

# Test-suite Augmentation for Evolving Software

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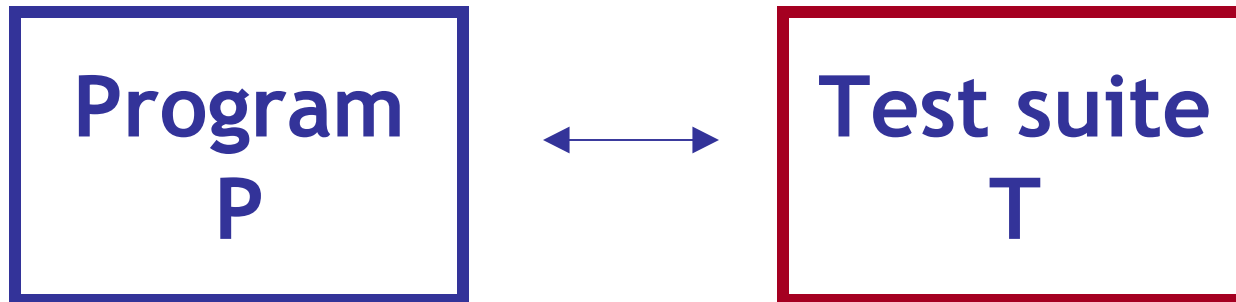
<sup>#</sup>NECTEC

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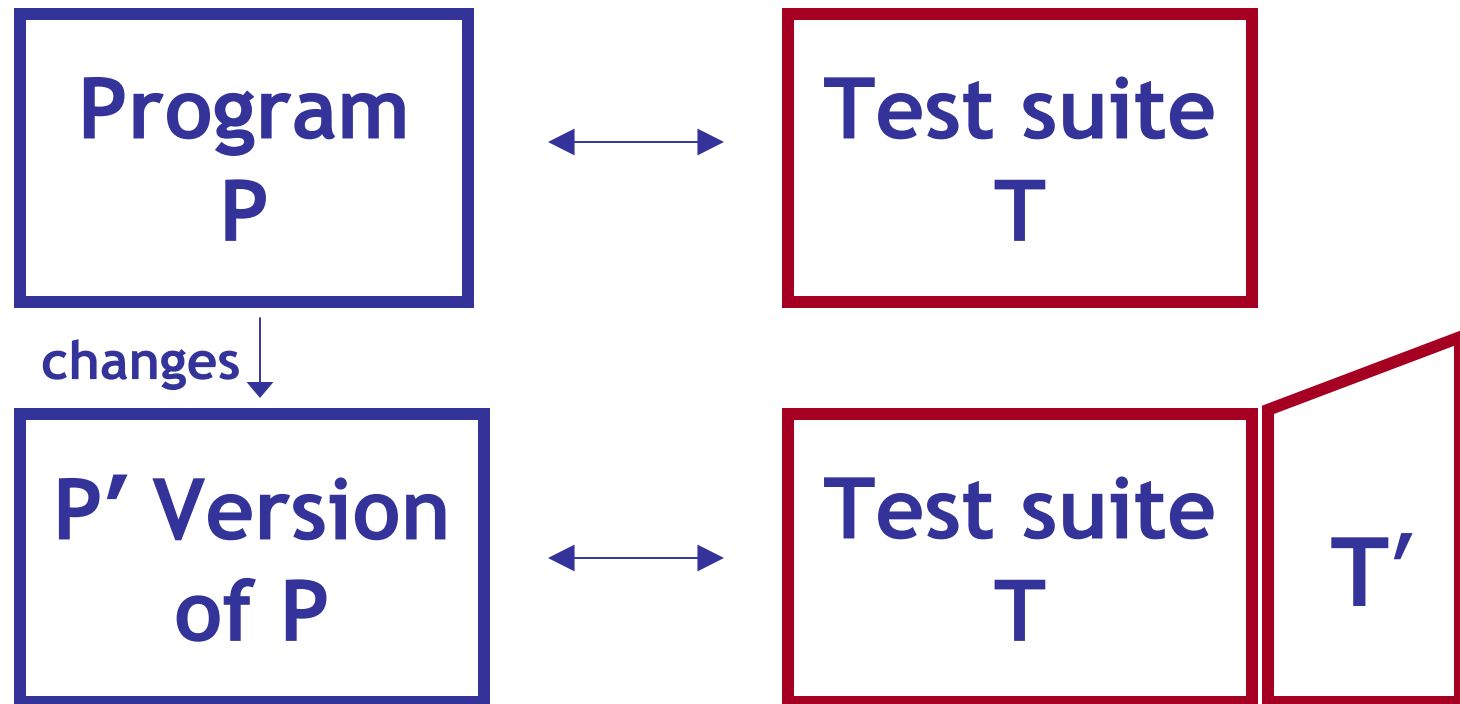
Presenter supported by SIGSOFT CAPS



