

Applications of membrane systems in population biology and ecological modelling

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- 1 Introduction
- 2 A P system based modeling framework
- 3 Example: Tritrophic Interactions
- 4 Simulation algorithms
- 5 Simulation results
- 6 Conclusions and future work



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Membrane computing

New modeling framework

- P Systems → modeling framework
 - **Ecosystems**
- Randomness → probabilistic strategies

Simulation algorithms

- Reproduce the behaviour of the models
- Validation
- Virtual experimentation

Software

- Implements the algorithms
- GUI for the end-user

Modeling Biological Phenomena

Diferents aproximations

- Ordinary Differential Equations (ODEs)
- Petri Nets (Goss, 1998)
- Agent Based Systems (Holcombe, 2003)
- Process Algebra, π -calculus (A. Regev, E. Shapiro, 2004)
- Bioambients (L. Cardelli, E.M. Panina, A. Regev, W. Silvermann, E. Shapiro, 2004)
- Brane Calculus (L. Cardelli, 2005)



Modeling Biological Phenomena

Desirable properties

- Relevance
- Understandability
- Extensibility
- Computability / treatability

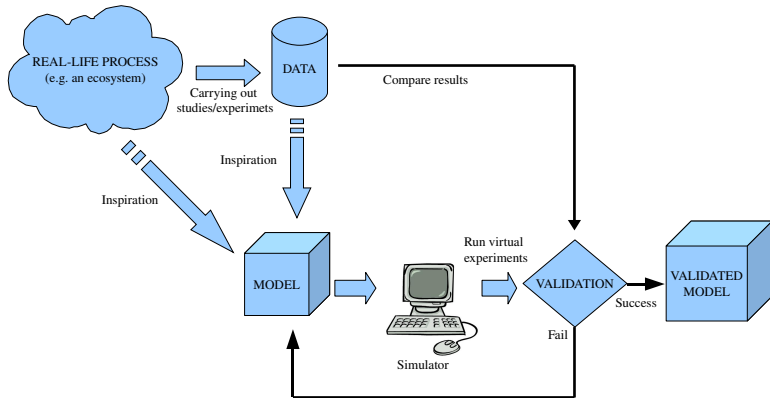
P systems

Modeling framework that meets these requirements.



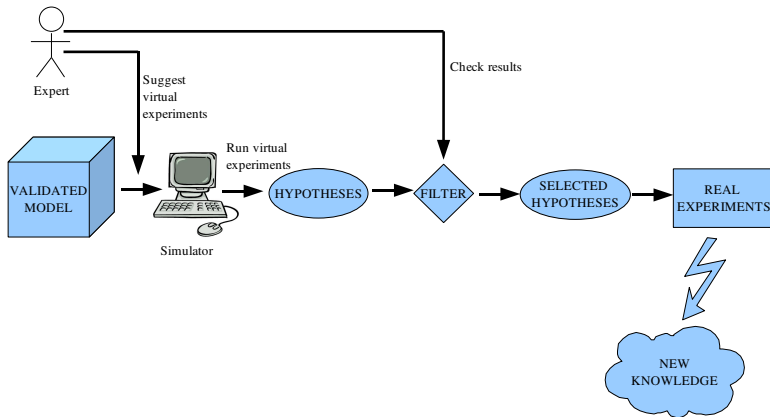
Modeling ecosystems

Validation process



Modeling ecosystems

Virtual Experiments



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Need to define a new variant of P Systems

- Cooperation
- Randomness
- Communication between environments
- Membrane polarization

A P system based modeling framework

- A skeleton of an extended P system with active membranes of degree $q \geq 1$,

$$(\Gamma, \mu, R)$$

- A probabilistic functional extended P system with active membranes of degree $q \geq 1$, taking T time units,

$$\Pi = (\Gamma, \mu, R, T, \{f_r : r \in R\}, M_0, \dots, M_{q-1})$$

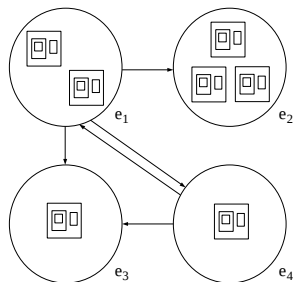
- A multienvironment probabilistic functional extended P system with active membranes of degree (m, q) taking T time units,

$$(\Sigma, G, R_E, \Gamma, \mu, R, T, \{f_{ij} : r \in R_\Pi, 1 \leq j \leq m\}, M_{ij} : 0 \leq i \leq q-1, 1 \leq j \leq m)$$

