

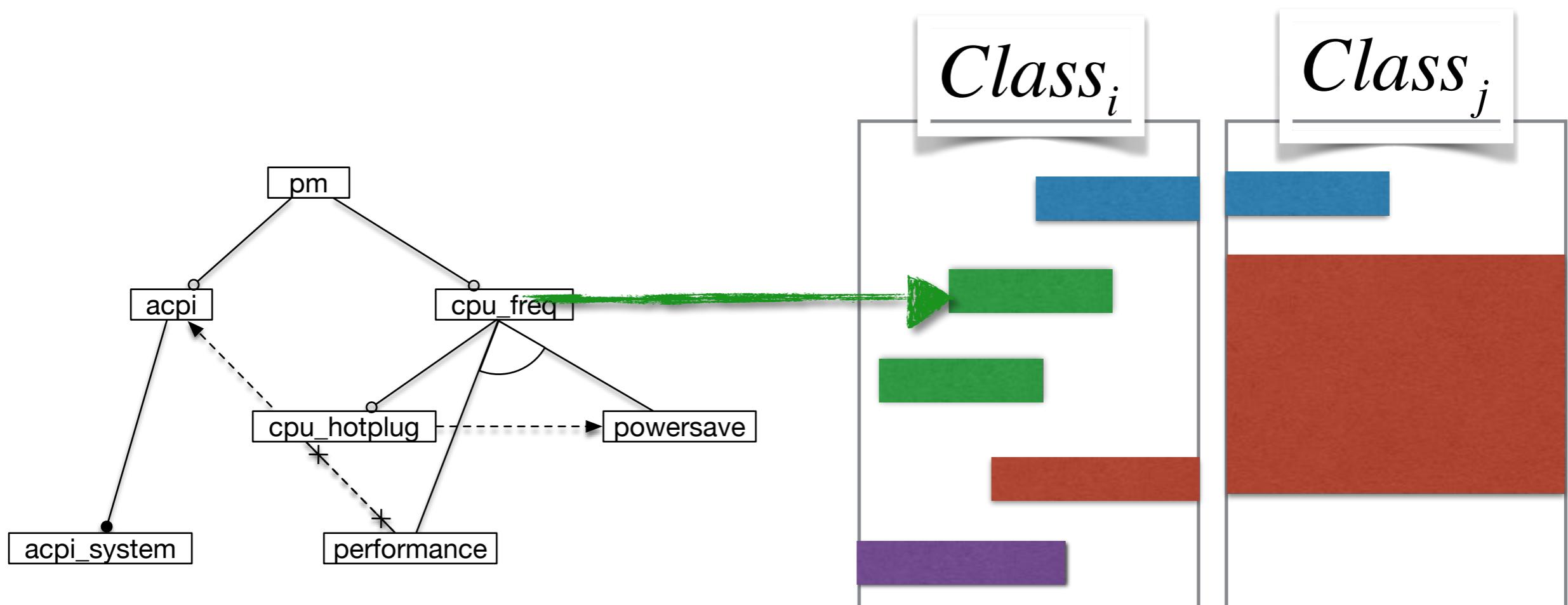
# StiCProb: A Novel Feature Mining Approach using Conditional Probability

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more: [www.chrisyttang.org/loong](http://www.chrisyttang.org/loong)