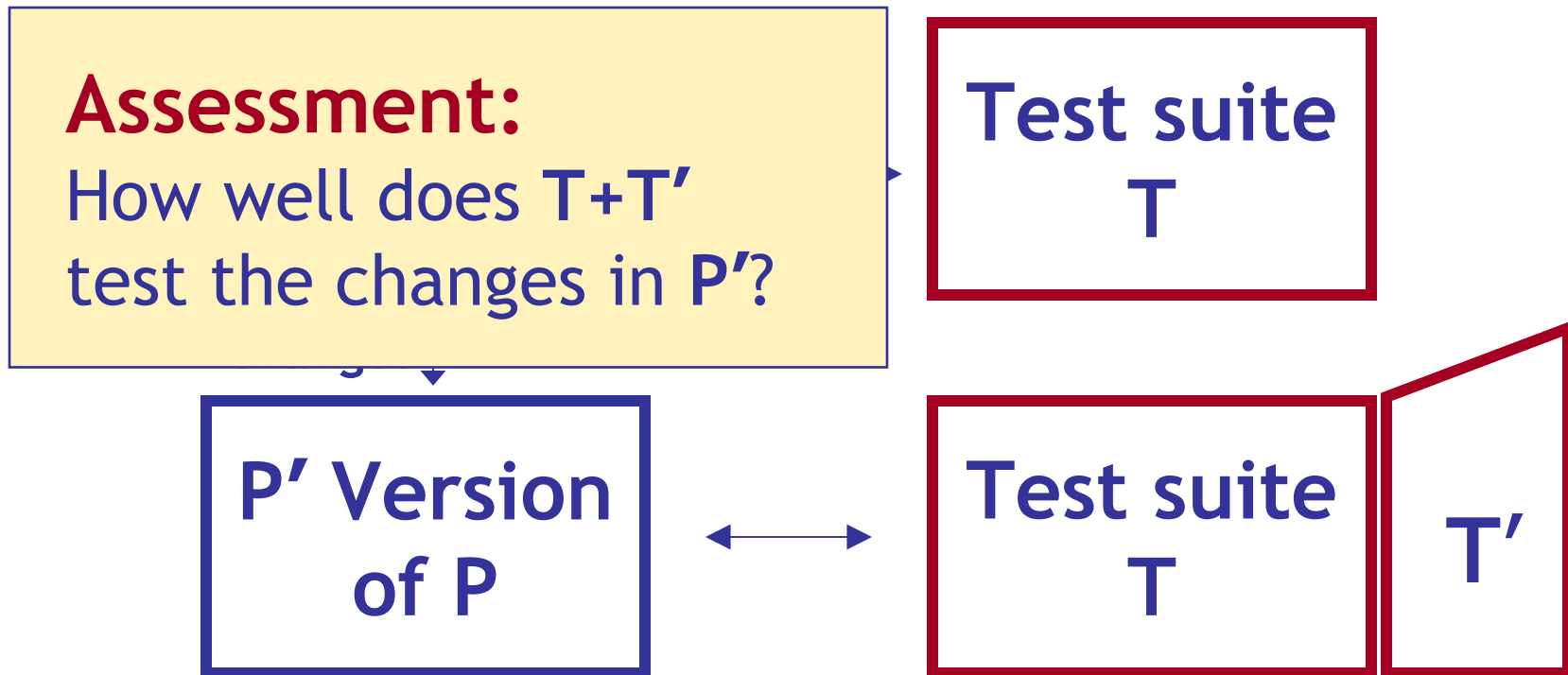
# Regression Testing



# Regression Testing



# Regression Testing



# Regression Testing

## Assessment:

How well does  $T+T'$  test the changes in  $P'$ ?

## Addition:

What else needs to be tested for those changes?

Test suite  
 $T$

Test suite  
 $T$

$T'$

$T''$

# Regression Testing

**Assessment:** —————→  
How well does  $T+T'$   
test the changes in  $P'$ ?

**Addition:** —————→  
What else needs to be  
tested for those  
changes?

**Test-suite  
Augmentation**

↑  
**need to compute  
requirements**

# Outline

- Background
- Our Technique
- Study
- Future Work and Contributions

# Example

100 inputs

```
program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2)
2.   y = y + 1
   else
3.   y = y - 1
4. z = abs(x)
5. if (y > 2)
6.   print 1
   else
7.   print 0
8. ... = z
```

Test suite **T**

test	input		out
	x	y	<b>A</b>
t1	1	10	<b>1</b>
t2	10	1	<b>0</b>

# Example

change

```
program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4. z = abs(x)
5. if (y > 2)
6.   print 1
   else
7.   print 0
8. ... = z
```

Test suite T

test	input		out	out
	x	y	A	A'
t1	1	10	1	1
t2	10	1	0	0

no difference

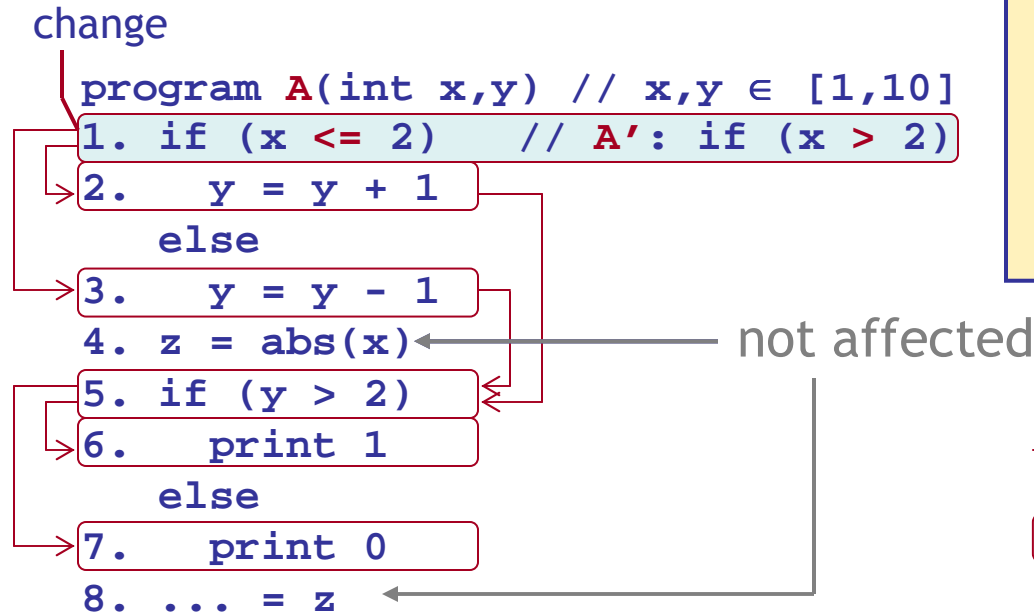
different output A → A'	x ≤ 2	x > 2
y = 2	1 → 0	0 → 1
y = 3	1 → 0	0 → 1

20/100 inputs have ≠ output!