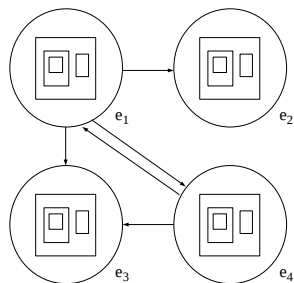


Multienvironment P systems

Functional extended with active membranes



Stochastic approximation



Probabilistic approximation

Multienvironment P systems

Functional extended with active membranes

Functions related to rules

Stochastic approximation

- Kinetic constants
- Stochastic constants
- Propensity

Probabilistic approximation

- Probabilistic functions



Multienvironment P systems

Functional extended with active membranes

Environment rules

$$(x)_{e_j} \xrightarrow{P(x,j,j')} (y)_{e_{j'}} \quad y \quad (\prod_k)_{e_j} \xrightarrow{P(k,j,j')} (\prod_k)_{e_{j'}}$$

Multienvironment P systems

Functional extended with active membrane

Simulation algorithms ← **Take the semantics of the model**

Stochastic approximation

- Multicompartmental Gillespie Algorithm ¹
- Time wait algorithm ¹

Probabilistic Approximation

- *Uniform Random Distribution Algorithm* ²
- *Binomial Random Distribution Algorithm* ²
- *DNDP: Direct non deterministic with probabilities algorithm* ³
- *BDD-CB: Block based direct distribution for consistent blocks*

- ¹ M.J. Pérez-Jiménez, F.J. Romero-Campero. P systems, a new computational modelling tool for Systems Biology. *Transactions on Computational Systems Biology VI*. Lecture Notes in Bioinformatics, 4220 (2006), 176–197
- ² M. Cardona, M.A. Colomer, A. Margalida, A. Palau, I. Pérez-Hurtado, M.J. Pérez-Jiménez, D. Sanuy. A Computational Modeling for Ecosystems Based on P Systems. *Natural Computing*, in press.
- ³ M.A. Martínez-del-Amor, I. Pérez-Hurtado, M.J. Pérez-Jiménez, A. Riscos-Núñez, M.A. Colomer. A new simulation algorithm for probabilistic P systems. *Proceedings of the 2010 IEEE Fifth International Conference on Bio-Inspired Computing: Theories and Applications BIC-TA*, 2010, 59–68



Applications

Stochastic approximation

- Quorum Sensing (*Vibrio Fischeri*)¹
- Signaling pathways (FAS, EGFR)^{2,3}
- Gene regulation in prokaryotes (Lac Operon)⁴

- 1 F.J. Romero, M.J. Pérez-Jiménez. A model of the Quorum Sensing System in *Vibrio Fischeri* using P systems. *Artificial Life*, 14, 1 (2008), 95-109.
- 2 S. Cheruku, A. Paun, F.J. Romero, M.J. Pérez-Jiménez, O.H. Ibarra. Simulating FAS-induced apoptosis by using P systems. *Progress in Natural Science*, 17, 4 (2007), 424-431.
- 3 M.J. Pérez-Jiménez, F.J. Romero. A study of the robustness of the EGFR signalling cascade using continuous membrane systems. *Lecture Notes in Computer Science*, 3561 (2005), 268-278.
- 4 F.J. Romero, M.J. Pérez-Jiménez. Modelling gene expression control using P systems: The Lac Operon, a case study. *BioSystems*, 91, 3 (2008), 438-457.



Applications

Probabilistic approximation

- A **real** ecosystem related to the bearded vulture ^{1,2}
- A **real** ecosystem related to the zebra mussel ³

- 1 M. Cardona, M. A. Colomer, M.J. Pérez-Jiménez, D. Sanuy, A. Margalida. Modelling ecosystems using P systems: The Bearded Vulture, a case of study. *Lecture Notes in Computer Science*, 5391 (2009), 137–156.
- 2 M. Cardona, M.A. Colomer, A. Margalida, I. Pérez-Hurtado, M.J. Pérez-Jiménez, D. Sanuy. A P system based model of an ecosystem of some scavenger birds. *Lecture Notes in Computer Science*, 5957 (2010), 182-195.
- 3 M. Cardona, M.A. Colomer, A. Margalida, A. Palau, I. Pérez-Hurtado, M.J. Pérez-Jiménez, D. Sanuy. A Computational Modeling for Ecosystems Based on P Systems. *Natural Computing*, in press.

Modeling real-life ecosystems

Some studies within the RGNC



- **Modeling Ecosystems using P systems: The Bearded Vulture, a case study.** Cardona et al. *LNCS*, 2009. Vol IV, 137–156.
- **P System Based Model of an Ecosystem of the Scavenger Birds.** Cardona et al. *LNCS*, 2010. Vol IV, 182–195.



