

# “Automatic Translation of MP<sup>+</sup>V Systems to Register Machines”

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# General Idea

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# Brief Background

# Register Machines

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- one *computationally universal/Turing complete* model of computation among the several existent
- similar to (real) computer architecture: von Neumann architecture, memory bank, cache memory, assembly, FPGA, ...
-  several descriptions; ours is the Shepderson & Sturgis<sup>1</sup> + subprograms
  1.  $\text{CPY}(R_1, R_2) \equiv R_2 \leftarrow R_1$
  2.  $\text{ADD}(R_1, R_2, R_3) \equiv R_3 \leftarrow R_1 + R_2$
  3.  $\text{SUB}(R_1, R_2, R_3) \equiv R_3 \leftarrow R_1 - R_2$

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<sup>1</sup>Shepherdson, J. C. and Sturgis, H. E. *Computability of Recursive Functions*. Journal of ACM, 10, pp. 217–255.

# Register Machine: Specification

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- triple  $\mathcal{R} = (R, O, P)$ 
  - $R = \{R_1, R_2, \dots, R_m\}$  is the finite set of *registers* (with infinite capacity)
  - $O' = \overbrace{\{\text{INC, DEC, CLR, JMP, JZ, JNZ, HALT}\}}^{\text{Instructions}} \cup \overbrace{\{\text{CPY, ADD, SUB}\}}^{\text{Subprograms}}$  is the extended, finite set of operations and subprograms;
  - $P = (I_1, I_2, \dots, I_n)$  is the (finite) program

# Metabolic P Systems

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- a model of membrane computing (P systems)
- with particular features:
  - generative grammar for temporal series
  - discrete dynamical systems
  - deterministic execution 🙌
- completely described and studied in Manca's book<sup>2</sup> (and tomorrow's talk! 📅)

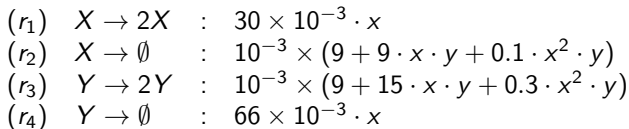
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<sup>2</sup>Manca, V. *Infobiotics: Information in Biotic Systems*. Springer Berlin Heidelberg. 2013.

# Metabolic P Systems: Example

## Lotka-Volterra

- Rules



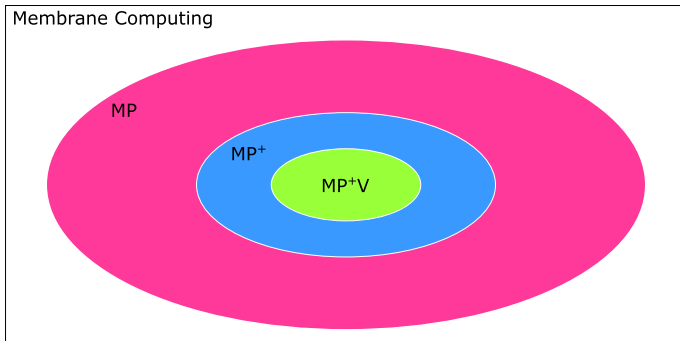
- Equation

$$\underbrace{\begin{bmatrix} x[t+1] \\ y[t+1] \end{bmatrix}}_{\vec{s}[t+1]} = \underbrace{\begin{bmatrix} x[t] \\ y[t] \end{bmatrix}}_{\vec{s}[t]} + \underbrace{\begin{bmatrix} 1 & -1 & 0 & 0 \\ 0 & 0 & 1 & -1 \end{bmatrix}}_{\text{Stoichiometric Matrix } \mathbb{A}} \times \underbrace{\begin{bmatrix} \varphi_1(\vec{s}[t]) \\ \varphi_2(\vec{s}[t]) \\ \varphi_3(\vec{s}[t]) \\ \varphi_4(\vec{s}[t]) \end{bmatrix}}_{\vec{U}[t]}$$



# The Universe of the Metabolic Systems

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## $MP^+V \subseteq MP^+$ Systems

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- $MP^+$  : subclass of MP that is semantically closer to biological metabolism
- restricts quantities and operations to positive numbers ( $\mathbb{N}, \mathbb{Q}^+, \mathbb{R}^+, \dots$ )
- and does it using two properties:
  1. every flux  $\geq 0$ , at any time
  2. fluxes cannot remove more quantities than available in the system, at each step