# Motivation

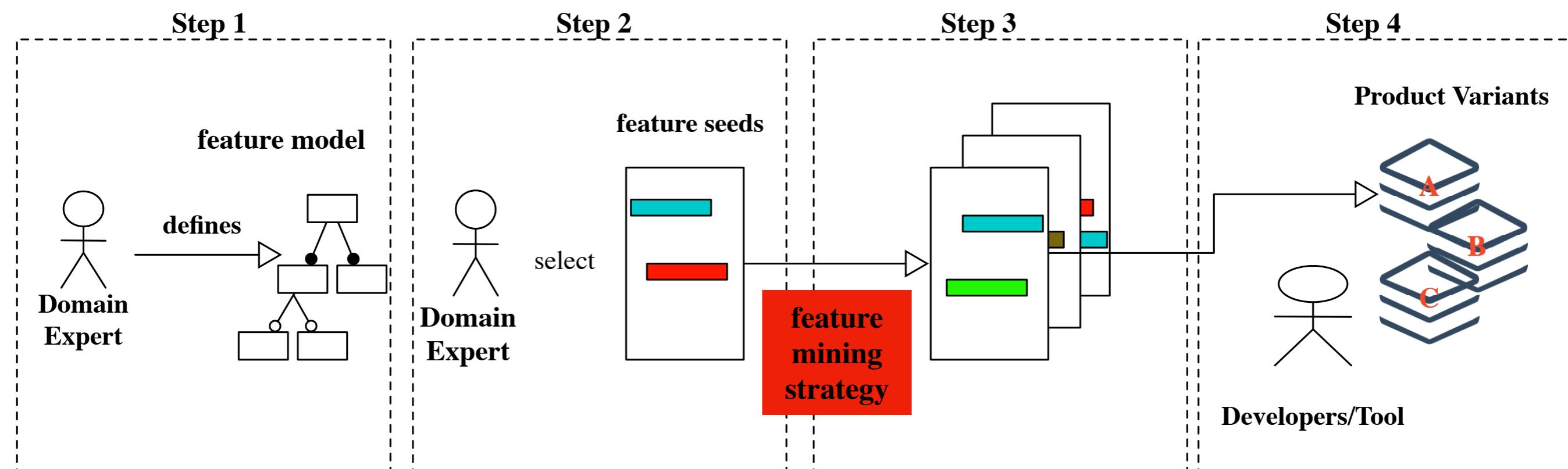
- legacy —> product line



1. How to locate the feature ?
2. How to measure ?

$feature \longleftrightarrow element$

# How to locate?



1. Select seeds
2. Annotate features

# Basic

Programming elements

$E$

Relationship

$R \subseteq E \times E$

Feature

$F$

(

Annotation

$A \subseteq E \times F$

# Basic (cond')

*mutual exclusion*

$$M \subseteq F \times F$$

*implications*

$$\Rightarrow \subseteq F \times F$$

*full annotation*

$$A \subseteq E \times F$$

$$A^* = \{(e, f) \mid (e, f) \in A, g \Rightarrow^* f\}$$

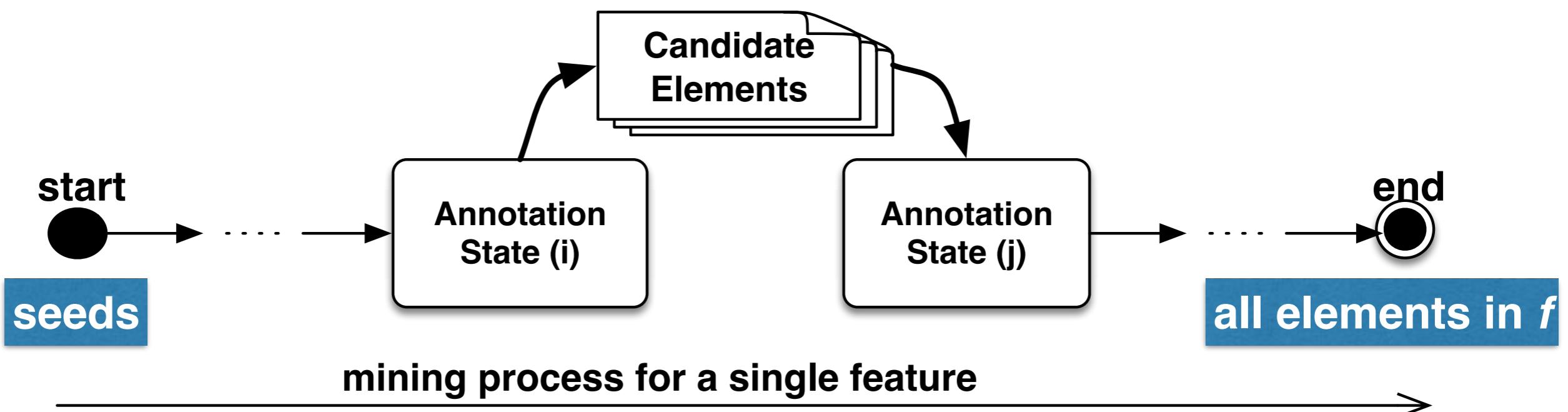
$(e, f)$

$(e, f) \mid (e, f) \in A, g \Rightarrow f$

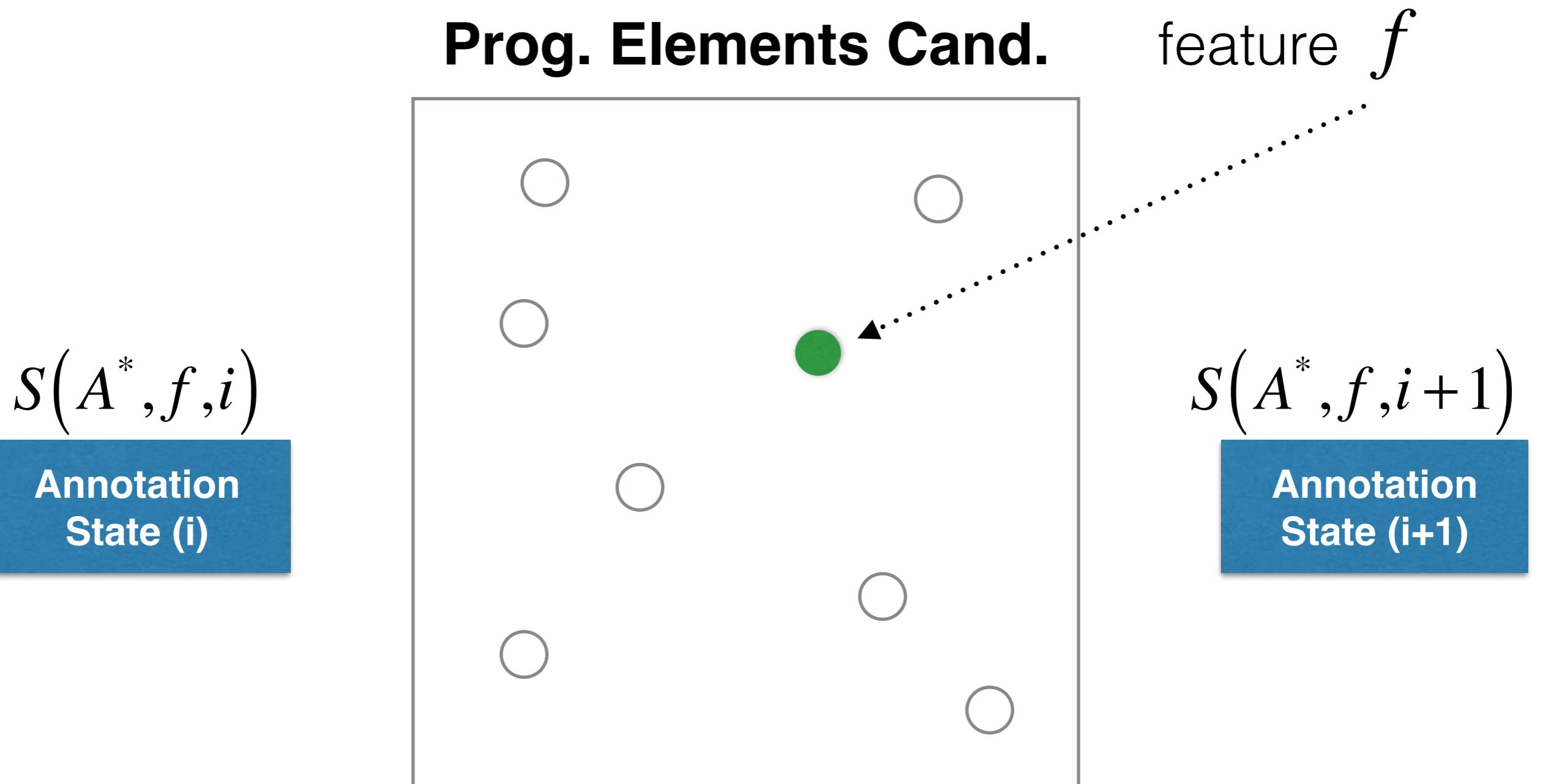
# Annotate Features

Annotation State

$$S(A^*, f, i) = \{e \mid (e, f) \in A^*\}$$



# Feature-Element Correlation Coefficient



# Feature-Element Correlation Coefficient(cond')

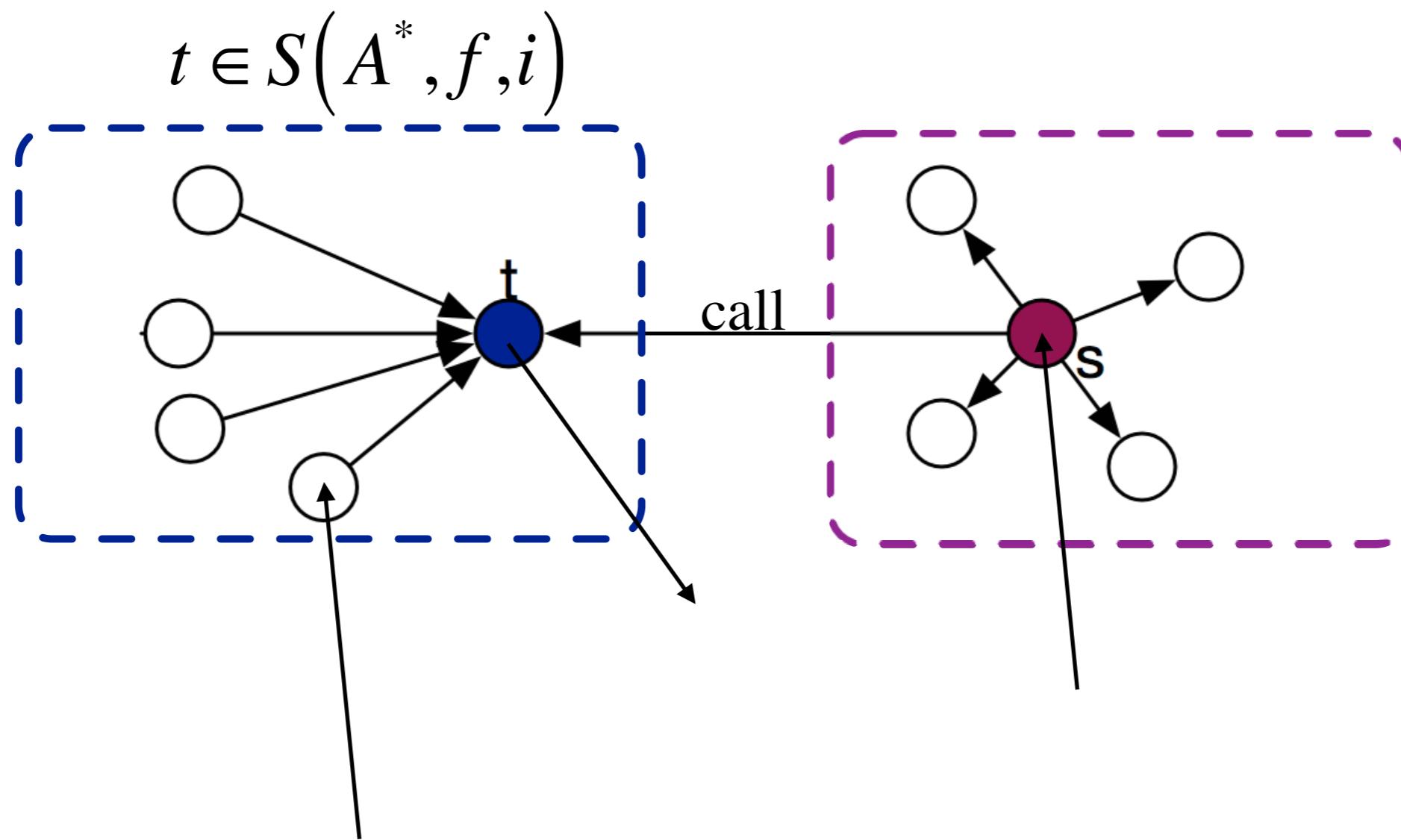
**Prog. Elements Cand.**

feature  $f$

$$S(A^*, f, i) \xrightarrow{\hspace{1cm}} \bullet \xrightarrow{\hspace{1cm}} S(A^*, f, i+1)$$