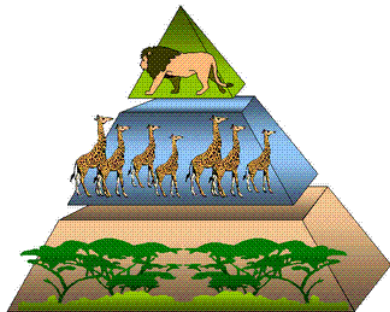
A Computational Modeling for real Ecosystems based on P systems.
Cardona et al. *Natural Computing*, 2010. on-line version.



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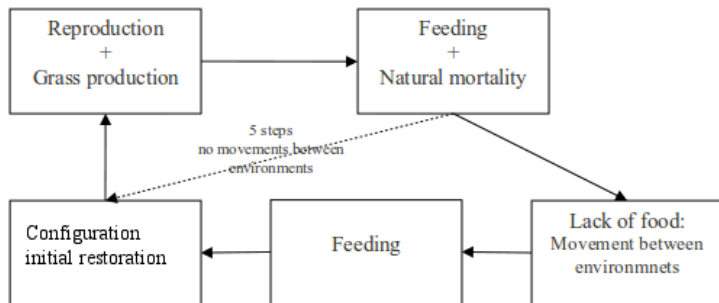


Example: Tritrophic Interactions



- Simplification of a real ecosystem
- Three trophic levels
 - (3) Carnivores
 - (2) Herbivores
 - (1) Grass
- The model
 - 5 modules
 - 9 steps per cycle
 - 10 areas

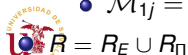
Tritrophic Interactions



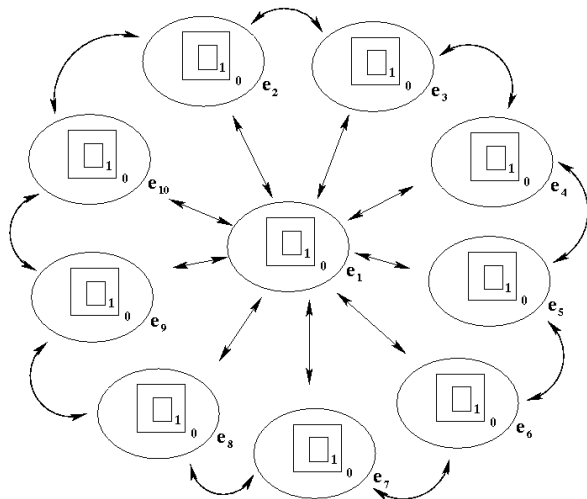
Tritrophic Interactions

$$(G, \Gamma, \Sigma, \Pi, R_E, \{f_{r,j} : r \in R_\Pi, 1 \leq j \leq 10\}, \\ \{\mathcal{M}_{ij} : 0 \leq i \leq 1, 1 \leq j \leq 10\})$$

- $G = (V, S)$
- $\Gamma = \{X_i : 1 \leq i \leq 7\} \cup \{X'_i, Y_i : 2 \leq i \leq 7\} \cup \{R_i : 0 \leq i \leq 6\} \cup G.$
- $\Sigma = \{X_i, X'_i : 2 \leq i \leq 7\}.$
- $\Pi = (\Gamma, [[]_1]_0, R_\Pi).$
- $R_E = \{(X_i)_{e_k} \xrightarrow{p_{k,s,i}} (X'_i)_{e_s} : 1 \leq k \leq 10, 1 \leq s \leq 10, 2 \leq i \leq 7\}$
- $\{f_{r,j} : r \in R_\Pi, 1 \leq j \leq 10\}$
- $\{\mathcal{M}_{ij} : 0 \leq i \leq 1, 1 \leq j \leq 10\}$
 - $\mathcal{M}_{0j} = \{X_1^{q_{1,j}}, R_0 : 1 \leq j \leq 10\}.$
 - $\mathcal{M}_{1j} = \{X_2^{q_{2,j}}, \dots, X_7^{q_{7,j}} : 1 \leq j \leq 10\}.$



Tritrophic Interactions



Tritrophic Interactions

Reproduction + Grass production

- Grass production

$$r_{1,j} \equiv X_1 []_1^0 \xrightarrow{m_j} [X_1, G^{h_j}]_1^+, \quad 1 \leq j \leq 3$$

- Females which reproduce and generate d_i offsprings.

$$r_{2,i} \equiv [X_i]_1^0 \xrightarrow{k_{i,1} \cdot 0,5} [X_i^{1+d_i}]_1^+, \quad 2 \leq i \leq 7$$