# MP<sup>+</sup> : Mathematically

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## Definition (MP<sup>+</sup> Grammar)

A MP<sup>+</sup> grammar  $G' = (M, R, I', \Phi')$  is a derivation from a (standard) MP grammar  $G = (M, R, I, \Phi)$  if its vector of initial values for substances  $I'$  has all components greater than zero and  $G'$  respects the following restrictions at every computational step  $t_i$ :

1.  $\varphi'(t_i) = \begin{cases} \varphi(t_i) & , \text{ if } \varphi(t_i) \geq 0 \\ 0 & , \text{ otherwise} \end{cases}$ , for all  $\varphi' \in \Phi'$  and their correspondents  $\varphi \in \Phi$ ;
2.  $\sum_{\varphi' \in \Phi'_x} \varphi'(t_i) \leq x$ , where the set of consuming fluxes of the metabolite  $x$  is defined as  $\Phi'_x = \{\varphi'_j : \text{mult}^-(x, r_j) > 0, \forall r_j \in R\}$ ; otherwise,  $\varphi'(t_i) = 0, \forall \varphi' \in \Phi'_x$  at the execution step  $t_i$ .

# MP<sup>+</sup>V Systems

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- MP<sup>+</sup>V : a minimalist MP<sup>+</sup> , **still** Turing complete!
- arose as a pattern on the equivalence between MP<sup>+</sup> and register machines<sup>3</sup>
- rule: at most **one single variable** at each side of it
- flux: either **one single variable** or a **subtraction of two variables**



# MP<sup>+</sup>V Systems: Formally Speaking

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## Definition (MP<sup>+</sup>V Grammar)

A MP<sup>+</sup>V grammar  $G = (M, R, I, \Phi)$  is a MP<sup>+</sup> one in which:

1.  $\forall r \in R$  and  $v', v'' \in M$ ,  $r$  must have one of the following shapes:
  - 1.1  $\emptyset \rightarrow v''$ ;
  - 1.2  $v' \rightarrow \emptyset$ ; or
  - 1.3  $v' \rightarrow v''$ ;
2.  $\forall \varphi \in \Phi$  and  $m', m'' \in M$ , the flux has either the form  $\varphi = m'$  or  $\varphi = m' - m''$ .

# The Translation

# Translation: Disclaimer

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- no surprises:  $MP^+V \equiv$  register machine  $\equiv$  Turing machine  $\Rightarrow \exists$  compilation 😊
- however, the current focus is **compilation**, not computational power<sup>3</sup>. 😊

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<sup>3</sup>Guiraldelli, R. and Manca, V. *The Computational Universality of Metabolic Computing*. arXiv:1505.02420. 2015.

# Not So Fast: Problems

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MP systems differ from register machine in three main ways:

1. unordered application of rules
2. parallel application of rules
3. positivity control



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Solution 1–2

**Command block** and/or Monad

# Not So Fast: Problems

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MP systems differ from register machine in two main ways:

1. application of rules
2. positivity control

## Solution 1

**Command block** and/or Monad

# Not So Fast: Problems

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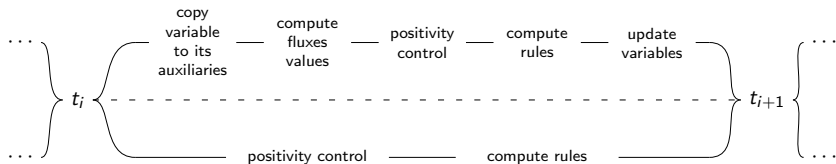
1. application of rules
2. positivity control

## Solution 2

Inclusion of a **subprogram**

## Runtime of a Computational Step

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**Figure:** Representation of a computation step MP<sup>+</sup>V systems (lower part) and its equivalent register machine (upper part).

# Standard $MP^+V$ Rules

## Rule $\emptyset \rightarrow V_1$

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$\emptyset \rightarrow V_1 : \varphi$

- 1 ADD( $R_{V_1}, R_\varphi, R_{aux}$ )
- 2 CPY( $R_{aux}, R_{V_1}$ )

## Rule $V_1 \rightarrow \emptyset$

---

$V_1 \rightarrow \emptyset : \varphi$

- 1 SUB( $R_{V_1}, R_\varphi, R_{aux}$ )
- 2 CPY( $R_{aux}, R_{V_1}$ )