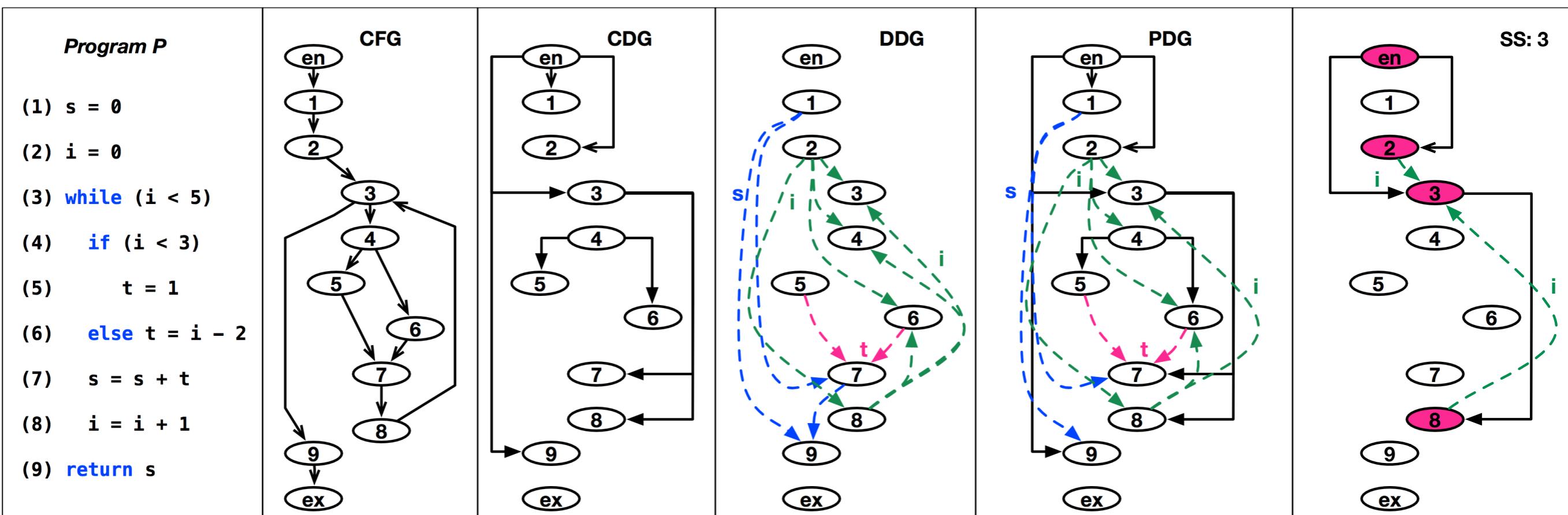
$p(e|S(A^*, f, i))$

# Feature-Element Correlation Coefficient(cond')



# Feature-Element Correlation

## Coefficient(cond')



Slicing Scope

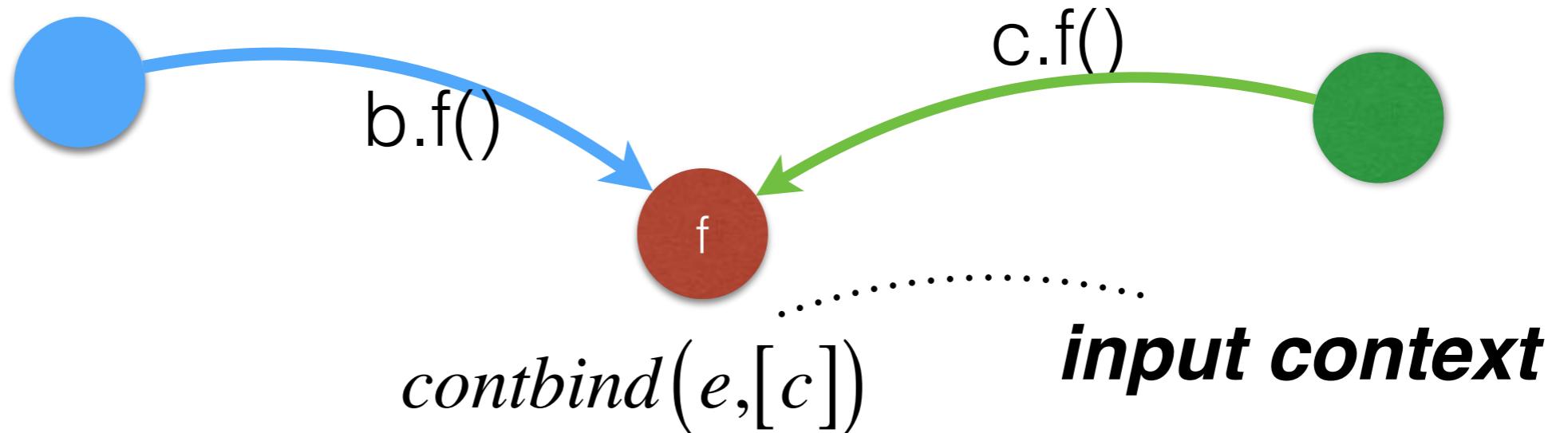
$$sscope(e) = e \cup \left\{ s \mid s \xrightarrow{\frac{df}{cf}} e, s \in E \right\}$$

# Feature-Element Correlation Coefficient(cond')

Binding

$$bind(e) = def(e) \cup use(e)^* \quad \text{in} \quad scope(e)$$

Context Binding



\* All def and use within e

# Context Binding

Context Binding @ Method Invocation

callsite

$$l = r_0.m(r_1, r_2, \dots, r_n)$$

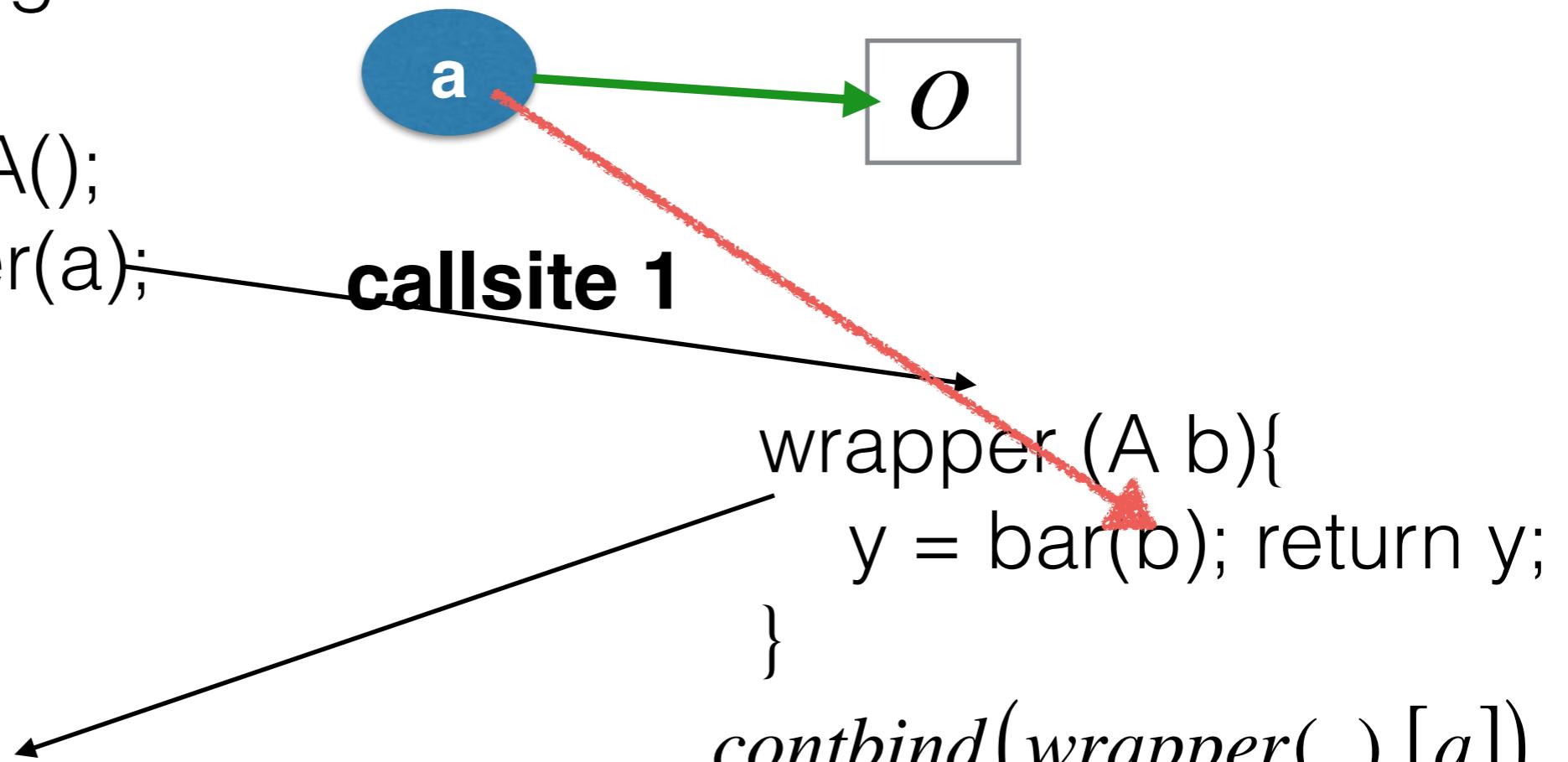
$$\text{contbind}(m, [r_1, \dots, r_n]) = \text{dispatch}(p_i = r_i) \rightarrow \text{bind}(m)$$