- Females and males which don't reproduce.

$$r_{3,i} \equiv [X_i]_1^0 \xrightarrow{1-k_{i,1} \cdot 0,5} [X_i]_1^+, \quad 2 \leq i \leq 7$$

- P system synchronization.

$$r_4 \equiv R_0 []_1^0 \rightarrow [R_0]_1^+$$



Tritrophic Interactions

Feeding + Natural mortality

- Animals which feed and survive.

$$r_{5,i} \equiv [X_i G_i^f]_1^+ \xrightarrow{1-k_{i,2}} [Y_i]_1^-, 2 \leq i \leq 6$$

$$r_{6,i} \equiv [X_7 X_i^{f_7}]_1^+ \xrightarrow{1-k_{7,2}} [Y_7]_1^-, 2 \leq i \leq 6$$

- Animals which feed and don't survive.

$$r_{7,i} \equiv [X_i G_i^f]_1^+ \xrightarrow{k_{i,2}} []_1^-, 2 \leq i \leq 6$$

$$r_{8,i} \equiv [X_7 X_i^{f_7}]_1^+ \xrightarrow{k_{i,2}} []_1^-, 2 \leq i \leq 6$$

- P system synchronization.

$$r_9 \equiv [R_0]_1^+ \rightarrow [R_0]_1^-$$



Tritrophic Interactions

Lack of food: movement between environments

- The objects related to species which have not eaten go to the skin membrane.

$$r_{10,i} \equiv [X_i]_1^- \longrightarrow X_i[]_1^-, 2 \leq i \leq 7$$

- The objects in the skin go to the environment.

$$r_{11,i} \equiv [X_i]_0^0 \longrightarrow X_i[]_0^0, 2 \leq i \leq 7$$

- Movement of objects between environments.

$$r_{12,k,s,i} \equiv (X_i)_{e_k} \xrightarrow{p_{k,s,i}} (X_i')_{e_s}, \\ 1 \leq k \leq 10, 1 \leq s \leq 10, 2 \leq i \leq 7$$

- The object X' goes into the skin membrane.

$$r_{13,i} \equiv X_i'[]_0^0 \longrightarrow [X_i']_0^0, 2 \leq i \leq 7$$

- The object X' goes into the inner membrane.

$$r_{14,i} \equiv X_i'[]_1^- \longrightarrow [X_i']_1^-, 2 \leq i \leq 7$$

- P system synchronization.

$$r_{15,l} \equiv [R_l]_1^- \longrightarrow [R_{l+1}]_1^-, 0 \leq l \leq 4$$



Tritrophic Interactions

Feeding

resources in the new area \rightarrow possibility to feed and survive.

$$r_{16} \equiv [X'_i G^f_i]_1^- \xrightarrow{1-k_{i,2}} [Y_i]_1^0, 2 \leq i \leq 6$$

$$r_{17} \equiv [X'_7 X'^f_i]_1^- \xrightarrow{1-k_{7,2}} [Y_7]_1^0, 2 \leq i \leq 6$$

$$r_{18} \equiv [X'_7 Y_i^f]_1^- \xrightarrow{1-k_{7,2}} [Y_7]_1^0, 2 \leq i \leq 6$$

$$r_{19} \equiv [X'_i G^f_i]_1^- \xrightarrow{k_{i,2}} []_1^0, 2 \leq i \leq 6$$

$$r_{20} \equiv [X'_7 X'^f_i]_1^- \xrightarrow{k_{7,2}} []_1^0, 2 \leq i \leq 6$$

$$r_{21} \equiv [X'_7 Y_i^f]_1^- \xrightarrow{k_{7,2}} []_1^0, 2 \leq i \leq 6$$

$$r_{22} \equiv [R_5]_1^- \longrightarrow [R_6]_1^0$$



Tritrophic Interactions

Reinit of the cycle

$$r_{23,i} \equiv [Y_i]_1^0 \longrightarrow [X_i]_1^0, 2 \leq i \leq 7$$

$$r_{24} \equiv [R_6]_1^0 \longrightarrow [R_0]_1^0$$

$$r_{25} \equiv [X_1]_1^0 \longrightarrow X_1 []_1^0$$

$$r_{26,i} \equiv [X'_i]_1^0 \longrightarrow []_1^0, 2 \leq i \leq 7$$

$$r_{27} \equiv [G]_1^0 \longrightarrow []_1^0$$



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Simulation algorithms

Binomial block based simulation algorithm

- Strategy based on the binomial distribution
- Blocks of rules with the same left-hand side
 - Probabilities summing 1
 - Consistent charges in right-hand side
- Each simulation step is composed by
 - (1) Selection micro-step
 - (2) Execution micro-step



Simulation algorithms

Binomial block based simulation algorithm

This simulation algorithm is useful for the most of the cases but it has the next disadvantages:

- It needs to classify the rules by its left-hand-side.
- It does not handle rules with intersections on their left-hand-sides.
- It does not check the consistency of charges in the selection of rules.
- It does not evaluate probabilistic functions related to rules.



