

The Impact of Equivalent Mutants

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Equivalent Mutants

A mutation may not change the semantics of a program. Then it is equivalent.

- What is the problem with them?
- Is it hard to find them?
- How frequent are these?
- How can we get rid of them?

Is this mutation equivalent?

```
if(!map.containsKey(key)) {  
    Integer value = key.length();  
    map.put(key, value);  
}
```

Is this mutation equivalent?

```
if(!map.containsKey(key)) {  
    Integer value = key.length();  
    map.put(key, value);  
}
```

Note: A yellow line is drawn through the condition `!map.containsKey(key)` in the original image, with the word `true` written above it.

Is this mutation equivalent?

```
if(!map.containsKey(key)) {  
    Integer value = key.length();  
    map.put(key, value);  
}
```

Equivalent

Is this mutation equivalent?

```
counter++;  
if(!map.containsKey(key)) {  
    Integer value = counter;  
    map.put(key, value);  
}
```

Is this mutation equivalent?

```
counter++;  
if(!map.containsKey(key)) {  
    Integer value = counter;  
    map.put(key, value);  
}
```

Is this mutation equivalent?

```
counter++;  
if(!map.containsKey(key)) {  
    Integer value = counter;  
    map.put(key, value);  
}
```

Not Equivalent

Background

- **Baldwin and Seyward: Heuristic Approach**
 - Semantic-preserving compiler optimization
 - Offutt and Craft: 10 % of equivalent mutants
- **Offutt and Pan: Path Conditions**
 - If constraint solver shows subsequent states are equivalent then mutant is equivalent
 - 48 % of equivalent mutants
 - 11 Fortran-77 Programs (10-30 Statements)

Javalanche

Javalanche

An efficient mutation framework for Java

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- Manipulates byte code directly

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An efficient mutation framework for Java

- Manipulates byte code directly
- Implements *selective mutation*:
 - replace constant C by $C \pm 1$, or 0
 - negate branch condition
 - replace operators (+ by -, * by /, etc.)

Javalanche

An efficient mutation framework for Java

- Manipulates byte code directly
- Implements *selective mutation*:
 - replace constant C by $C \pm 1$, or 0
 - negate branch condition
 - replace operators (+ by -, * by /, etc.)
- Uses coverage and mutant schemata

Javalanche: Efficiency

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Jaxen	12,449 LOC	6,626 mutations	6.5h
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Publicly available this Summer

Equivalent Mutants

Experimental base – the *Jaxen XQuery Engine*:

- 6626 mutations applied by Javalanche
- 29% undetected by test suite

Random Mutants

**20 randomly chosen mutations
from 20 different classes**

non-equivalent

equivalent

undecided

Random Mutants

20 randomly chosen mutations from 20 different classes	
non-equivalent	50% (10)
equivalent	40% (8)
undecided	10% (2)

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Manual Classification

- About 15 minutes per mutant to classify.
 - Sometimes up to 2 hours per mutant.
- For 1900 (29%) mutations in Jaxen:
 - Nearly 500 hours!

Manual Classification



Idea: Measuring Impact



Coverage Impact

Original

Mutant

Coverage Impact

A mutated loop

Original

Mutant

```
for (int i = 0; i < 10; i++)
```

Coverage Impact

A mutated loop

Original

```
|  
|  
| for (int i = 0; i < 10; i++)  
10  
10  
10  
10  
10  
|  
|
```

Mutant

Coverage Impact

A mutated loop

Original

```
| .....  
| .....  
| for (int i = 0; i < 10; i++)  
10 .....  
10 .....  
10 .....  
10 .....  
10 .....  
| .....  
| .....
```