[13] L. O. Anderson, “Program analysis and specification for c programming language”

# Context Binding

Context Binding @ Method Invocation

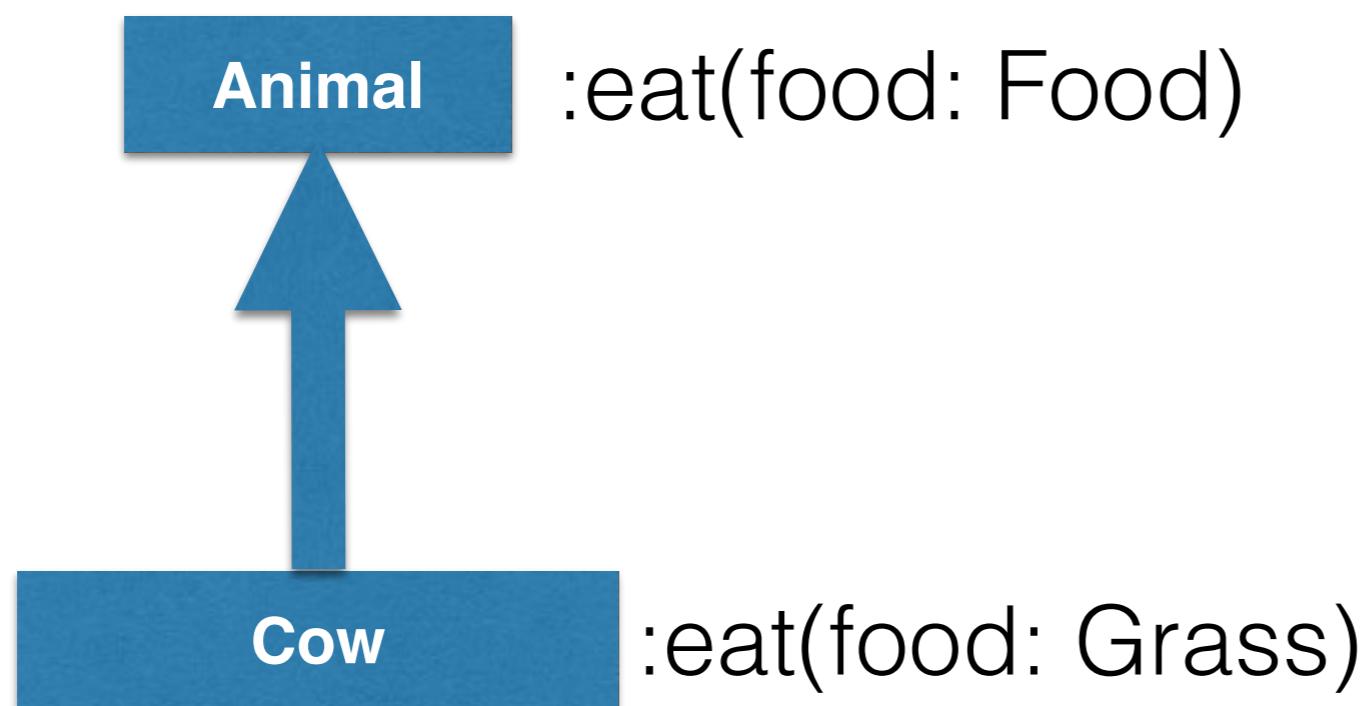
```
main(){  
    A a = new A();  
csite 1: z = wrapper(a);  
}  
  
bar (A c){  
    x = c.f; return x;  
}
```



[13] L. O. Anderson, "Program analysis and specification for c programming language"

# Context Binding

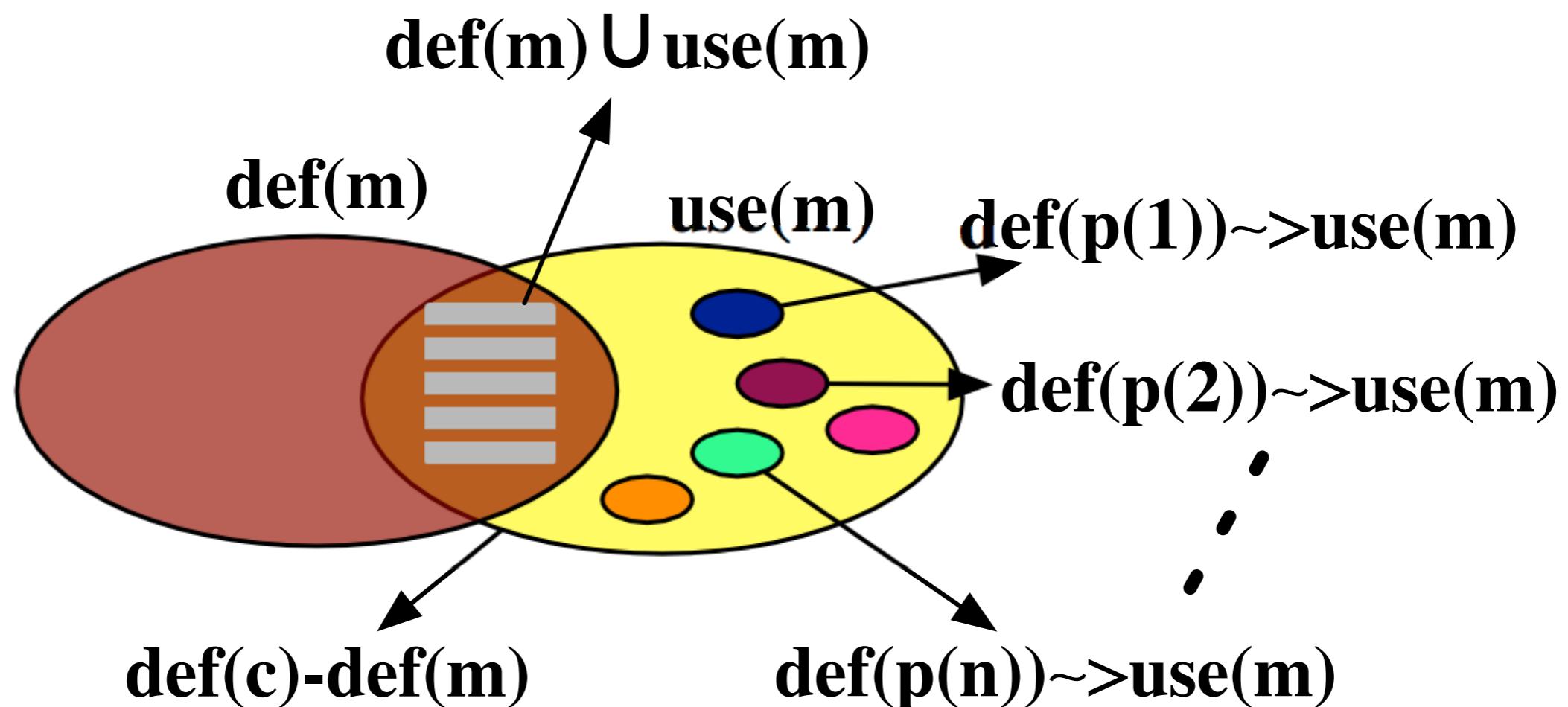
Context Binding @ Overriding



- m 1. def and use in m
- 2. def in parent class/interface, used in m

# Context Binding

Context Binding @ Overriding



# Context Binding

Context Binding @ Overriding

defined in m:  $\text{def}(m)$

used in m:  $\text{use}(m)$

1. used in m and defined in m
2. used in m and not defined in m

$$\text{contbind}(m, [p_1, \dots, p_n]) = \text{def}(m) \cup \bigcup_{i=1}^n (\text{use}(m) \rightsquigarrow \text{def}(p_i))$$