Simulation algorithms

Direct non-deterministic distribution algorithm with probabilities (DNDP)

Input: A multienvironment functional P system with active membranes of degree (q, m) with $q \geq 1$, $m \geq 1$, taking T time units, $T \geq 1$, and a natural number $K \geq 1$.

- 1: **for** $t \leftarrow 0$ to $T - 1$ **do**
- 2: $C_t \leftarrow$ configuration of the system at the moment t
- 3: $C'_t \leftarrow C_t$
- 4: initialization
- 5: First selection phase. It generates a multiset of *consistent* applicable rules.
- 6: Second selection phase. It generates a multiset of *maximally consistent* applicable rules.
- 7: Execution of selected rules.
- 8: $C_{t+1} \leftarrow C'_t$
- 9: **end for**



Simulation algorithms

DNDP Algorithm: Initialization

- 1: $R_{\Pi} \leftarrow$ ordered set of rules of Π
- 2: **for** $j \leftarrow 1$ to m **do**
- 3: $R_{E,j} \leftarrow$ ordered set of rules from R_E related to the environment j
- 4: $A_j \leftarrow$ ordered set of rules from $R_{E,j}$ whose probability at the moment t is > 0
- 5: $M_j \leftarrow$ ordered set of pairs $\langle label, charge \rangle$ for all the membranes from C_t contained in the environment j
- 6: $B_j \leftarrow \emptyset$
- 7: **for each** $\langle h, \alpha \rangle \in M_j$ (following the considered order) **do**
- 8: $B_j \leftarrow B_j \cup$ ordered set of rules $u[v]_h^\alpha \leftarrow u'[v']_h^\beta$ from R_{Π} whose probability at the moment t is > 0 for the environment j
- 9: **end for**
- 10: **end for**



Simulation algorithms

DNDP Algorithm: First selection phase (*consistency*)

```
1: for  $j \leftarrow 1$  to  $m$  do
2:    $R_{sel,j}^1 \leftarrow$  the empty multiset
3:    $R_{sel,j}^2 \leftarrow$  the empty multiset
4:   for  $k \leftarrow 1$  to  $K$  do
5:      $D_j \leftarrow A_j \cup B_j$  with a random order
6:     for each  $r \in D_j$  (following the considered order) do
7:       if  $r$  is consistent with the rules in  $R_{sel,j}^1$  then
8:          $n \leftarrow$  applications( $r,j$ )
9:         if  $n > 0$  then
10:           $C'_t \leftarrow C'_t - n \cdot I(r)$ 
11:           $R_{sel,j}^1 \leftarrow R_{sel,j}^1 \cup \{ \langle r, n \rangle \}$ 
12:        else
13:           $R_{sel,j}^2 \leftarrow R_{sel,j}^2 \cup \{ \langle r, n \rangle \}$ 
14:        end if
15:      end if
16:    end for
17:  end for
18: end for
```



Simulation algorithms

DNDP Algorithm: First selection phase (*applications* function)

```
1:  $n \leftarrow 0$ 
2:  $N' \leftarrow \max\{\text{number of times that } r \text{ is applicable to } C_t'\}$ 
3: if  $N' > 0$  then
4:   if  $p_{r,j}(t) = 1$  then
5:      $n \leftarrow F_b(N', 0, 5)$ 
6:   else
7:      $N \leftarrow \max\{\text{number of times that } r \text{ is applicable to } C_t\}$ 
8:      $n \leftarrow F_b(N, p_{r,j}(t))$ 
9:     if  $n > N'$  then
10:       $n \leftarrow N'$ 
11:     end if
12:   end if
13: end if
14: return  $n$ 
```



