

# Membrane Computing: Power, Efficiency, Applications

## (A Quick Introduction)

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## Summary:

- generalities
- the basic idea
- examples
- classes of P systems
- types of results
- types of applications
  - applications in biology
  - Nishida's membrane algorithms

Goal: abstracting computing models/ideas from the structure and functioning of living cells (and from their organization in tissues, organs, organisms)

hence not producing models for biologists (although, this is now a tendency)

result:

- distributed, parallel computing model
- compartmentalization by means of membranes
- basic data structure: multisets (but also strings; recently, numerical variables)

## References:

- Gh. Păun, Computing with Membranes. *Journal of Computer and System Sciences*, 61, 1 (2000), 108–143, and *Turku Center for Computer Science-TUCS Report No 208*, 1998 ([www.tucs.fi](http://www.tucs.fi))  
ISI: “fast breaking paper”, “emerging research front in CS” (2003)  
<http://esi-topics.com>
- Gh. Păun, *Membrane Computing. An Introduction*, Springer, 2002
- G. Ciobanu, Gh. Păun, M.J. Pérez-Jiménez, eds., *Applications of Membrane Computing*, Springer, 2006
- Website: <http://psystems.disco.unimib.it>

(Yearly events: BWMC (February), WMC (summer), TAPS/WAPS (fall))

## SOFTWARE AND APPLICATIONS:

[http://www.dcs.shef.ac.uk/~marian/PSimulatorWeb/P\\_Systems\\_applications.htm](http://www.dcs.shef.ac.uk/~marian/PSimulatorWeb/P_Systems_applications.htm)

[www.cbmc.it](http://www.cbmc.it) – PSim2.X simulator

Verona (Vincenzo Manca: [vincenzo.manca@univr.it](mailto:vincenzo.manca@univr.it))

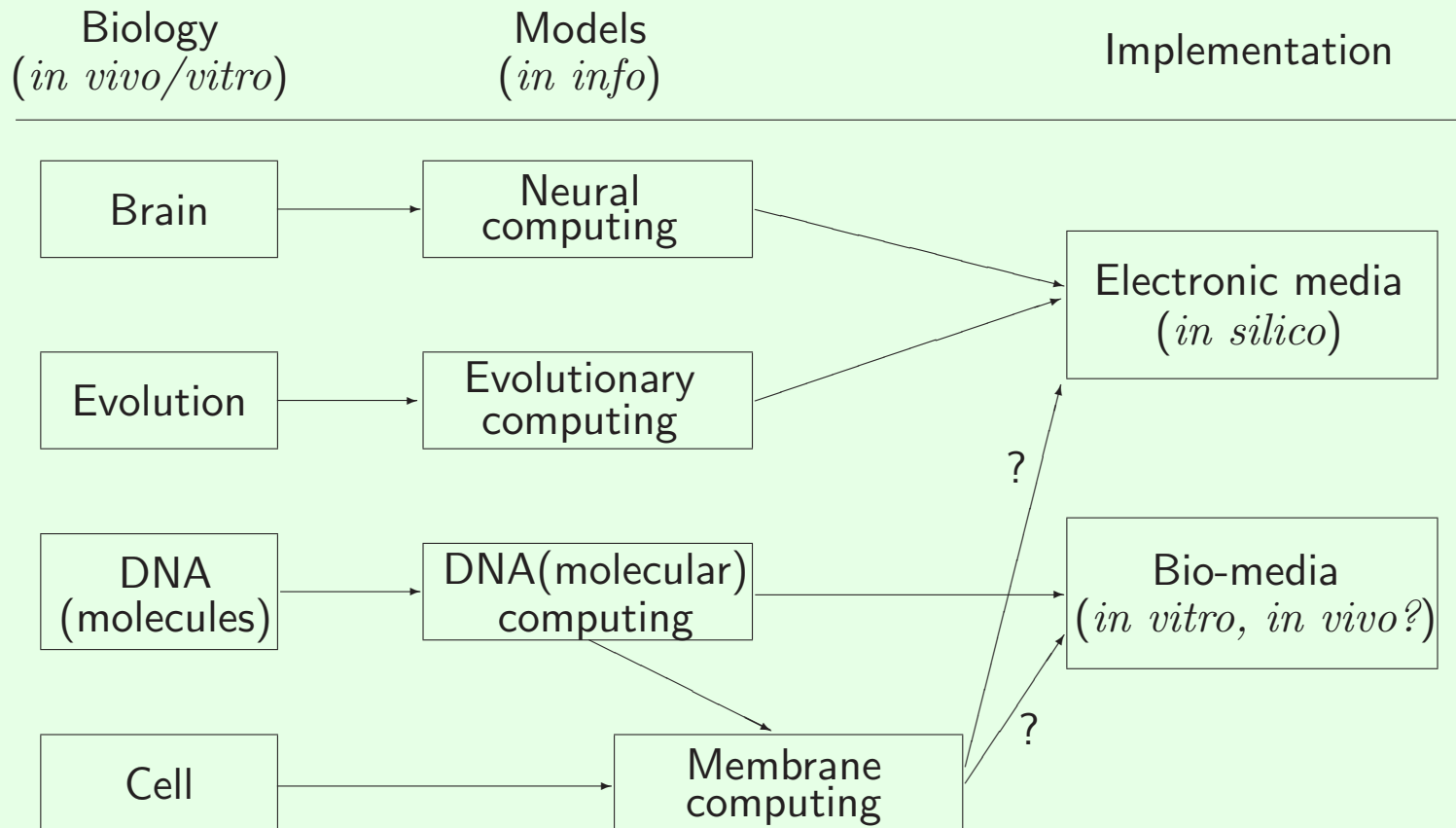
Sheffield (Marian Gheorghe: [M.Gheorghe@dcs.shef.ac.uk](mailto:M.Gheorghe@dcs.shef.ac.uk))