$\rightsquigarrow$  used to specify the source of the context

# Context Binding

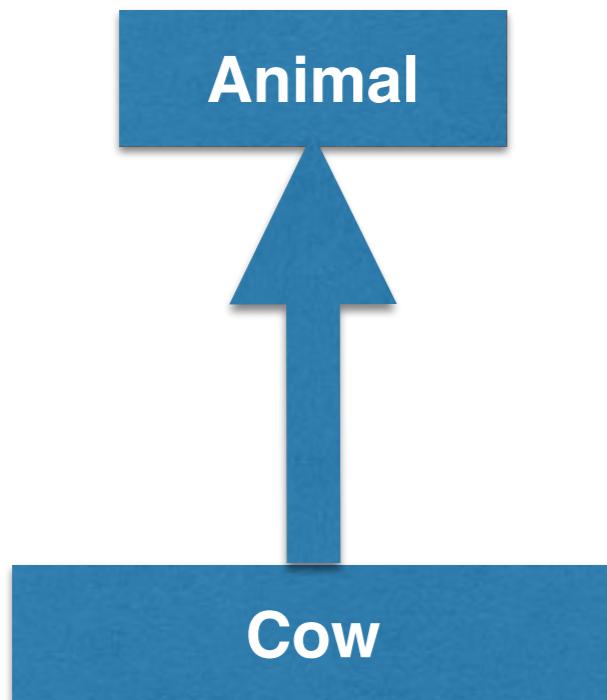
## Context Binding @ Overriding : Example

```
1 public class FlyingCar implements OperateCar {  
2     public int startEngine(int encryptedValue) {  
3         OperateCar.super.startEngine(OperateCar.encryptedValue);  
4     }  
5 }  
-----  
6 public interface OperateCar {  
7     int encryptedValue = 1;    ←  
8     default public int startEngine(int value) { ... }  
9 }
```

*contextbind(startEngine) = {encryptedValue, OperateCar.encryptedValue}*

# Context Binding

Context Binding @ Inheritance

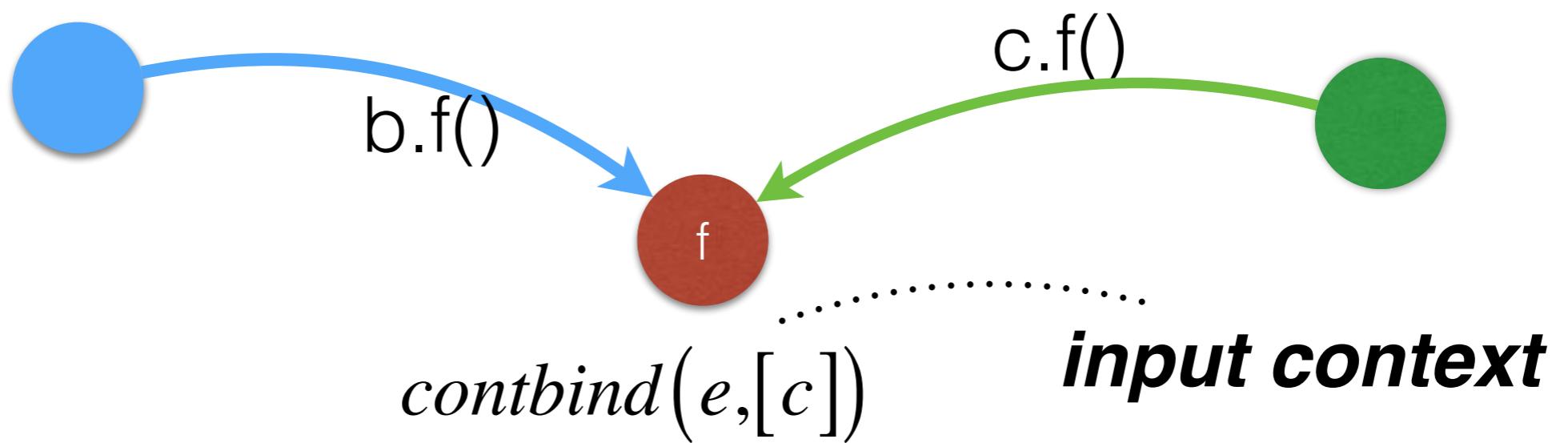


$$contbind(c, [p_1, \dots, p_n]) = bind(c) \cup \bigcup_{i=1}^n def(p_i)$$

# Feature-Element Correlation

## Coefficient(cond')

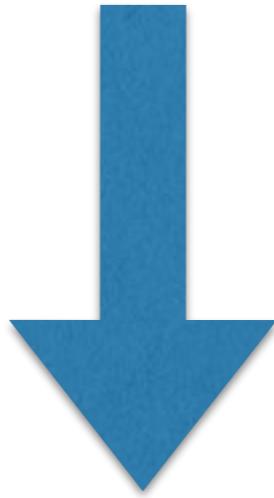
Context Binding



by default: ***context-aware points-to analysis***

# Feature-Element Correlation Coefficient(cond')

$$S(A^*, f, i) = \{e | (e, f) \in A^*\}$$



$$S(A^*, f, i) = \bigcup_{a \in S(A^*, f, i)} contextbind(a)$$

# StiCProb

1. Build Program DB
2. Build uniqueness table
3. Annotate features

# StiCProb: Uniqueness Table

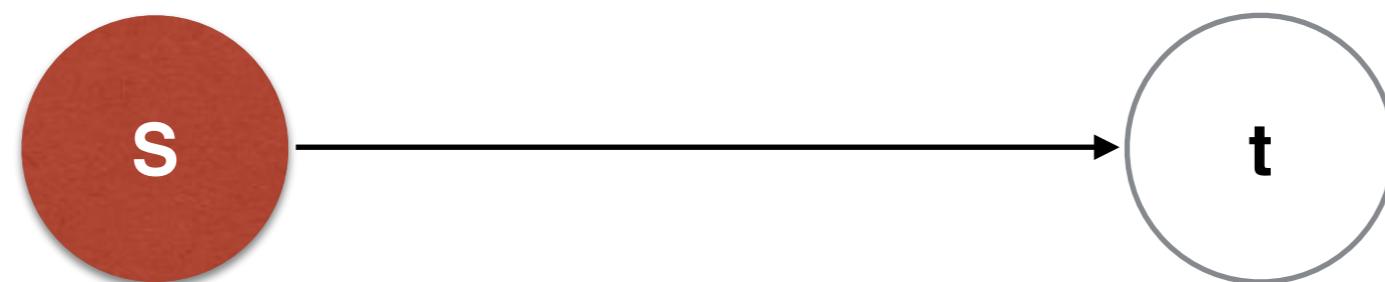
element  $s$  and  $t$  with a relation  $r$

