

# On the Use of Sequence Mining within Spectrum Based Fault Localisation

Gulsher Laghari and Serge Demeyer

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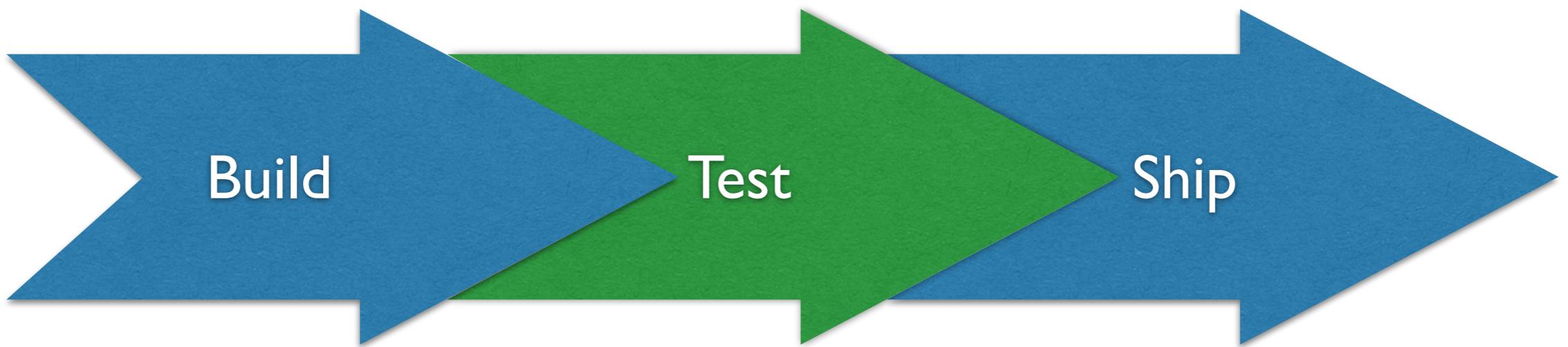


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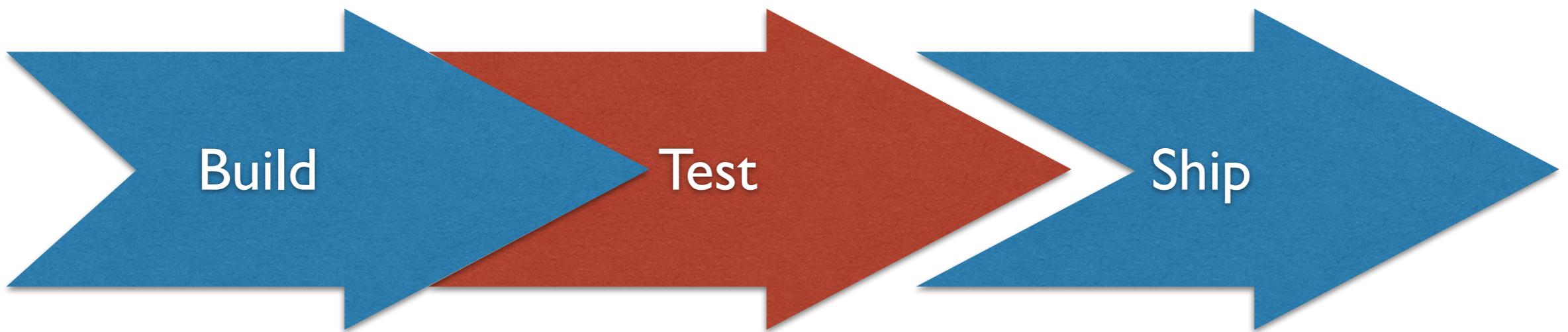
Antwerp Systems & Software Modelling  
University of Antwerp



# Overview



# Overview



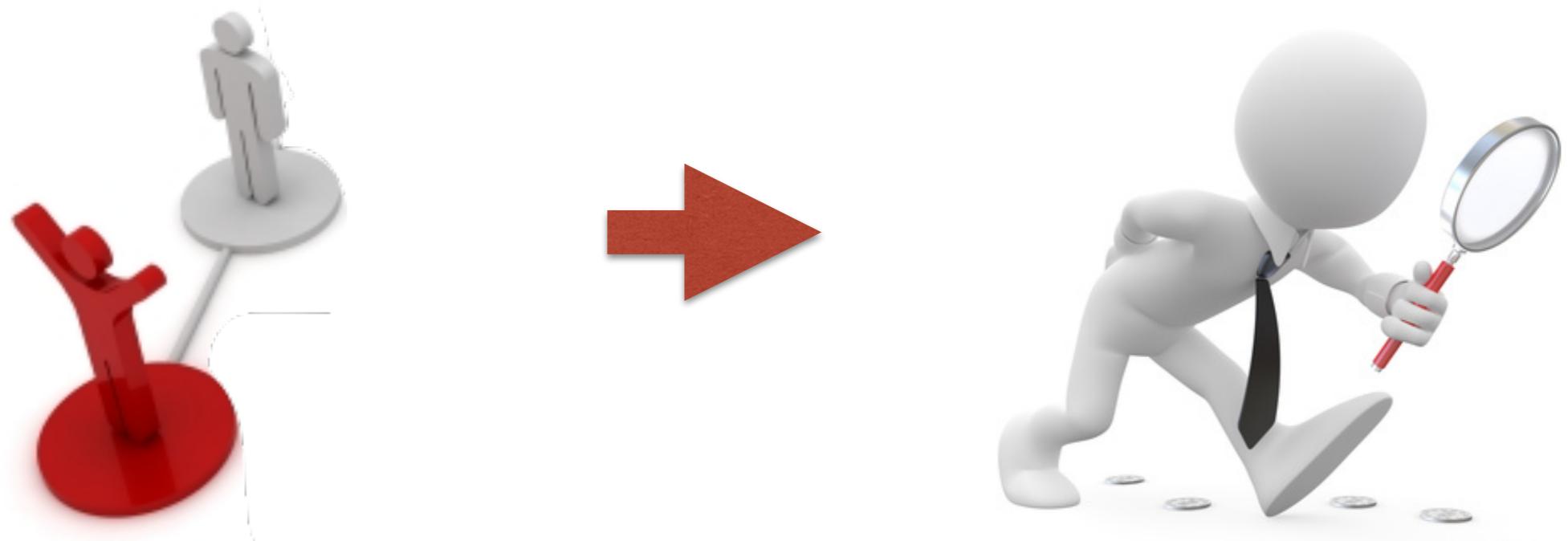
# Fault Localisation



Fault Localisation an important step in debugging process

# Fault Localisation

Test to code mapping - 1:1



# Fault Localisation

Test to code mapping - 1:N



# Fault Localisation

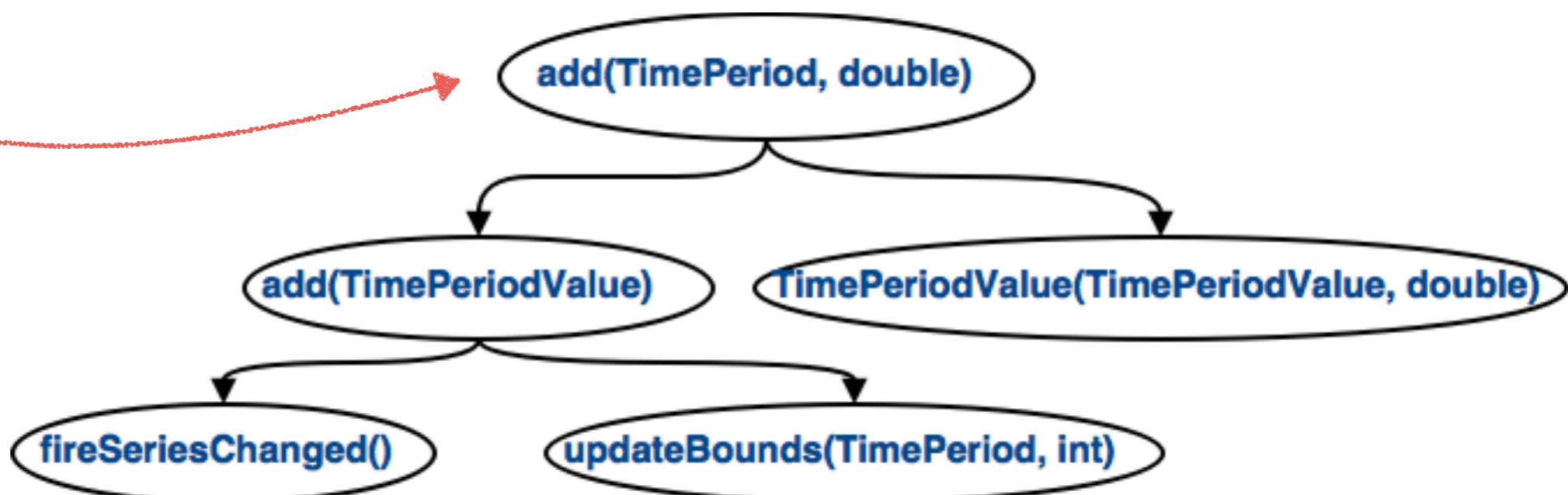
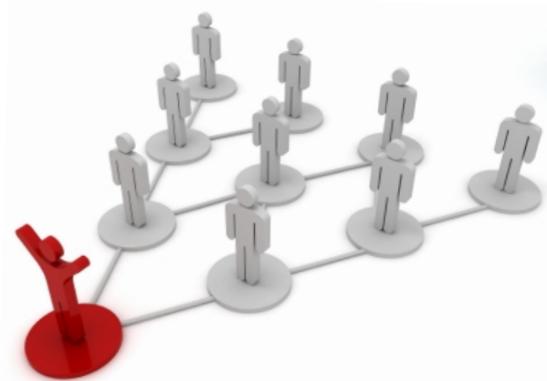
```
public void testGetMaxMiddleIndex() {  
    TimePeriodValues s = new TimePeriodValues("Test");  
    assertEquals(-1, s.getMaxMiddleIndex());  
    s.add(new SimpleTimePeriod(100L, 200L), 1.0);  
    assertEquals(0, s.getMaxMiddleIndex());  
    s.add(new SimpleTimePeriod(300L, 400L), 2.0);  
    assertEquals(1, s.getMaxMiddleIndex());  
    s.add(new SimpleTimePeriod(0L, 50L), 3.0);  
    assertEquals(1, s.getMaxMiddleIndex());  
    s.add(new SimpleTimePeriod(150L, 200L), 4.0);  
    assertEquals(1, s.getMaxMiddleIndex());  
}
```



