

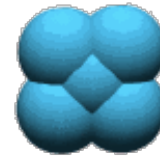
MGS: a declarative spatial computing programming language

Jean-Louis Giavitto^a

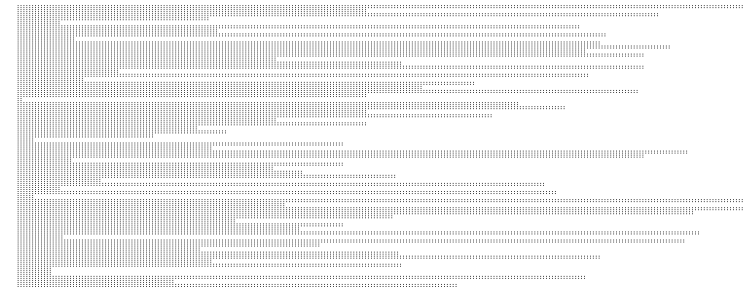
Antoine Spicher^b

^a IRCAM – CNRS

^b LACL – Université de Paris Est



<http://mgs.spatial-computing.org>

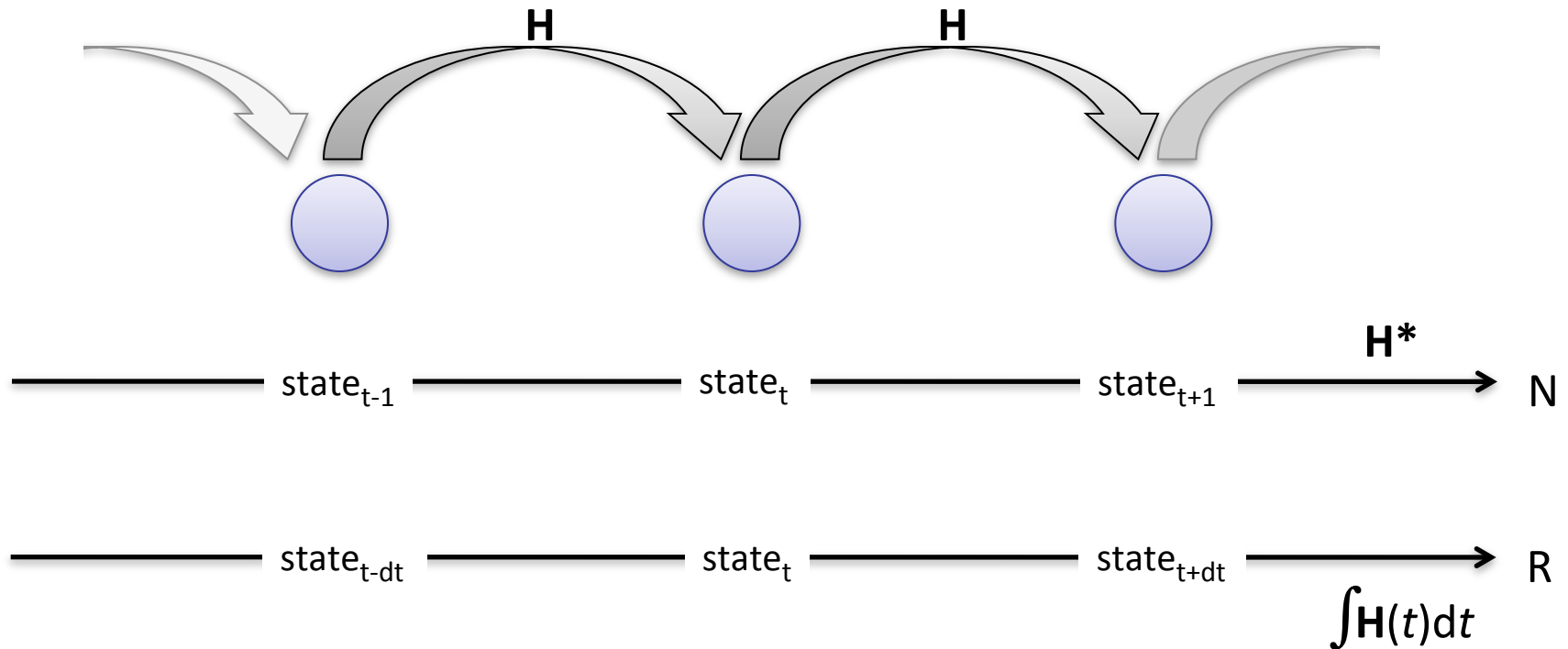


We sketch the rationals of the MGS programming language. MGS is an experimental programming language developed to study the application of several spatial computing concepts to the specification and simulation of dynamical systems with a dynamical structure. MGS extends the notion of rewriting by considering more general structure than terms. The basic computation step in MGS replaces in a topological collection A , some subcollection B , by another topological collection C . A topological collection is a set of element structured by a neighborhood relationships describing an underlying space representing a data structure or constraints that must be fulfilled by the computation. This process proposes a unified view on several computational mechanisms initially inspired by biological or chemical processes (Gamma and the CHAM, Lindenmayer systems, Paun systems and cellular automata).