$$s \xrightarrow{r} t$$

$$U(E, T, R, P_{forward}, P_{backward})$$

$$P_{forward}$$

the uniqueness of  $t$  to  $s$  if  $s$  has been annotated to a feature  $f$



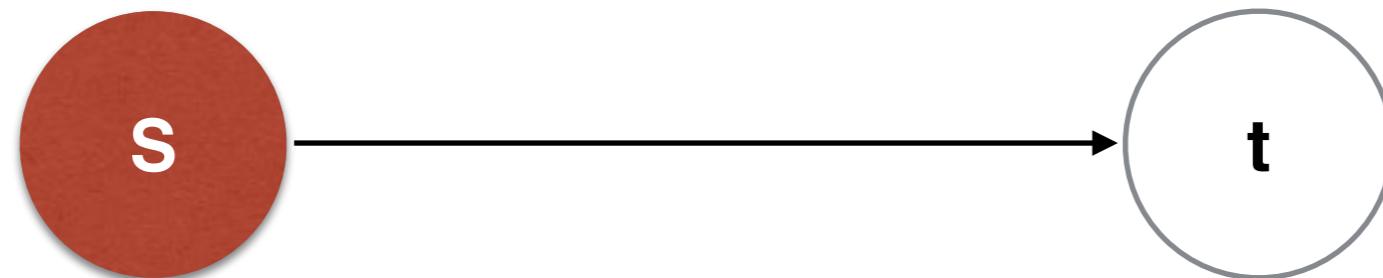
$$s \in S(A^*, f, i)$$

$$S(A^*, f, i) = \{e \mid (e, f) \in A^*\}$$

# StiCProb: Uniquess Table(cond')

$P_{forward}$

the uniqueness of t to s if s has been annotated to a feature **f**



$$s \in S(A^*, f, i)$$

$$S(A^*, f, i) = \{e \mid (e, f) \in A^*\}$$

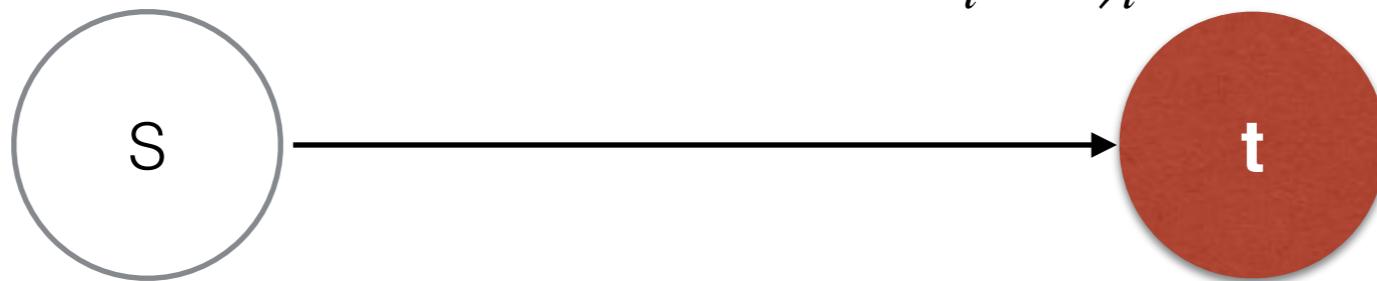
$$p_{forward}(s \xrightarrow{r} t \mid (s, f) \in A^*) = \frac{\text{contbind}(t, [s])}{\text{contbind}(s)}$$

# StiCProb: Uniquess Table(cond')

$P_{backward}$

the uniqueness of s to t if t has been annotated to a feature **f**

$$p_{backward}(s \xrightarrow{r} t | (t, f) \in A^*) = \frac{\text{contbind}(t, [s])}{\bigcup_{i \xrightarrow{r} t} \text{contbind}(t, [i])}$$

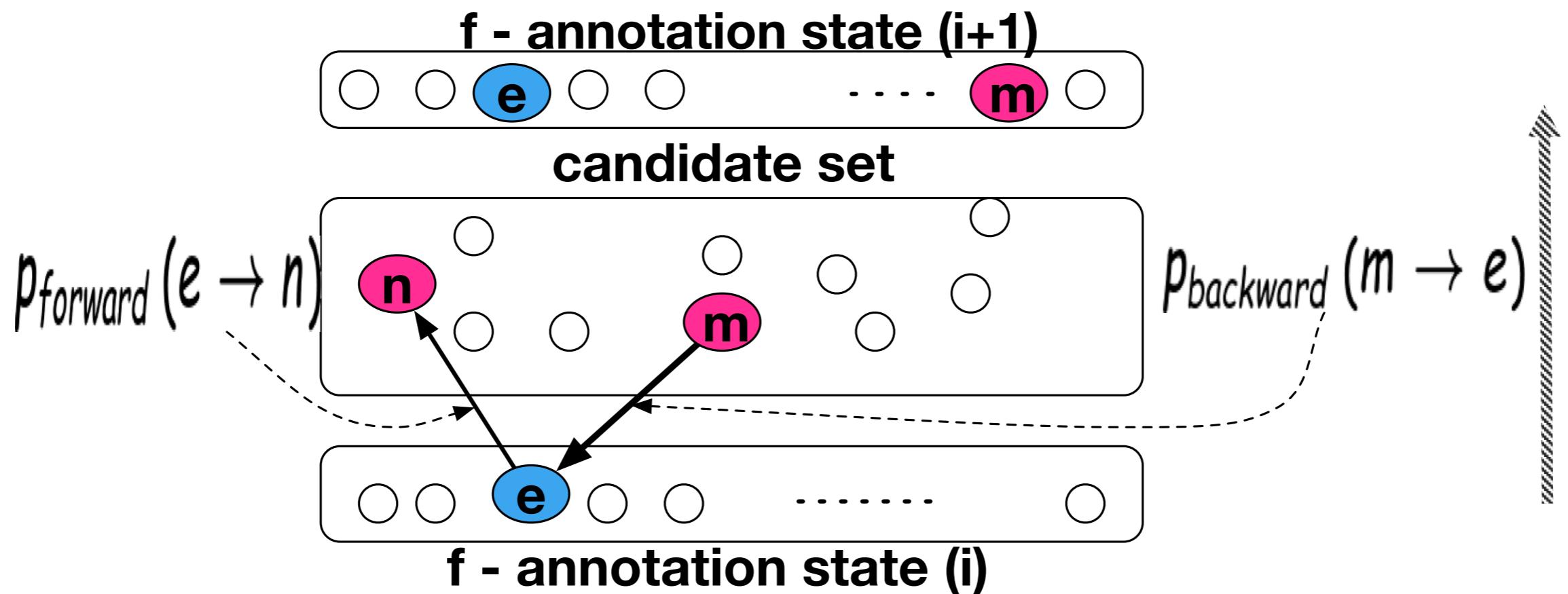


$$s \in S(A^*, f, i)$$

$$S(A^*, f, i) = \{e | (e, f) \in A^*\}$$

$\bigcup_{i \xrightarrow{r} t} \text{contbind}(t, [i])$  a collection of context binding from all prog.  
elements, which have relation  $r$  with  $t$ .

# StiCProb



# StiCProb

---

**Algorithm 1:** StiCProb feature mining approach

---

**Input:**  $seeds$ ,  $fm$ ,  $threshold$ ,  $U$

**Output:** all annotation states for features  $Sset$  in  $fm$

1 Create a set of annotation states as

$$Sset = \bigcup_f^{f \in features} S(A^*, f);$$

2 Assign seeds to each feature as  $S(A^*, f) = seeds(f)$ ;

3 Create feature set  $features$  with all features in  $fm$ ;

4 **while**  $features$  not *NULL* **do**

5   **for** feature  $f$  in  $features$  **do**

6     Create set  $waitList = \emptyset$ ;

7     Create candidate set  $C(S, f) = \emptyset$  for  $f$ ;

8     Add all elements have relations with elements in  
 $S(A^*, f)$  to  $C(S, f)$ ; // initialize  $C(S, f)$

9     **for** element  $m$  in  $C(S, f)$  **do**

10       **if** there is a relation  $r$  from  $m$  to the element  
 $e$  in  $S(A^*, f)$  **then**

11           Let  $value =$

$$p_{backward}(m \xrightarrow{r} e | (e, f) \in A^*);$$

12       **else**

13           Let  $value =$

$$p_{forward}(e \xrightarrow{r} m | (e, f) \in A^*);$$

14       **if**  $value > threshold$  **then**

15           Add  $m$  to  $waitList$ ;

16     Update  $S(A^*, f) \leftarrow S(A^*, f) \cup waitList$ ;

17     **if**  $StopCheck(f)$  is TRUE **then**

18       Remove  $f$  from  $features$ ;

19 **return**  $Sset$ ;

Feature model(fm)

Seeds(seeds)

threshold

Uniqueness Table(U)

# Case Study

Projects	LOC	#features	domain
Prevalyer	8,009	5	object persistence library
MobileMedia	4,653	6	mobile
Lampiro	44,584	*2	message client
ArgoUML	~120K	7	modeling tool