# Previous Work on Test-suite Augmentation Criteria

- Identify entities **affected** by (dependent on) changes
  - statements
  - control dependencies (branches)
  - data dependencies (du-pairs)
- Require testing (coverage) of those entities  
[Rothermel & Harrold 94, Binkley 97]

# Example: Affected Dependencies



Dependencies are affected by a change directly or transitively

→ affected dependence  
 □ affected statement

Dependencies:  $(1,2), (1,3), (2,5), (3,5), (5,6), (5,7)$   
 direct on 1                      transitive on 1

# Example: Augmentation with Affected Dependencies

```

change
program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
  2.   y = y + 1
  else
  3.   y = y - 1
  4.   z = abs(x)
5. if (y > 2)
  6.   print 1
  else
  7.   print 0
8.   ... = z
  
```

Test suite T

test	input		out	out
	x	y	A	A'
t1	1	10	1	1
t2	10	1	0	0

assess

no difference


add <nothing>

Coverage of dependencies on A':

test	(1,2)	(1,3)	(2,5)	(3,5)	(5,6)	(5,7)
t1		•		•		•
t2	•		•		•	

# Analyzing Augmentation Criteria

change

```
program A(int x,y)  x,y ∈ [1, 100]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
```

## PIE model [Voas 92]:

- Execute change
- Infect state
- Propagate infection to output

# Analyzing Augmentation Criteria

E

```
program A(int x,y) // x,y ∈ [1,
1. if (x ≤ 2) // A': if (x >
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
```

## PIE model [Voas 92]:

- Execute change
- Infect state
- Propagate infection to output

# Analyzing Augmentation Criteria

E  
!

```
program A(int x,y) // x,y ∈ [1,
1. if (x ≤ 2) // A': if (x >
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
```

## PIE model [Voas 92]:

- Execute change
- Infect state
- Propagate infection to output

# Analyzing Augmentation Criteria

```
E
program A(int x,y) // x,y ∈ [1,
1. if (x ≤ 2) // A': if (x >
P 2.   y = y + 1
   else
   → 3.   y = y - 1
     4. z = abs(x)
     ← P
   5. if (y > 2)
   P 6.   print 1
     else
     → 7.   print 0 (!)
     8. ... = z
```

output infected ✓

**PIE** model [Voas 92]:

- Execute change
- Infect state
- Propagate infection to output

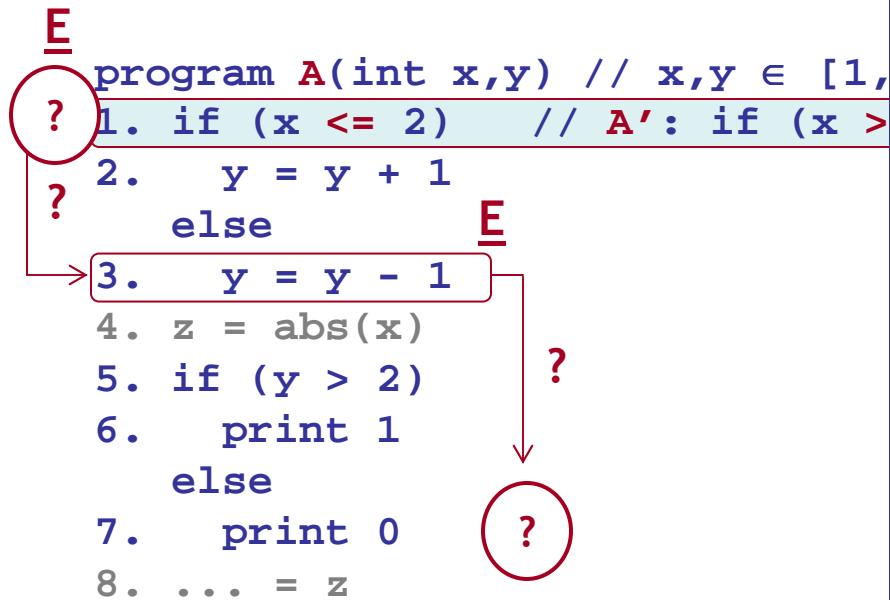
# Limitations of Previous Work

```
E  
program A(int x,y) // x,y ∈ [1,  
1. if (x ≤ 2) // A': if (x >  
2.   y = y + 1  
   else  
3.   y = y - 1 E  
4.   z = abs(x)  
5.   if (y > 2)  
6.     print 1  
   else  
7.     print 0  
8.   ... = z
```

## PIE model [Voas 92]:

- Execute change
- Infect state
- Propagate infection to output

# Limitations of Previous Work



## PIE model [Voas 92]:

- Execute change
- Infect state
- Propagate infection to output

Problems: State might not be infected  
Infection might not propagate to output

# Our Technique

E  
program **A**(int x,y) /  
1. if (x <= 2) //  
2. y = y + 1  
   else  
3. y = y - 1  
4. z = abs(x)  
5. if (y > 2)  
6. print 1  
   else  
7. print 0  
8. ... = z

Goal: requirements for **i**nfecting state at change and **p**ropagating infection to **o**utput