## Rule $V_1 \rightarrow V_2$

---

$$V_1 \rightarrow V_2 : \varphi$$
$$\equiv$$
$$\begin{cases} V_1 \rightarrow \emptyset : \varphi \\ \emptyset \rightarrow V_2 : \varphi \end{cases}$$

- 1 SUB( $R_{V_1}, R_\varphi, R_{aux}$ )
- 2 CPY( $R_{aux}, R_{V_1}$ )
- 3 ADD( $R_{V_2}, R_\varphi, R_{aux}$ )
- 4 CPY( $R_{aux}, R_{V_2}$ )



## Rule $V_1 \rightarrow HALT$

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$V_1 \rightarrow HALT : \varphi$

- 1 JNZ( $R_{HALT}$ , 3)
- 2 JMP(4)
- 3 HALT
- 4 SUB( $R_{V_1}$ ,  $R_\varphi$ ,  $R_{aux}$ )
- 5 CPY( $R_{aux}$ ,  $R_{V_1}$ )
- 6 ADD( $R_{HALT}$ ,  $R_\varphi$ ,  $R_{aux}$ )
- 7 CPY( $R_{aux}$ ,  $R_{HALT}$ )

# The “Exoskeleton” of MP<sup>+</sup>V

# Exoskeleton Is **The** Important Part

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- With experience, basic rules arises fast
  - HALT requires some reasoning
- Exoskeleton is the tricky part
  - always there to ensure the proper/correct overall execution
  - the hidden dynamics of the system
  - 4/5 of the process, most of the generated source code

## The Exoskeleton Rules

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1. copy variables values to auxiliaries registers
2. compute fluxes values for current computational step
3. perform *positivity control* on every rule
4. update the variables values with computed ones
5. loop the systems up to fixed-point  $\text{HALT} \neq 0$

# The Exoskeleton Rules

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5. *loop the systems up to fixed-point HALT  $\neq 0$*

## Values Update in Computational Step

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- copy variables values to auxiliaries registers
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## Values Update in Computational Step

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- copy variables values to auxiliaries registers

$$\begin{aligned} &\forall V \in M, \\ &\forall t \in \mathbb{N}, \\ &\exists V_{aux} : V_{aux}|_{t-} \leftarrow V|_t \end{aligned} \quad 1 \quad \text{CPY}(R_V, R_{V_{aux}})$$

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## Values Update in Computational Step

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- copy variables values to auxiliaries registers
- compute fluxes values for current computational step

$$\begin{aligned} &\forall \varphi \in \Phi, \\ &\forall t \in \mathbb{N}, \\ &\varphi[t] = \varphi(\vec{V}|_t) \end{aligned}$$

```
if  $\varphi = V$  then
  1  CPY( $R_V, R_\varphi$ )
else       $\triangleright$  Hence,  $\varphi = V_1 - V_2$ 
  1  SUB( $R_{V_1}, R_{V_2}, R_\varphi$ )
end if
```

- update the variables values with computed ones



## Values Update in Computational Step

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- copy variables values to auxiliaries registers
- compute fluxes values for current computational step
- update the variables values with computed ones

$$\begin{aligned} &\forall V \in M, \\ &\forall t \in \mathbb{N}, \\ &\exists V_{aux} : V|_{t+1} \leftarrow V_{aux}|_{t+} \end{aligned} \quad 1 \quad \text{CPY}(R_{V_{aux}}, R_V)$$

# Positivity Control

---

- must satisfy two constraints
  1. fluxes must always belong to the set of positive number

$$\varphi'(t_i) = \begin{cases} \varphi(t_i) & , \text{ if } \varphi(t_i) \geq 0 \\ 0 & , \text{ otherwise} \end{cases}$$

2. the sum of all consuming fluxes for a given variable must be smaller or equal to the amount of the variable

