# **Simplification and Computation**

Eugenio Moggi

moggi@disi.unige.it

DISI, Univ. of Genova

# **Overall Goal**

A framework for operational semantics based on

- ideas from monadic metalanguages [MF03]
  - simplification  $e \longrightarrow e'$  compatible&confluent relation on terms
  - computation  $Cfg_1 \mapsto Cfg_2$  relation on configurations
- ideas from CHAM [BB92] and related calculi (Join [FG96], Kell [Ste03,BS03])
  - configurations as multisets of terms (and reaction rules)
  - computation as chemical reaction (and heating)
- expressive patterns
  - patterns p for simplification subsume ML& PMC [Kah03] Do we need more? LINDA/KLAIM, XML
  - join patterns J with weaken linearity assumptions Kell patterns a[!x] address a different issue: match for active kells.
  - distinguish atoms a from varaibles y (as in FreshML [GP99,SGP03])

Use framework for multi-lingual extensions and for defining monadic interpreters.

# **Overview of Key Properties**

- we distinguish atoms a, name variables y and term variables x
- a term e may have free occurrences of atoms FN(e) and variables FV(e)
- a configuration Cfg is a finite multiset of closed terms (i.e. no free variables)

# **Overview of Key Properties**

- we distinguish atoms a, name variables y and term variables x
- a term e may have free occurrences of atoms FN(e) and variables FV(e)
- a configuration Cfg is a finite multiset of closed terms (i.e. no free variables)

The SIMPLIFICATION relation  $e \longrightarrow e'$  is

- name/variable preserving, i.e.  $FN(e') \subseteq FN(e)$  and  $FV(e') \subseteq FV(e)$
- compatible, i.e. can be performed in any context
- confluent, i.e. can be performed in any order
- Invariant w.r.t. permutations  $\pi$  of atoms and substitutions  $\rho$  of name/term  $e \longrightarrow e'$

variables with names/terms, i.e.

$$\begin{array}{c} e \longrightarrow e' \\ \hline e[\pi] \longrightarrow e'[\pi] \end{array} \begin{array}{c} e \longrightarrow e' \\ \hline e[\rho] \longrightarrow e'[\rho] \end{array}$$

# **Overview of Key Properties**

 $Cfg_1 \longmapsto Cfg_2$ 

 $Cfq'_1 \mid --- > Cfq'_2$ 

- we distinguish atoms a, name variables y and term variables x
- a term e may have free occurrences of atoms FN(e) and variables FV(e)
- a configuration Cfg is a finite multiset of closed terms (i.e. no free variables)

The COMPUTATION relation  $Cfg_1 \mapsto Cfg_2$  is

invariant w.r.t. permutations  $\pi$  of atoms

preserved by simplification, i.e. \*

• preserved by extension, i.e.  $Cfg \uplus Cfg_1 \longmapsto Cfg \uplus Cfg_2$  when  $Cfg_1 \longmapsto Cfg_2$ and  $FN(Cfg) \# (FN(Cfg_2) - FN(Cfg_1))$ . Thus atomic broadcast not allowed.

## **Syntax**

- atom  $a \in A$ , name variable y, term variable x, Name  $u \in N ::= a \mid y$
- **J** Term  $e \in E$ , Pattern p, Join pattern J, Match m

Meaning of matches and related constructs:

- $\bullet$  ok e succeeds and returns e
- $e: p \Rightarrow m$  if e matches p try instance of m, otherwise fail
- (m; e') returns e when m succeeds and returns e, and e' when m fails
- $(p \Rightarrow e_1 | e_2) @ e \longrightarrow (e: p \Rightarrow e_1; e_2 @ e) \text{ is } \beta \text{-reduction}$

## **Syntax**

- atom  $a \in A$ , name variable y, term variable x, Name  $u \in N : := a \mid y$
- **J** Term  $e \in E$ , Pattern p, Join pattern J, Match m

Linearity constrains and binding in patterns:

- In p a variable !x or !y can be declared at most once
  - \* the occurrences of y after !y are bound
- In J a term variable !x can be declared at most one  $u_i \overline{p}_i$ 
  - \* while a name variable y can be declared in several  $u_i \overline{p}_i$ .

#### **Examples of patterns** *p* **and** *J* **without free variables**

- $p_0 \equiv c \ (c \ !x)$  matches  $c \ (c \ e)$ , where  $c \in A$
- $p_1 \equiv c \; ! y \; y$  matches  $c \; a \; a$  for any  $a \in A$
- $p_2 \equiv !y \; (y \; !x)$  matches  $a \; (a \; e)$  for any  $a \in A$

How to express function eq to test equality of names (e.g. references)

$$eq = (!y \Rightarrow (y \Rightarrow true | !y' \Rightarrow false | fail) | fail)$$

eq fails when an argument is not an atom (does not simplify to an atom)

#### **Examples of patterns** *p* **and** *J* **without free variables**

We define reaction rules for the operation on ML-references, represented by atoms newR, getR and setR (term constructor names). Program represented by molecule named prog, store represented by molecules named ref.