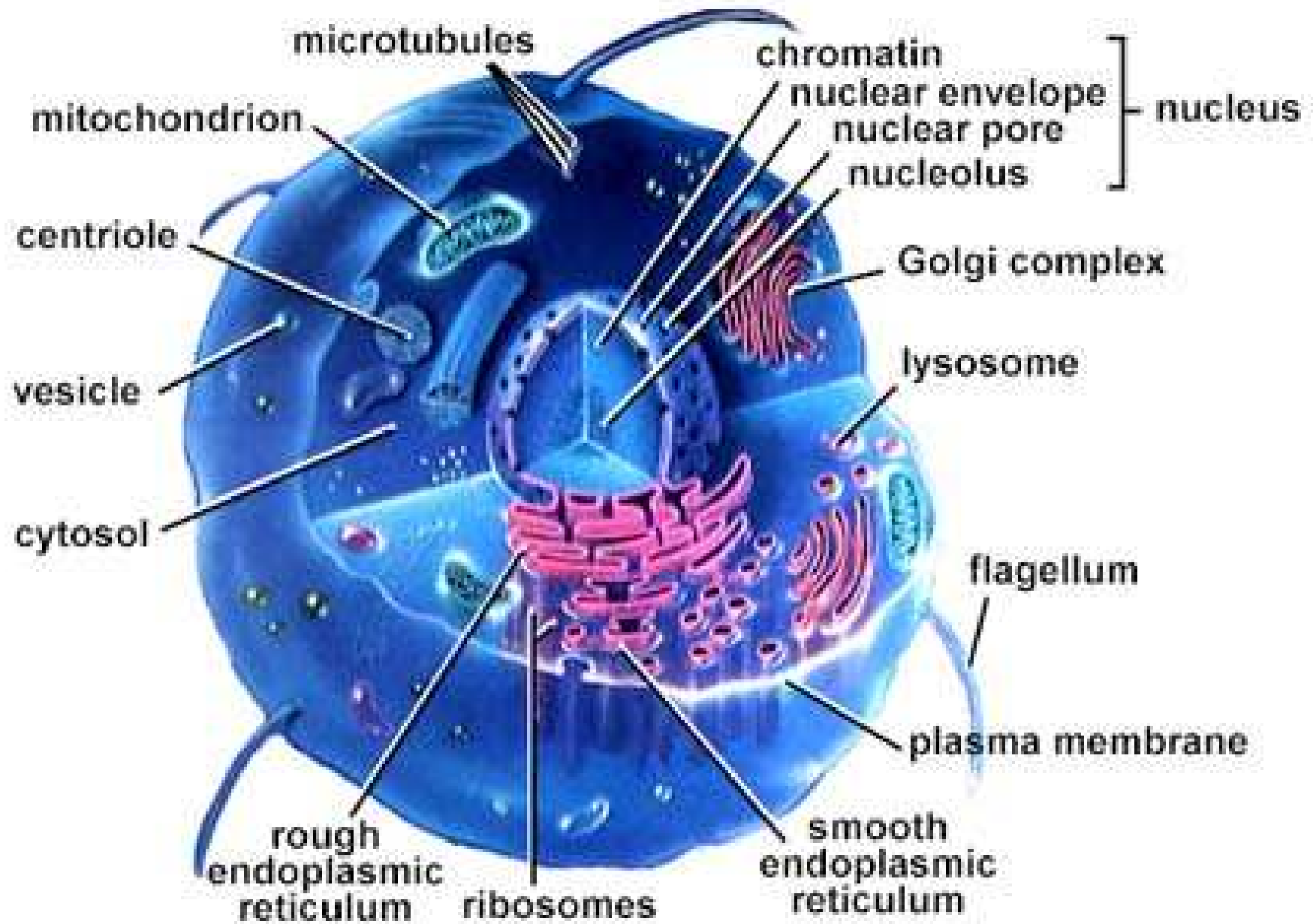
Sevilla (Mario Pérez-Jiménez: [marper@us.es](mailto:marper@us.es))

Milano (Giancarlo Mauri: [mauri@disco.unimib.it](mailto:mauri@disco.unimib.it))