$$\sum_{\varphi' \in \Phi'_x^-} \varphi'(t_i) \leq x$$

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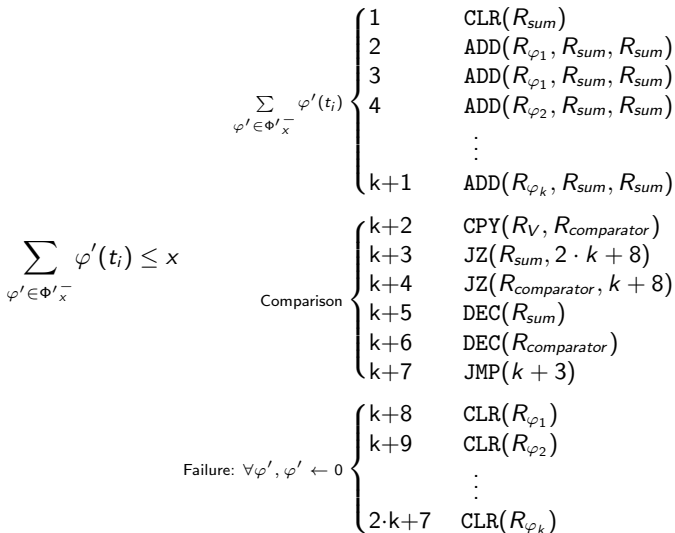
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$$\sum_{\varphi' \in \Phi'_x} \varphi'(t_j) \leq x$$

# Positivity Control



## Perpetuum Mobile, or Not

---

- as dynamics, it shouldn't stop—unless it stucks in a fixed-point<sup>3</sup> 😊
- as computational process, it **must** stop—unless of halting problem ⚠️
- is it possible to differ `while(true)` from `for(i = 0; i < limit; i++)` at *compile time*?

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- is it possible to differ `while(true)` from `for(i = 0; i < limit; i++)` at *compile time*?
  - Yes, it is!
  - Trick #1: require HALT variable and  $V_i \rightarrow HALT$  rule! 🎉
  - Trick #2:  $HALT \neq 0$  is the *signal to halt*—and since there aren't any  $HALT \rightarrow V_i$  rules, it is guaranteed

## Perpetuum Mobile, or Not

---

$\nexists$ HALT variable

$\vee$

$\nexists V_i \rightarrow HALT$

1

CPY(..., ...)

$\vdots$

$\ell-1$

JMP(1)

$\ell$

HALT

---

$\exists$ HALT variable

$\wedge$

$\exists V_i \rightarrow HALT$

1

JNZ( $R_{HALT}, \ell$ )

$\vdots$

$\ell-1$

JMP(1)

$\ell$

HALT





# Pseudo-code of a Translation from $MP^+V$ to Register Machine

```
while  $R_{HALT} = 0$  do
  for all variable  $v \in M$  do           ▷ copy variables to auxiliaries
     $R_{v'} \leftarrow R_v$ 
  end for
  for all flux  $\varphi \in \Phi$  do             ▷ compute fluxes
     $R_\varphi \leftarrow \varphi(t_i)$ 
  end for
  for all variable  $v \in M$  do           ▷ positivity control property
    for all flux  $\varphi_v^- \in \Phi_v^-$  do
       $R_{sum} \leftarrow R_{sum} + R_{\varphi_v^-}$ 
    end for
    if  $R_{sum} > v$  then
      for all flux  $\varphi_v^- \in \Phi_v^-$  do
         $R_{\varphi_v^-} \leftarrow 0$ 
      end for
    end if
  end for
  for all rule  $r$  do                   ▷ compute rules
    if  $r$  is of the form  $\emptyset \rightarrow v : \varphi$  then
       $R_{v'} \leftarrow R_{v'} + \varphi$ 
    else if  $r$  is of the form  $v \rightarrow \emptyset : \varphi$  then
       $R_{v'} \leftarrow R_{v'} - \varphi$ 
    else                               ▷ hence, it must be of the form  $v_1 \rightarrow v_2 : \varphi$ 
       $R_{v_1'} \leftarrow R_{v_1'} + \varphi$ 
       $R_{v_2'} \leftarrow R_{v_2'} - \varphi$ 
    end if
  end for
  for all variable  $v \in M$  do           ▷ update variables
     $R_v \leftarrow R_{v'}$ 
  end for
end while
```

# Conclusions



# Conclusions

---

- paradigm change possibly brings *big exoskeleton*
  -  *critical insight* over the failures of the past
  - metabolic  $\mapsto$  computational
  - parallel  $\mapsto$  sequential
  - local (pair substances)  $\mapsto$  global (execution control)
- $MP \supseteq MP^+ \supseteq MP^+V$  , and all computationally universal 


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# Conclusions

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- bidirectional, automatic translation between von Neumann architecture (register machine) and metabolic computing (MP+V systems)
  - Can we extend it to real metabolism? Yes, we can. . . Theoretically.<sup>3</sup>
- open door  for new translations, including *hardware description languages*, programming languages, visual representations, etc
- lead the way to implementation of circuits based on metabolic (MP) systems

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Thank you!



Obrigado!



¡Gracias!



Grazie!



Ačiū!