\* Failing test in Apache Commons Math

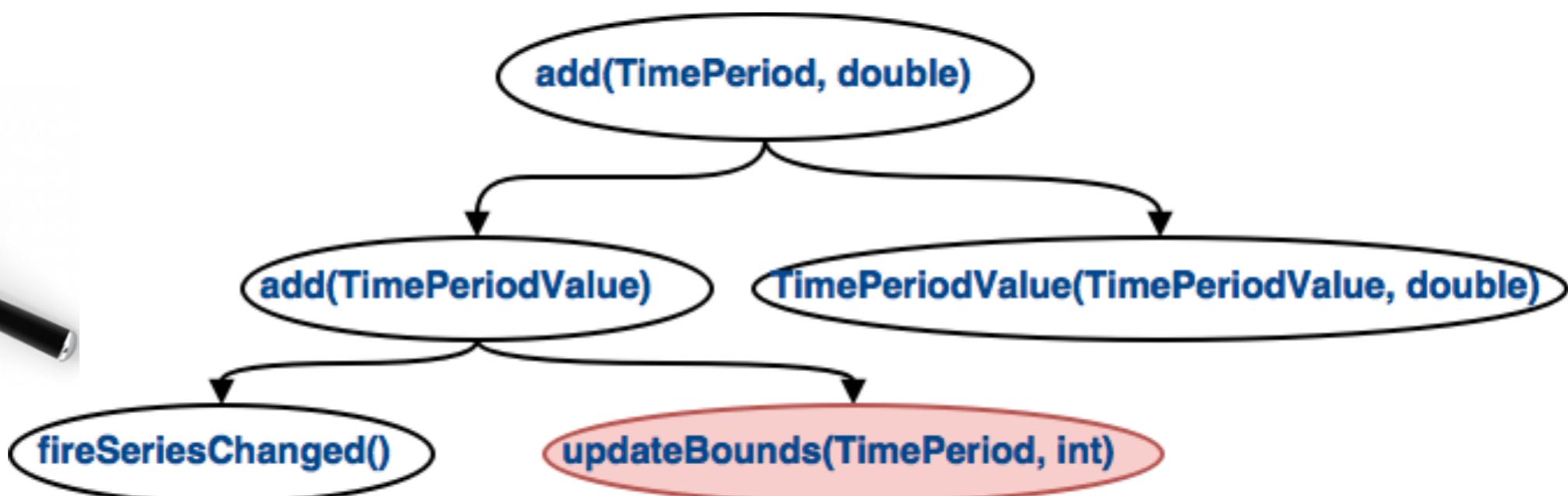
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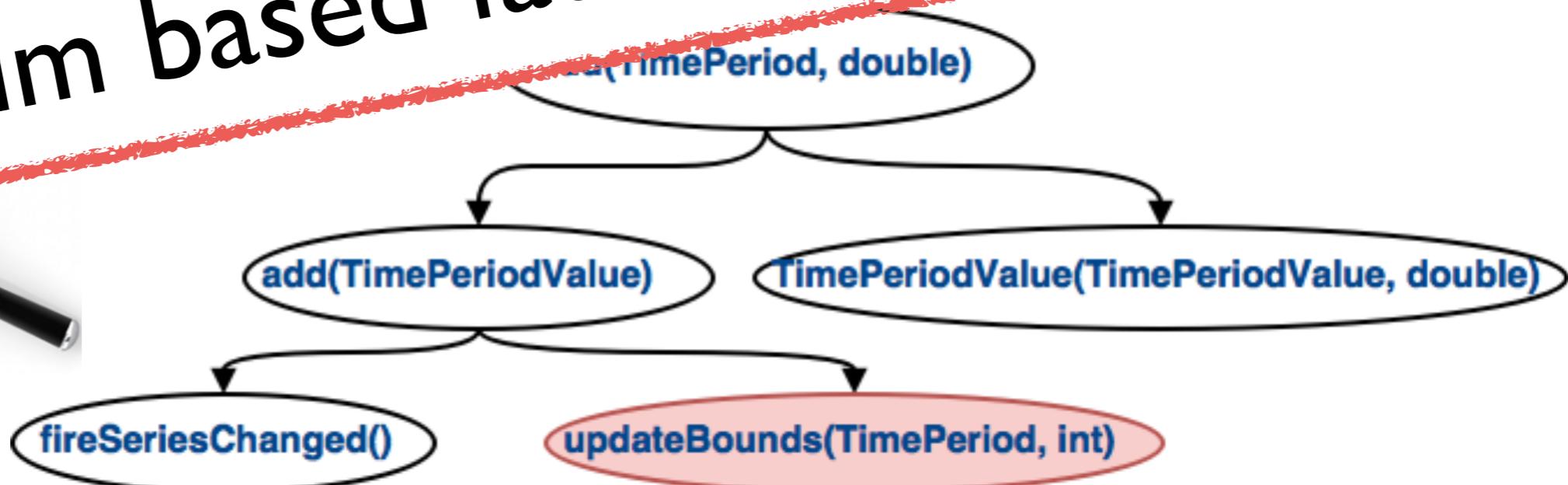


# Fault Localisation

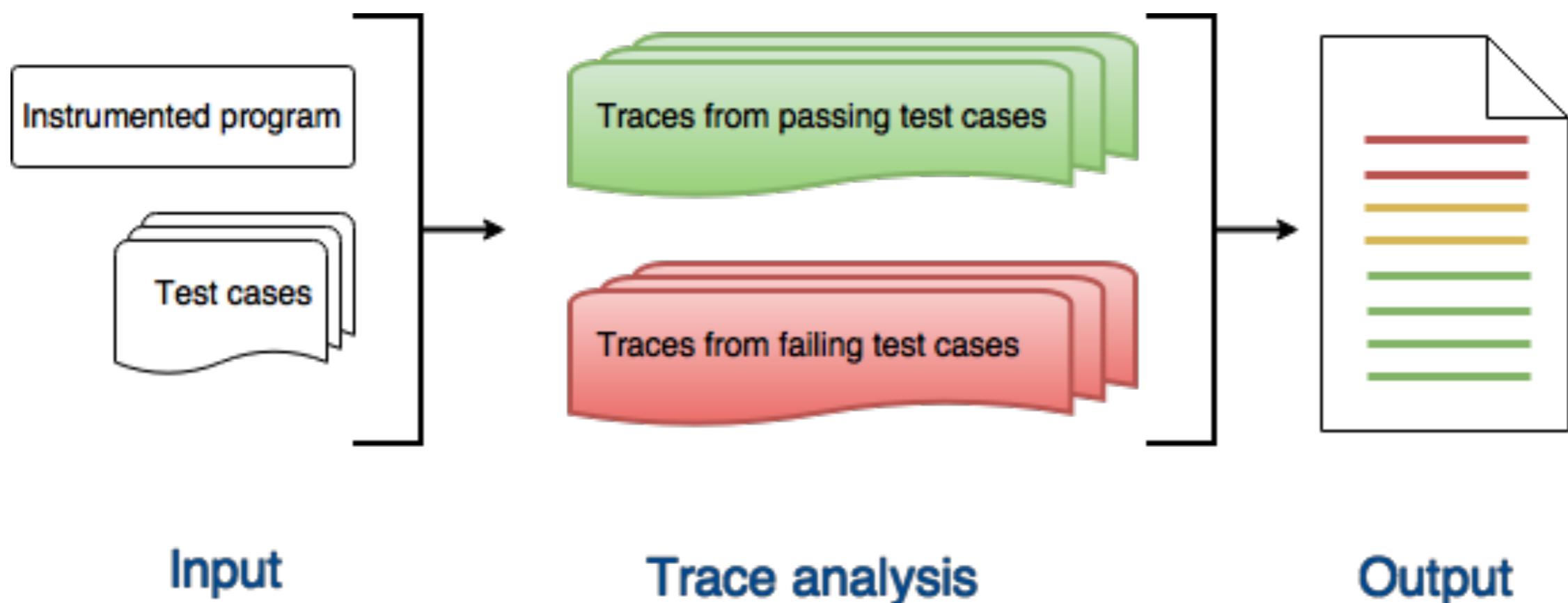
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```