Mutant

```
for (int i = 0; i < 0; i++)
```


Coverage Impact

A mutated loop

Original

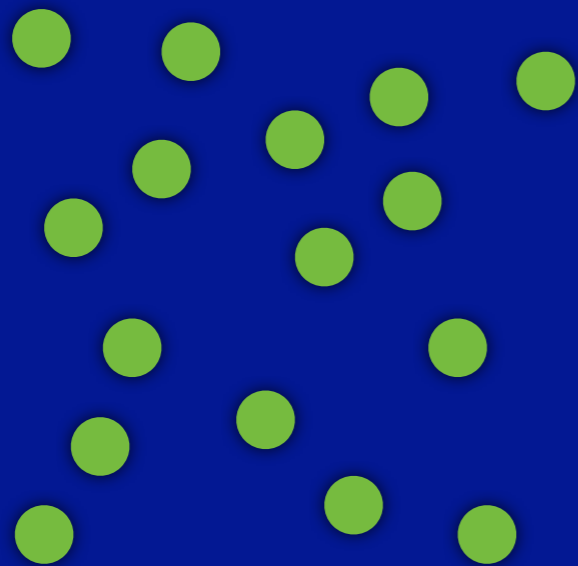
```
|  
|  
|  
| for (int i = 0; i < 10; i++)  
10  
10  
10  
10  
10  
|  
|
```

Mutant

```
|  
|  
| for (int i = 0; i < 0; i++)  
0  
0  
0  
0  
0  
0  
|  
|
```

