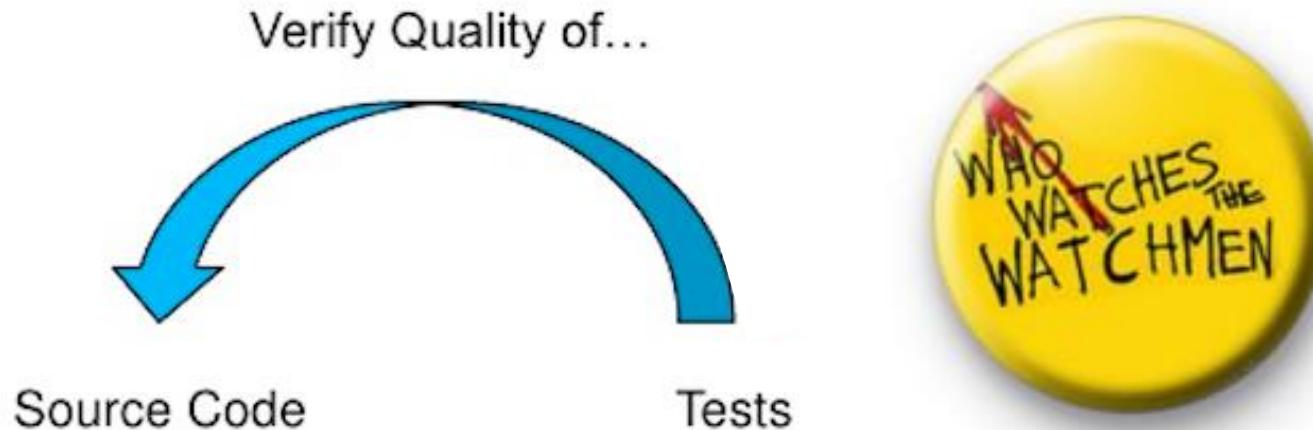


Mutant Subsumption in First- and Second-Order Mutation Testing

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Introduction



Mutation Testing

- Introducing **artificial syntactic changes (mutations)** into original source code
 - Intending to represent real common programming bugs
 - Changed programs are called **mutants**
- Running the test suite on mutants
 - Result different from original: mutant **killed**
 - Otherwise: **alive**

Example of a mutant

Mutation place:

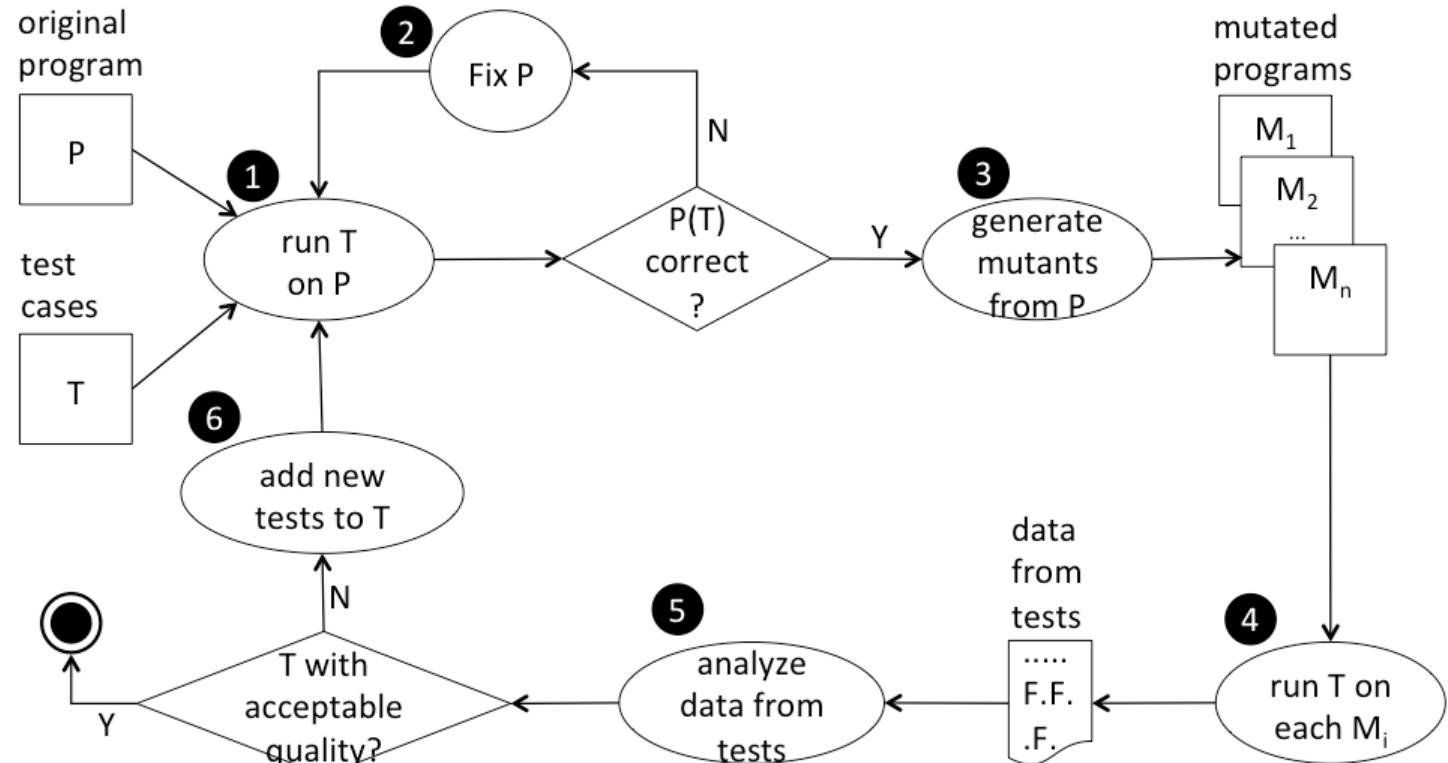
```
public class Taxes {  
  
    double simpleTax(double amount) {  
  
        return amount * 0.2;  
    }  
}
```