Nottingham, Leiden, Vienna, Evry, Iași

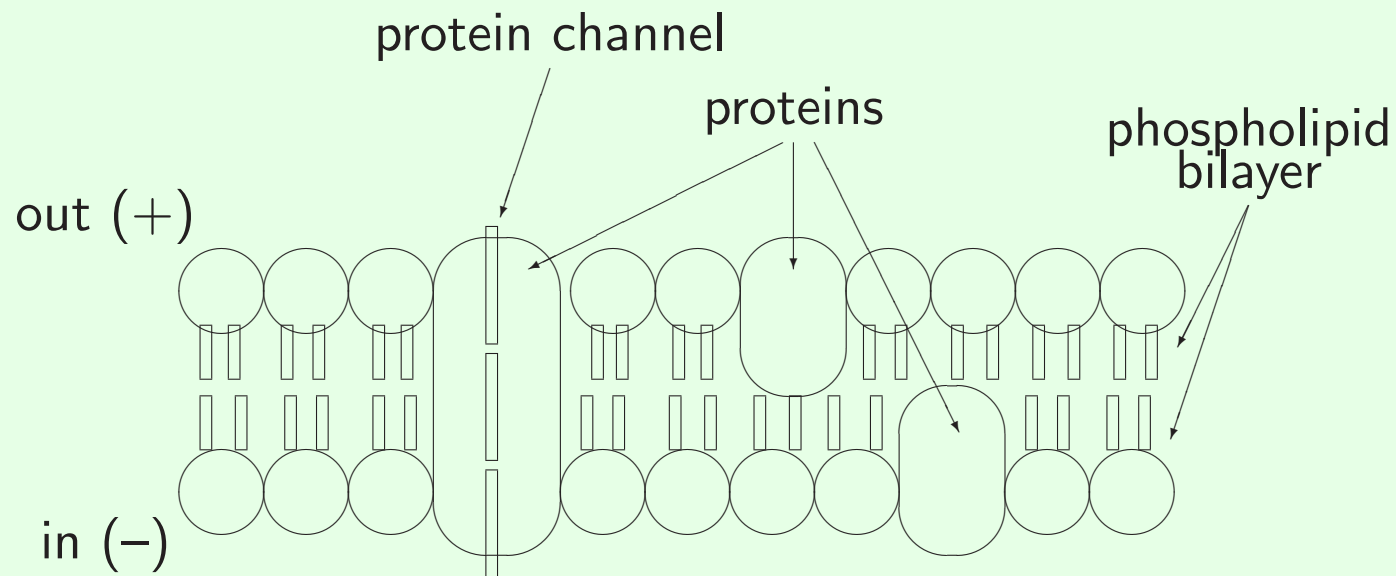
## Natural computing



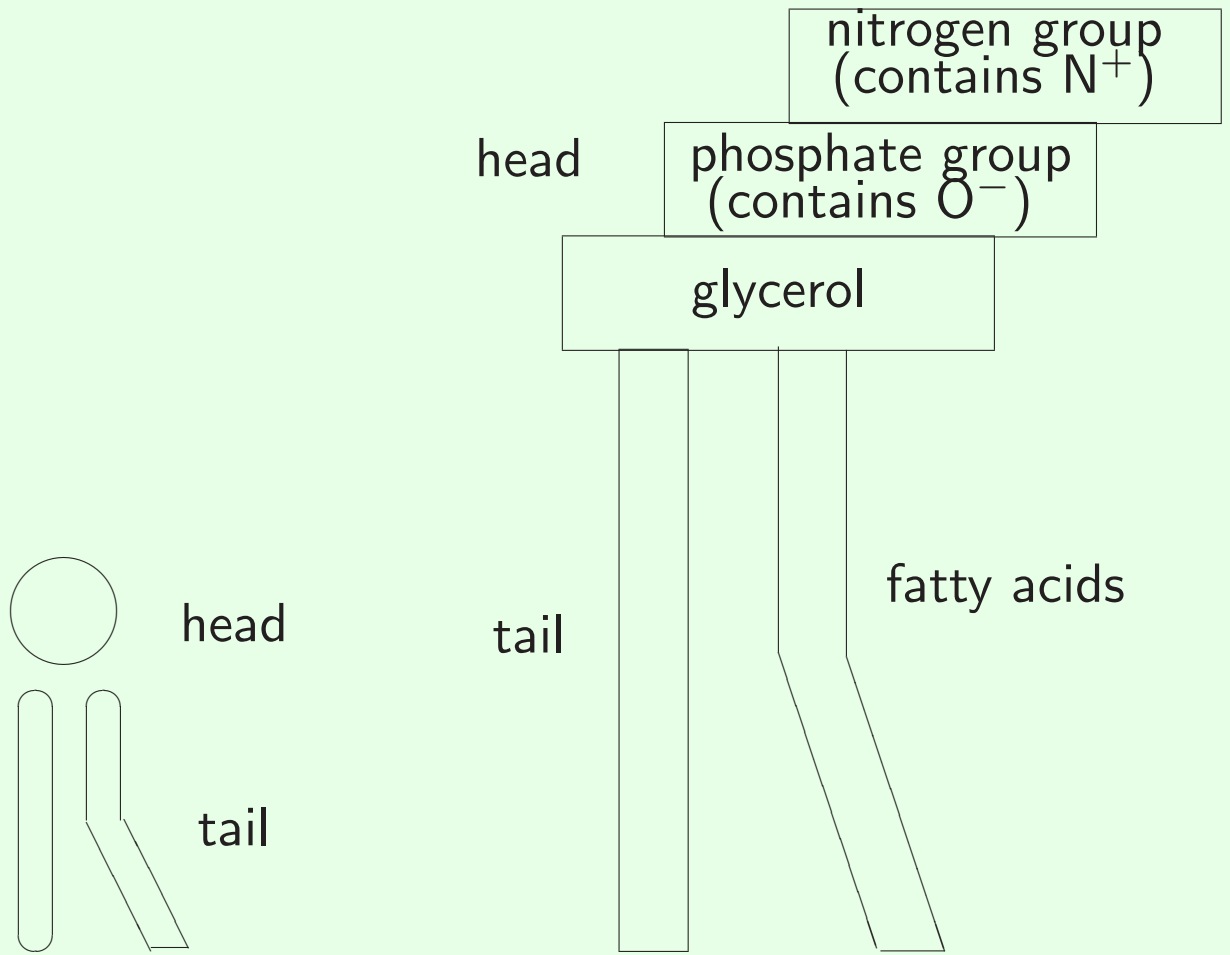


## THE STRUCTURE OF PLASMA MEMBRANE

The *fluid-mosaic model*, S. Singer and G. Nicolson, 1972

