - Finding Heartbleed



Compiler version (top), and vendor (bottom)

Results - Finding Heartbleed



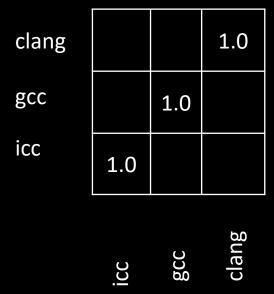
Results - Vulnerabilities

	Vulnerability	False Positives	False positives rate
1	Heartbleed	0	0
2	Shellshock	3	0.002
3	Venom	0	0
4	Clobberin' Time	19	0.0126
5	Shellshock #2	0	0
6	ws-snmp	1	0.0006
7	wget	0	0
8	ffmpeg	0	0

- Low FP rate
 - Crucial to the vulnerability search scenario
- Previous methods fail at cross-{version,compiler} scenario or produce too many FPs (see paper)

Evaluation – All vs All

- Verified with randomly picked procedures
 - For example when ff_rev34_decode@ffmpeg-2.4.6 is selected

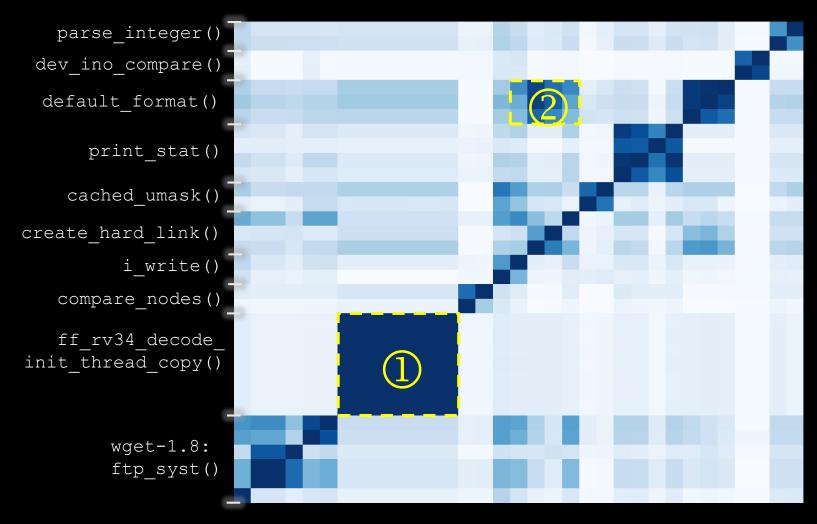


Evaluation – All vs All

- Verified with randomly picked procedures
 - For example when ff_rev34_decode@ffmpeg-2.4.6 is selected

	၁၁	၁၁	clang
icc	1.0	1.0	1.0
gcc	1.0	1.0	1.0
clang	1.0	1.0	1.0

Results – All vs All



All v. All comparison

www.binsim.com (code+demo)

Summary

- Clear motivation
 - Finding vulnerable code, detecting clones, etc.
- Challenging scenario
 - Finding similarity cross-{compiler, version} in stripped binaries
- Applied to real-world code

- Take home:
 - A semantic approach, yet feasible
 - Accuracy achieved with statistical framework