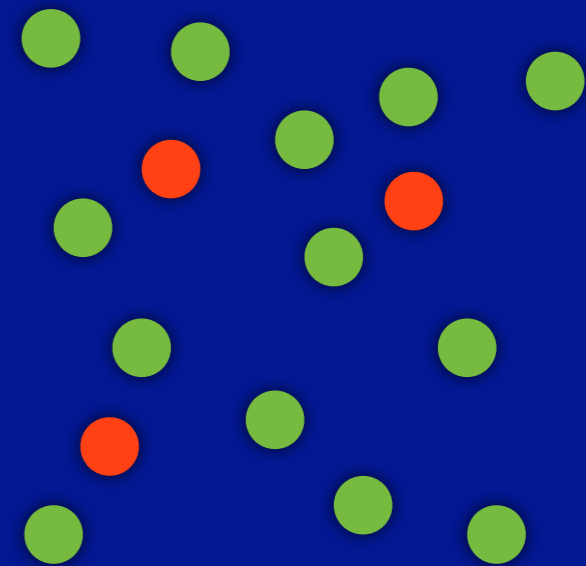
Coverage Impact

Original



- Class without Change
- Class with Change

Mutant



Coverage Impact

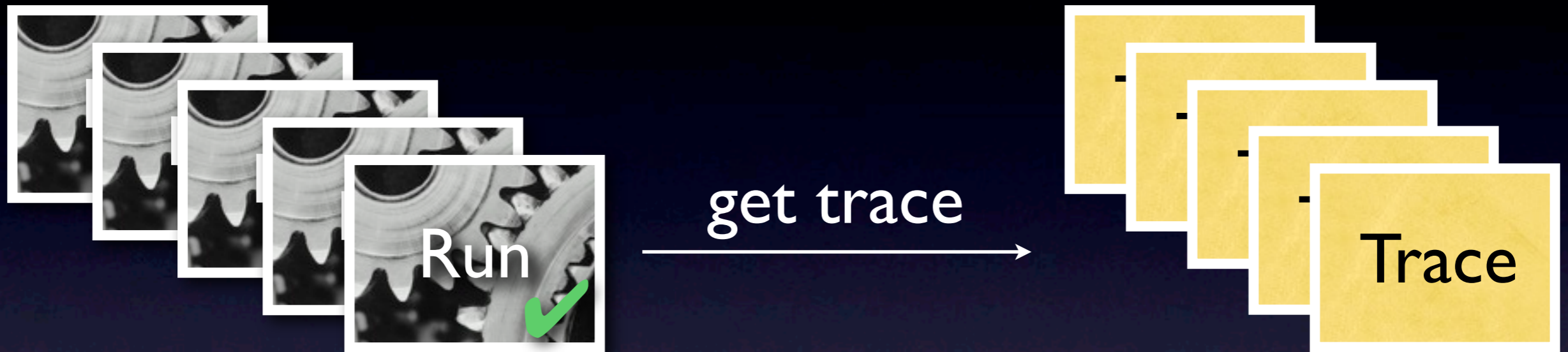
Original

- Class without Change
- Class with Change

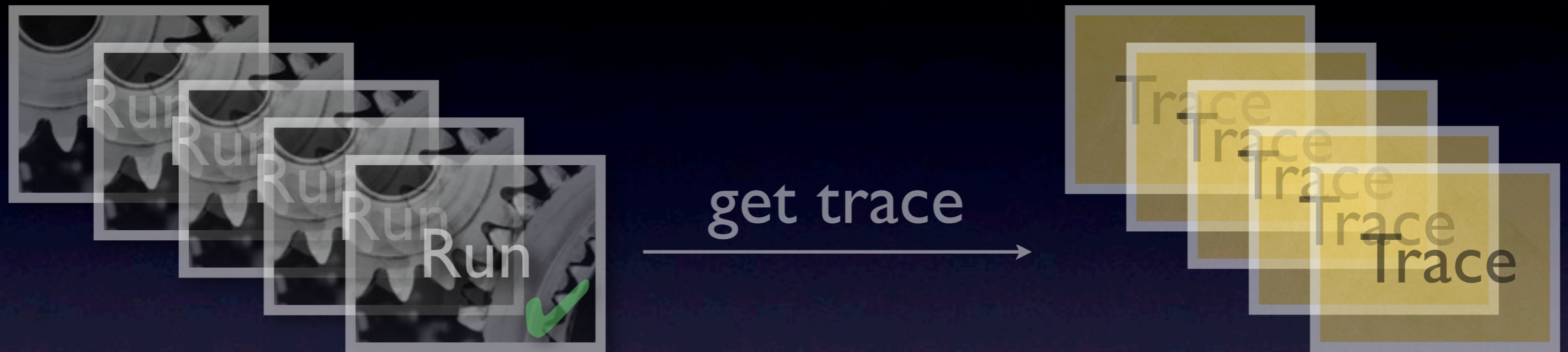
Mutant

Impact = # of Classes with Coverage Changes

Tracing Programs



Tracing Programs



Original

compare traces

Mutant

Quality of Impact

- Using the same 20 random mutations