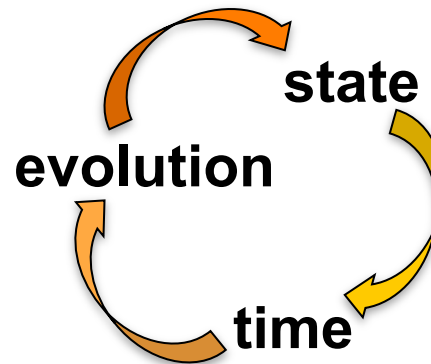
1. $(DS)^2$
2. Gamma, P systems, L systems, cellular automata...
3. Spatial generalization
4. MGS
5. Algorithmic examples
6. Biological modeling

Dynamical systems and Dynamical Structures

Specifying a dynamical system (for simulation)



- Specification of**
- **structure of state**
 - **structure of time**
 - **evolution function**



- State : often structured by space (e.g. fields)
- Time
- Evolution function

C : continuous, D : discrete	PDE	Coupled ODE	Iteration of functions	Cellular automata	...
<i>state</i>	C	C	C	D	...
<i>time</i>	C	C	D	D	...
<i>space</i>	C	D	D	D	...

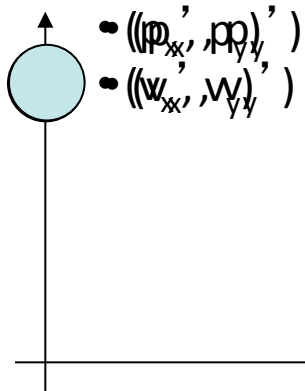


THE CHEMICAL BASIS OF MORPHOGENESIS

BY A. M. TURING, F.R.S. *University of Manchester*

(Received 9 November 1951—Revised 15 March 1952)

a falling ball



at any time a state is a position and a speed

A dynamical system (DS)



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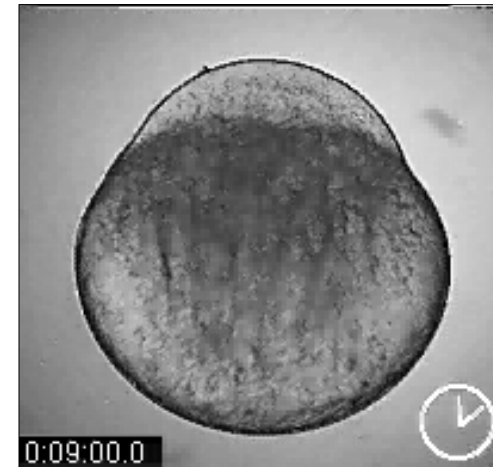
a falling ball



at any time a state is a position and a speed

A dynamical system (DS)

a developing embryo

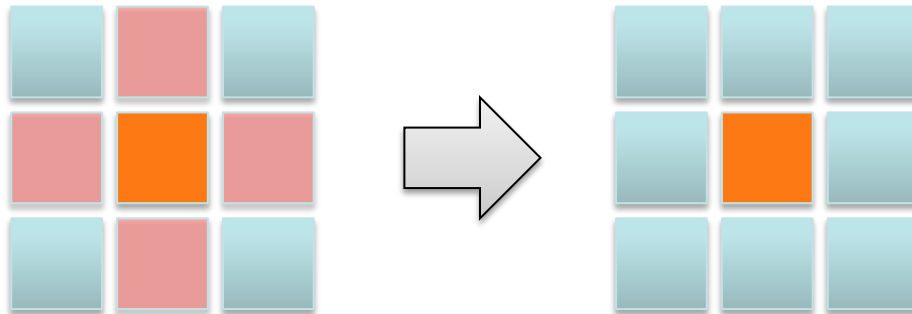


the structure of the state is changing in time
(chemical and mechanical state of each cell)