# Our Technique

```
E  
program A(int x,y) //  
1. if (x <= 2) //  
P 2. y = y + 1  
   else  
   → 3. y = y - 1  
     4. z = abs(x)  
     P 5. if (y > 2)  
       6. print 1  
       else  
       → 7. print 0  
     8. ... = z
```

Goal: requirements for **i**nfecting state at change and **p**ropagating infection to **o**utput

stage

**1**

identify *chains* of dependencies (propagation paths)

Example chain: (1,3),(3,5),(5,7)

# Our Technique

```
E
program A(int x,y) //
1. if (x <= 2) //
P 2.   y = y + 1
   else
   → 3.   y = y - 1
     4.   z = abs(x)
     ← P
     5.   if (y > 2)
     P 6.   print 1
       else
       → 7.   print 0 (!)
     8.   ... = z
```

Goal: requirements for **i**nfecting state at change and **p**ropagating infection to **o**utput

stage

1

identify *chains* of dependencies (propagation paths)

Example chain: (1,3),(3,5),(5,7)

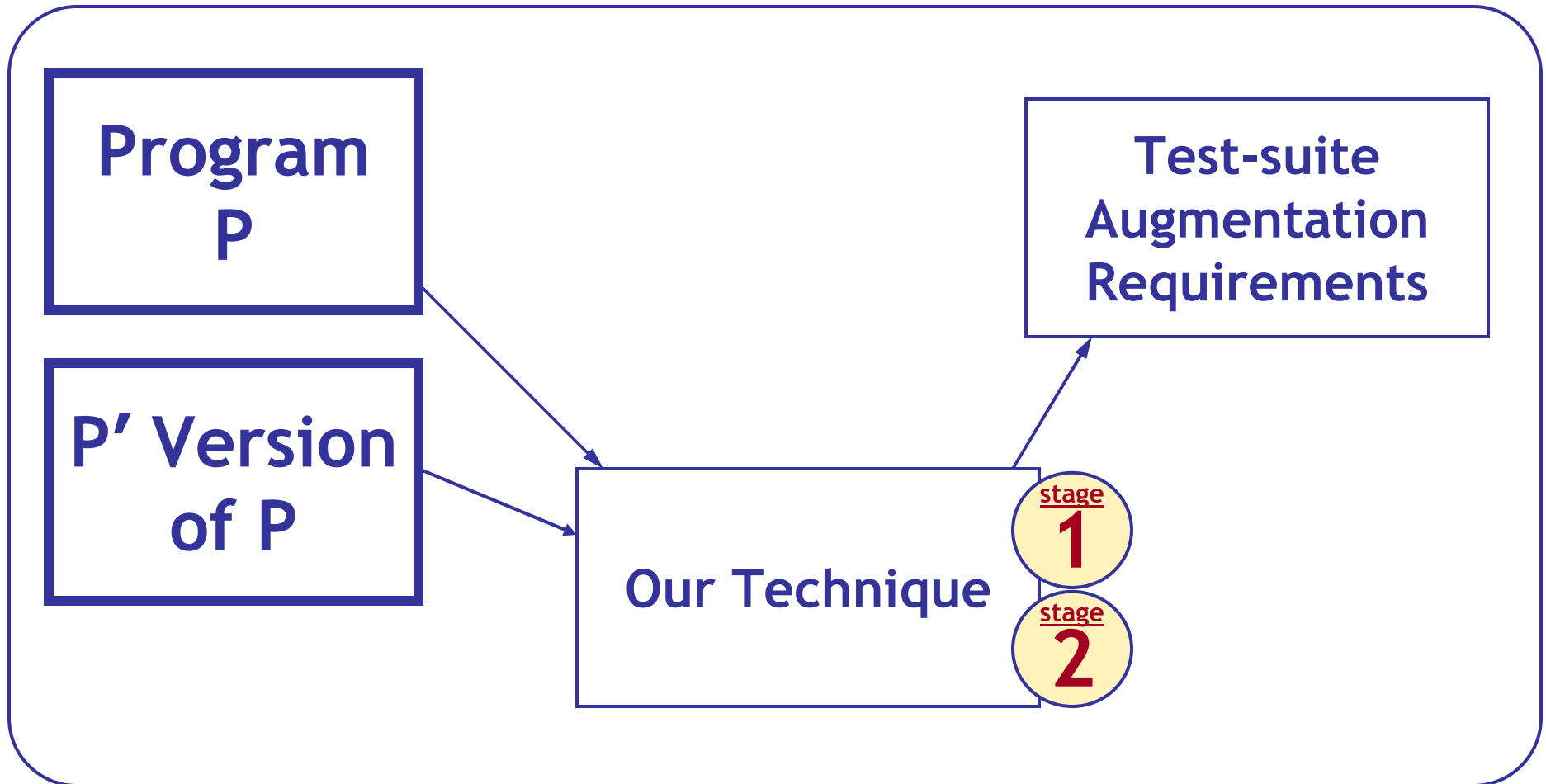
stage

2

compute *infection conditions* on chains (using *partial* symbolic execution)

infection propagates to end of chain!