Example of a mutant

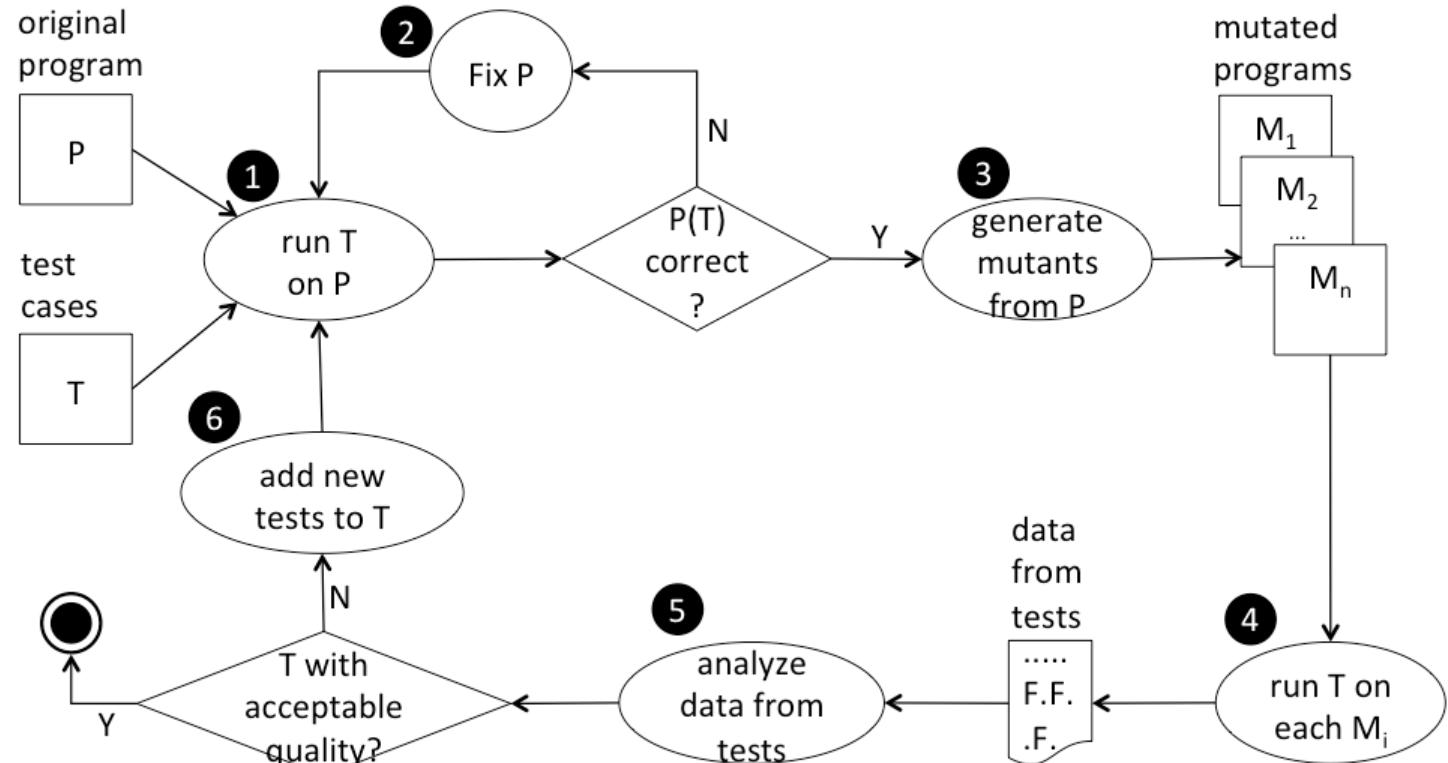
Mutation: * → +

```
public class Taxes {  
  
    double simpleTax(double amount) {  
  
        return amount + 0.2;  
    }  
}
```