**A dynamical system
with a dynamical structure
(DS)²**

Bio-inspired models of $(DS)^2$

- Von Neumann
- Avoids the dynamic structure problems
 - Predefined underlying (unbounded) space
- Replace a cell X
in an NEWS grid
by another one (with a new state)



It might be possible, however, to treat a few particular cases in detail with the aid of a digital computer. This method has the advantage that it is not so necessary to make simplifying assumptions as it is when doing a more theoretical type of analysis. It might even be possible to take the mechanical aspects of the problem into account as well as the chemical, when applying this type of method. The essential disadvantage of the method is that one only gets results for particular cases. But this disadvantage is probably of comparatively little importance.

Lindenmayer systems

- The structure of a tree can be coded by a string of parenthesised symbols
- A symbol is an elementary part of the plant
- The symbol between [and] represents a sub-tree
- Additional conventions are used to represent main axis, orientation, depth, etc.

- A rule

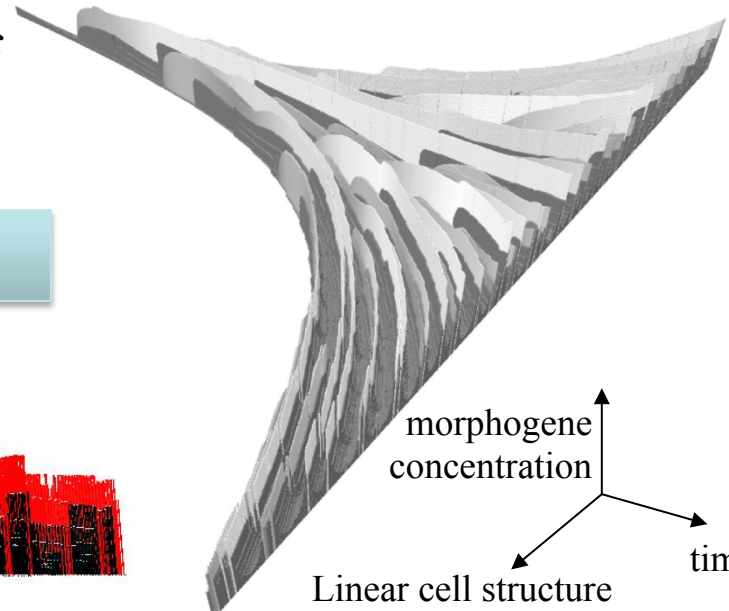
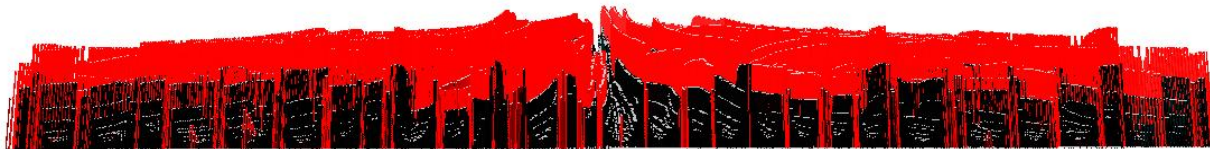
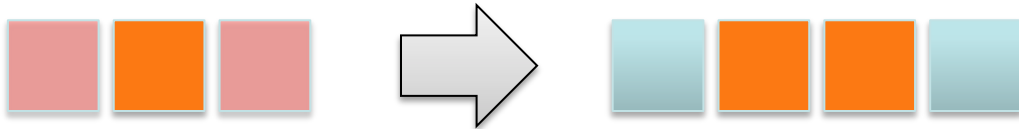
$$s_0 \rightarrow s_1 s_2 s_3 \dots$$
represents the evolution of s_0