Simulation algorithms

DNDP Algorithm: Second phase of rules selection (*maximality*)

```
1: for  $j \leftarrow 1$  to  $m$  do
2:    $R_{sel,j} \leftarrow R_{sel,j}^1 + R_{sel,j}^2$  with an order by the rule probabilities, from highest to lowest
3:   for each  $\langle r, n \rangle \in R_{sel,j}$  (following the selected order) do
4:     if  $n > 0 \vee (r$  is consistent with the rules in  $R_{sel,j}^1)$  then
5:        $N' \leftarrow \max\{\text{number of times that } r \text{ is applicable to } C'_t\}$ 
6:       if  $N' > 0$  then
7:          $R_{sel,j}^1 \leftarrow R_{sel,j}^1 \cup \{\langle r, N' \rangle\}$ 
8:          $C'_t \leftarrow C'_t - N' \cdot I(r)$ 
9:       end if
10:    end if
11:  end for
12: end for
```



Simulation algorithms

DNDP Algorithm: Execution of selected rules

- 1: **for each** $\langle r, n \rangle \in R_{sel,j}^1$ **do**
- 2: $C'_t \leftarrow C'_t + n \cdot r(r)$
- 3: Update the electrical charges of C'_t according to $r(r)$
- 4: **end for**

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Simulation results

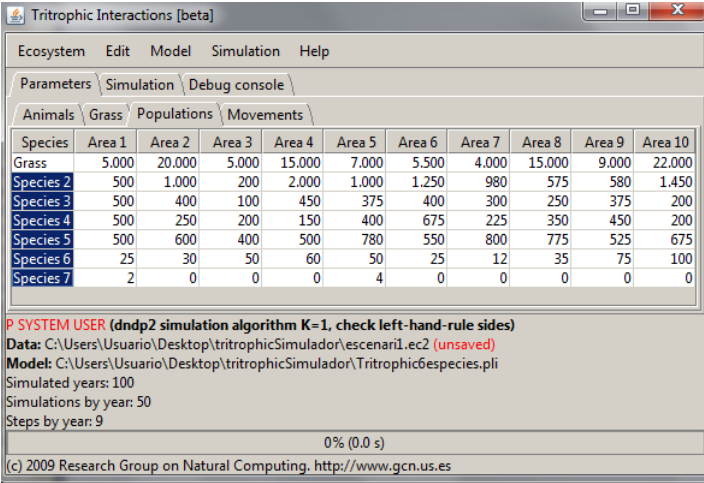
Software used

- P-Lingua: programming language to define P systems
<http://www.p-lingua.org>
- pLinguaCore: Java library → P-Lingua parser + simulation algorithms
- A specific Java GUI over pLinguaCore
 - Input
 - Initial ecosystem parameters
 - Number of years (complete cycles) to simulate
 - Number of simulations per year
 - Output
 - Evolution of the populations
 - Tables and graphs



Simulation results

Number of animals of each species and grass surface



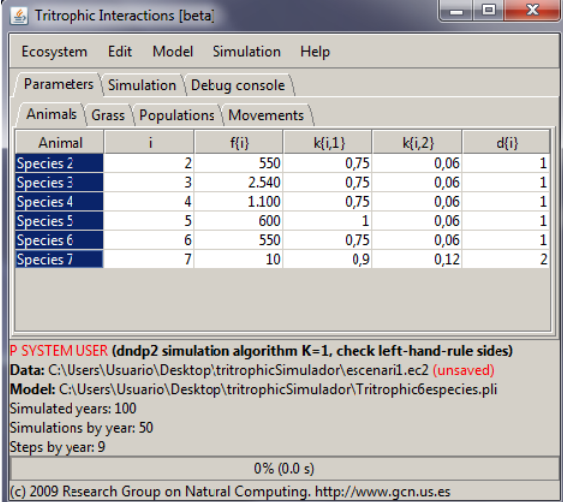
The screenshot shows the 'Tritrophic Interactions [beta]' application window. The 'Parameters' tab is active, and the 'Populations' sub-tab is selected. A table displays the number of animals for each species across ten different areas. Below the table, system parameters and simulation progress are shown.

Species	Area 1	Area 2	Area 3	Area 4	Area 5	Area 6	Area 7	Area 8	Area 9	Area 10
Grass	5.000	20.000	5.000	15.000	7.000	5.500	4.000	15.000	9.000	22.000
Species 2	500	1.000	200	2.000	1.000	1.250	980	575	580	1.450
Species 3	500	400	100	450	375	400	300	250	375	200
Species 4	500	250	200	150	400	675	225	350	450	200
Species 5	500	600	400	500	780	550	800	775	525	675
Species 6	25	30	50	60	50	25	12	35	75	100
Species 7	2	0	0	0	4	0	0	0	0	0