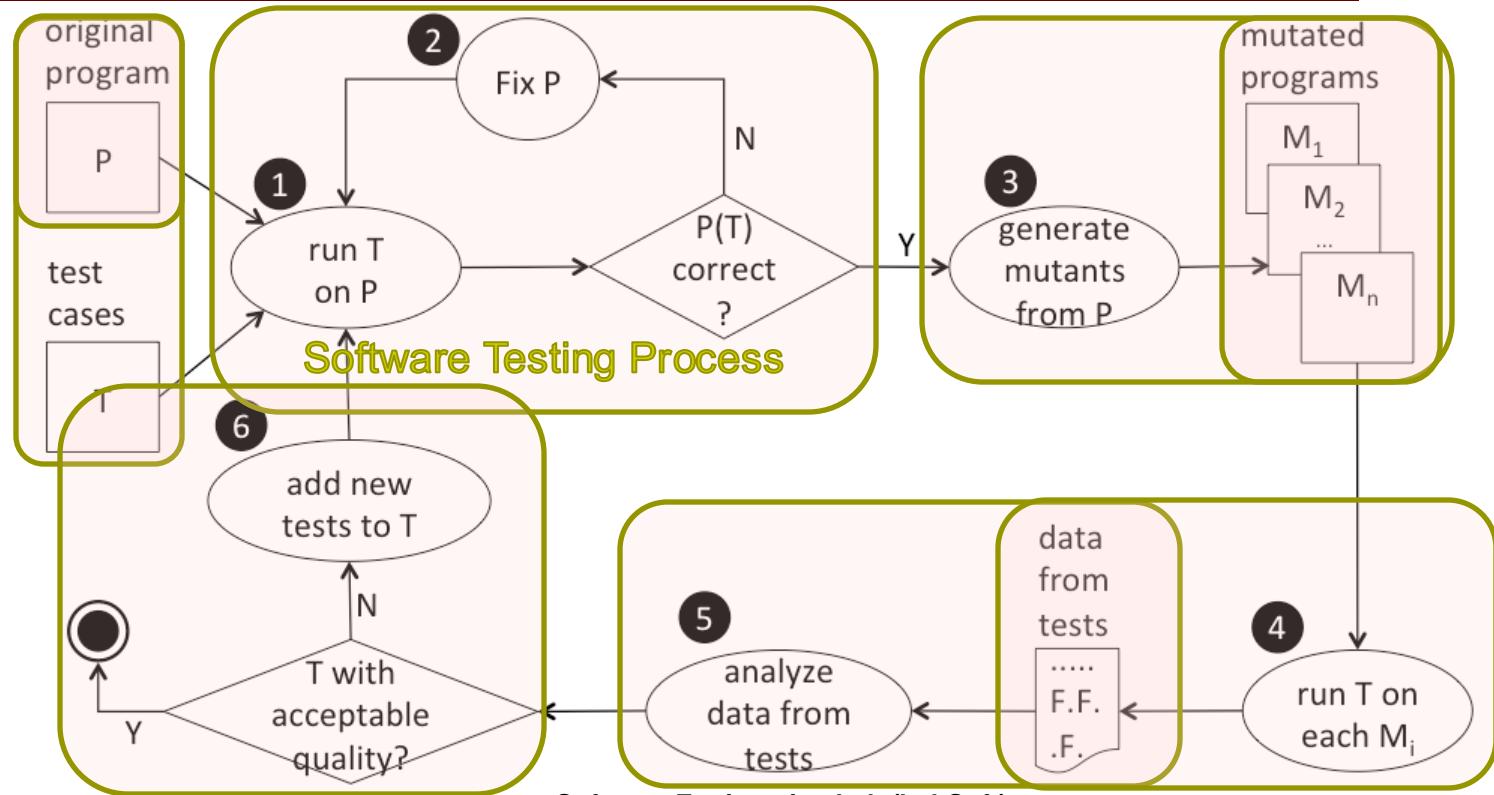
Mutation testing process



Mutation testing process



Mutation testing process



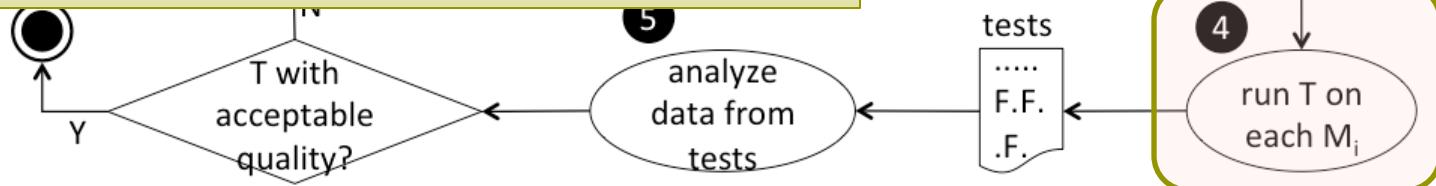
Mutation testing drawbacks

3. Many mutable places

- Mutants generation
- Compilation

4. High computational cost

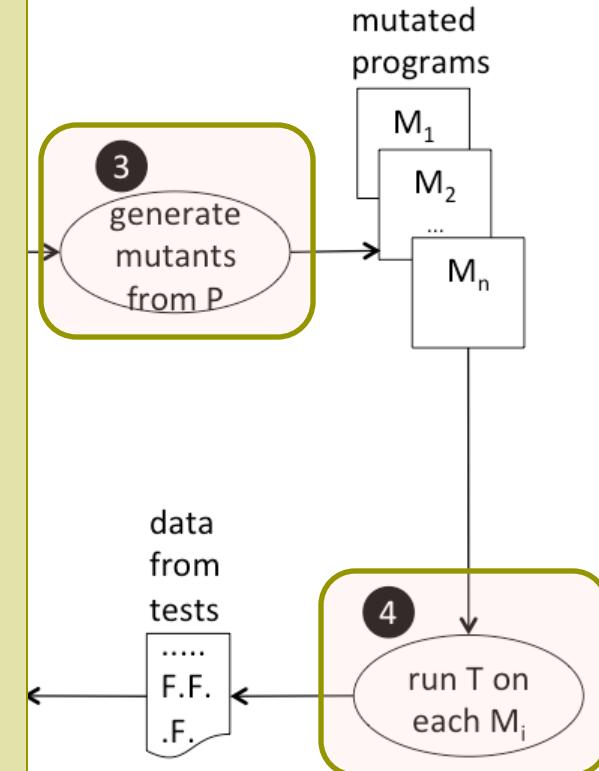
- Running tests



Mutation testing drawbacks

- Cost reduction techniques

- Number of test cases
- Test case prioritization
- Number of mutants
 - **subsumption**



Mutants subsumption

Contextualizing

```
def greaterThan(a, b):  
    return a > b # original
```

```
def greaterThan(a, b):  
    return a >= b # mutant 1
```

```
def greaterThan(a, b):  
    return a <= b # mutant 2
```