Spectrum based fault localisation

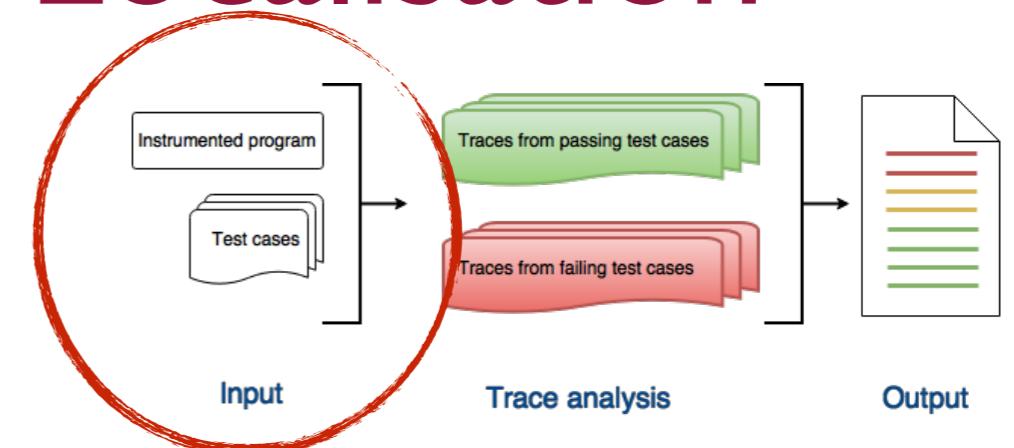


# Spectrum Based Fault Localisation

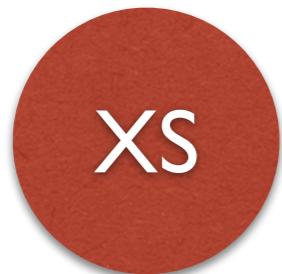


# Spectrum Based Fault Localisation

## Granularity



Statement



Block



Method

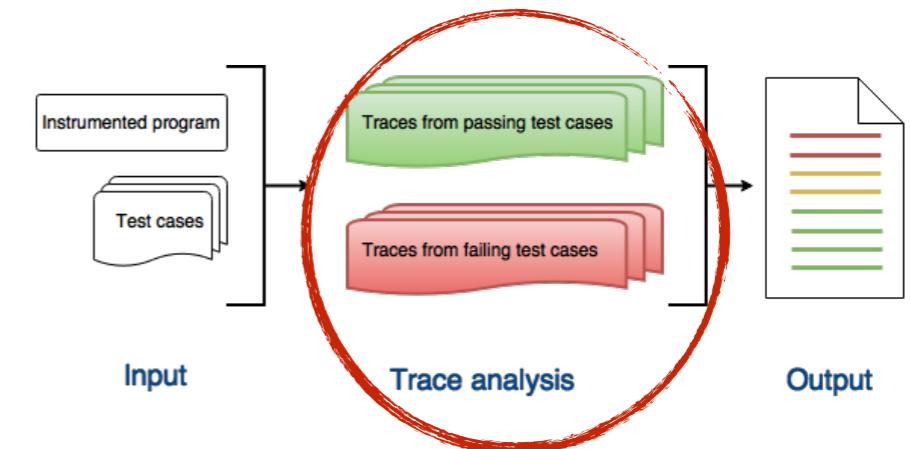


Class

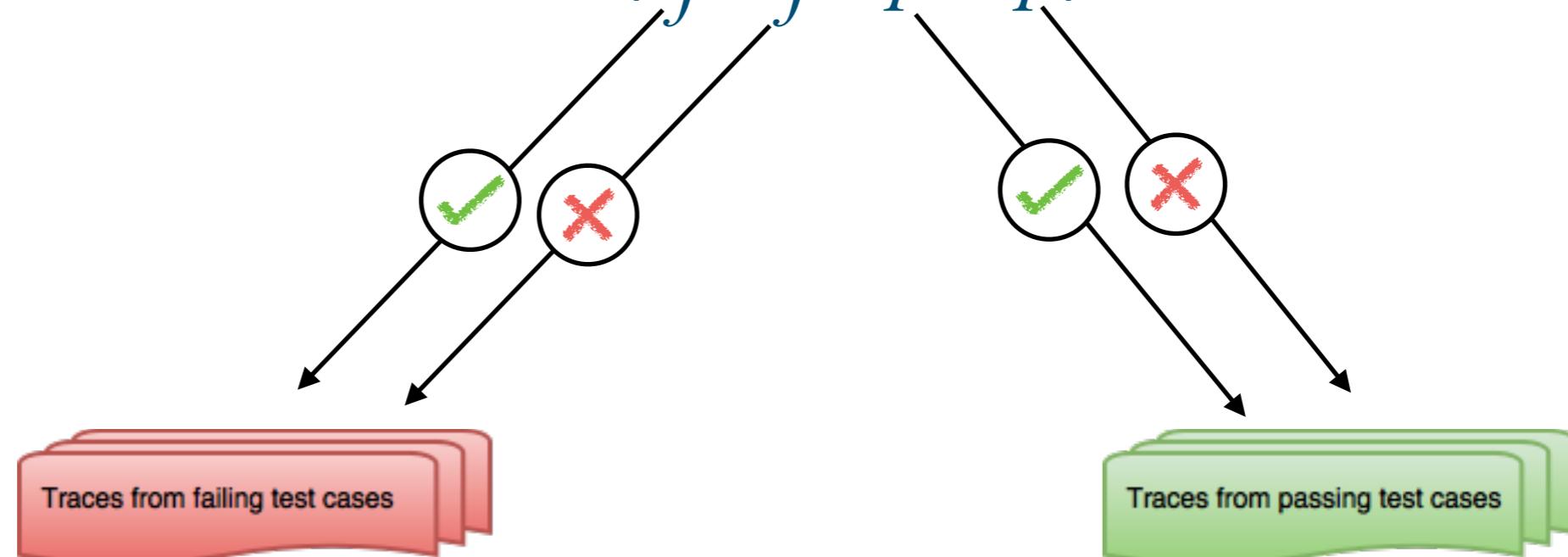


# Spectrum Based Fault Localisation

## Hit Spectrum



$$\text{Method} = (e_f, n_f, e_p, n_p)$$

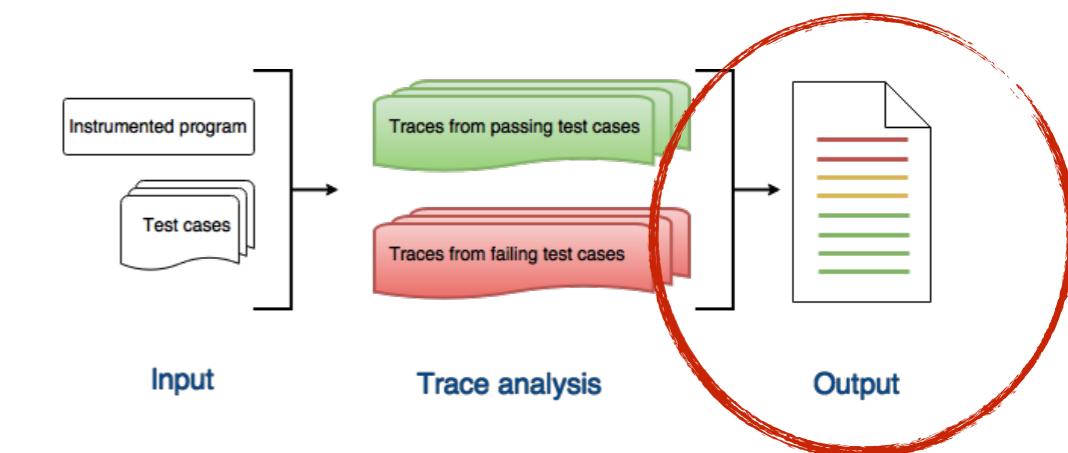


Number of traces that contain UUT

Number of traces that do not contain UUT

# Spectrum Based Fault Localisation

## Fault Locator



$$f(x)$$

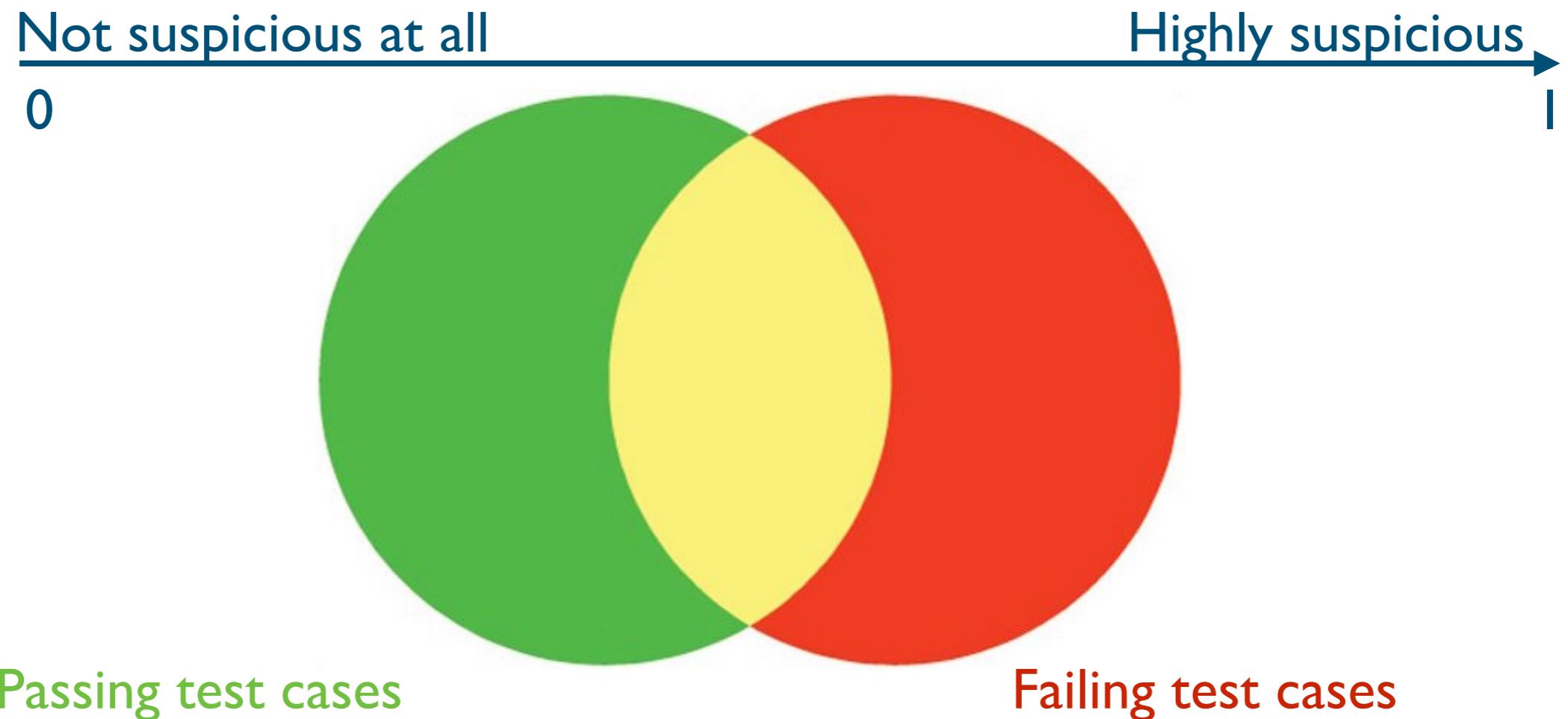
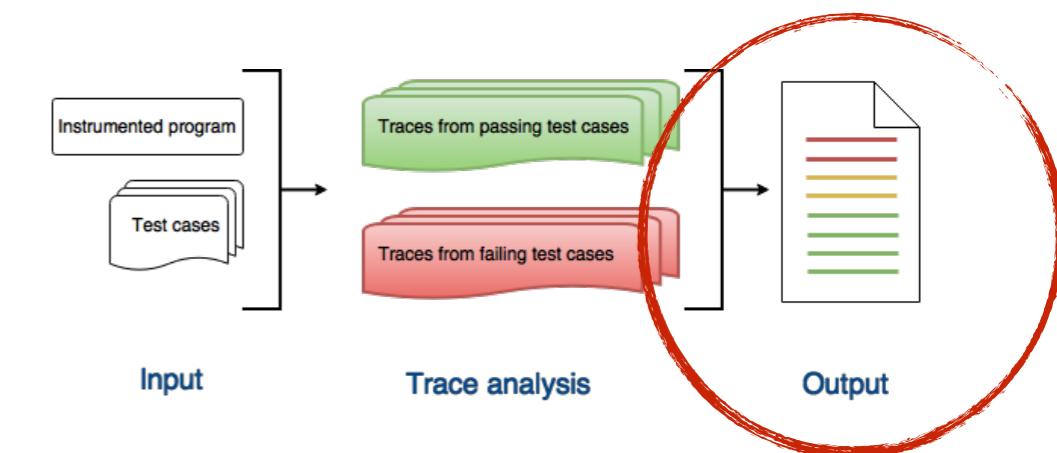
Method =  $(e_f, n_f, e_p, n_p)$

Method = Suspiciousness [0,1]

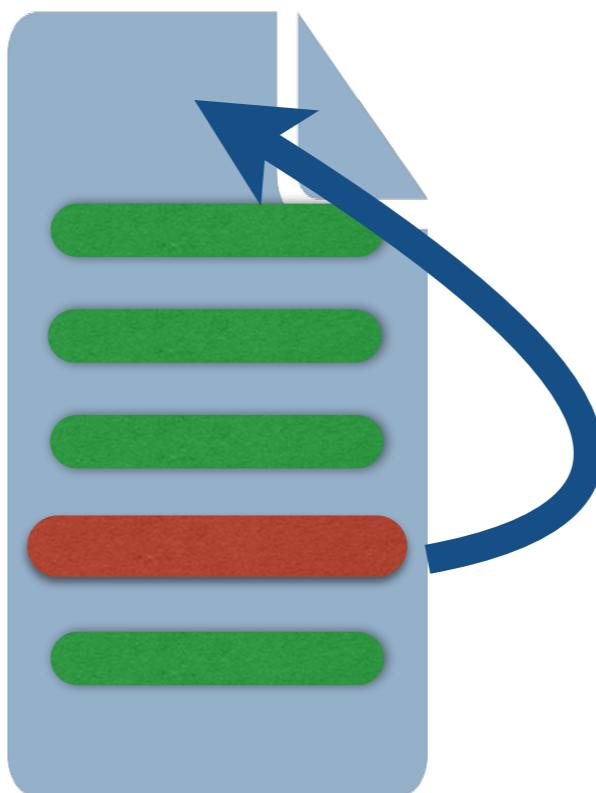
$$suspiciousness = \frac{e_f}{\sqrt{(e_f + n_f)(e_f + e_p)}}$$

# Spectrum Based Fault Localisation

## Fault Locator



# Spectrum Based Fault Localisation



# Spectrum Based Fault Localisation



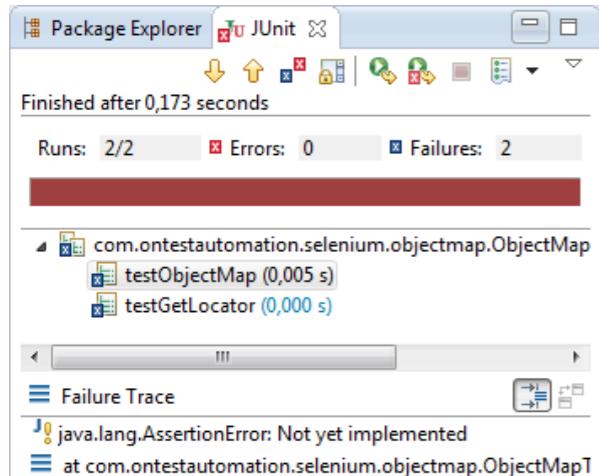
# Raw Spectrum Analysis

Missing

```
method1( ) {  
    methodA( )  
    methodB( )  
    if (condition) {  
        return  
    }  
    methodC( )  
}
```

# Raw Spectrum Analysis

## Missing

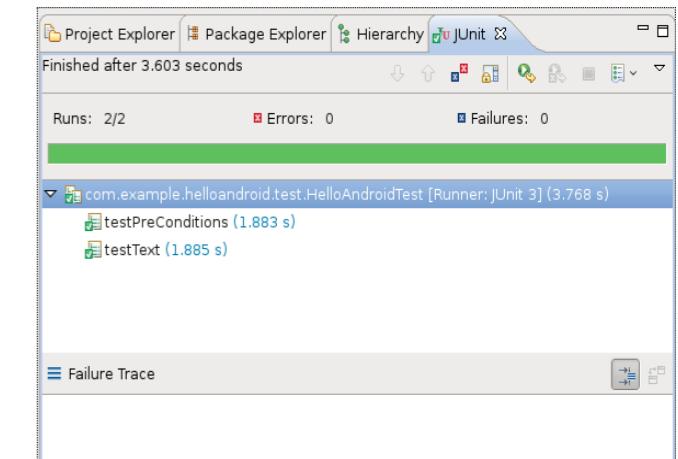
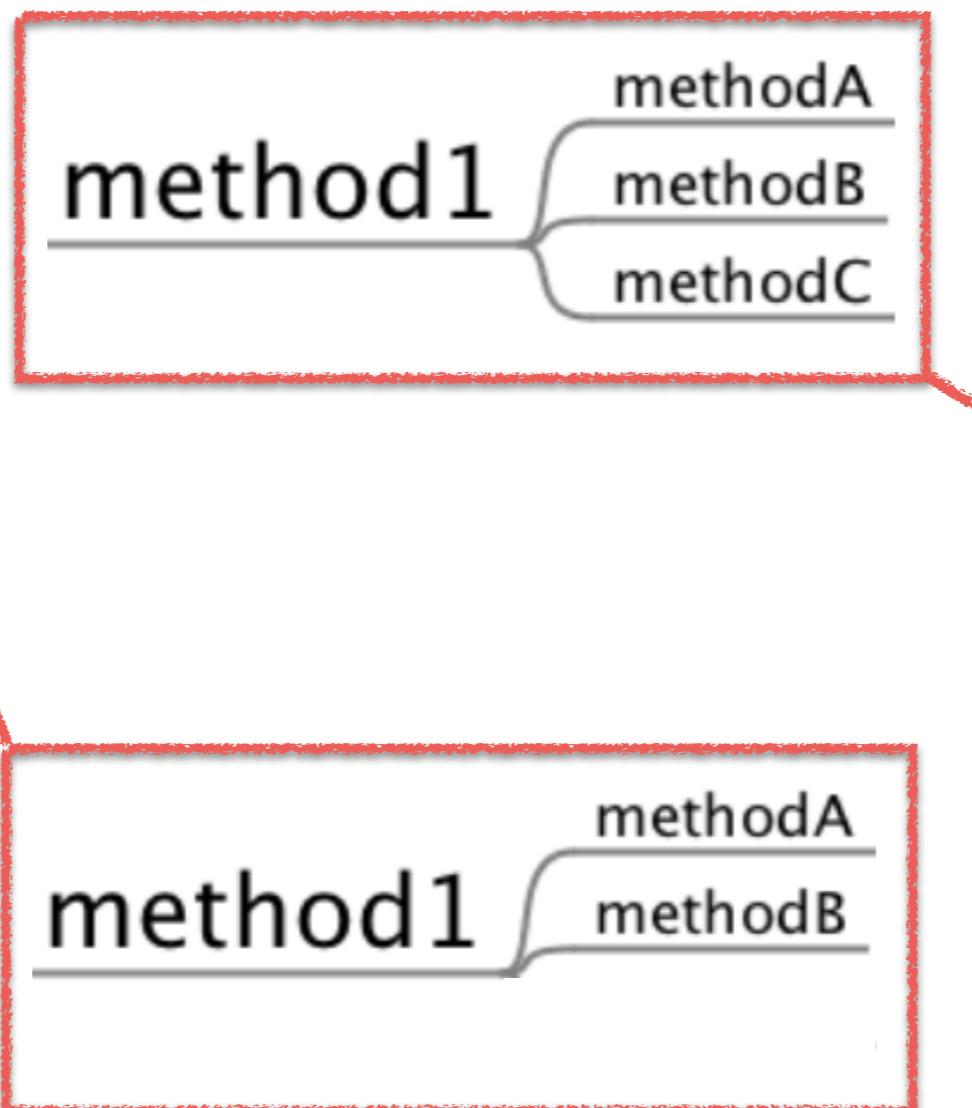
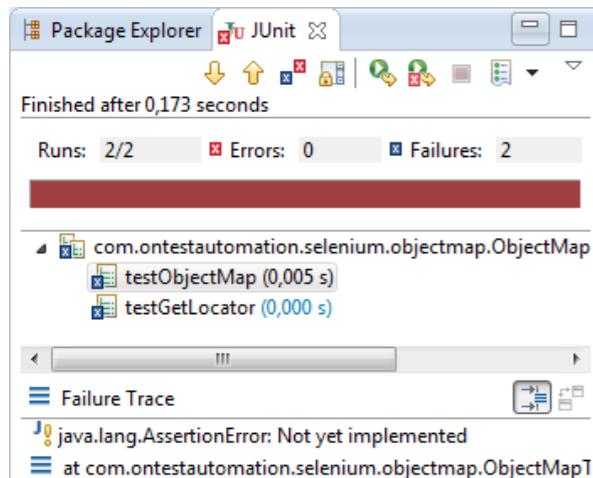