- Replace a substring X in a string by another one

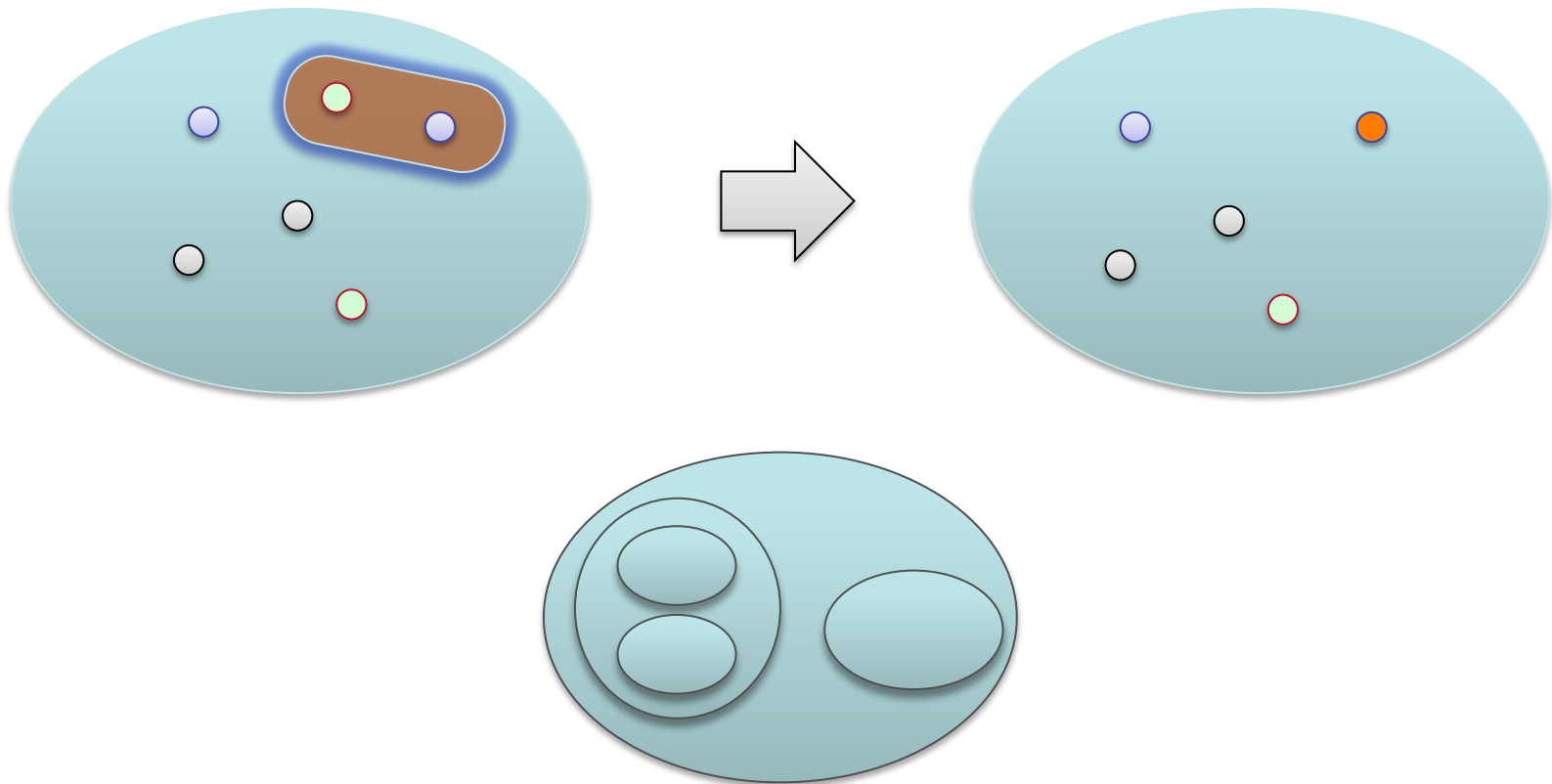
```
e / (C(e) & (e.x >= lm) & (e.p == L))
```

```
=> {type="C", a=e.a, h=e.h, x=e.x*shorter, p=L},
```

```
{type="C", a=e.a, h=e.h, x=e.x*longer
```