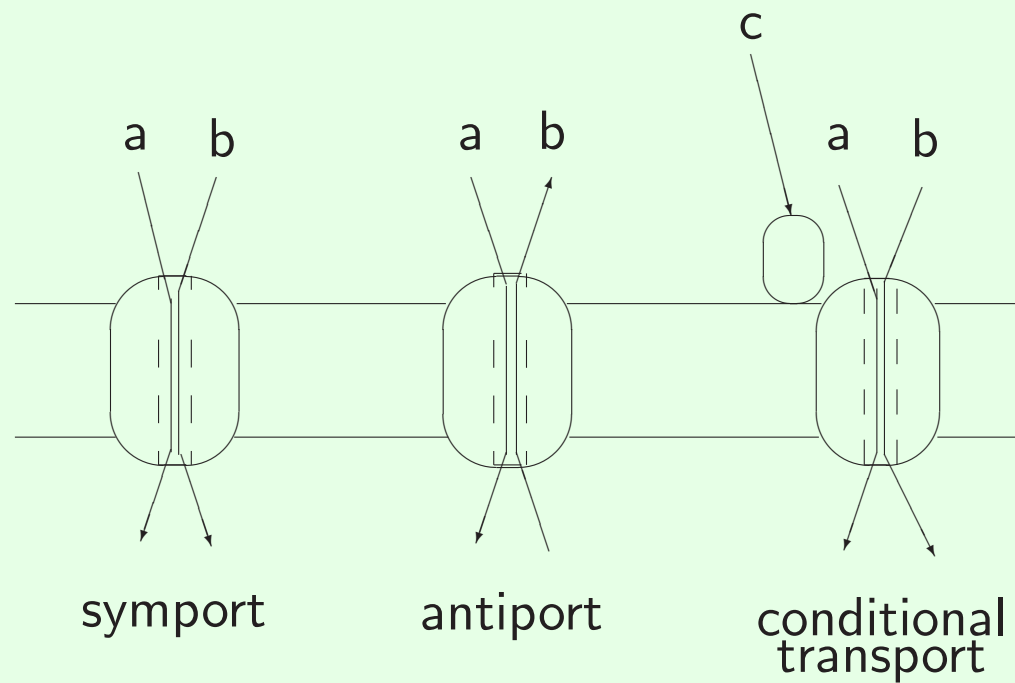




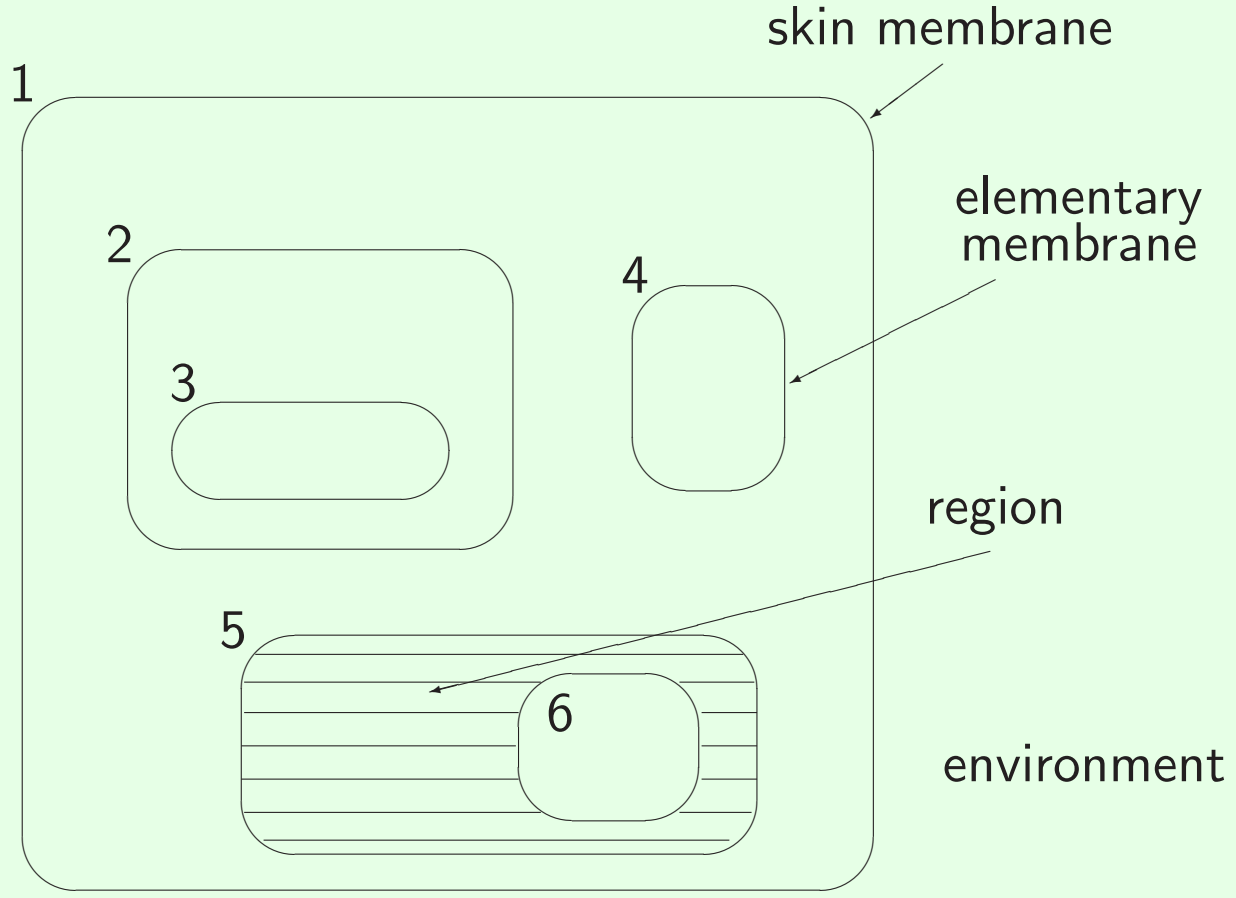


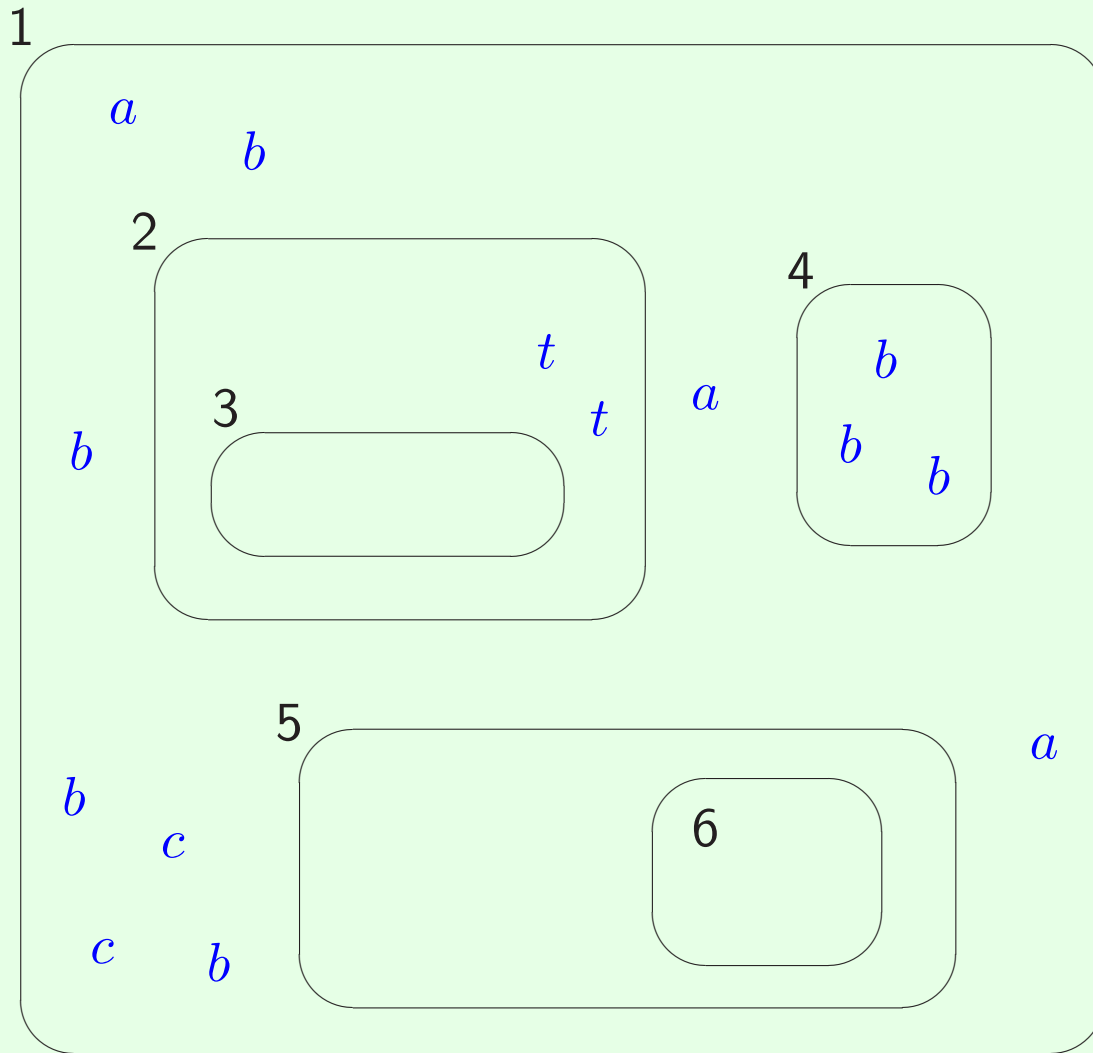
Trans-Membrane Transport  
passive (diffusion – concentration based)  
active (protein channels)  
vesicle-mediated