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    methodB()  
    if(condition) {  
        return  
    }  
    methodC()  
}
```



# Raw Spectrum Analysis

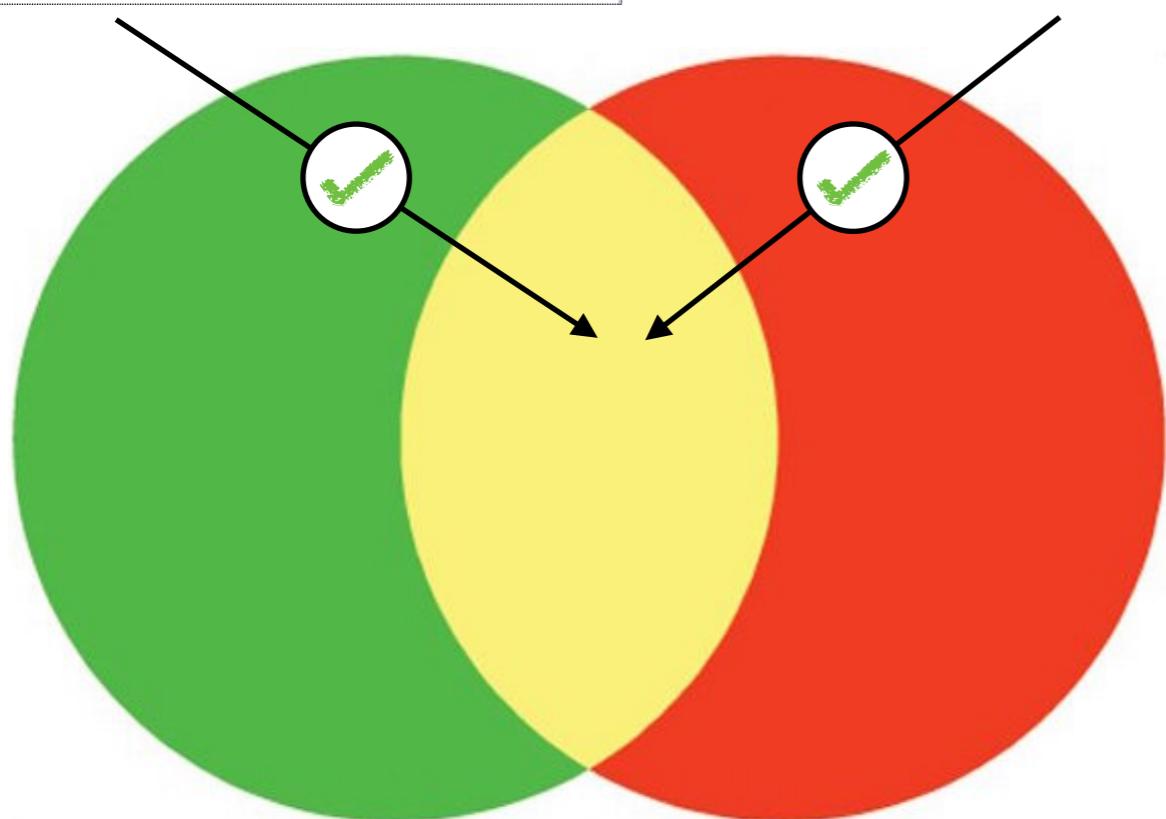
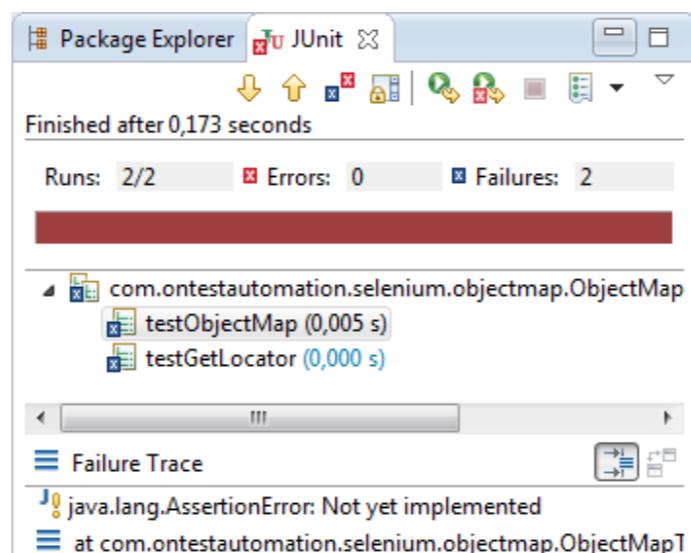
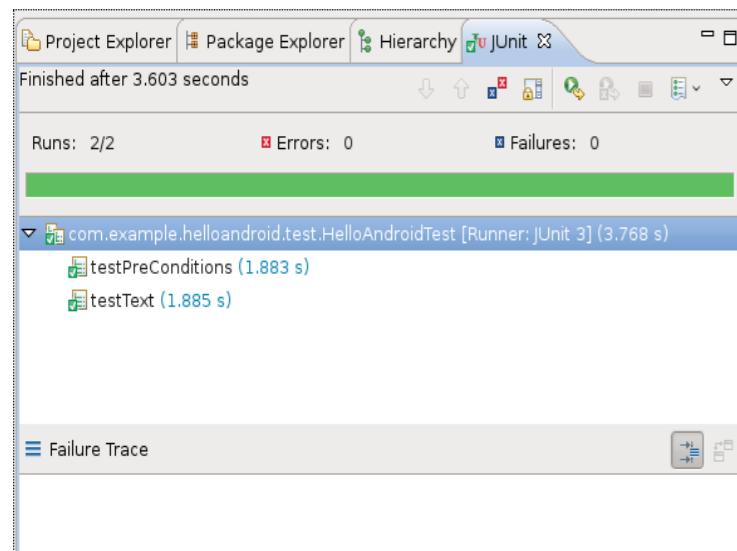
## Missing



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method1() {
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# Raw Spectrum Analysis

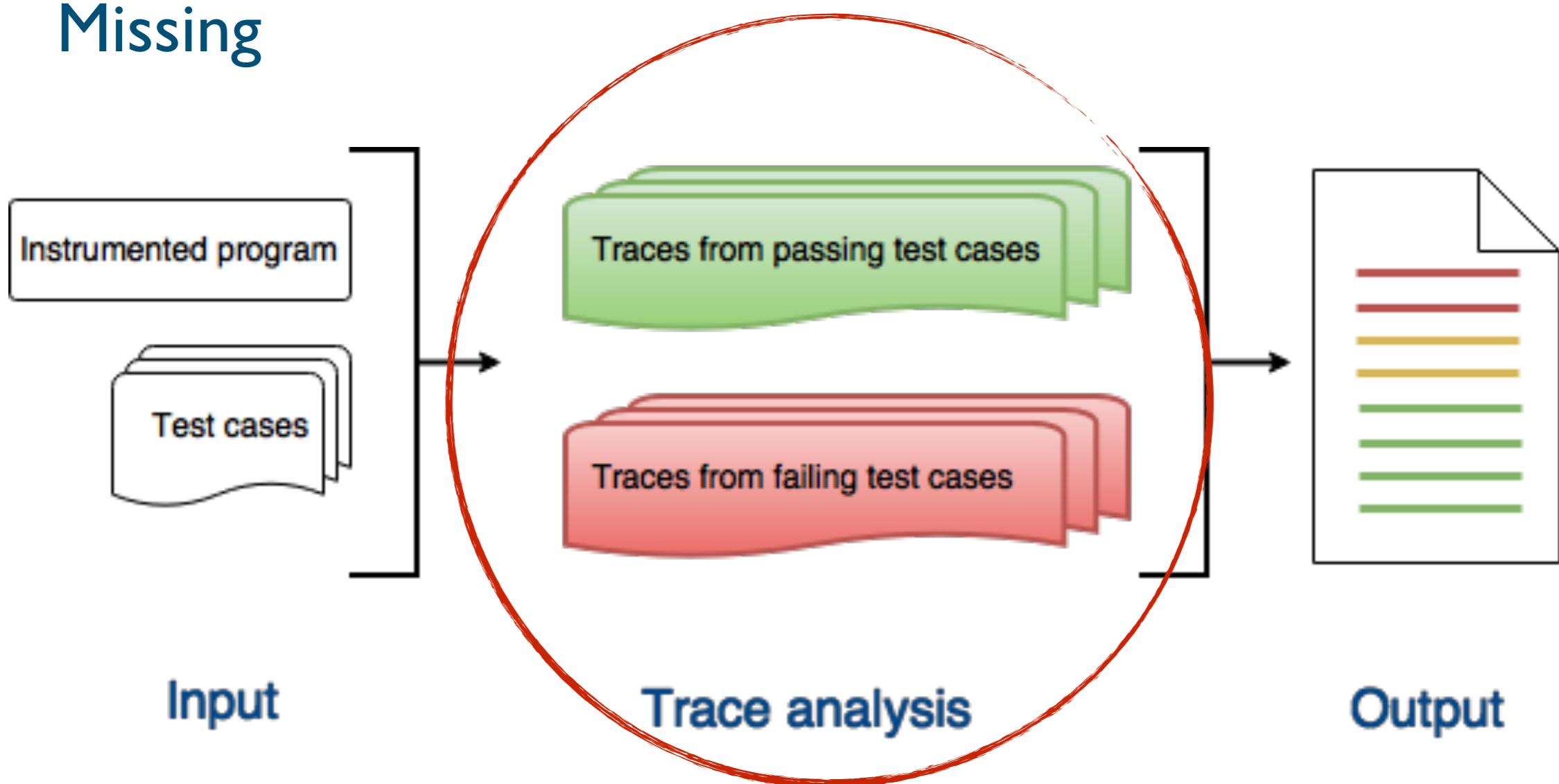
## Missing