Contextualizing

```
def greaterThan(a, b):  
    return a > b  # original
```

	Test	orig
t1	assertTrue(greaterThan(6, 5))	✓
t2	assertFalse(greaterThan(5, 5))	✓
t3	assertFalse(greaterThan(5, 6))	✓

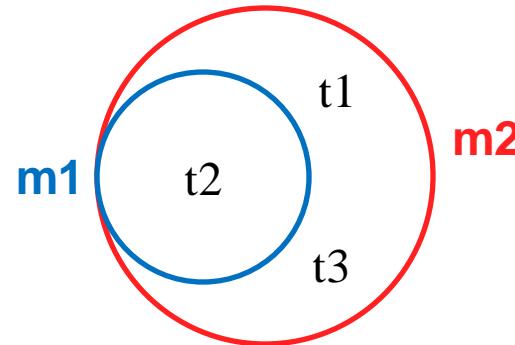
Contextualizing

```
def greaterThan(a, b):  
    return a > b # original  
    return a >= b # mutant 1  
    return a <= b # mutant 2
```

	Test	orig	m1	m2
t1	assertTrue(greaterThan(6, 5))	✓	✓	✗
t2	assertFalse(greaterThan(5, 5))	✓	✗	✗
t3	assertFalse(greaterThan(5, 6))	✓	✓	✗

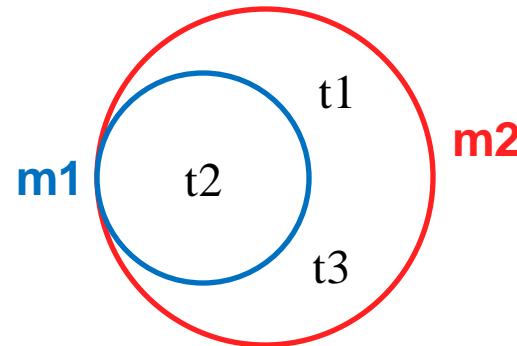
Contextualizing

□ Killing tests



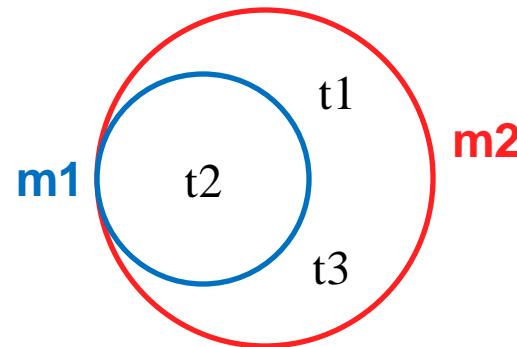
Contextualizing

- All test sets that kill **m1** also kill **m2**



Definition

- **m1 subsumes m2**



In summary

- If we know beforehand that
 - **m₁** subsumes **m₂**
- Therefore,
 - **m₂** should not have been generated