- Replace a sub-multiset X in a multiset by another one



A general device

A general rewriting mechanism

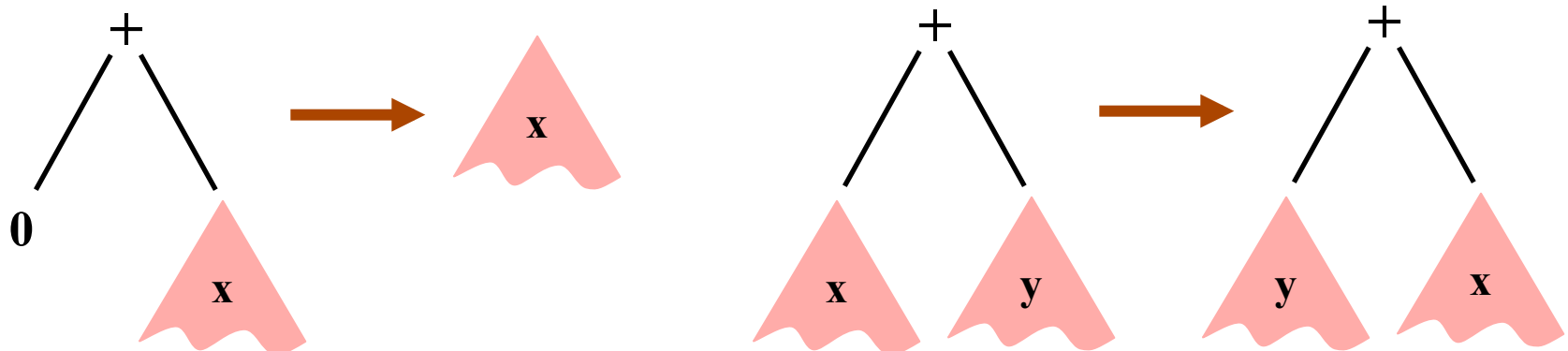
1. In a *collection* of elements
2. Replace a *subcollection* X
3. With a collection Y computed from X and its *neighbors*

	Collection	Neighborhood	data structure
	– Term	father/son	free
monoidal	– Set	all	A + C + Idempotent
	– Multiset	all	A + Commutative
	– Sequence	left and right (with jump for trees)	Associative
	Commutative		
	– Grid	NEWS	

- Rewriting system

- Used to formalize equational reasoning
- A generative device (grammar)
- Replace a sub-part of an entity by another
- Set of rewriting rules $\alpha \rightarrow \beta$
 - α : pattern specifying a sub-part
 - β : expression evaluating a new sub-part

- Example: arithmetic expressions simplification



$$1 + 2 \rightarrow \dots$$

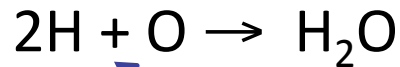
arithmetic operation

(arithmetic) term rewriting

$$a . b \rightarrow \dots$$

string concatenation: « . » is a formal associative operation

string rewriting (\sim L systems)



multiset concatenation (= the chemical soup): « . » is AC

multiset rewriting (\sim chemistry)

A general rewriting mechanism

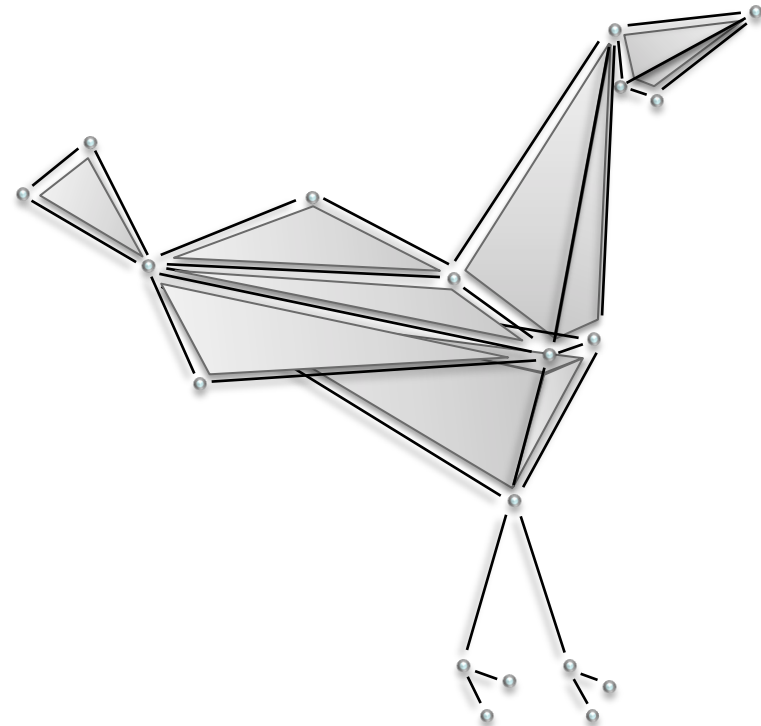
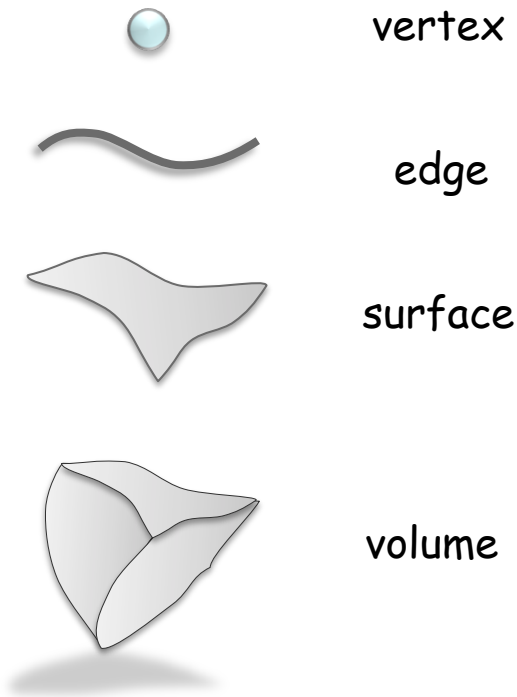
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	Commutative		
	– Grid	NEWS	