```
method1 () {  
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    methodB ()  
    if (condition) {  
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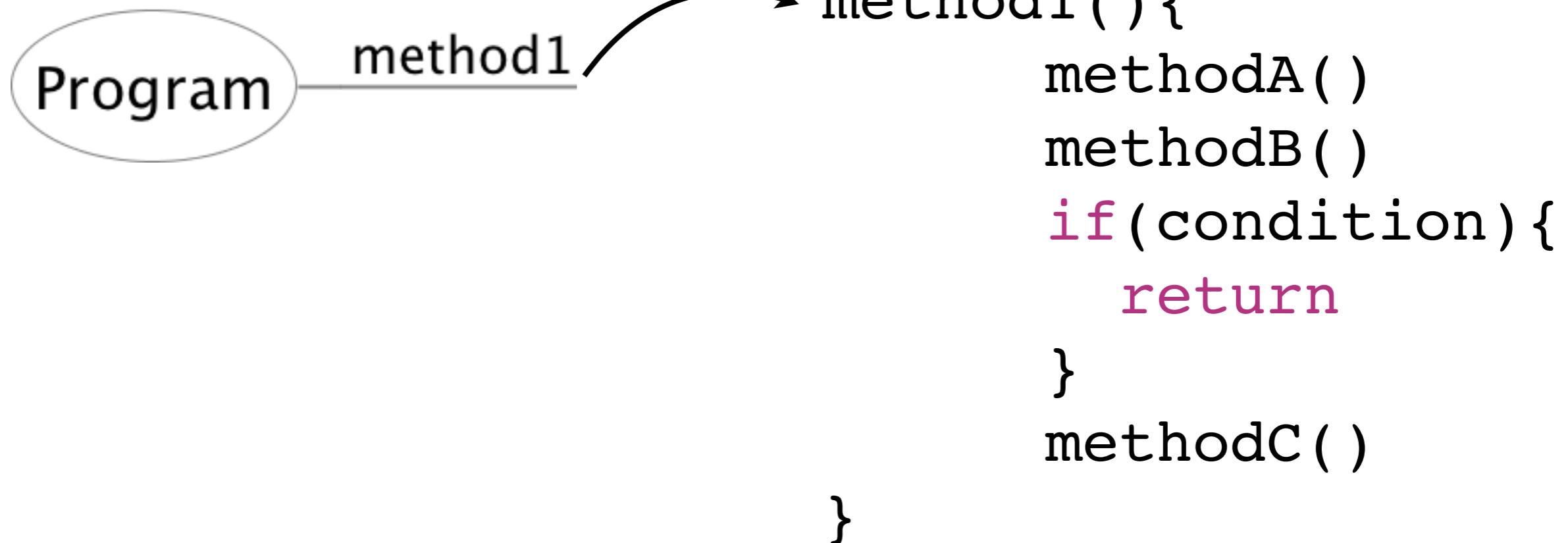
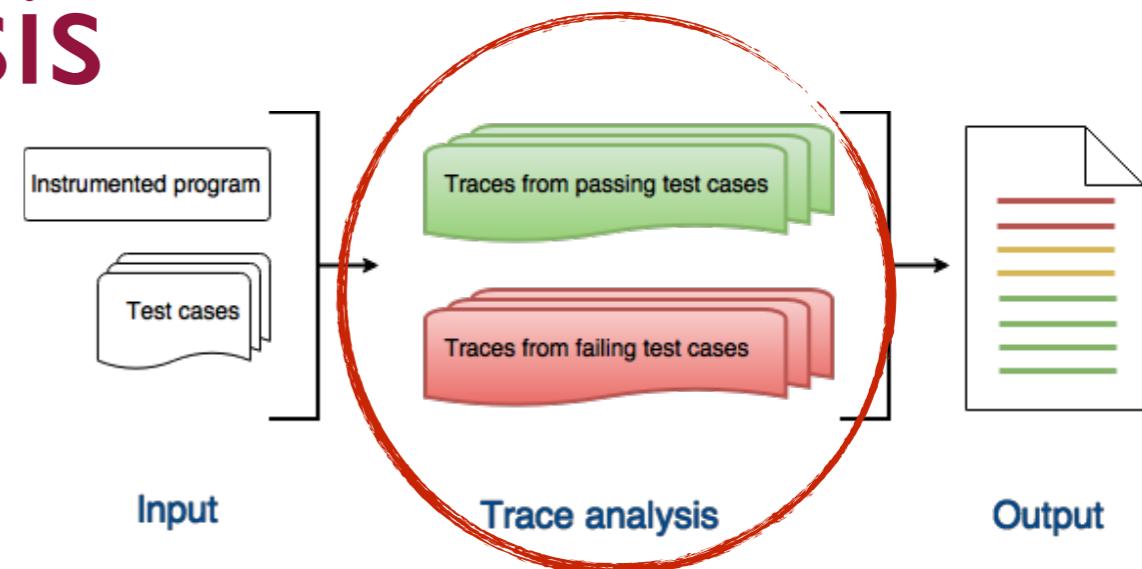
# Raw Spectrum Analysis