# Our Technique



# stage 1 Dependence Chains

## Problem

- Chains to output can be too long (or infinite)
- Number of chains might grow exponentially

## Solution

Set limit  $d$  to length of chains

## Intuition

Covering all dependence chains of length  $d$  increases chances of propagation to output

stage  
**1**

# Coverage of Chains

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

Chain length: 1

chain on A'	x	y	# inputs	≠ output
ch <sub>1</sub> = (1,2)	x > 2	[1,10]	80	
ch <sub>2</sub> = (1,3)	x ≤ 2	[1,10]	20	

different output A → A'	x ≤ 2	x > 2
y = 2	1 → 0	0 → 1
y = 3	1 → 0	0 → 1

stage  
**1**

# Coverage of Chains

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
   2.   y = y + 1
   else
   3.   y = y - 1
4. z = abs(x)
5. if (y > 2)
6.   print 1
   else
7.   print 0
8. ... = z
  
```

Chain length: 1

chain on A'	x	y	# inputs	≠ output
ch <sub>1</sub> = (1,2)	x > 2	[1,10]	80	16
ch <sub>2</sub> = (1,3)	x ≤ 2	[1,10]	20	

different output A → A'	x ≤ 2	x > 2
y = 2	1 → 0	0 → 1
y = 3	1 → 0	0 → 1

8 × 2 = 16

# stage 1 Coverage of Chains

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

Chain length: 1

chain on A'	x	y	# inputs	≠ output
ch <sub>1</sub> = (1,2)	x > 2	[1,10]	80	16
ch <sub>2</sub> = (1,3)	x ≤ 2	[1,10]	20	4

different output A → A'	x ≤ 2	x > 2
y = 2	1 → 0	0 → 1
y = 3	1 → 0	0 → 1

2 × 2 = 4

# stage 1 Coverage of Chains

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

Chain length: 1

chain on A'	x	y	# inputs	≠ output
ch <sub>1</sub> = (1,2)	x > 2	[1,10]	80	16
ch <sub>2</sub> = (1,3)	x ≤ 2	[1,10]	20	4

Probability of test suite finding a difference =

$$p_{\text{diff}}(\text{ch}_1) + (1 - p_{\text{diff}}(\text{ch}_1)) \times p_{\text{diff}}(\text{ch}_2) = 16/80 + (1 - 16/80) \times 4/20 = \mathbf{36\%}$$

# stage 1 Coverage of Chains

```

change
program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

Chain length: 3

chain on A'	x	y	# inputs	≠ output
(1,2),(2,5),(5,6)	$x > 2$	$y > 1$	72	16
(1,2),(2,5),(5,7)	$x > 2$	$y \leq 1$	8	0
(1,3),(3,5),(5,6)	$x \leq 2$	$y > 3$	14	0
(1,3),(3,5),(5,7)	$x \leq 2$	$y \leq 3$	6	4

Probability of test suite finding a difference = **83%** ← much better

different output A → A'	$x \leq 2$	$x > 2$
$y = 2$	<b>1 → 0</b>	<b>0 → 1</b>
$y = 3$	<b>1 → 0</b>	<b>0 → 1</b>

stage  
**2**

# Infection Conditions

change

```
program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.
```

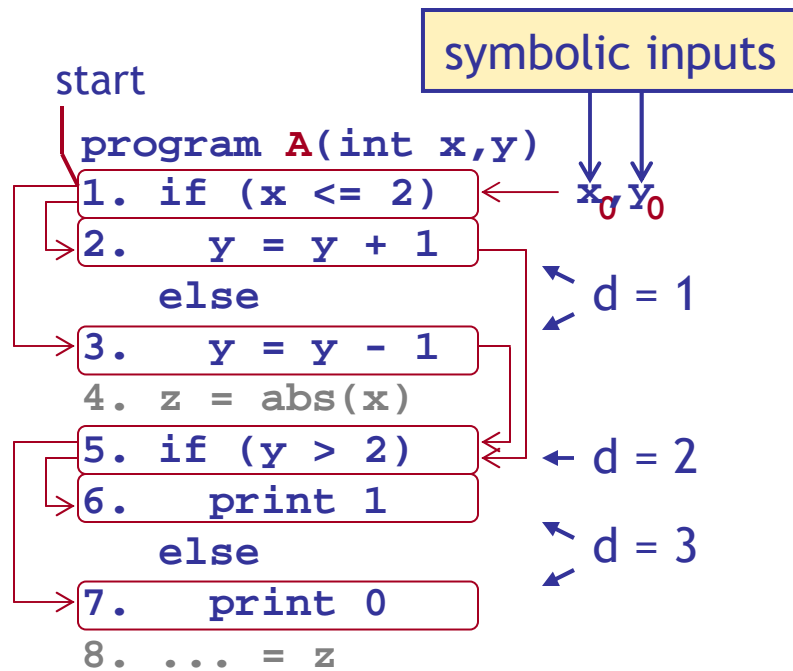
Our technique adds conditions to **guarantee** infection of state and propagation of infection to end of each chain.

## Intuition

Propagating infection to end of chain increases chances of propagating to the **output**.

stage  
**2**

# Partial Symbolic Execution



For program **A**:

always  
 $x = x_0$

statement	pc	state
1	true	$y = y_0$
2	$x_0 \leq 2$	$y = y_0$
3	$x_0 > 2$	$y = y_0$
5	$x_0 \leq 2$	$y = y_0 + 1$
	$x_0 > 2$	$y = y_0 - 1$
6	$x_0 \leq 2 \wedge y_0 + 1 > 2$	$y = y_0 + 1$
	$x_0 > 2 \wedge y_0 - 1 > 2$	$y = y_0 - 1$
7	$x_0 \leq 2 \wedge y_0 + 1 \leq 2$	$y = y_0 + 1$
	$x_0 > 2 \wedge y_0 - 1 \leq 2$	$y = y_0 - 1$