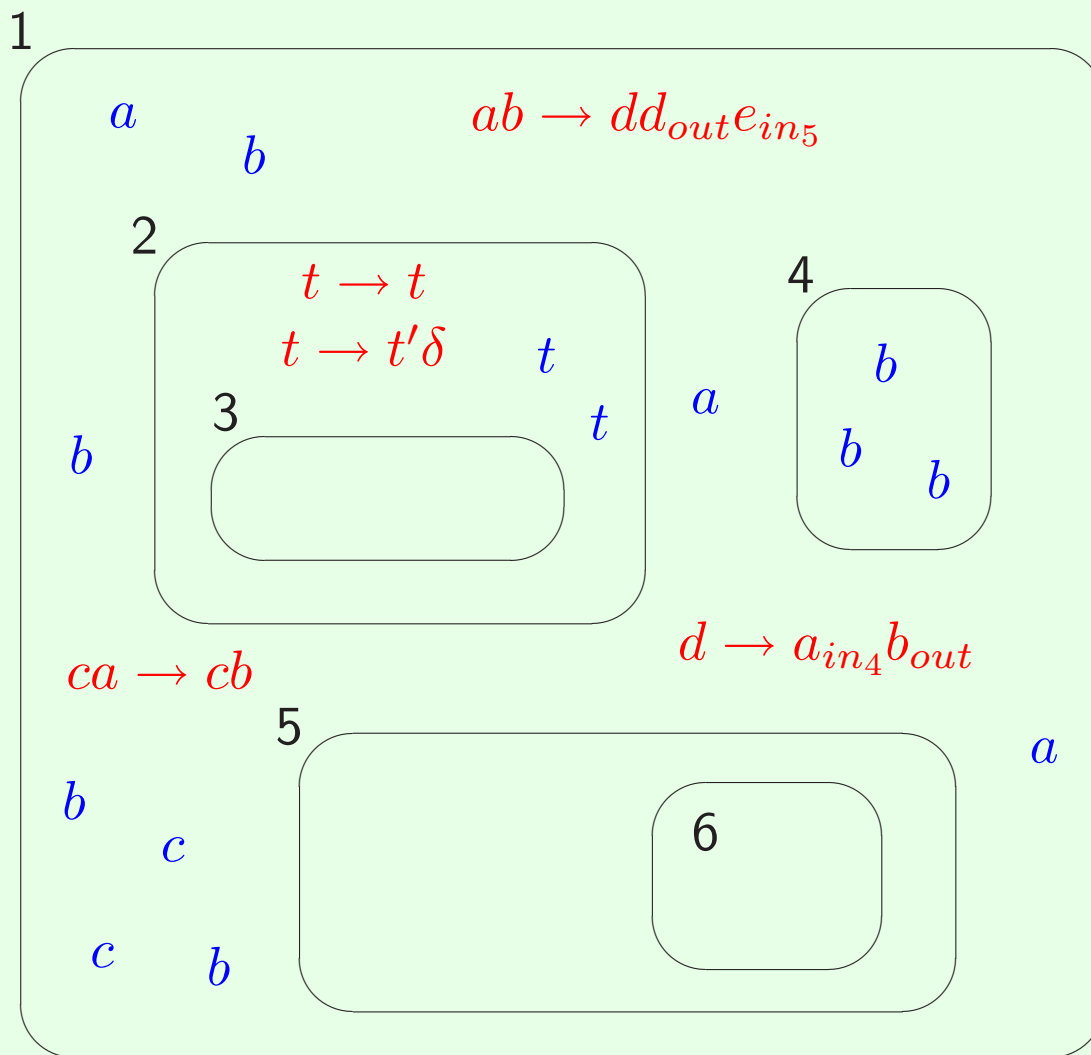
Important case of active transport: coupled transport  
symport  
antiport



# THE BASIC IDEA



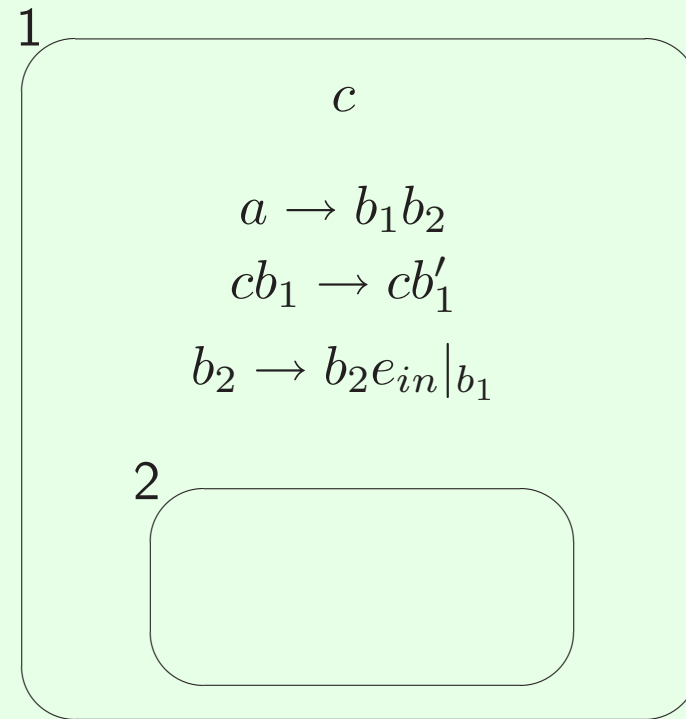




## Functioning (basic ingredients):

- nondeterministic choice of rules and objects
- maximal parallelism
- transition, computation, halting
- internal output, external output, traces