	impact	no impact
non-equivalent		
equivalent		
undecided		

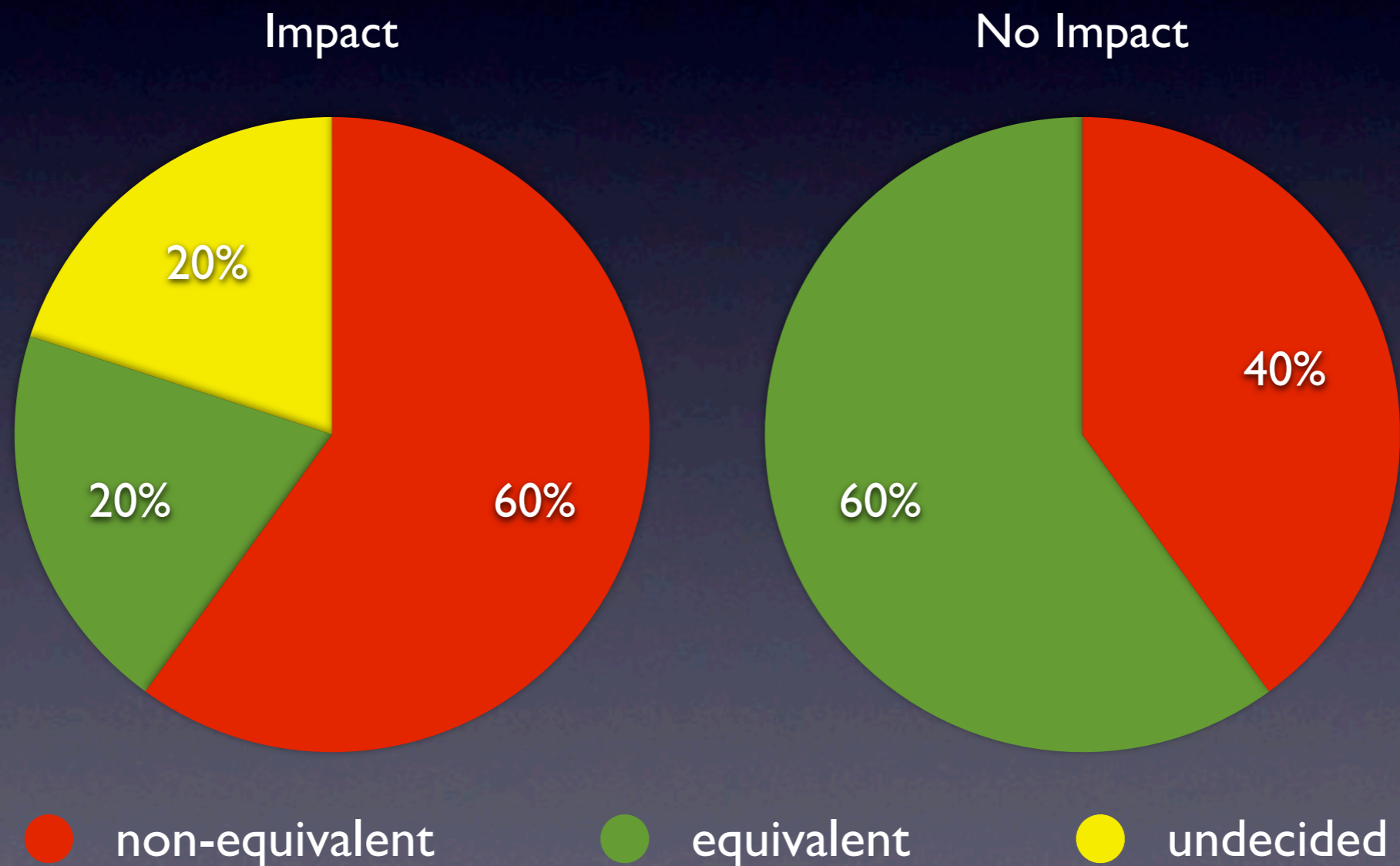
Quality of Impact

- Using the same 20 random mutations

	impact	no impact
non-equivalent	6	4
equivalent	2	6
undecided	2	0

Quality of Impact

- Using the same 20 random mutations



Ranking along Impact

- Hypothesis:
 - Higher Impact means more likely non-equivalent.
- Experiment:
 - 20 mutations with the *most impact*.
 - 20 mutations with the *least impact*.