# Case Study

## Experimental Setting:

**seeds**: FLAT3 tool

**tool**: Loong Eclipse plugin

**feature model**: benchmark  
**benchmark**

## Related Approaches:

Type system, Topology analysis, Text comparison

## Measurement:

precision recall f-score

# Case Study

## **Other Settings:**

**# seeds:** 3

**threshold:** 0.6

# Experimental Result

StiCProb with threshold  $t = 0.6$

Project	Feature	Feature Size			Mining Results		
		LOC	FR	FI	IT	Recall	Prec.
Prevayler	Censor	105	10	5	3	17%	60%
	Gzip	165	4	4	3	16%	100%
	Monitor	240	19	8	2	17%	82%
	Replication	1487	37	28	26	79%	98%
	Snapshot	263	29	5	9	42%	99%
MobileM.	CopyMedia	79	18	6	4	43%	95%
	Sorting	85	20	6	4	32%	100%
	Favorites	63	18	6	12	20%	100%
	SMS Trans.	714	26	14	23	91%	49%
	Music	709	38	16	4	39%	90%
	Photo	493	35	13	5	63%	61%

**LOC:** line of code, **FR:** count of distinct code fragments, **IT:** number of iteration,

**Prec.:** precision

# Experimental Result (cond')

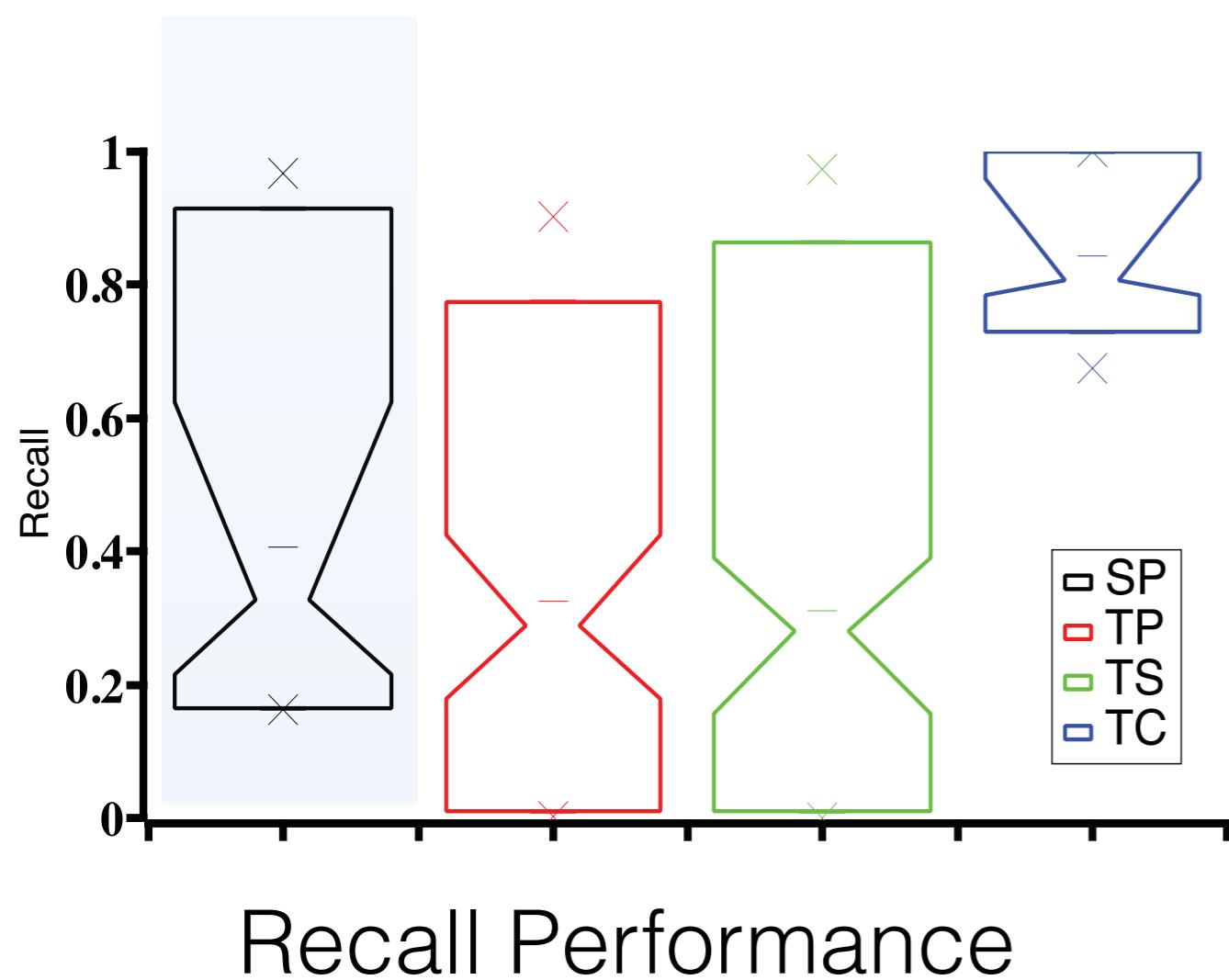
StiCProb with threshold  $t = 0.6$

Project	Feature	Feature Size			IT	Mining Results	
		LOC	FR	FI		Recall	Prec.
MobileM.	M.Transfer	153	4	3	14	97%	94%
Lampiro	Compre.	5155	33	20	34	40%	82%
ArgoUML	Cognitive	16319	285	233	127	70%	92%
	Activity	2282	115	80	17	26%	74%
	State	3917	115	88	18	33%	82%
	Collab.	1579	53	40	40	17%	72%
	Sequence	5379	65	53	98	33%	89%
	Use-Case	2712	59	49	39	19%	70%
	Deployment	3147	57	47	36	22%	67%

**LOC:** line of code, **FR:** count of distinct code fragments, **IT:** number of iteration,

**Prec.:** precision

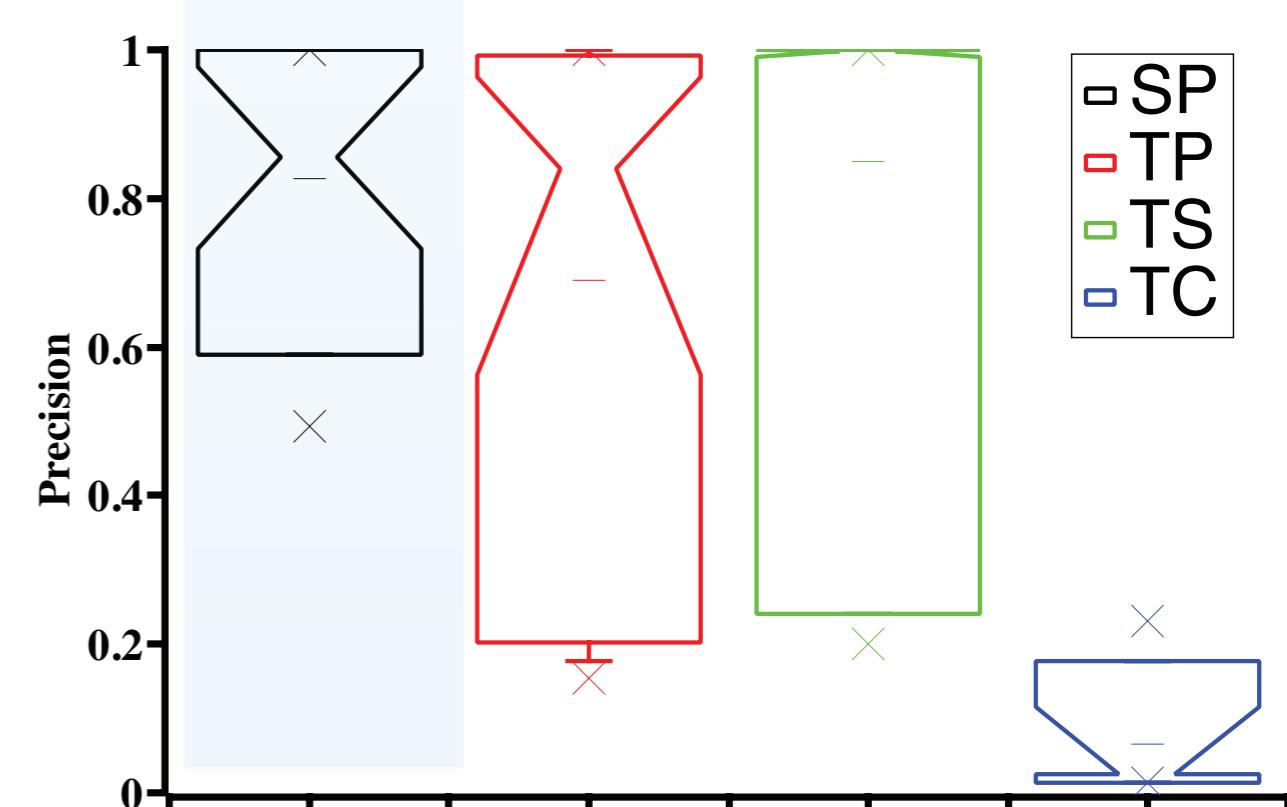
# Experimental Result (cond')