P SYSTEM USER (dndp2 simulation algorithm K=1, check left-hand-rule sides)
Data: C:\Users\Usuario\Desktop\tritrophicSimulador\escenari1.ec2 (unsaved)
Model: C:\Users\Usuario\Desktop\tritrophicSimulador\Tritrophic6species.pli
Simulated years: 100
Simulations by year: 50
Steps by year: 9
0% (0.0 s)
(c) 2009 Research Group on Natural Computing. <http://www.gcn.us.es>

Simulation results

Biological parameters



The screenshot shows a software window titled "Tritrophic Interactions [beta]". The window has a menu bar with "Ecosystem", "Edit", "Model", "Simulation", and "Help". Below the menu bar are several tabs: "Parameters", "Simulation", "Debug console", "Animals", "Grass", "Populations", and "Movements". The "Animals" tab is active, displaying a table with the following data:

Animal	i	f(i)	k{i,1}	k{i,2}	d(i)
Species 2	2	550	0,75	0,06	1
Species 3	3	2.540	0,75	0,06	1
Species 4	4	1.100	0,75	0,06	1
Species 5	5	600	1	0,06	1
Species 6	6	550	0,75	0,06	1
Species 7	7	10	0,9	0,12	2

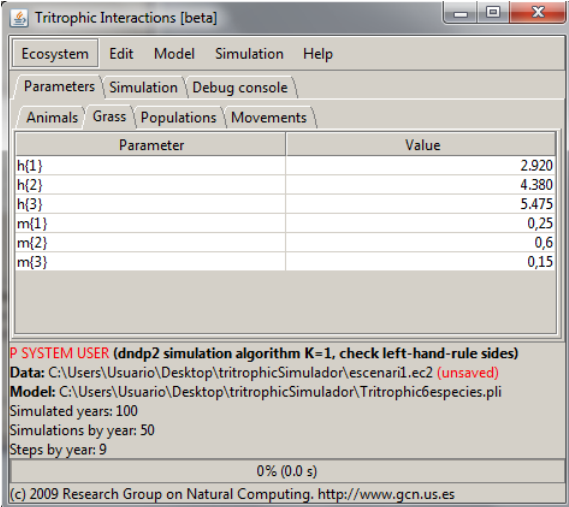
Below the table, there is a status bar with the following text:

P SYSTEM USER (dndp2 simulation algorithm K=1, check left-hand-rule sides)
Data: C:\Users\Usuario\Desktop\tritrophicSimulador\escenari1.ec2 (unsaved)
Model: C:\Users\Usuario\Desktop\tritrophicSimulador\Tritrophic6species.pli
Simulated years: 100
Simulations by year: 50
Steps by year: 9
0% (0.0 s)
(c) 2009 Research Group on Natural Computing. <http://www.gcn.us.es>



Simulation results

Parameters related to grass



The screenshot shows a software window titled "Tritrophic Interactions [beta]". The interface includes a menu bar with "Ecosystem", "Edit", "Model", "Simulation", and "Help". Below the menu bar are several tabs: "Parameters", "Simulation", "Debug console", "Animals", "Grass", "Populations", and "Movements". The "Grass" tab is active, displaying a table of parameters and their values.

Parameter	Value
h{1}	2.920
h{2}	4.380
h{3}	5.475
m{1}	0,25
m{2}	0,6
m{3}	0,15

Below the table, the following system information is displayed:

P SYSTEM USER (dndp2 simulation algorithm K=1, check left-hand-rule sides)
Data: C:\Users\Usuario\Desktop\tritrophicSimulador\escenari1.ec2 (unsaved)
Model: C:\Users\Usuario\Desktop\tritrophicSimulador\Tritrophic6especies.pli
Simulated years: 100
Simulations by year: 50
Steps by year: 9

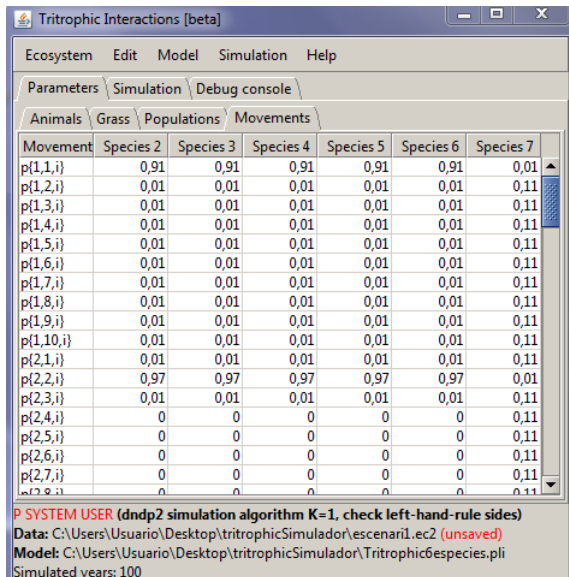
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(c) 2009 Research Group on Natural Computing. <http://www.gcn.us.es>