Cost reduction: fewer mutants to run the test suite against

Dynamic mutant subsumption graphs

Example

test	m1	m2	m3	m4	m5
t1	✗	✗		✗	✗
t2	✗		✗	✗	
t3				✗	
t4		✗		✗	✗

Subsumption relationships

test	m1	m2	m3	m4	m5
t1	✗	✗		✗	✗
t2	✗		✗	✗	
t3				✗	
t4		✗		✗	✗

$m1 \rightarrow m4$

$m2 \rightarrow m4$

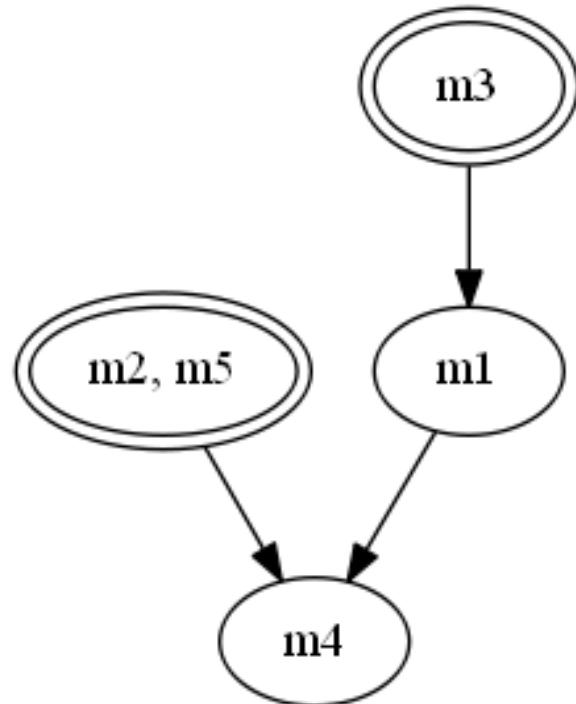
$m3 \rightarrow m1$

$m3 \rightarrow m4$

$m5 \rightarrow m4$

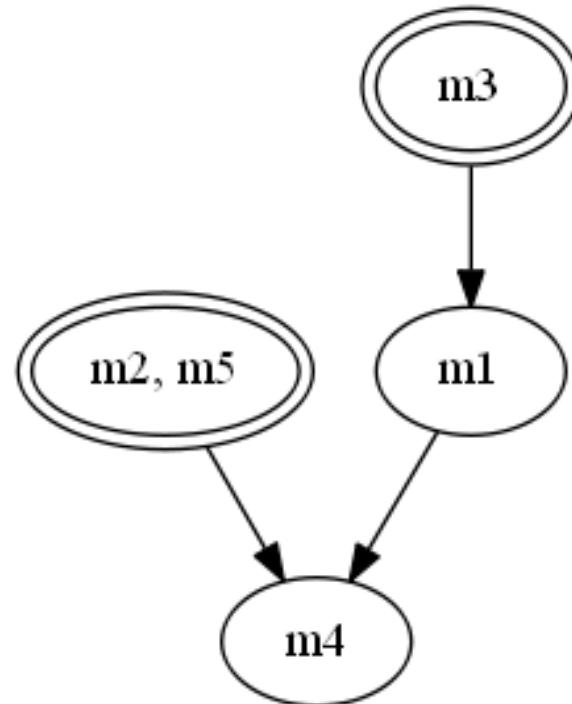
Subsumption graph

Test	m1	m2	m3	m4	m5
t1	✗	✗		✗	✗
t2	✗		✗	✗	
t3				✗	
t4		✗		✗	✗



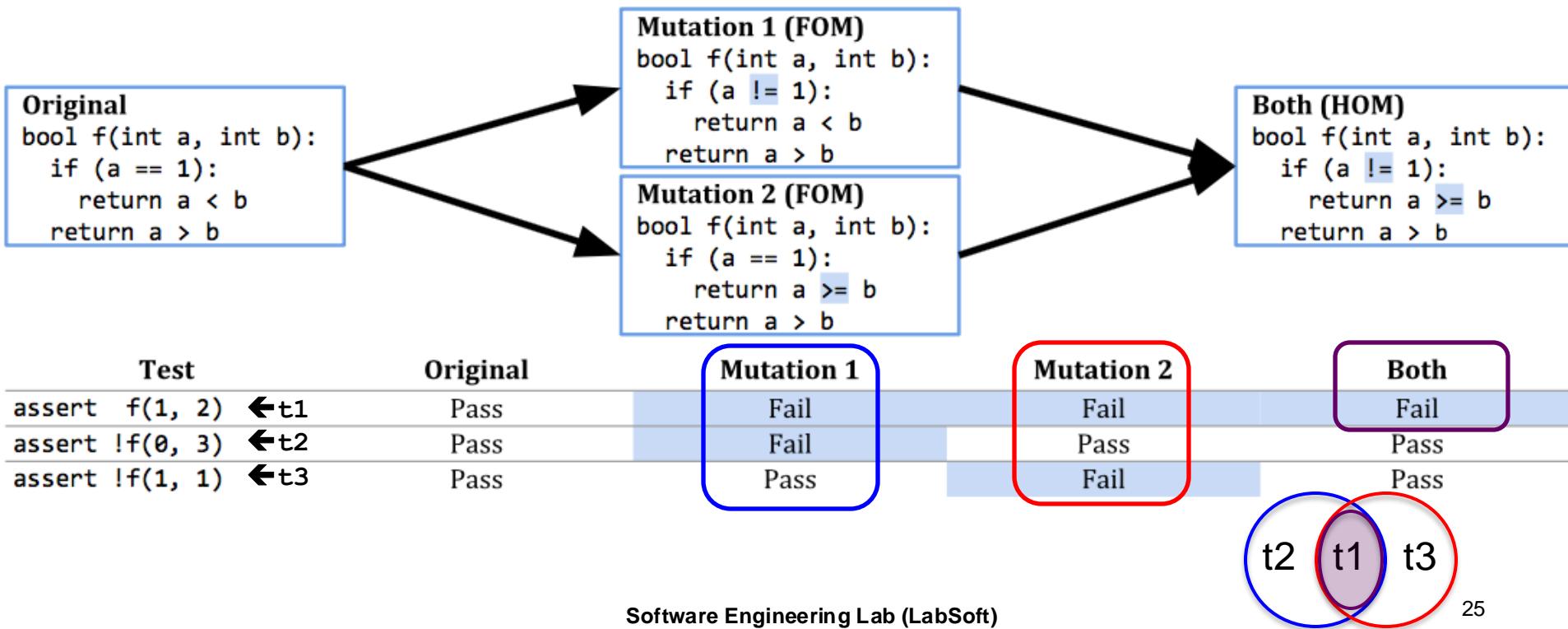
Conclusion

- Root nodes are kept
 - 2 minimal
 - 3 mutants
- Remaining nodes
 - are disregarded
 - (redundants)



Second-order mutants subsumption

A 2OM subsumption example



A 2OM subsumption example

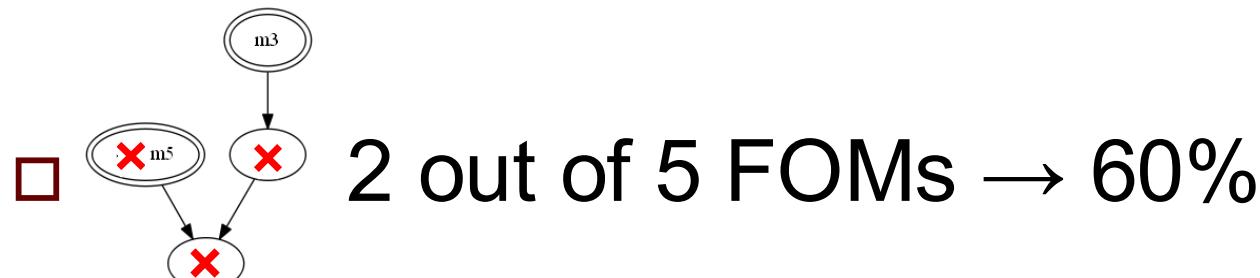
Strongly subsuming second-order mutant
(SS2OM)