Recall Performance

SP: StiCProb ( $t = 0.6$ )

TP: topology analysis



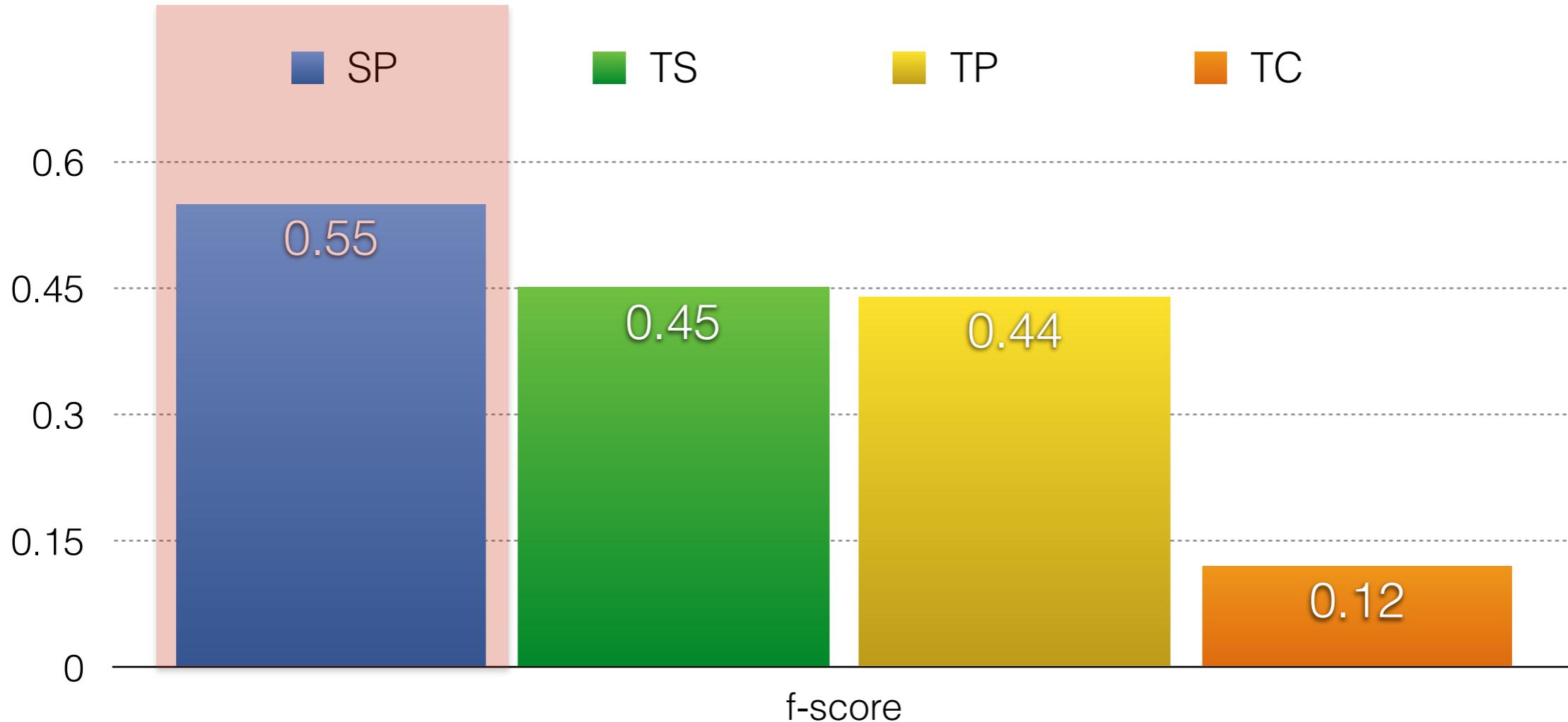
Precision Performance

TS: type system

TC: text comparison

# Experimental Result

## *f-score*



SP: StiCProb ( $t = 0.6$ )

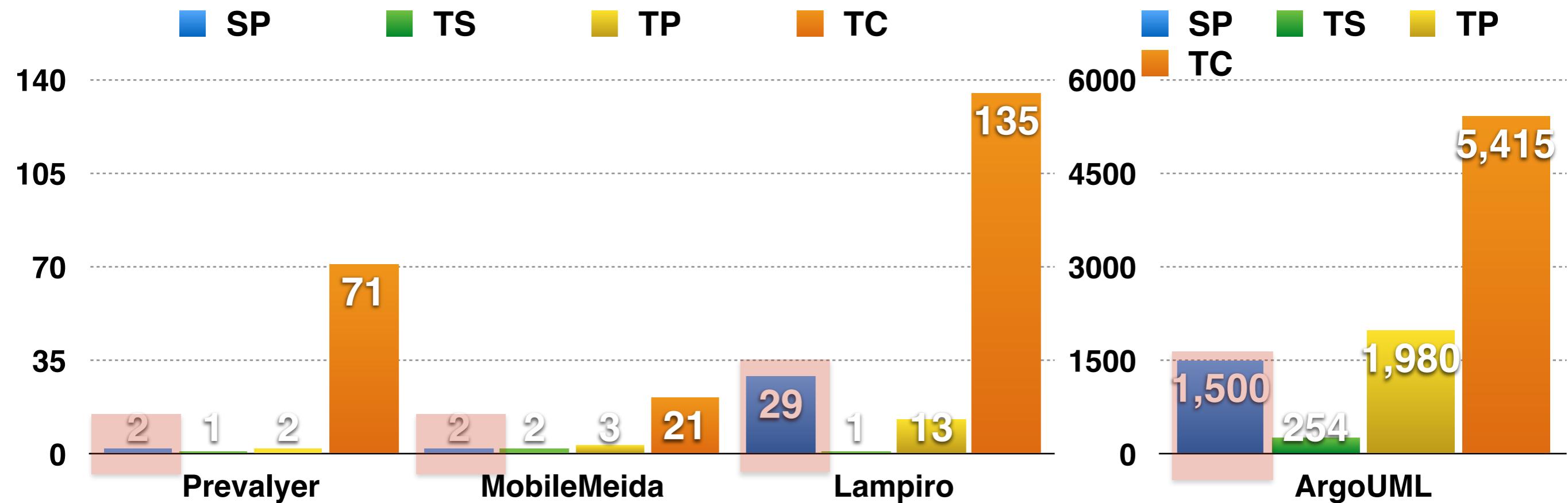
TP: topology analysis

TS: type system

TC: text comparison

# Experimental Result

## Runtime



SP: StiCProb ( $t = 0.6$ )

TP: topology analysis

TS: type system

TC: text comparison

# Discussion

## **Seeds:**

1. seeds provided by FLAT3 might be not correct
2. number of seeds
3. granularity of seeds: coarse granularity could improve the recall performance, but sometimes at the cost of precision.

# Discussion

## Thresholds:

threshold: 0.6 —> 0.8

precision: 83% —> 85%

The *threshold* contributes less to the performance.

Structure of the program

Thanks

# Loong Plugin

- Download: <http://www.chrisyttang.org/loong/>
- Source code: <https://github.com/csytang/Loong>
- Experimental results: <https://drive.google.com/folderview?id=0B9l0qvk6pnW0ZDRYMmxIQVhRb0U&usp=sharing>
- Online Tutorial: <http://www.chrisyttang.org/loong/>

# Discussion

- Need of req. specification <—> seeds selection/poor naming
- Variants of our approach? or better solutions ?
- - weighted graph —> graph clustering
- ?