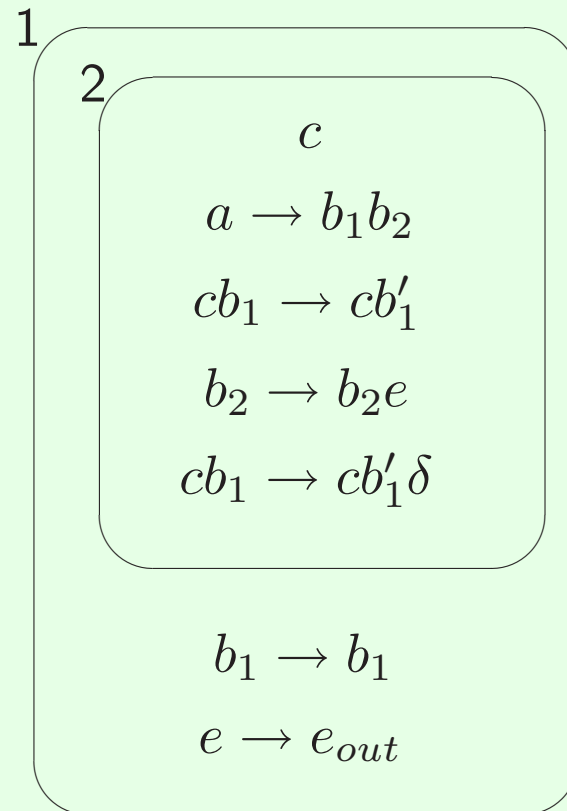
## EXAMPLES



Computing system:  $n \longrightarrow n^2$  (catalyst, promoter, determinism, internal output)

Input (in membrane 1):  $a^n$

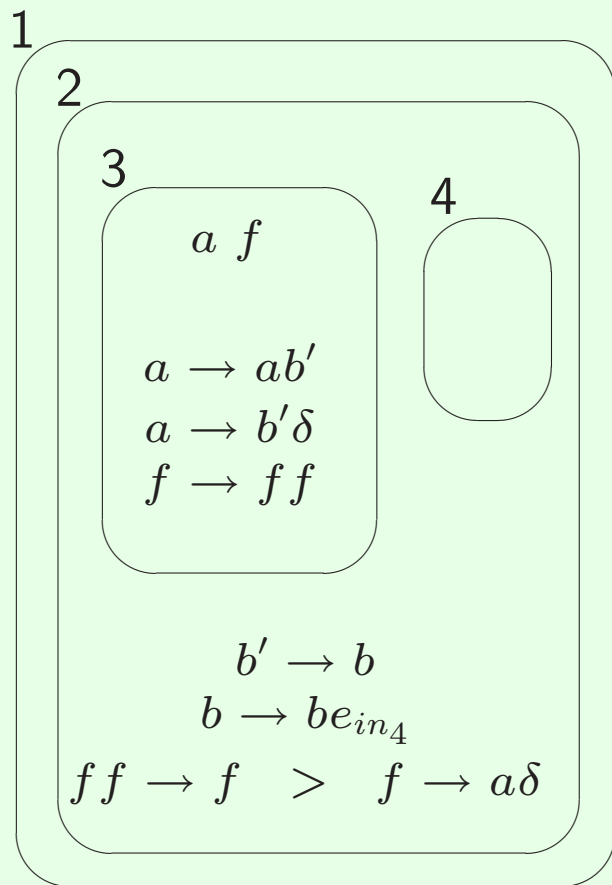
Output (in membrane 2):  $e^{n^2}$



The same function ( $n \longrightarrow n^2$ ), with catalyst, dissolution, nondeterminism, external output



Generative mode :  $\{n^2 \mid n \geq 1\}$



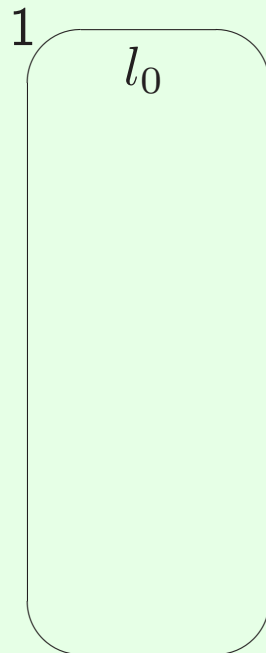
0	$af$	
1	$ab'ff$	
...	...	
$m \geq 0$	$ab'^m f^{2^m}$	
$m + 1$	$b'^{m+1} f^{2^{m+1}}$	$\delta$
$m + 2$	$b^{m+1} f^{2^m}$	
$m + 3$	$b^{m+1} f^{2^{m-1}}$	$e_{in_4}^{m+1}$
...	...	...
$2m + 1$	$b^{m+1} f^2$	$e_{in_4}^{m+1}$
$2m + 2$	$b^{m+1} f$	$e_{in_4}^{m+1}$
$2m + 3$	$b^{m+1} a\delta$	$e_{in_4}^{m+1}$
$m + 1$ times	<b>HALT!</b>	

$$(m + 1) \times (m + 1)$$

$$N(\Pi) = \{n^2 \mid n \geq 1\}$$

# SIMULATING A REGISTER MACHINE $M = (m, B, l_0, l_h, R)$

$$E = \{a_r \mid 1 \leq r \leq m\} \cup \{l, l', l'', l''', l^{iv} \mid l \in B\}$$



$$\left. \begin{array}{l} (l_1, out; a_r l_2, in) \\ (l_1, out; a_r l_3, in) \end{array} \right\} \text{ for } l_1 : (\text{add}(r), l_2, l_3)$$

$$\left. \begin{array}{l} (l_1, out; l'_1 l''_1, in) \\ (l'_1 a_r, out; l'''_1, in) \\ (l''_1, out; l^{iv}_1, in) \\ (l^{iv} l'''_1, out; l_2, in) \\ (l^{iv} l'_1, out; l_3, in) \end{array} \right\} \text{ for } l_1 : (\text{sub}(r), l_2, l_3)$$

$$(l_h, out)$$

Symport/antiport rules (of weight 2)

## A bird eye view to the MC jungle:

- cell-like, tissue-like, neural-like (spiking neural) systems
- symbols, strings, arrays, numerical variables, etc
- multisets, sets, fuzzy
- multiset rewriting, symport/antiport, membrane evolving, combinations
- controls: priority, promoters, inhibitors,  $\delta$ ,  $\tau$ , activators,
- maximal, bounded, minimal parallelism, sequential
- synchronized, time-, clock-free
- generating, accepting, computing/translating, dynamical system
- computing power, computing efficiency, others
- implementations/simulations
- applications: biology/medicine, economics, optimization, computer graphics, linguistics, computer science, cryptography, etc
- quantum
- etc (e.g., brane-membrane bridge)