- {(prog (newR !x !x'))} >  $\nu y$ . {(prog (x'@y) | ref y x)} semantics of let val y=ref x in x'y

### **Simplification rules: left-linear and non-overlapping**

$v ::= u \ \overline{e} \   \ fail \   \ (p \Rightarrow e_1   e_2) \   \ \nu y.e \   \ \{(e_i   i \in n)\} \   \ J > e$	$p ::= !x \mid !y \ \overline{p} \mid u \ \overline{p}$
$e ::= x \mid v \mid e_1 @e_2 \mid (m; e) \mid let \{ x_i = e_i \mid i \in n \}  in  e$	$m ::= ok \ e \mid e : p \Rightarrow m$

Unfolding of recursive definitions

let 
$$\{x_i = e_i | i \in n\}$$
 in  $e \longrightarrow e[x_i: \text{let } \{x_i = e_i | i \in n\}$  in  $e_i | i \in n]$ 

Application

$$\begin{array}{rccc} fail@e & \longrightarrow & fail\\ (p \Rightarrow e_1 | e_2)@e & \longrightarrow & (e: p \Rightarrow ok \ e_1; e_2@e) \end{array}$$

# **Simplification rules: left-linear and non-overlapping**

$v ::= u \overline{e}   fail   (p \Rightarrow e_1   e_2)   \nu y.e   \{(e_i   i \in n)\}   J > e$	$p ::= !x \mid !y \ \overline{p} \mid u \ \overline{p}$
$e ::= x \mid v \mid e_1 @e_2 \mid (m; e) \mid let \{ x_i = e_i \mid i \in n \}  in  e$	$m ::= ok \ e \mid e : p \Rightarrow m$

### Simplification of matching

$$\begin{array}{rcl} (ok \; e; e') & \longrightarrow & e \\ (e: !x \Rightarrow m; e') & \longrightarrow & (m[x:e]; e') \\ (u \; \overline{e}: !y \; \overline{p} \Rightarrow m; e') & \longrightarrow & (\overline{e}: \overline{p}[y:u] \Rightarrow m[y:u]; e') \quad \text{when } |\overline{e}| = |\overline{p}| \\ (v: !y \; \overline{p} \Rightarrow m; e') & \longrightarrow & e' \quad \text{when } v \not\equiv u \; \overline{e} \; \text{with } |\overline{e}| = |\overline{p}| \\ (a_1 \; \overline{e}: a_2 \; \overline{p} \Rightarrow m; e') & \longrightarrow & \begin{cases} (\overline{e}: \overline{p} \Rightarrow m; e') & \text{if } a_1 = a_2 \\ e' & \text{if } a_1 \neq a_2 \end{cases} \quad \text{when } |\overline{e}| = |\overline{p}| \\ (v: u \; \overline{p} \Rightarrow m; e') & \longrightarrow & e' \quad \text{when } v \not\equiv u \; \overline{e} \; \text{with } |\overline{e}| = |\overline{p}| \end{array}$$

#### **Computation rules: heating and chemical reaction**

$$v ::= u \overline{e} | fail | (p \Rightarrow e_1 | e_2) | \nu y.e | \{(e_i | i \in n)\} | J > e$$
  
$$p ::= !x | !y \overline{p} | u \overline{p} \qquad J ::= \{(u_i \overline{p}_i | i \in n)\}$$

Heating

$$\begin{array}{cccc} \textit{Cfg}, \ \{(e_i | i \in n)\} & \longmapsto & \textit{Cfg}, \ \{e_i | i \in n\} \\ & \textit{Cfg}, \ \nu y.e & \longmapsto & \textit{Cfg}, \ e[y:a] & \textit{with } a \notin \mathrm{FN}(\textit{Cfg}, \nu y.e) \end{array}$$

Reaction a la Join-calculus

Cfg, 
$$J > e$$
,  $J\rho \longrightarrow Cfg$ ,  $e[\rho]$ ,  $J > e \rho$  closed substitution

 $J\rho$  is the multiset obtained by replacing the only occurrence of !x in J with  $\rho(x)$ , and occurrences of !y and y in J with  $\rho(y)$  (each occurrence of y in  $u_i \overline{p}_i$  is bound by a !y)

### **Conclusion: multi-lingual extensions and interpreters**

• new term constructors encoded as fresh atoms new term destructors (and their simplification rules) defined using let-binding encoding of natural numbers: zero z and successor s are atoms, iterator  $it: X \to (X \to X) \to N \to X$  is a variable defined recursively

 $\nu z. \ \nu s. \text{ let } it = (!x \Rightarrow !f \Rightarrow (z \Rightarrow x \mid s !n \Rightarrow it@x@f@n \mid fail)) \text{ in } \dots$ 

interpreter for existing term constructors as reaction rules for new molecules interpreter for operation on references newR, getR and more

 $\begin{array}{l} \nu p. \ \nu r. \quad \text{molecule names for interpreted programs and local store} \\ \left\{ \left( \begin{array}{c} \left\{ \left( p \ (newR \ !x \ !x') \right) \right\} > \nu y. \left\{ \left( p \ (x'@y) \ \mid r \ y \ x \right) \right\}, \\ \left\{ \left( p \ (getR \ !y \ !x') \ \mid r \ !y \ !x \right) \right\} > \left\{ \left( p \ (x'@x) \ \mid r \ y \ x \right) \right\}, \\ \dots \end{array} \right) \right\} \end{array} \right\}$ 

restrict visibility of r to ensure that store is manipulated only by the interpreter

### **Conclusion: multi-lingual extensions and interpreters**

- new term constructors encoded as fresh atoms new term destructors (and their simplification rules) defined using let-binding
- Interpreter for existing term constructors as reaction rules for new molecules interpreter for operation on references newR, getR and more

 $\begin{array}{l} \nu p. \ \nu r. \quad \text{molecule names for interpreted programs and local store} \\ \left\{ \left( \begin{array}{c} \left\{ \left( p \ (newR \ !x \ !x') \right) \right\} > \nu y. \left\{ \left( p \ (x'@y) \ \mid r \ y \ x \right) \right\}, \\ \left\{ \left( p \ (getR \ !y \ !x') \ \mid r \ !y \ !x \right) \right\} > \left\{ \left( p \ (x'@x) \ \mid r \ y \ x \right) \right\}, \\ \dots \end{array} \right) \right\} \end{array} \right\}$ 

restrict visibility of r to ensure that store is manipulated only by the interpreter