An SS2OM can replace their constituent
FOMs without loss of effectiveness

Mutants reduction via subsumption

From FOMs

- Remove all FOMs in subsumed nodes
- Randomly pick one FOM in each minimal node



From SS2OMs

FOMs

m1

m2

m3

m4

m5

m6

m7

m8

From SS2OMs

FOMs SS2OMs

m1 [m1,m2]

m2 [m3,m4]

m3 [m5,m6]

m4

m5

m6

m7

m8

From SS2OMs

FOMs	SS2OMs	Resulting mutants
m1	[m1,m2]	[m1,m2]
m2	[m3,m4]	[m3,m4]
m3	[m5,m6]	[m5,m6]
m4		m7
m5		m8
m6		
m7		
m8		

non-subsumed FOMs

From SS2OMs

FOMs	SS2OMs	Resulting mutants	Reduction
m1	[m1,m2]	[m1,m2]	3 out of 8 mutants
m2	[m3,m4]	[m3,m4]	(37.5%)
m3	[m5,m6]	[m5,m6]	
m4		m7	
m5		m8	
m6			
m7			
m8			

non-subsumed FOMs

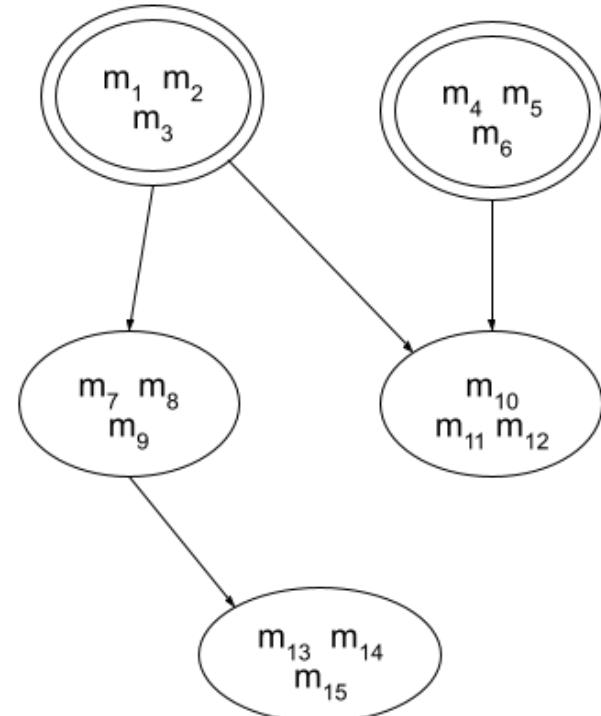
Study design

RQ1

- Which *subsumption* can reduce more mutants?
 - From FOMs
 - From SS2OMs

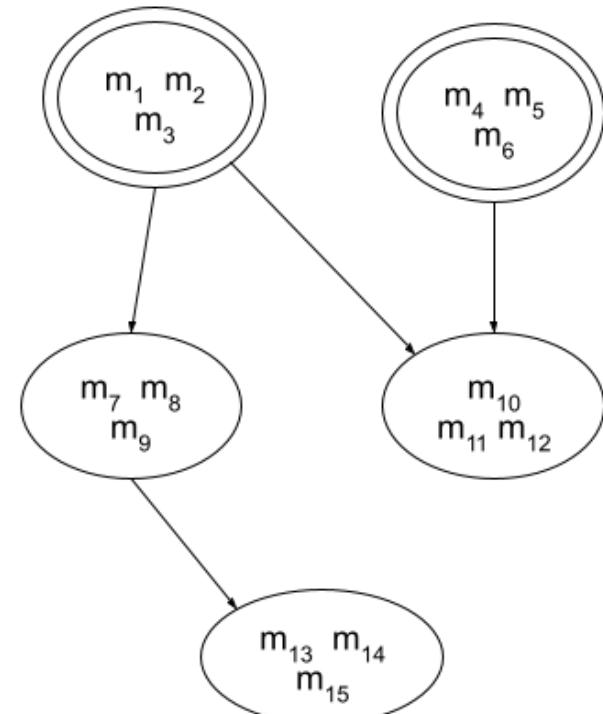
RQ2

- What is the distribution of the SS2OMs **constituent FOMs** in the subsumption graphs?



RQ2 – example of SS2OMs

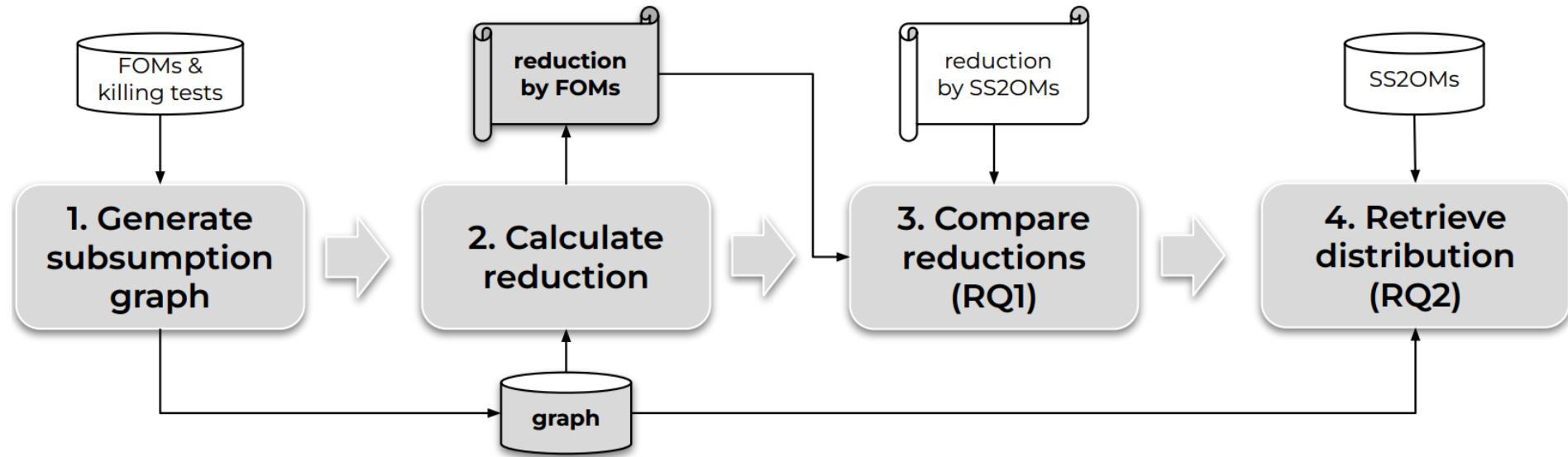
- $[m_1, m_3]$: same minimal node
- $[m_1, m_4]$: \neq minimal nodes
- $[m_4, m_7]$: 1 minimal, 1 subsumed
- $[m_{10}, m_{11}]$: same subsumed nodes
- $[m_{10}, m_{13}]$: \neq subsumed nodes