Missing



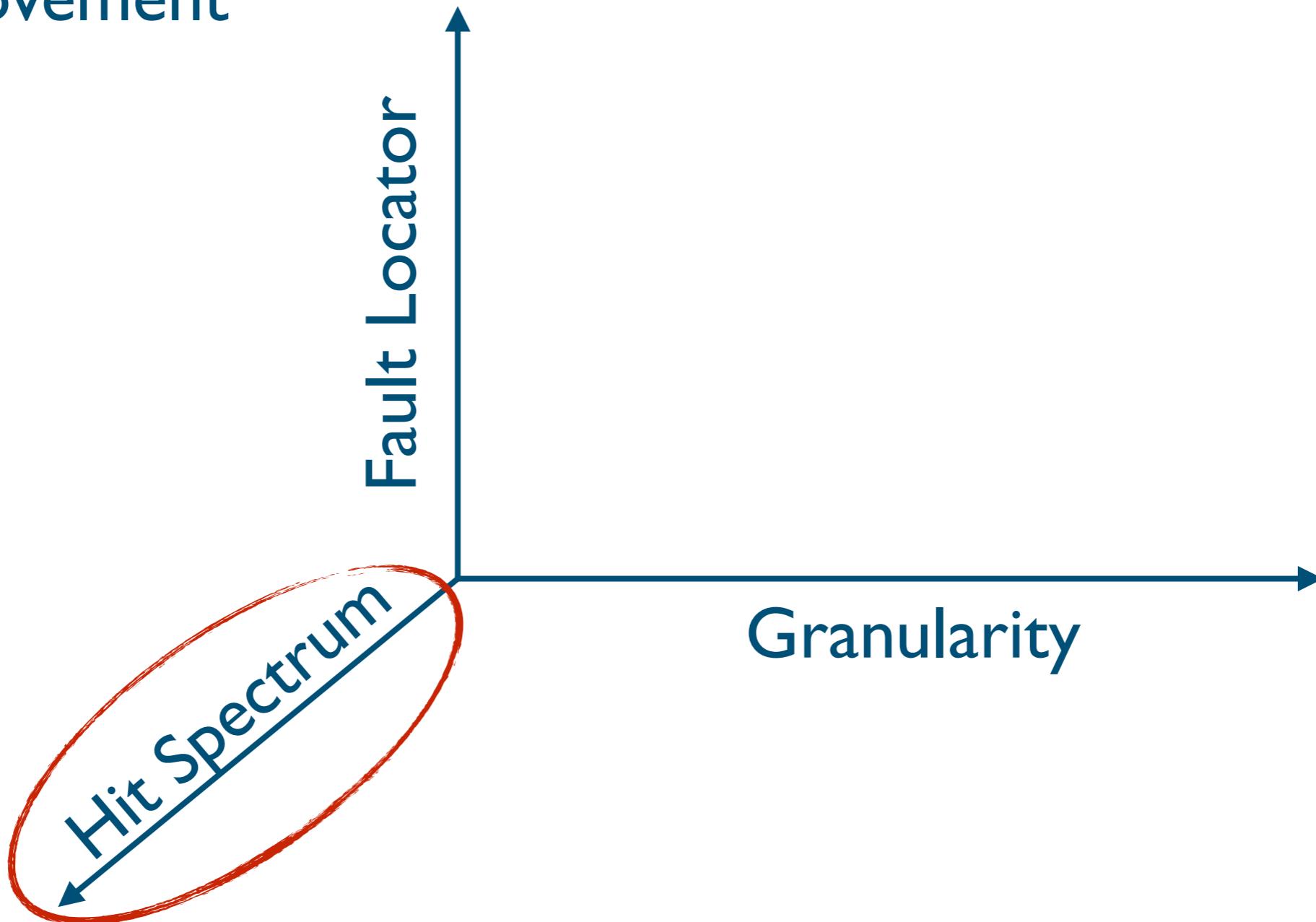
# Raw Spectrum Analysis

Missing



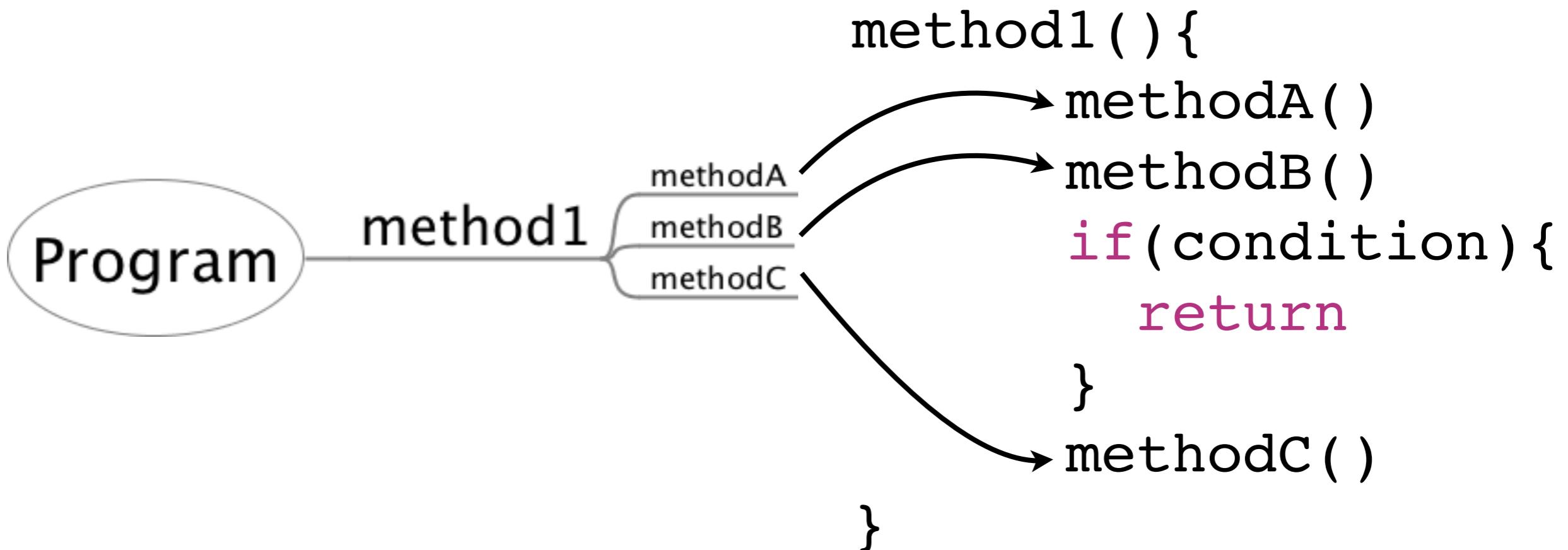
# Raw Spectrum Analysis

Improvement



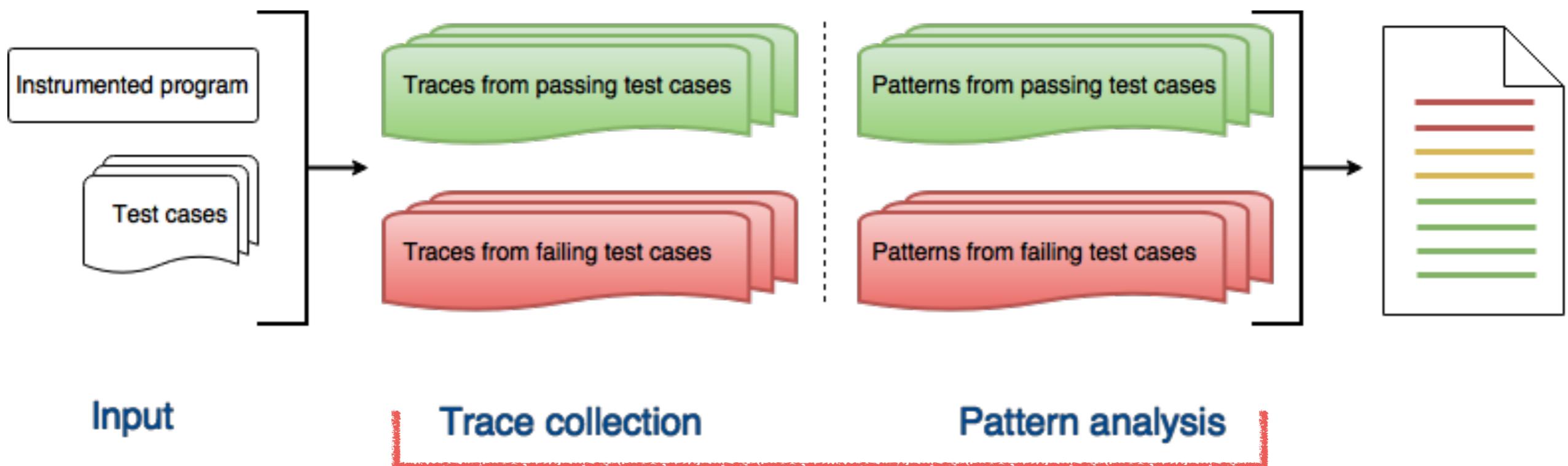
# Raw Spectrum Analysis

## Improvement



# Raw Spectrum Analysis

## Improvement

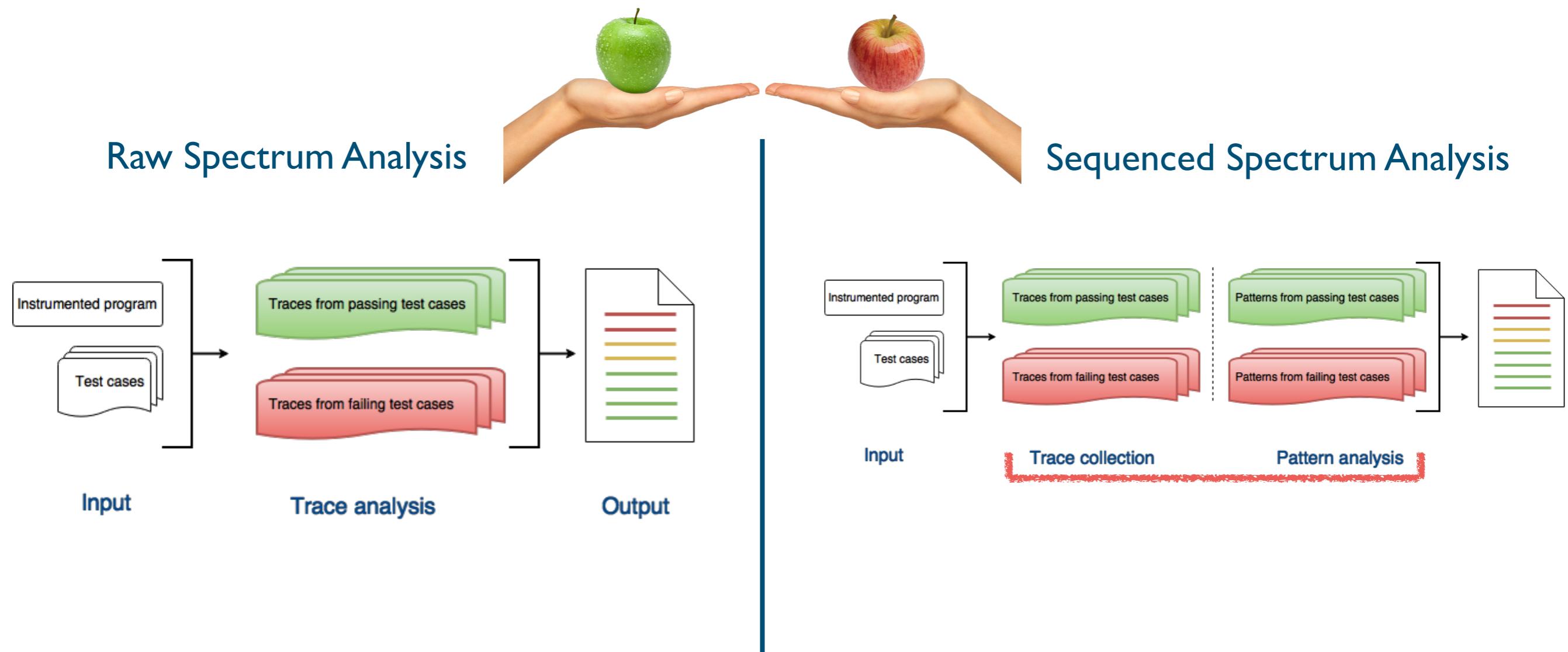


# Sequenced Spectrum Analysis

## Improvement



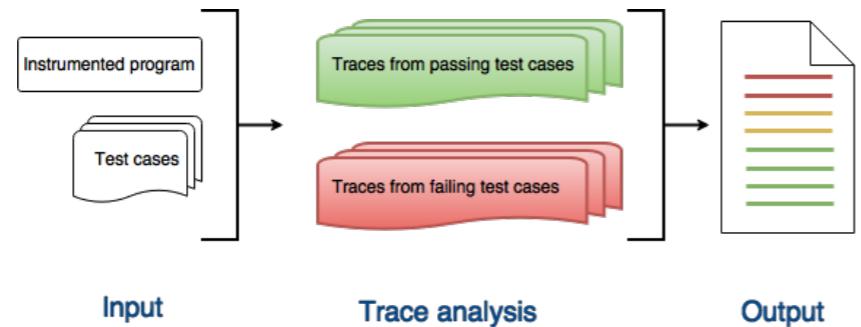
# Case Study



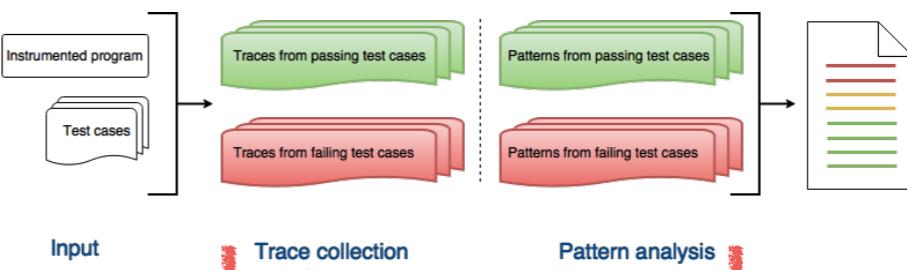
# Case Study



## Raw Spectrum Analysis



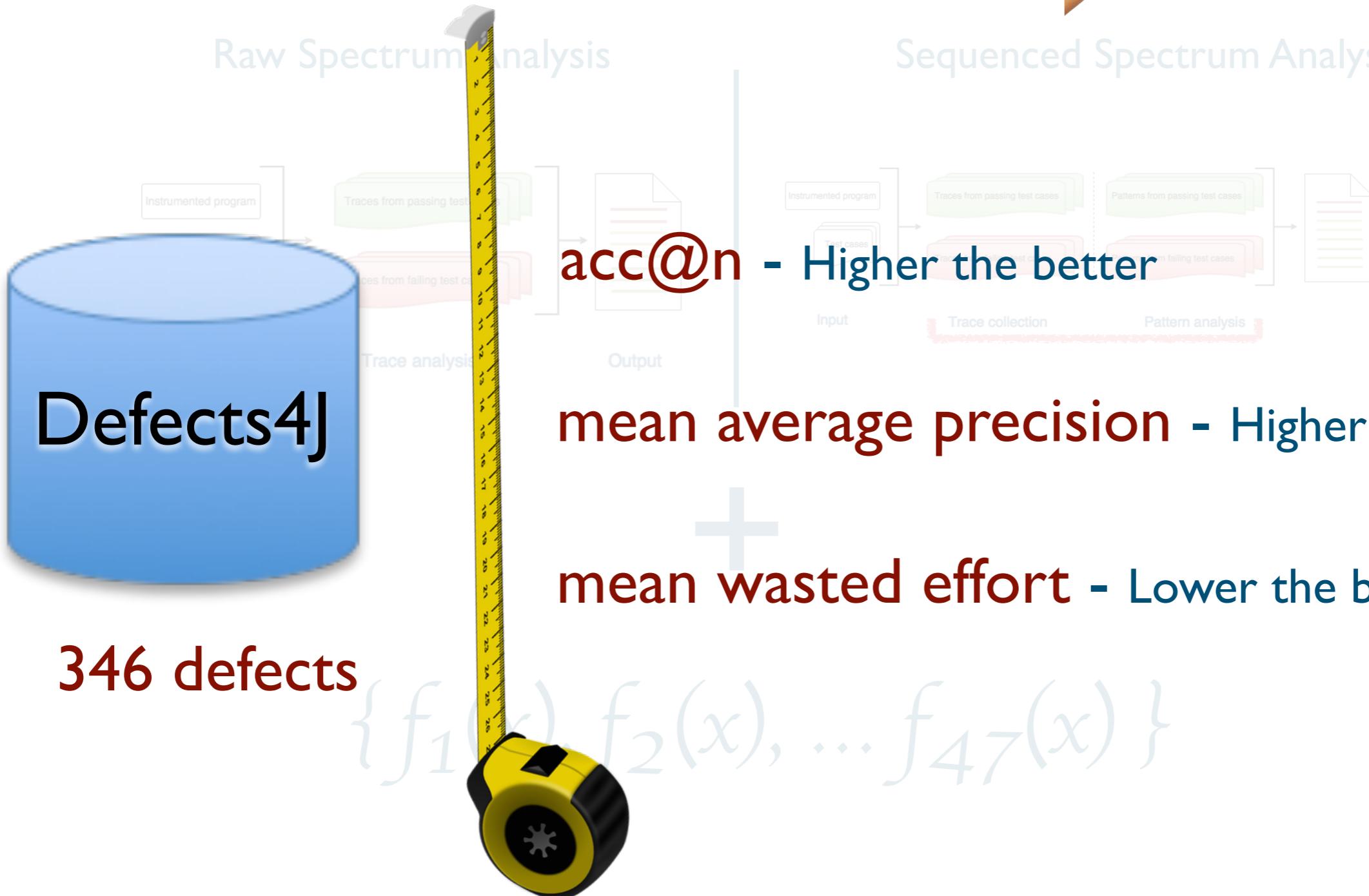
## Sequenced Spectrum Analysis



+

$$\left\{ f_1(x), f_2(x), \dots, f_{47}(x) \right\}$$

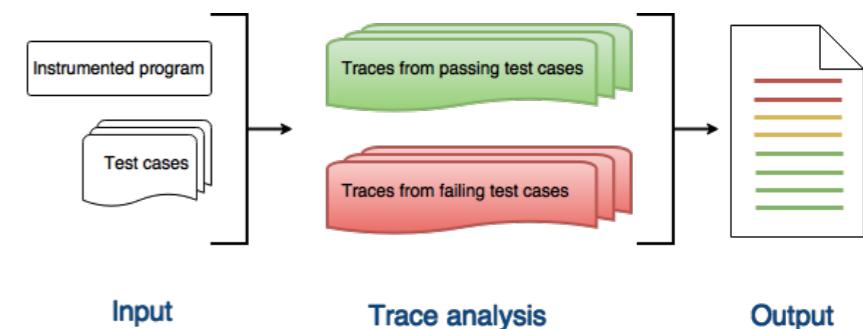
# Case Study



# Results

RQ1. What is the baseline performance of raw spectrum analysis?

## Raw Spectrum Analysis



Rank	Fault Locator	acc@1	acc@3	acc@5	MAP	MWE
1	GP13	63	120	142	0.2780349	96.73
1	Naish2	63	120	142	0.2776756	96.64
2	M2	62	118	141	0.2753030	96.32
3	Goodman	61	116	138	0.2695181	16.68
3	Ample2	64	120	140	0.2764775	101.24
4	T*	62	119	139	0.2744910	96.37
5	Zoltar	61	118	138	0.2735461	96.14

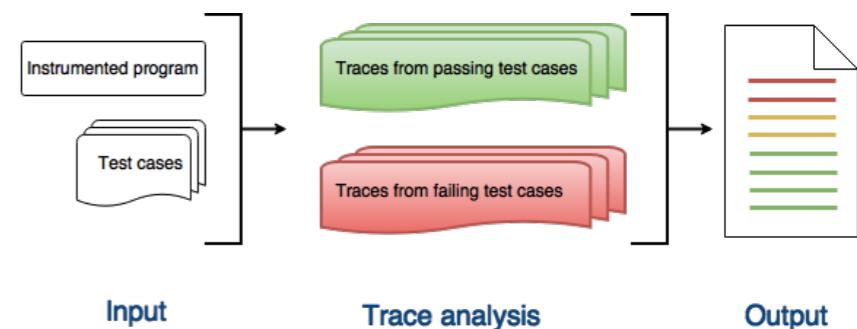
MAP = mean average precision

MWE = mean wasted effort

# Results

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## Raw Spectrum Analysis



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1	GP13	63	120	142	0.2780349	96.73
1	Naish2	63	120	142	0.2776756	95.64
2						
3		18% exact hit			average non-faulty methods on top of the faulty method	
3	Ample2	64	120	140	0.2764775	101.24
4	T*	62	119	139	0.2744910	96.37
5	Zoltar	61	118	138	0.2735461	96.14

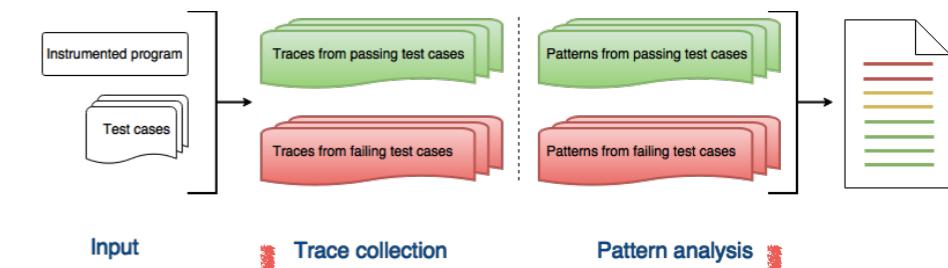
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# Results

RQ2. How much can sequenced spectrum analysis improve upon raw spectrum analysis?