[Apiwattanapong et al., 2006]

stage

2

# Partial Symbolic Execution

## Summary

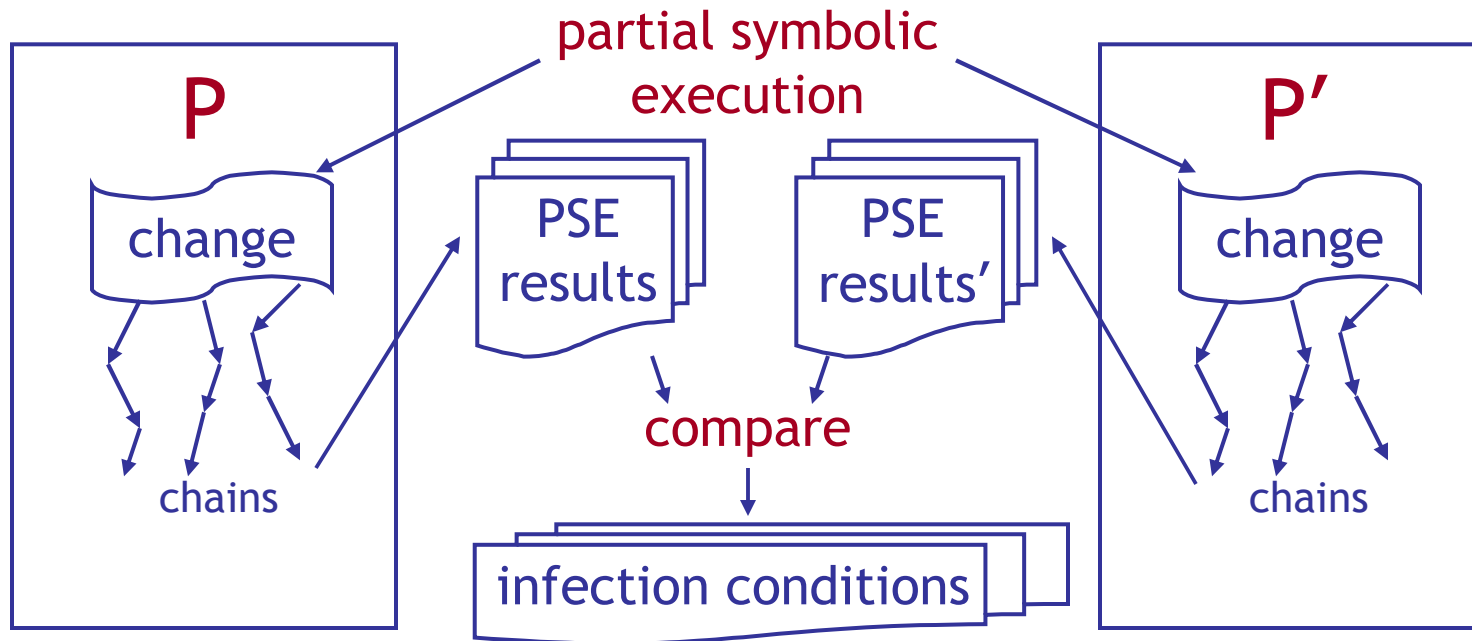
1. Starts at arbitrary point (change) in program (symbolic inputs are variables at that point)
2. Stops at given distance  $d$

Our technique uses partial symbolic execution to compute **infection conditions**.

**[Apiwattanapong et al., 2006]**

stage  
**2**

# Infection Conditions



For each chain in  $P'$ , compare results on  $P$  and  $P'$ :

1. *Live* states of  $P'$  and  $P$  differ at *end* of the chain
2.  $P'$  covers the chain, but  $P$  does not reach its *end*

# stage 2 Concrete Example

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

chain  $h = (1,2),(2,5),(5,6)$

program	pc	live state
<b>A</b>	$x_0 < 2 \wedge y_0 + 1 > 2$ $x_0 \geq 2 \wedge y_0 - 1 > 2$	- -
<b>A'</b>	$x_0 \geq 2 \wedge y_0 + 1 > 2$	-

conditions for chain  
 $(1,2),(2,5),(5,6)$

Types of infection for a chain  $h$  in  $A'$ :

1. Live states of  $A'$  and  $A$  differ at end of  $h \rightarrow \text{state}(A',h) \neq \text{state}(A,h)$
2.  $A'$  covers  $h$ , but  $A$  does not reach  $\text{end}(h) \rightarrow \text{pc}(A',h) \wedge \neg \text{pc}(A,\text{end}(h))$

# stage 2 Concrete Example

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

chain  $h = (1,2),(2,5),(5,6)$

program	pc	live state
A	$x_0 < 2 \wedge y_0 + 1 > 2$	-
	$x_0 \geq 2 \wedge y_0 - 1 > 2$	-
A'	$x_0 \geq 2 \wedge y_0 + 1 > 2$	-

conditions for chain  
 $(1,2),(2,5),(5,6)$

Types of infection for a chain  $h$  in  $A'$ :

1. Live states of  $A'$  and  $A$  differ at end of  $h \rightarrow$  *false*
2.  $A'$  covers  $h$ , but  $A$  does not reach  $end(h) \rightarrow pc(A',h) \wedge \neg pc(A,end(h))$

# stage 2 Concrete Example

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

chain  $h = (1,2),(2,5),(5,6)$

program	pc	live state
A	$x_0 < 2 \wedge y_0 + 1 > 2$	-
	$x_0 \geq 2 \wedge y_0 - 1 > 2$	-
A'	$x_0 \geq 2 \wedge y_0 + 1 > 2$	-

conditions for chain  
 $(1,2),(2,5),(5,6)$

Types of infection for a chain  $h$  in  $A'$ :