Dataset: 9 Java systems

System	LOC	# Tests	#FOMs	#SS2OMs
Vending Machine	~100	35	57	25
Triangle	34	12	138	393
Monopoly	1,181	124	866	3,324
Commons CSV	~2k	325	925	4,430
Commons CLI	2,699	318	1,082	1,852
ECal	3,626	224	1,207	1,421
Commons Validator	7,409	536	3,197	17,546
Gson	> 10k	1,089	3,712	6,970
Chess	4,924	930	5,287	8,959
Overall			16,471	44,960

Study steps



Results

Reduction comparison (RQ1)

RQ1 answer

Overall	#FOMs	#minimal nodes	#remaining FOMs
9 systems	16,471	1,115	3,376

Highlight on Triangle

{91}

{65, 67, 68, 70, 75, 77, 80, 52, 85, 56, 57, 58, 59, 63}

{35, 37, 38, 39, 108, 112, 114, 115, 116, 118, 119}

{129, 130, 131, 46, 122, 123, 124, 127}

{11}

{62}

{79}

{5}

{96, 132, 136, 121, 106, 111}

{69}

{97, 99, 100, 101, 103, 104, 23, 24, 25, 26, 27, 28, 93}

{0}

	#FOMs	#minimal nodes	#remaining FOMs
	138	12	59

RQ1 answer

Overall	FOMs	Via subsumption graph	Via SS2OMs
9 systems	16,471	77.35%	22.37%

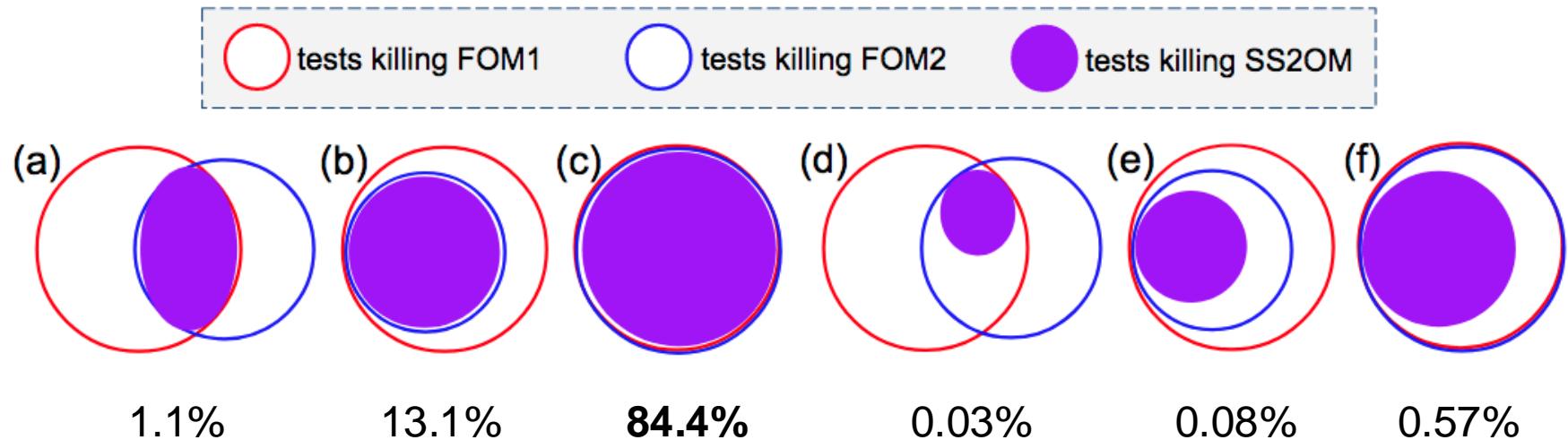
SS2OMs distribution (RQ2)

RQ2 answer

**44,960 SS2OMs &
their constituent FOMs distribution over
the subsumption graphs**

Overall	Same minimal	1 minimal 1 subsumed	Same subsumed	\neq subsumed
9 systems	20,368	1,968	18,298	4,286
	45.3%	4.4%	40.7%	9.6%

A previous result



RQ2 answer

**44,960 SS2OMs &
their constituent FOMs distribution over
the subsumption graphs**

Overall	Same minimal	\neq minimal	1 minimal 1 subsumed	Same subsumed	\neq subsumed
9 systems	20,368	0	1,968	18,298	4,286
	45.3%	0%	4.4%	40.7%	9.6%