## Sequenced Spectrum Analysis



Rank	Fault Locator	acc@1	acc@3	acc@5	MAP	MWE
1	Ample2	103	159	191	0.3925699	25.88
2	Fleiss	103	157	191	0.3874628	16.01
3	T*	102	160	190	0.3936098	31.18
4	M2	102	160	189	0.3933028	31.10
5	Arithmetic Mean	103	157	189	0.3875244	26.67

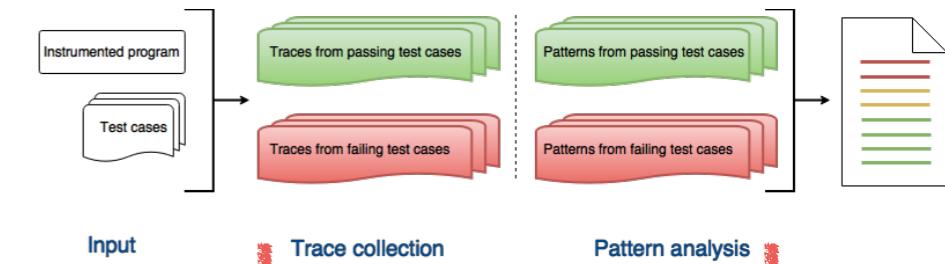
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2	Fleiss	103	157	191	0.3874628	10.01
3	T*	30% exact hit			average non-faulty methods on top of the faulty method	
4	M2	102	160	189	0.3933028	31.10
5	Arithmetic Mean	103	157	189	0.3875244	26.67

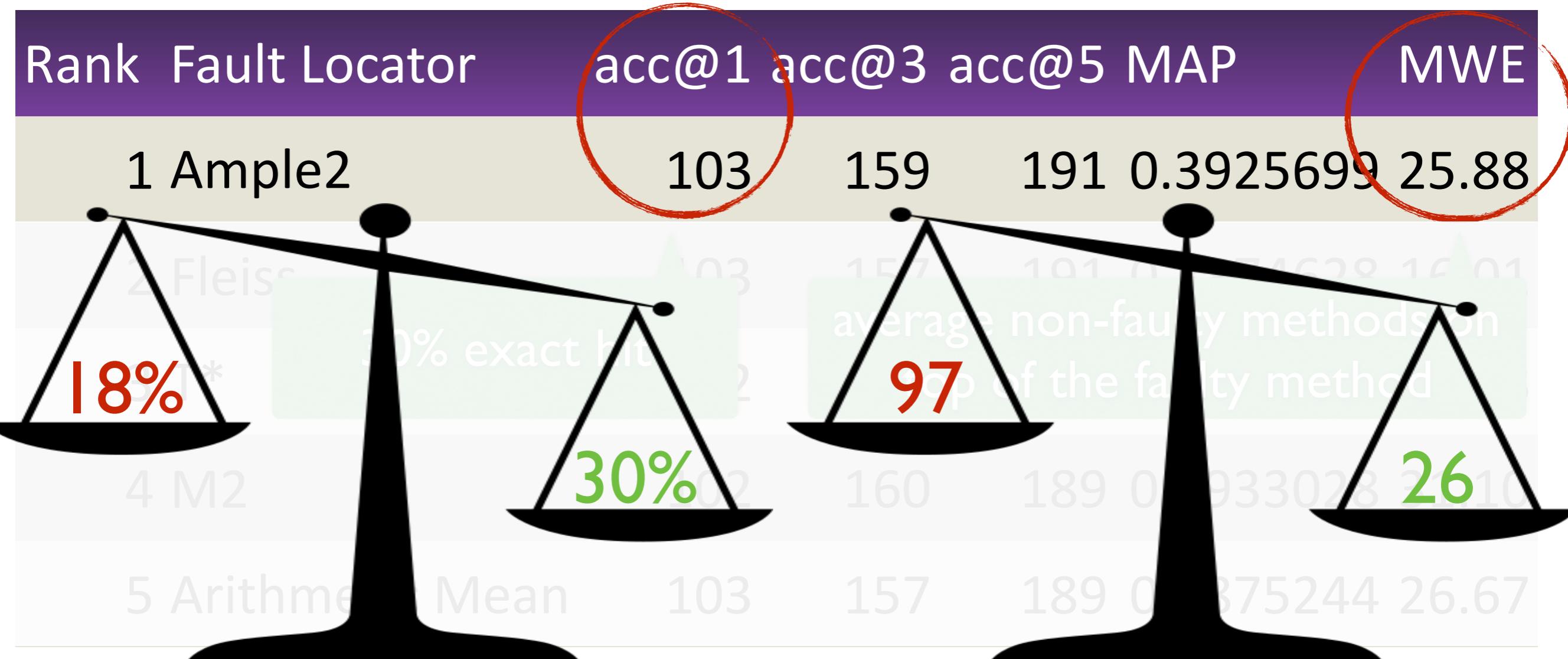
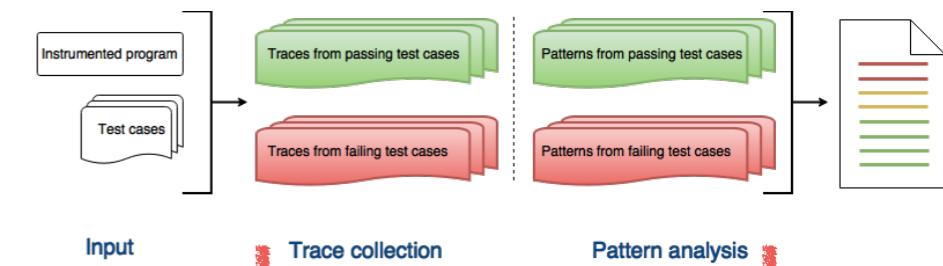
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## Sequenced Spectrum Analysis

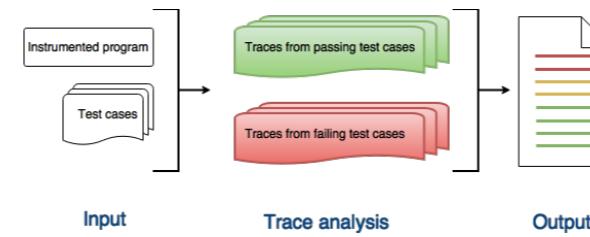


MAP = mean average precision

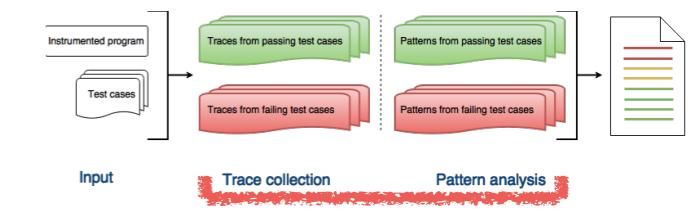
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# Results