- ~~1. Live states of  $A'$  and  $A$  differ at end of  $h \rightarrow false$~~
- $A'$  covers  $h$ , but  $A$  does not reach  $end(h) \rightarrow (x_0 \geq 2 \wedge y_0 + 1 > 2) \wedge \neg ((x_0 < 2 \wedge y_0 + 1 > 2) \vee (x_0 \geq 2 \wedge y_0 - 1 > 2))$

# stage 2 Concrete Example

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
  
```

chain  $h = (1,2),(2,5),(5,6)$

program	pc	live state
A	$x_0 < 2 \wedge y_0 + 1 > 2$	-
	$x_0 \geq 2 \wedge y_0 - 1 > 2$	-
A'	$x_0 \geq 2 \wedge y_0 + 1 > 2$	-

conditions for chain  
 $(1,2),(2,5),(5,6)$

Types of infection for a chain  $h$  in  $A'$ :

- ~~1. Live states of  $A'$  and  $A$  differ at end of  $h \rightarrow false$~~
- $A'$  covers  $h$ , but  $A$  does not reach  $end(h) \rightarrow x_0 \geq 2 \wedge y_0 \in \{2,3\}$   
 $\underbrace{\hspace{10em}}_{8 \times 2 = 16 \text{ inputs}}$

# stage 1 Chains + stage 2 Infection Conditions

change

```

program A(int x,y) // x,y ∈ [1,10]
1. if (x ≤ 2) // A': if (x > 2)
2.   y = y + 1
   else
3.   y = y - 1
4.   z = abs(x)
5.   if (y > 2)
6.     print 1
   else
7.     print 0
8.   ... = z
    