Threats to validity

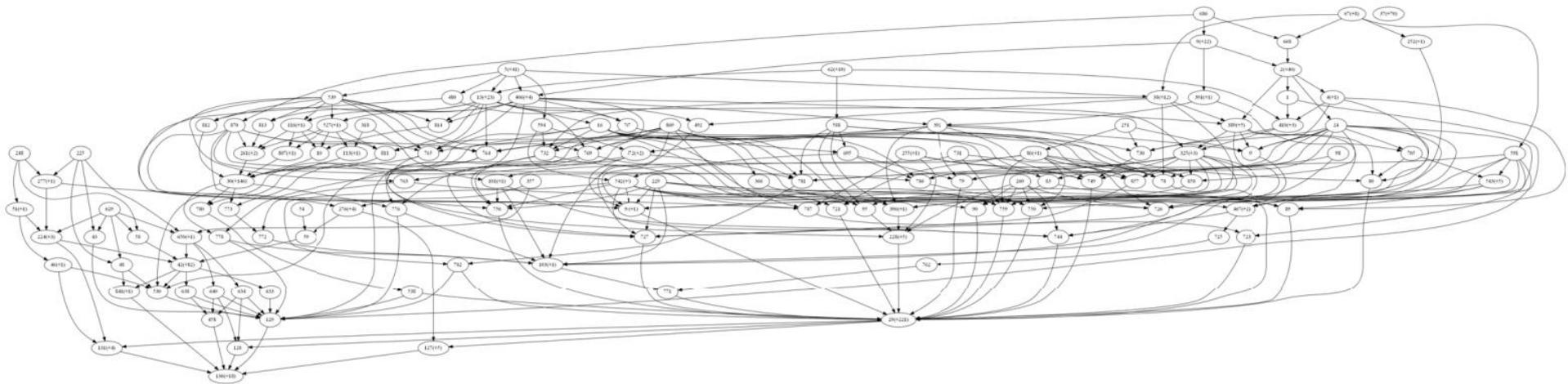
Threats to validity

- External
 - Results may not generalize
- Internal
 - Possible incorrect data from dataset
- Construction
 - Subsumption graphs correct?

Related work

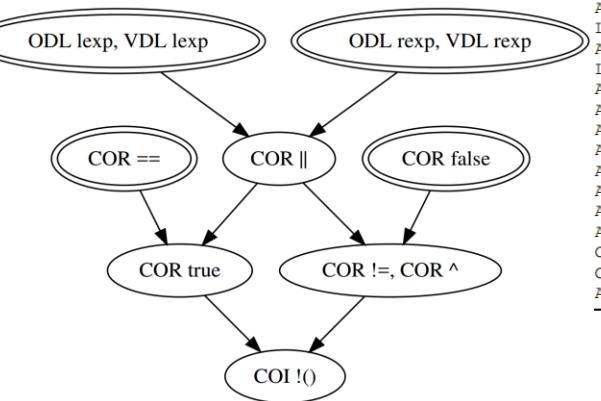
Kurtz *et al.* 2014

□ Mutant Subsumption Graphs



Guimarães *et al.* 2020

□ Optimizing Mutation Testing by Discovering Dynamic Mutant Subsumption Relations



Target	Minimal Set	Reduction
lexp + rexp	AORB %, ODL lexp, ODL rexp	62.5%
lexp + rexp (obj)	ODL lexp, ODL rexp	50.0%
lexp - rexp	AORB %, ODL lexp, ODL rexp	62.5%
lexp * rexp	AORB /, ODL lexp, ODL rexp	62.5%
lexp / rexp	AORB %, AOIS *, ODL rexp	62.5%
lexp % rexp	AORB +, AORB -, AORB /, ODL lexp	50.0%
lexp > rexp	ROR false, ROR !=, ROR >=	62.5%
lexp >= rexp	ROR true, ROR ==, ROR >	62.5%
lexp < rexp	ROR false, ROR !=, ROR <=	62.5%
lexp <= rexp	ROR true, ROR ==, ROR <	62.5%
lexp == rexp	ROR !=	62.5%
lexp == rexp (obj)	ROR !=	50.0%
lexp == rexp (bool)	ROR !=, ODL lexp, ODL rexp	50.0%
lexp != rexp	ROR true, ROR <, ROR >	62.5%
lexp != rexp (obj)	ROR ==	50.0%
lexp != rexp (bool)	ROR ==, ODL lexp, ODL rexp	50.0%
lexp && rexp	COR false, COR ==	81.8%
lexp rexp	COR true, COR !=	81.8%
lexp & rexp	ODL lexp, ODL rexp	66.7%
lexp rexp	ODL lexp, ODL rexp, LOR ^	66.7%
lexp ^ rexp	LOR	83.3%
lexp ^ rexp (bool)	COR false, COR	80.0%
exp	AOIU -exp	83.3%
+exp	LOI exp	75.0%
-exp	COD exp	50.0%
~exp	AODU exp	66.7%
	LOD exp	75.0%
	AODS exp	75.0%
	LOI exp	75.0%
	AODS exp	75.0%
	LOI exp	75.0%
	ASRS -=, ASRS %=, ODL =	50.0%
	ASRS +=, ASRS %=, ODL =	50.0%
	ASRS /=, ASRS %=, ODL =	50.0%
	ASRS *=, ASRS %=, ODL =	50.0%
	ASRS +=, ASRS -=, ASRS *=, ASRS /=	33.3%
	ASRS >=,	66.7%
	ASRS <=, ODL =, SDL	0.0%
	ASRS >=, ASRS <=	50.0%
	ODL =, SDL	50.0%
	ODL =, ASRS ^=, SDL	25.0%
	ASRS =	75.0%

Conclusion

Main finding

SS2OMs do not seem to contribute to any further reduction than the one achieved by the subsuming FOMs.

Future work

- Analyze SS3OMs
- Formal proof
 - SS2OMs constituent FOMs cannot belong to same minimal node