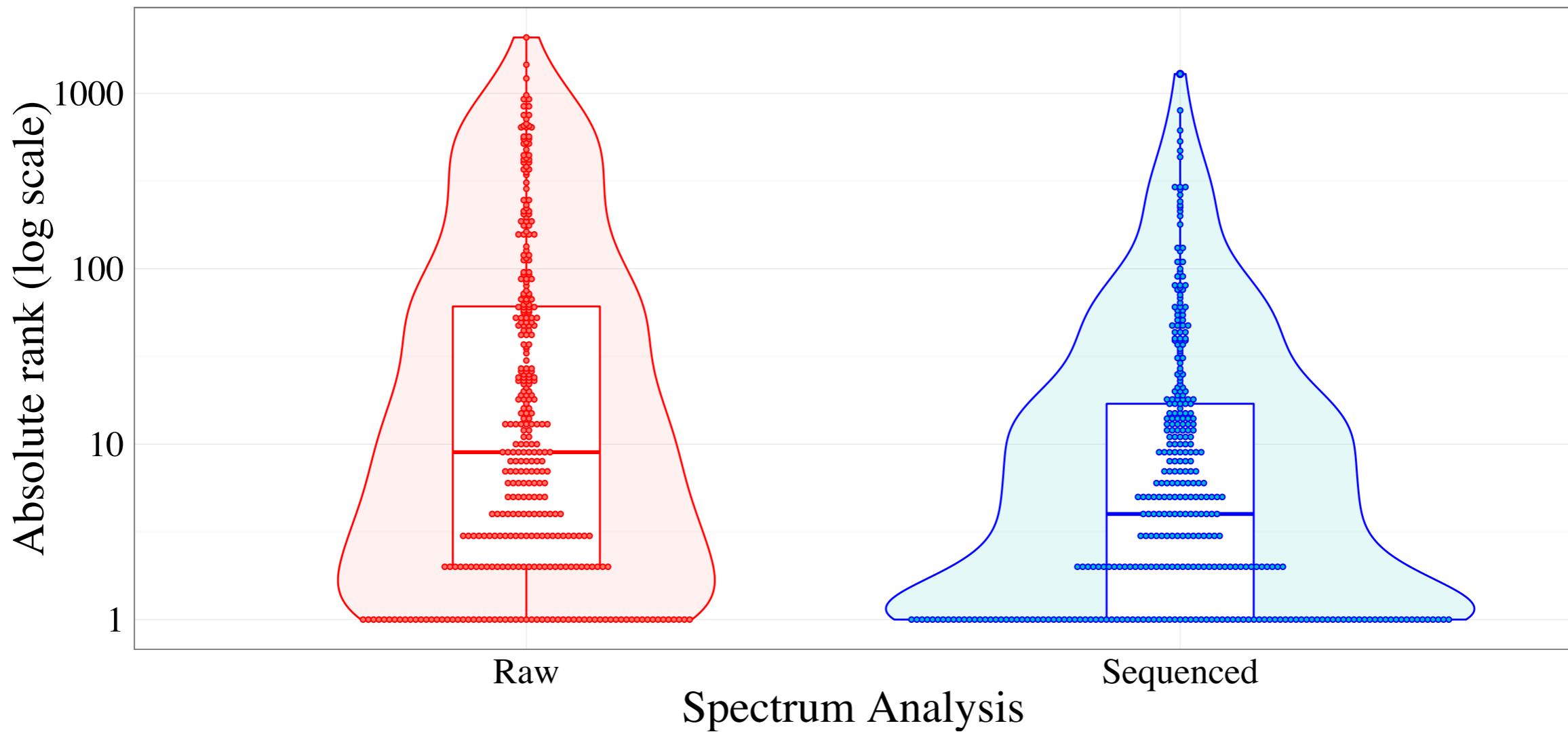
## Raw Spectrum Analysis



## Sequenced Spectrum Analysis

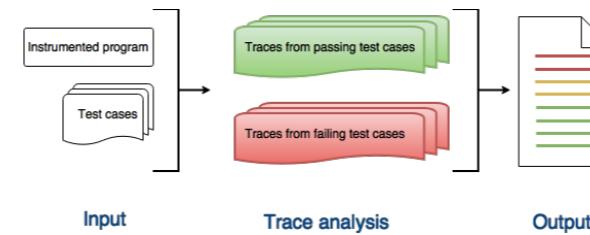


Spectrum Analysis: Raw Spectrum Analysis Sequenced Spectrum Analysis

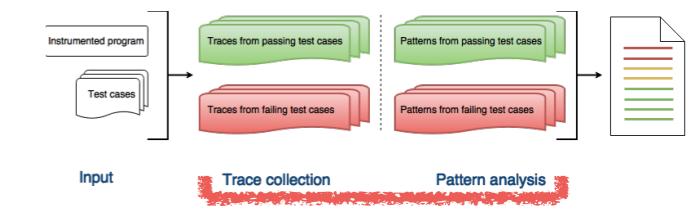


# Results

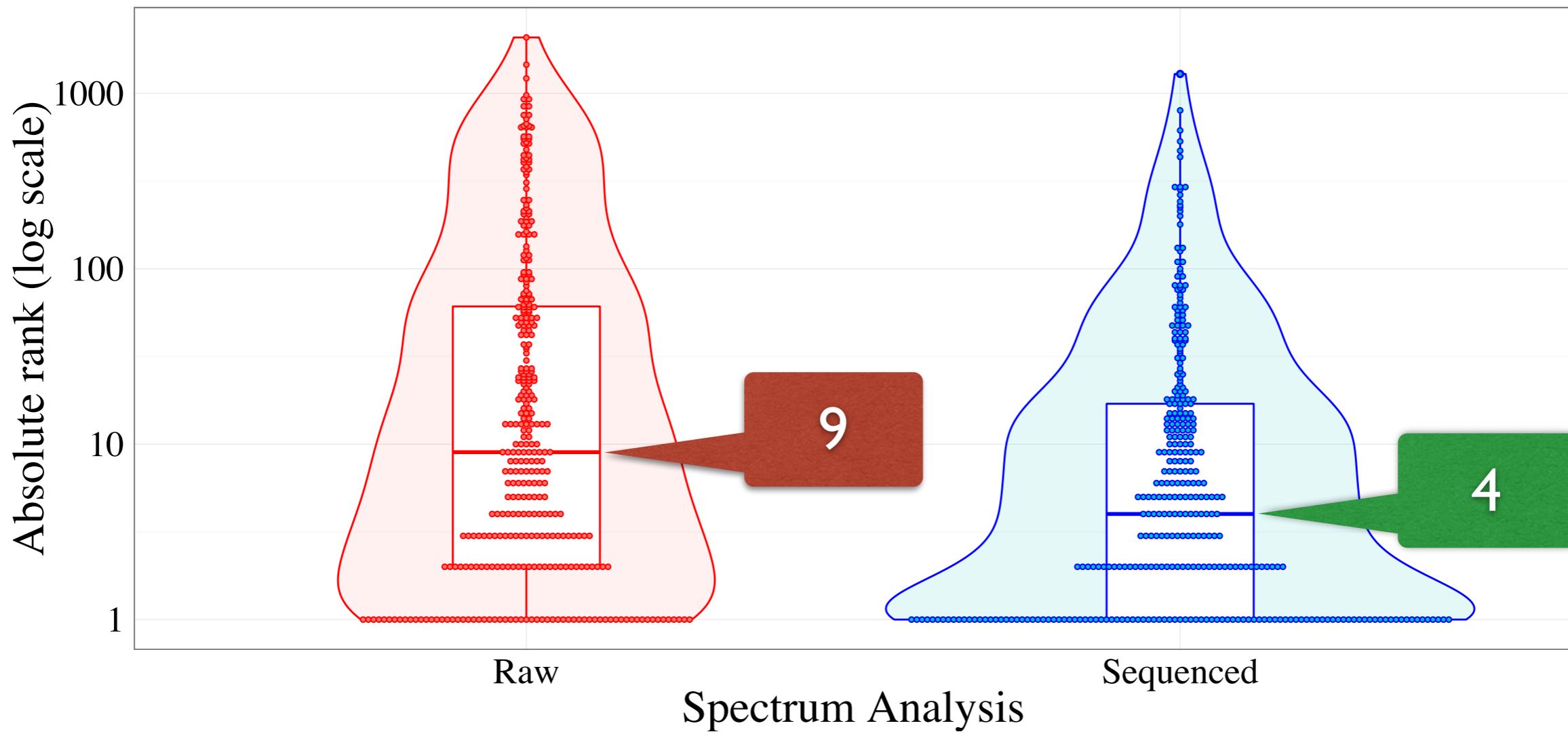
## Raw Spectrum Analysis



## Sequenced Spectrum Analysis

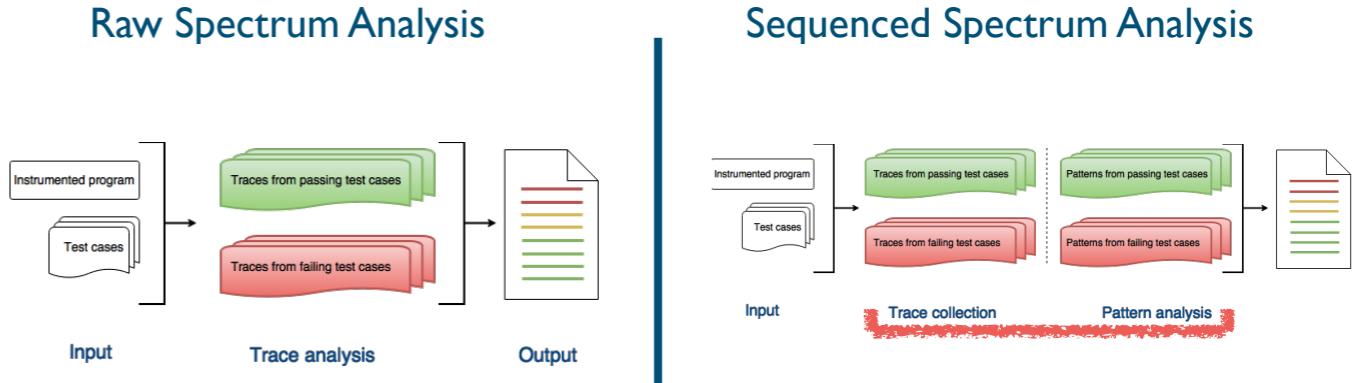


Spectrum Analysis: Raw Spectrum Analysis Sequenced Spectrum Analysis



# Results

RQ3. Are there project specific differences between the rankings?



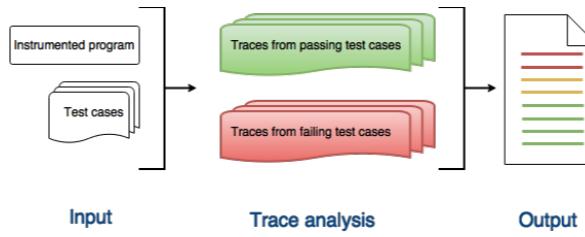
## Top Fault Locators

Project	Sequenced	Raw
Closure	Fleiss	GP13
Math	Goodman	Goodman
Lang	Geometric Mean	GP13
Time	Ample2	GP13
Chart	CBISqrt	Tarantula

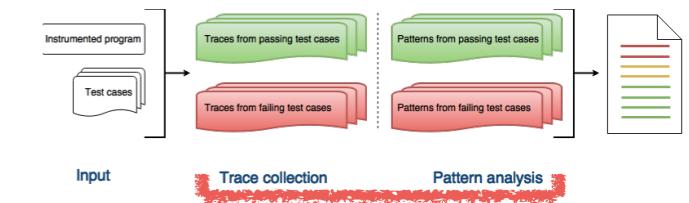
# Results

RQ3. Are there project specific differences between the rankings?

Raw Spectrum Analysis



Sequenced Spectrum Analysis



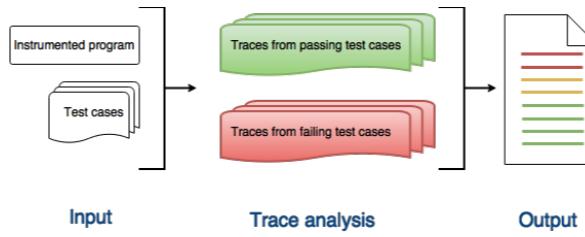
acc@1

Project	Sequenced	Raw
Closure	17 / 133	7 / 133
Math	33 / 106	21 / 106
Lang	41 / 65	21 / 65
Time	6 / 27	5 / 27
Chart	9 / 26	10 / 26

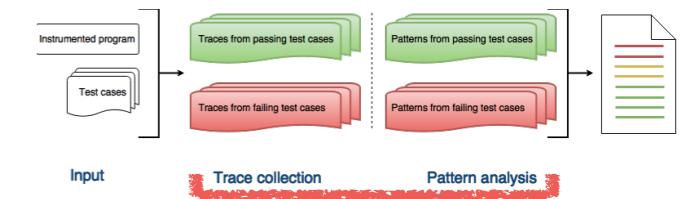
# Results

RQ3. Are there project specific differences between the rankings?

Raw Spectrum Analysis



Sequenced Spectrum Analysis



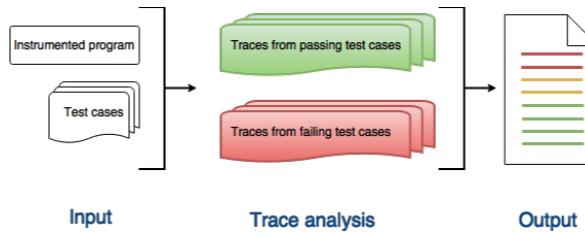
acc@I

Project	Sequenced	Raw
Closure	17 / 133	7 / 133
Math	33 / 106	21 / 106
Lang	41 / 65	21 / 65
Time	6 / 27	5 / 27
Chart	9 / 26	10 / 26

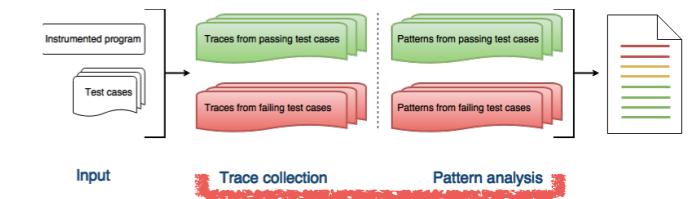
# Results

RQ3. Are there project specific differences between the rankings?

Raw Spectrum Analysis



Sequenced Spectrum Analysis

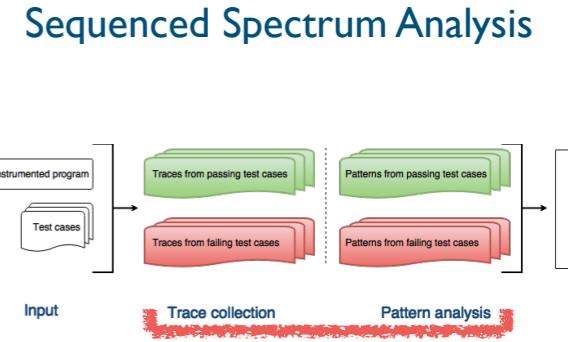
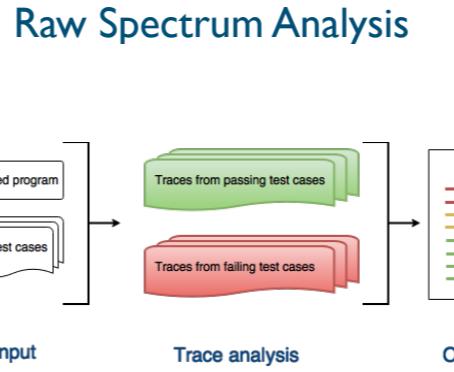


## Mean Wasted Effort

Project	Sequenced	Raw
Closure	31	223
Math	5	8
Lang	1	4
Time	16	39
Chart	13	27

# Results

RQ3. Are there project specific differences between the rankings?

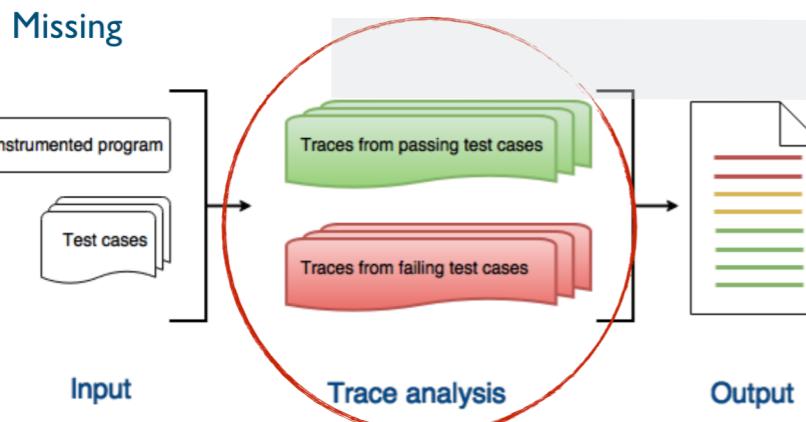


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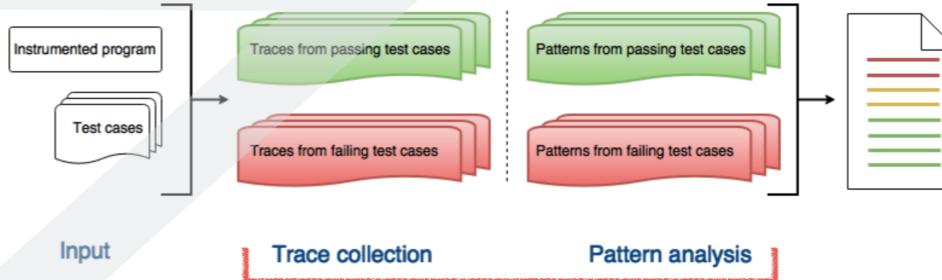
# Summary

## Raw Spectrum Analysis

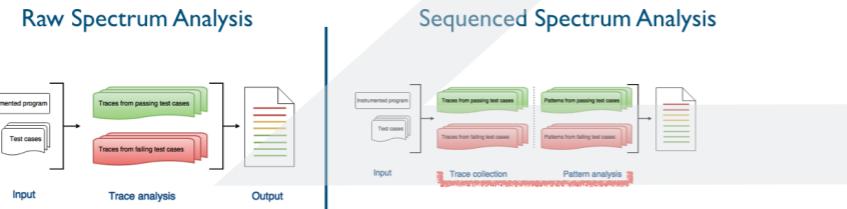


## Sequenced Spectrum Analysis

### Improvement



## Case Study



$$\{f_1(x), f_2(x), \dots, f_{47}(x)\}$$

## Results