Ranking Example

IMPACT	MUTATION
10	A
2	B
0	C
3	D
7	E
0	F
5	G

Ranking Example

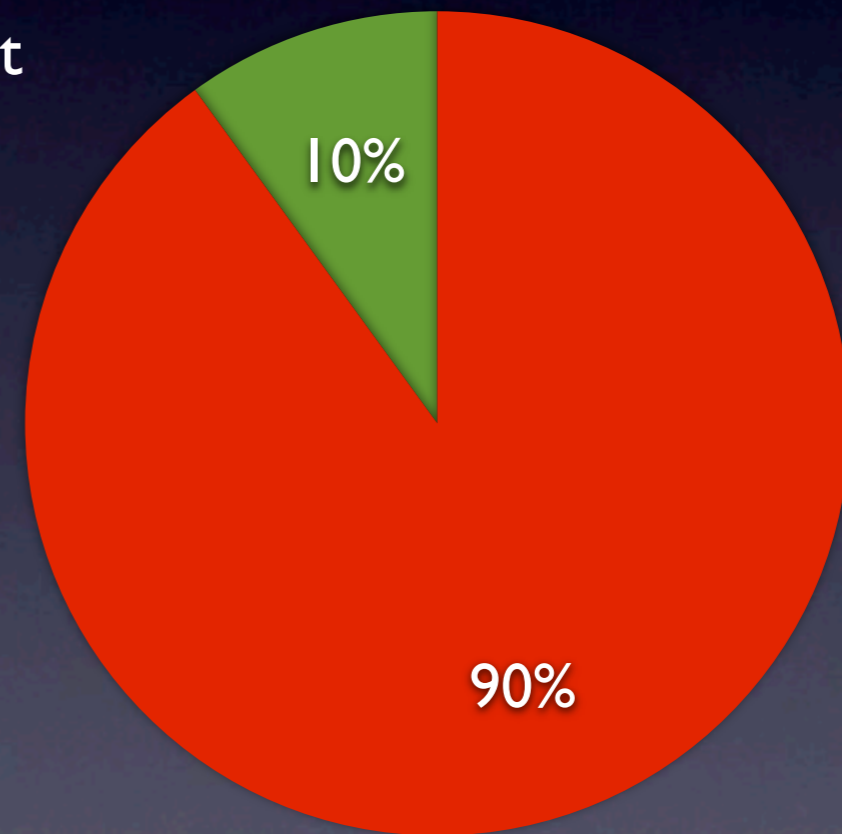
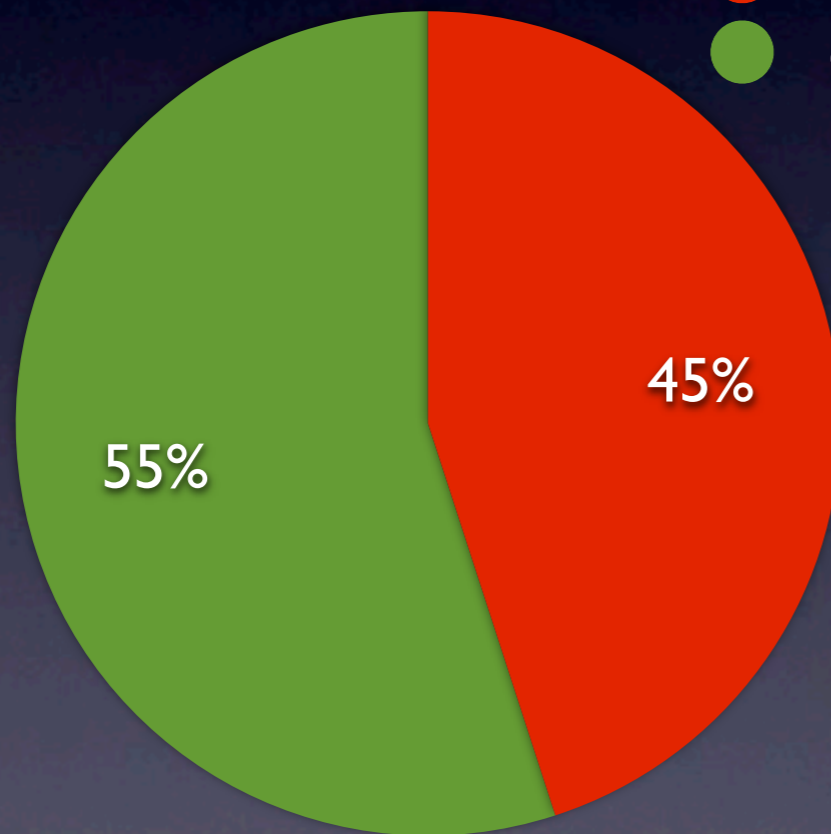
IMPACT	MUTATION
10	A
7	E
5	G
3	D
2	B
0	C
0	F

Ranking Results

BOTTOM 20

TOP 20

● non-equivalent
● equivalent



Ranking along Impact



String Alteration

```
// org.jaxen.XPathSyntaxException, Line 140
public String getPositionMarker() {
    StringBuffer buf = new StringBuffer();
    int pos = getPosition();
    for (int i = 0; i < pos; i++) {
        buf.append(" ");
    }
    buf.append("^");
    return buf.toString();
}
```