```

	stage 1	+	stage 2	
chain	# inputs covering chain		# inputs infecting chain	P <sub>diff</sub> before → after
(1,2),(2,5),(5,6)	72		16	16/72 → 16/16
(1,2),(2,5),(5,7)	8		0	0 → 0
(1,3),(3,5),(5,6)	14		0	0 → 0
(1,3),(3,5),(5,7)	6		4	4/6 → 4/4

Technique guarantees that test suite finds a difference in this example!

# Work Related to Our Technique

Theory of fault-based testing [Morell 84]

↑  
refines

RELAY framework [Richardson & Thompson 88]

↑  
complements

PIE dynamic estimation technique [Voas 92]

# Study: Goal and Setup

Goal: evaluate effectiveness

Toolset: **MATRIX-RELOADED** for Java bytecode\*

- Successor of **MATRIX** [Apiwattanapong et al., 2006]

Subjects:

subject	SLOC	tests	changes
Tcas	131	1608	6
NanoXML	~ 4000	216	9

\* Based on the Soot Analysis Framework

# Experiment Design

## Augmentation criteria

Statement: changed statements

Data: chains of data-only dependencies

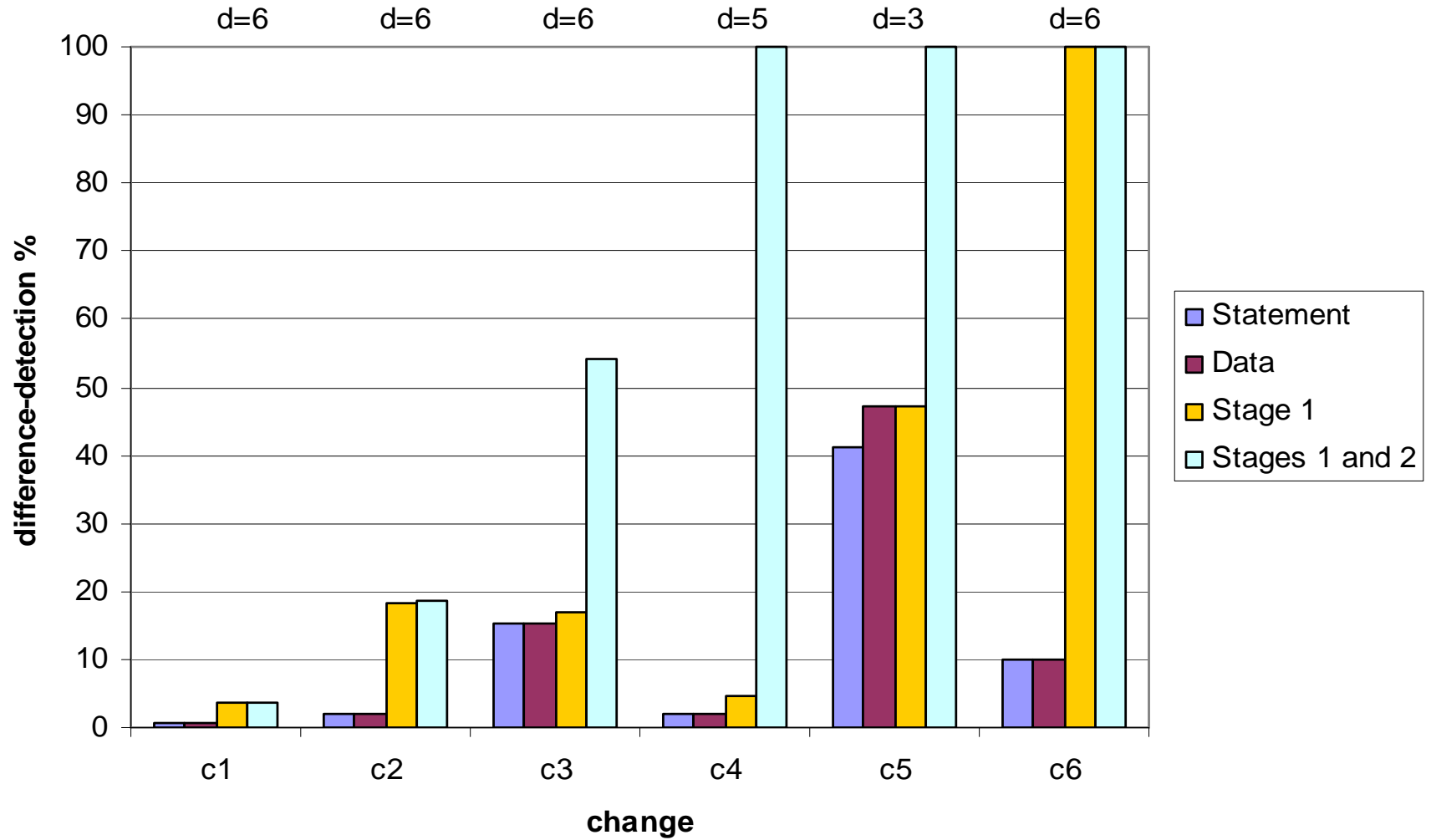
**stage 1** Stage 1: chains of data/control dependencies

**stage 1+2** Stages 1 and 2: chains with infection conditions

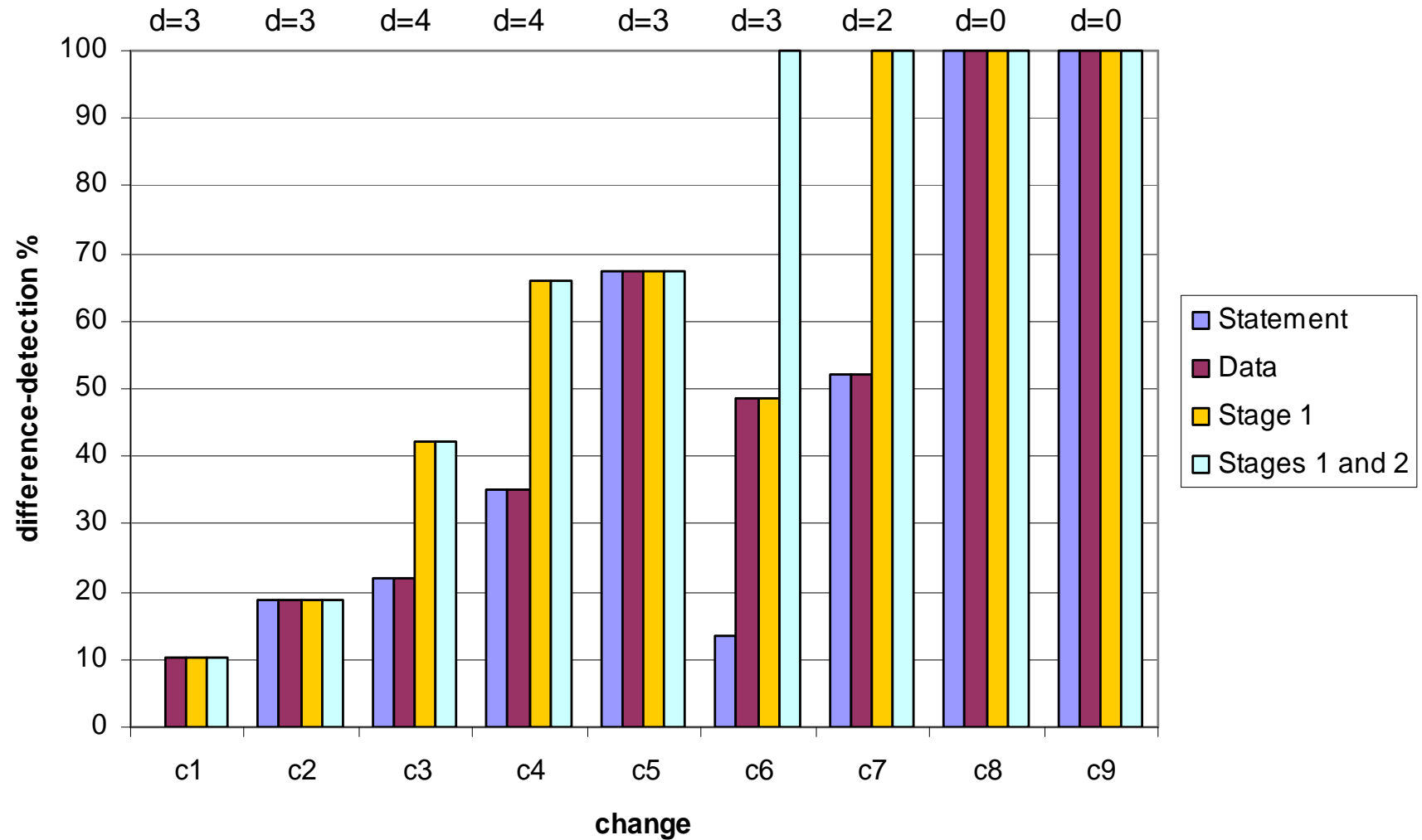
## For each criterion

- Generated 100 test suites satisfying criterion
- Computed % of test suites revealing output difference
- Used maximum distance our toolset reached

# Results for Tcas



# Results for NanoXML



# Future Work

1. Improve toolset and perform **more** experiments
2. Improve chances of propagating infection by finding “infection-killing” points **beyond** limit
3. Address the problem of large changes
4. Automate **generation** of test cases for our requirements

# Contributions

1. Approach is **propagation-based**, rather than just coverage-based
2. Technique increases probability of difference detection in two stages
  - Stage 1**: “propagation paths” (chains)
  - Stage 2**: infection conditions (partial symbolic execution)
3. Technique is **practical**
  - distance limit reduces **length** and **number** of chains
  - partial symbolic execution** controls infection conditions