Simulation results

Probabilities of species movement

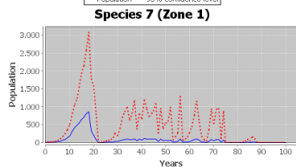
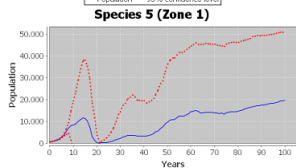
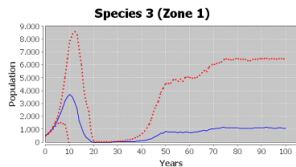


The screenshot shows the 'Tritrophic Interactions [beta]' software window. The 'Movements' tab is active, displaying a table of probabilities for 7 species. The table has columns for Movement, Species 2, Species 3, Species 4, Species 5, Species 6, and Species 7. The data is as follows:

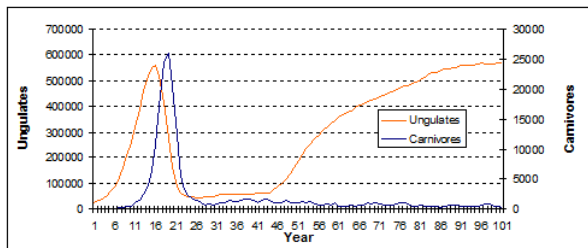
Movement	Species 2	Species 3	Species 4	Species 5	Species 6	Species 7
p(1,1,i)	0,91	0,91	0,91	0,91	0,91	0,01
p(1,2,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,3,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,4,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,5,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,6,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,7,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,8,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,9,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(1,10,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(2,1,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(2,2,i)	0,97	0,97	0,97	0,97	0,97	0,01
p(2,3,i)	0,01	0,01	0,01	0,01	0,01	0,11
p(2,4,i)	0	0	0	0	0	0,11
p(2,5,i)	0	0	0	0	0	0,11
p(2,6,i)	0	0	0	0	0	0,11
p(2,7,i)	0	0	0	0	0	0,11
p(2,8,i)	0	0	0	0	0	0,11

Below the table, the status bar shows: **P SYSTEM USER (dndp2 simulation algorithm K=1, check left-hand-rule sides)**
Data: C:\Users\Usuario\Desktop\tritrophicSimulador\escenari1.ec2 (unsaved)
Model: C:\Users\Usuario\Desktop\tritrophicSimulador\Tritrophic6species.pli
Simulated years: 100

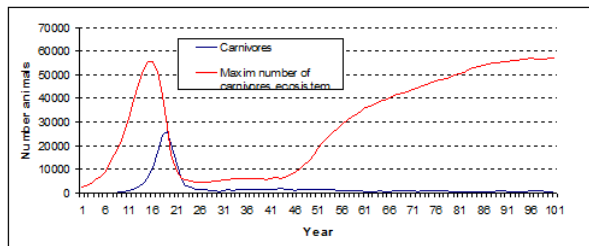
Simulation results



Simulation results



Simulation results



Simulation results

	Scenario 1		Scenario 2	
Algorithm	Binomial	Dndp	Binomial	Dndp
Simulation 1	58,41	54,94	63,62	57,81
Simulation 2	58,57	55,10	61,56	58,58
Simulation 3	58,29	56,05	61,39	57,22
Simulation 4	58,19	56,31	62,81	58,19
Simulation 5	58,75	55,21	61,20	58,75
Simulation 6	57,56	55,19	62,86	57,17
Simulation 7	58,13	54,62	61,92	58,68
Average	58,27	55,35	62,19	58,06
Deviation	0,38	0,61	0,91	0,67

- 1 Introduction
- 2 A P system based modeling framework
- 3 Example: Tritrophic Interactions
- 4 Simulation algorithms
- 5 Simulation results
- 6 Conclusions and future work**



Conclusions

- P systems provide a high-level modeling framework for ecosystems
- Software tools based on membrane computing can be used by ecologists
 - Bearded vulture
 - Zebra mussel
- It has been designed simulation algorithms
- A software framework based on P-Lingua has been provided
- A virtual ecosystem has been used as an example



- Design new simulation algorithms
- Develop simulators based on High Performance Computing (GPUs)
- Design a common protocol to communicate simulators and user interfaces
 - Using different platforms for simulators
 - Codyfing P systems on a standard format file
- Extend the software framework to cover more types of P systems
- Design more efficient and standard GUIs for final users

Thanks!