String Alteration

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Return Values

```
// org.jaxen.function.CeilingFunction, Line 129
public static Double evaluate(Object obj, Navigator nav) {
    Double value = NumberFunction.evaluate(obj, nav);

    return new Double( Math.ceil( value.doubleValue() ) );
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Return Values

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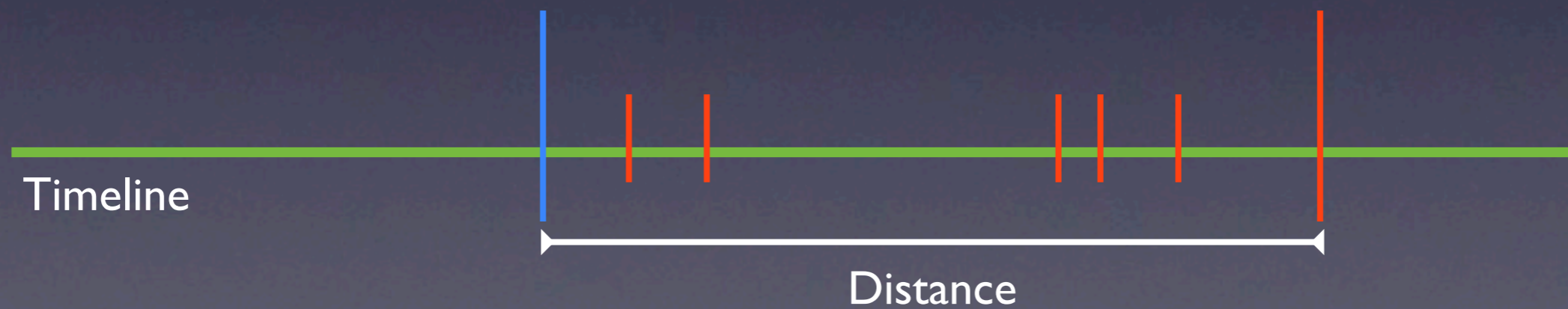
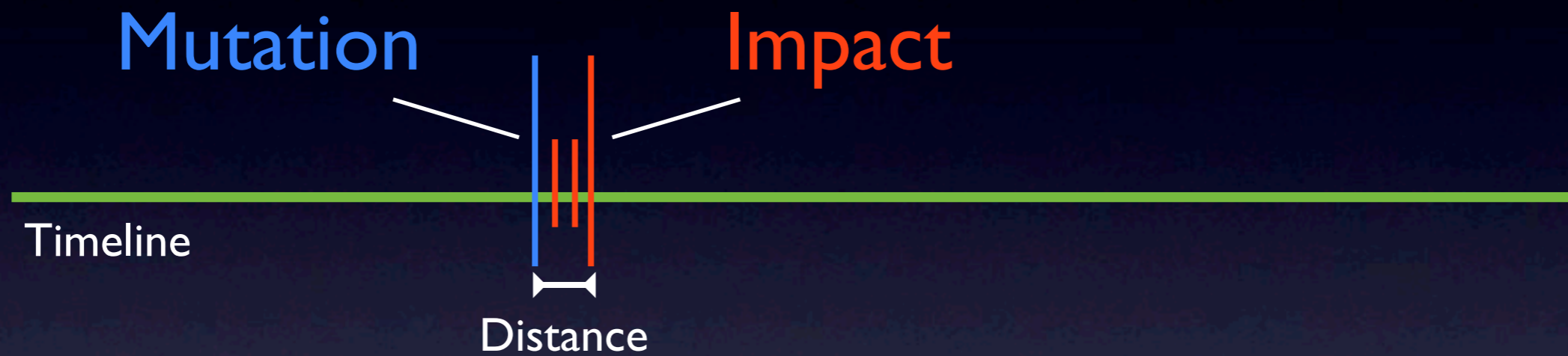
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0

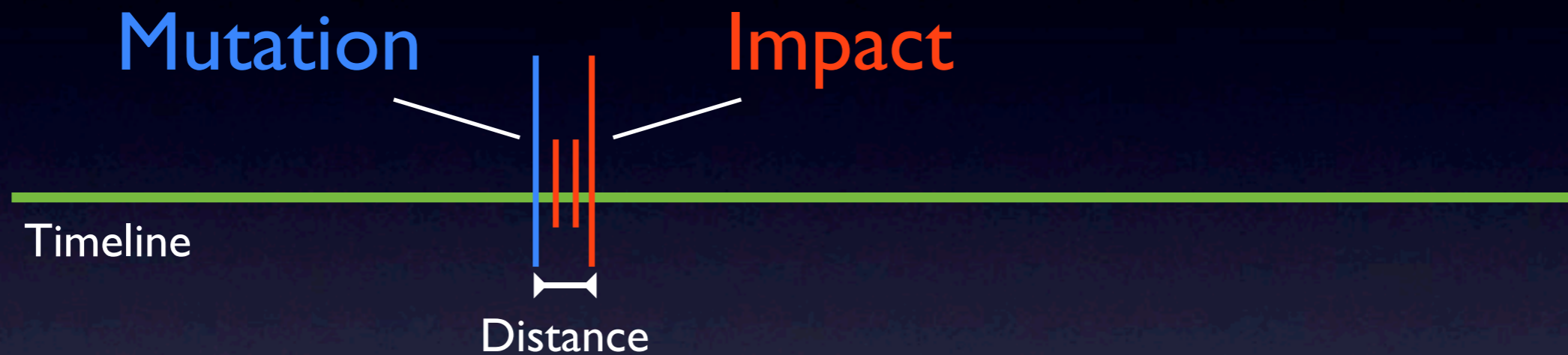
Future Work

- Find other impact measures:
 - Trace *method return values* and *invariants*
see our paper at ISSTA 2009
 - Count *methods* instead of classes
 - Use some *distance measure*