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- etc. (e.g., brane-membrane bridge)

## Types of rules:

$u \rightarrow v$  with targets in  $v$   
(possibly conditional: promoters or inhibitors)  
particular cases:  $ca \rightarrow cu$  (catalytic)  
 $a \rightarrow u$  (non-cooperative)

$(ab, in), (ab, out)$  – symport (in general,  $(x, in), (x, out)$ )  
 $(a, in; b, out)$  – antiport (in general,  $(u, in; v, out)$ )

$u]_i.v \rightarrow u']_i.v'$  – boundary (Manca, Bernardini)

$ab \rightarrow a_{tar_1}b_{tar_2}$  – communication (Sosik)

$ab \rightarrow a_{tar_1}b_{tar_2}c_{come}$

$a \rightarrow a_{tar}$

$$\begin{aligned}
a[ ]_i &\rightarrow [b]_i \\
[a]_i &\rightarrow b[ ]_i \\
[a]_i &\rightarrow b \\
a &\rightarrow [b]_i \\
[a]_i &\rightarrow [b]_j [c]_k \\
[a]_i [b]_j &\rightarrow [c]_k \\
[a]_i [ ]_j &\rightarrow [[b]_i]_j \\
[[a]_i]_j &\rightarrow [b]_i [ ]_j \\
[u]_i &\rightarrow [ ]_i [u]_{@j} \\
[Q]_i &\rightarrow [O - Q]_j [Q]_k
\end{aligned}$$

go in  
go out  
membrane dissolution  
membrane creation  
membrane division  
membrane merging  
endocytosis  
exocytosis  
gemmation  
separation

and others

## Results:

- characterization of **Turing computability** ( $RE$ ,  $NRE$ ,  $PsRE$ )  
Examples: by catalytic P systems (2 catalysts) [Sosik, Freund, Kari, Oswald]  
by (small) symport/antiport P systems [many]
- polynomial solutions to **NP-complete problems** (by using an exponential workspace created in a “biological way”: membrane division, membrane creation, string replication, etc) [Sevilla team], [Madras team], [Obtulowicz], [Alhazov, Pan] etc
- other types of **mathematical results** (normal forms, hierarchies, determinism versus nondeterminism, complexity) [Ibarra group]
- **connections** with ambient calculus, Petri nets, X-machines, quantum computing, lambda calculus, brane calculus, etc [many]
- **simulations** and implementations
- **applications**

## Open problems, research topics:

Many: see the P page

- borderlines: universality/non-universality, efficiency/non-efficiency (the power of 1 catalyst, the role of polarizations, dissolution, uniform versus semi-uniform, etc.)
- semantics (events, causality, etc.)
- neural-like systems (more biology, complexity, applications, etc.)
- user friendly, flexible software for bio-applications
- MC and economics



## Applications:

- biology, medicine (continuous versus discrete mathematics) [Sevilla, Verona, Milano, Sheffield, etc]
- computer science (computer graphics, sorting/ranking, 2D languages, cryptography, general model of distributed-parallel computing) [many]
- linguistics (modeling framework, parsing) [Tarragona]
- optimization (membrane algorithms [Nishida, 2004])
- economics ([Warsaw group], [R. Păun], [Vienna group])

## A typical application in biology/medicine:

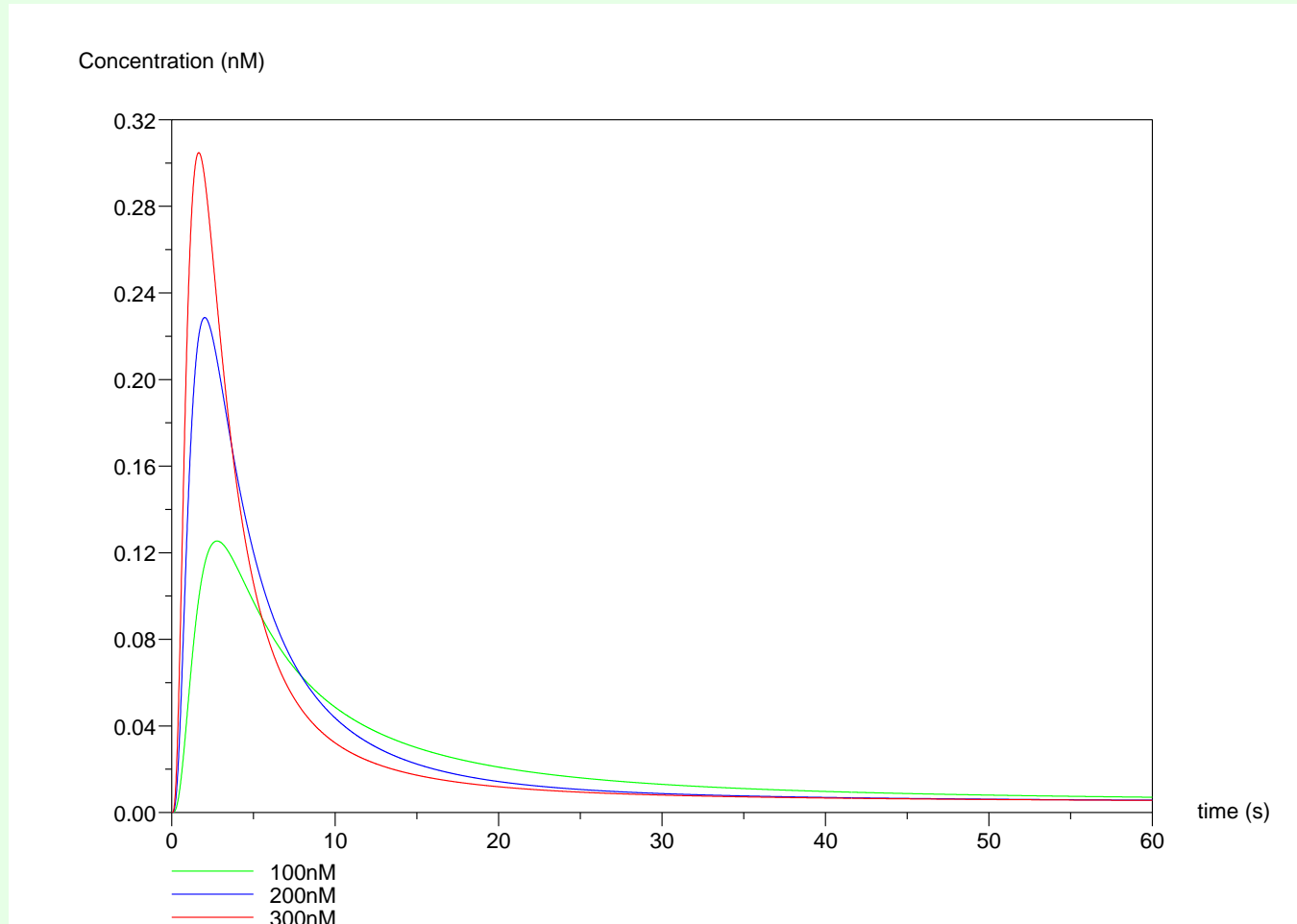
M.J. Pérez–Jiménez, F.J. Romero–Campero:

A Study of the Robustness of the EGFR Signalling Cascade Using Continuous Membrane Systems.

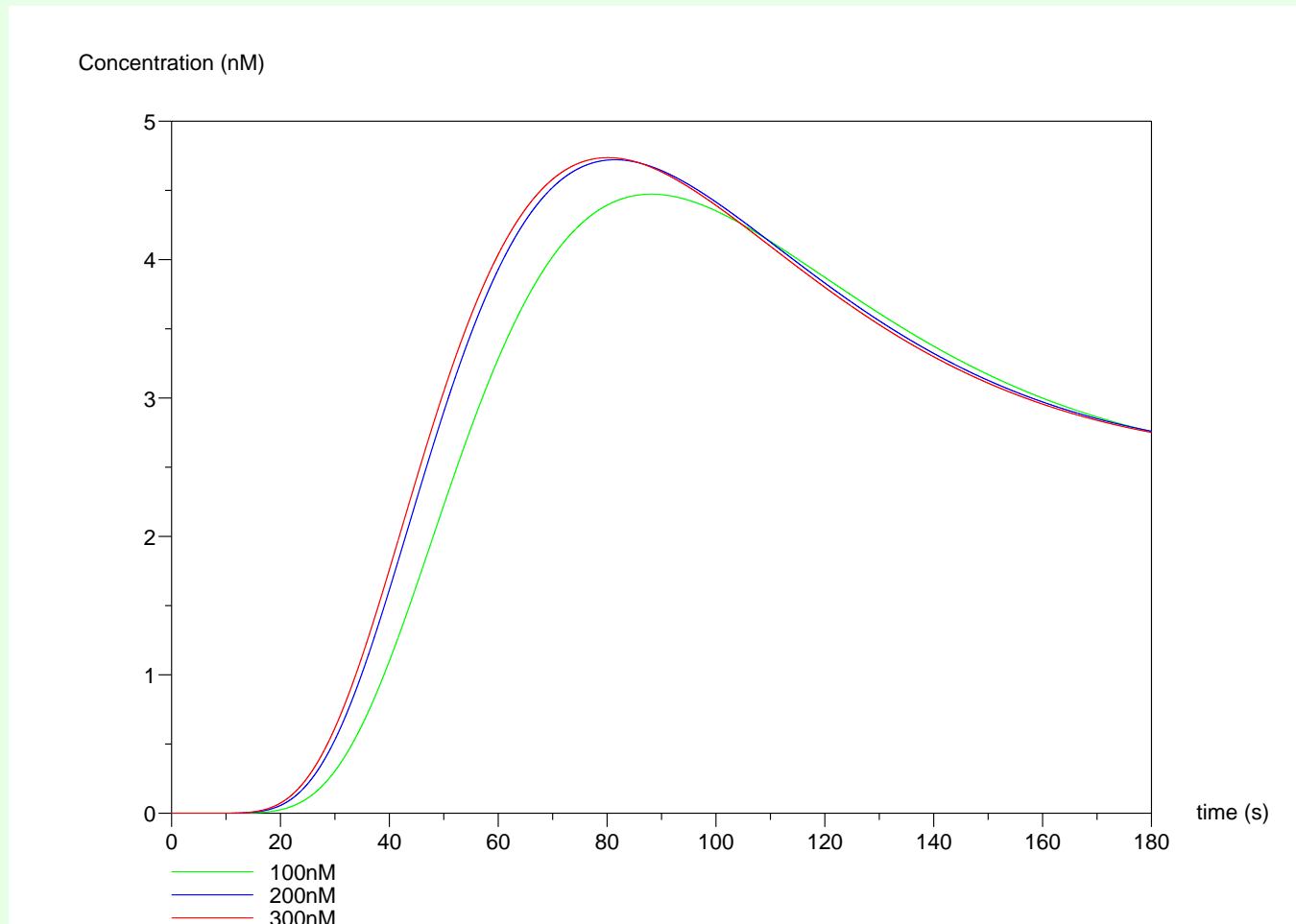
In *Mechanisms, Symbols, and Models Underlying Cognition. First International Work-Conference on the Interplay between Natural and Artificial Computation, IWINAC 2005* (J. Mira, J.R. Alvarez, eds.), LNCS 3561, Springer, Berlin, 2005, 268–278.

- 60 proteins, 160 reactions/rules
- reaction rates from literature
- results as in experiments

## Typical outputs:



The EGF receptor activation by auto-phosphorylation  
(with a rapid decay after a high peak in the first 5 seconds)



The evolution of the kinase MEK  
(proving a surprising robustness of the signalling cascade)

### Nishida's membrane algorithms:

- candidate solutions in regions, processed locally (local sub-algorithms)
- better solutions go down
- static membrane structure – dynamical membrane structure
- two-phases algorithms

Excellent solutions for Travelling Salesman Problem (benchmark instances)

- rapid convergence
- good average and worst solutions (hence reliable method)
- in most cases, better solutions than simulated annealing

Still, many problems remains: check for other problems, compare with sub-algorithms, more membrane computing features, parallel implementations (no free lunch theorem)

**Recent:** L. Huang, N. Wang, J. Tao; G. Ciobanu, D. Zaharie; A. Leporati, D. Pagani

# Thank you!

...and please do not forget: <http://psystems.disco.unimib.it>

(with mirrors in China: <http://bmc.hust.edu.cn/psystems>,  
<http://bmchust.3322.org/psystems>)