- Analyze more software projects

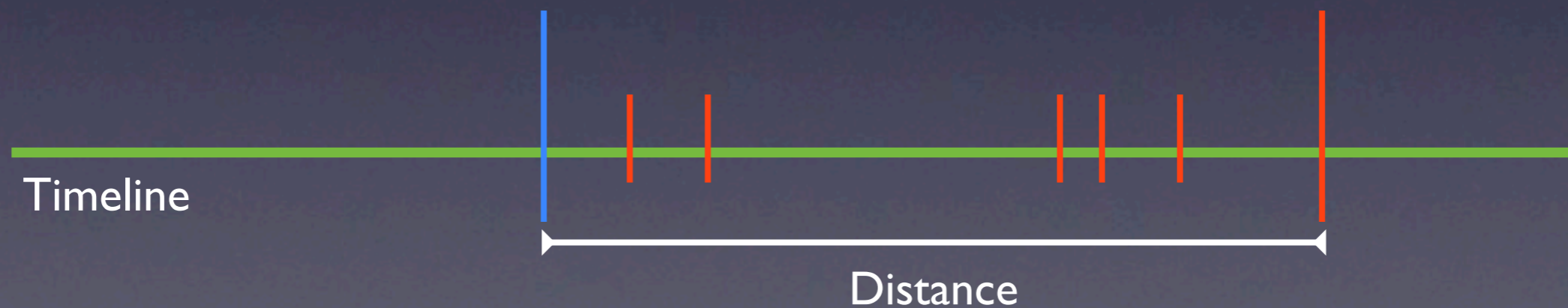
Idea: Distance Measure



Idea: Distance Measure



Early impact \Rightarrow equivalent



Late impact \Rightarrow non-equivalent

A person is swinging on a swing set. The background is dark and textured, possibly a wall or a large tree. The swing is in motion, and the person's hands are visible on the chains.

Conclusion

<http://www.st.cs.uni-saarland.de/mutation>

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Manual Classification



Manual Classification is not efficient!

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Random Mutants

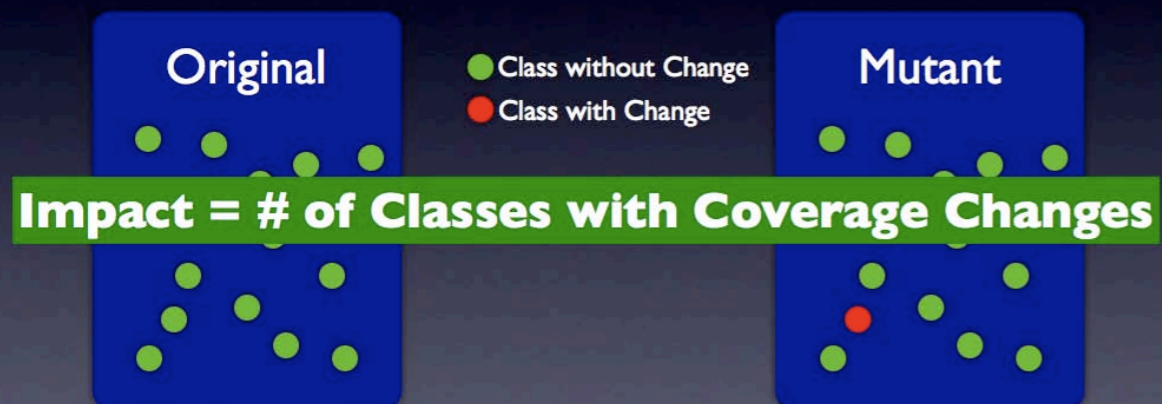
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Coverage Impact



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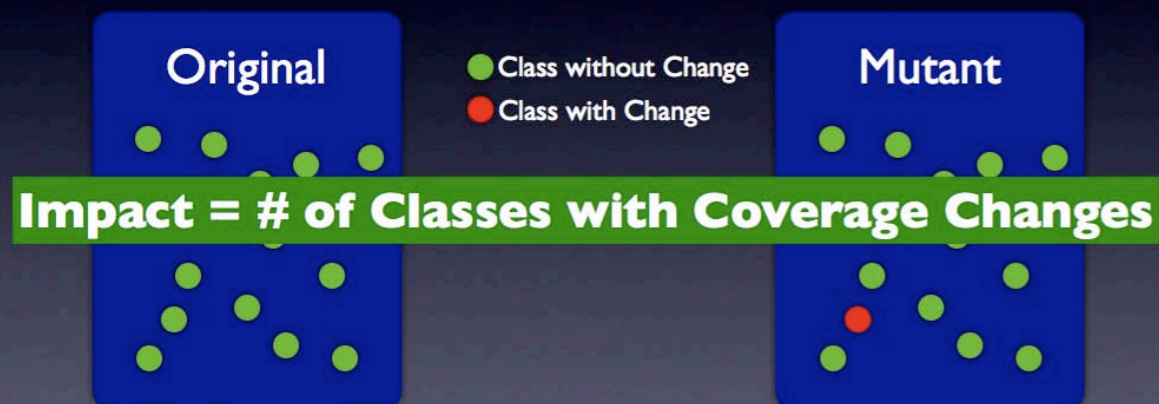
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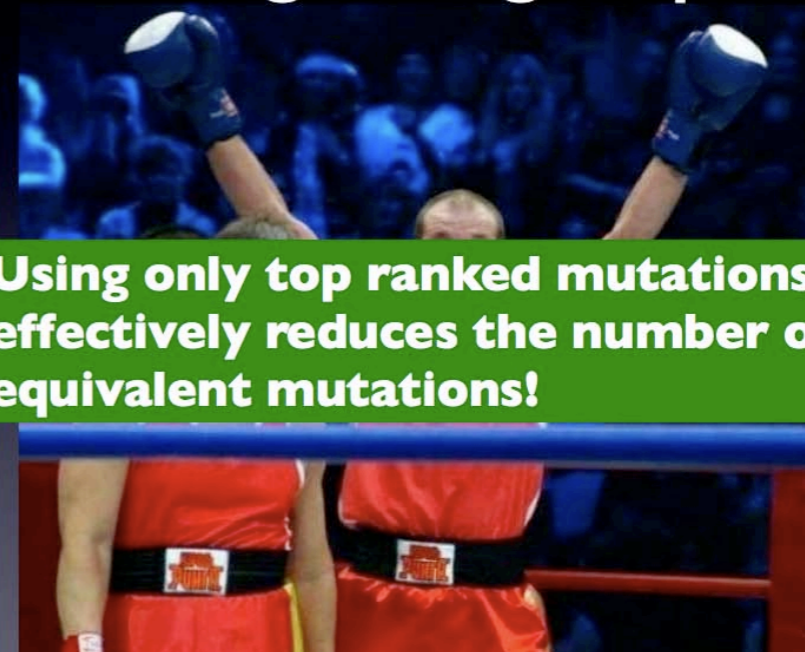


Manual Classification is not efficient!

Coverage Impact



Ranking along Impact



✓ Using only top ranked mutations effectively reduces the number of equivalent mutations!

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Manual Classification



Manual classification is not efficient!

Future Work

- Find other impact measures.
- Tracing of method return values.
- Based on methods instead of classes.
- Using some distance measure.
- Analyze more software projects.

Coverage



Impact = # of Classes with Coverage Changes

Along Impact

✓ Using only top ranked mutations effectively reduces the number of equivalent mutations!



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