- Use space (topology) to unify the various collection structures
 - space as a **resource**
 - space as a **constraint**
 - space as an **input/output**
- **Neighborhood** relationships:
 - the structure of the collection
 - the structure of the subcollection
 - the computation dependencies
- Substitution (replacement)
topological surgery

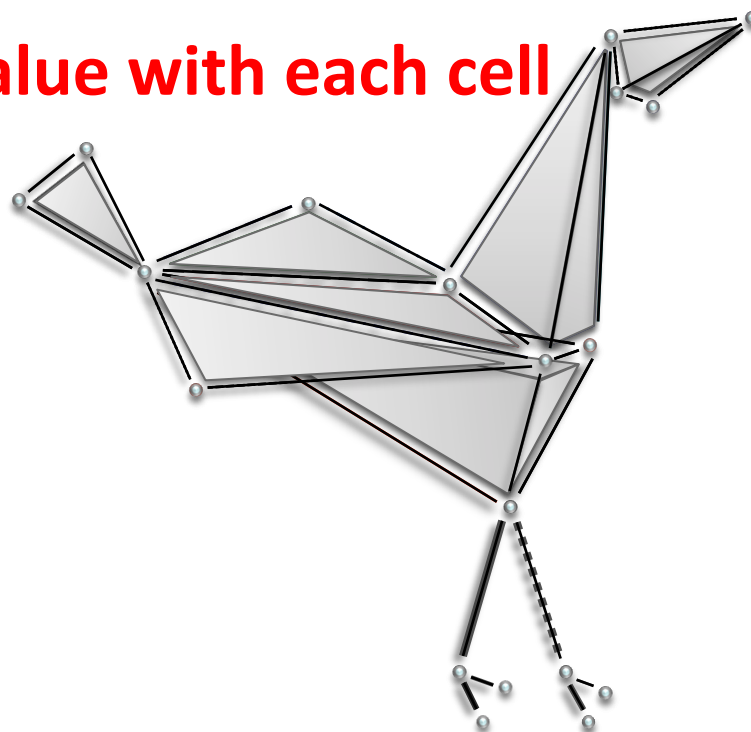
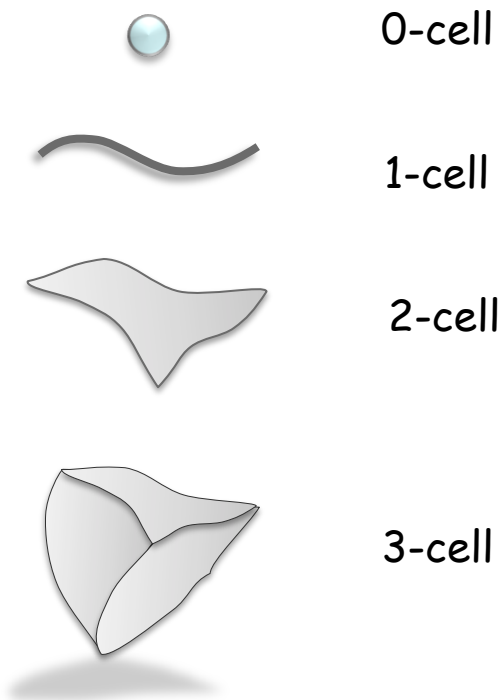
- *local evolution rules*
mandatory when you cannot express a global function/relation because the domain of the function/relation is changing in time
- *interaction based approach*
the l.h.s. of a rule specifies a set of elements in *interaction*,
the r.h.s. the result of the interaction
- *the phase space is well defined but not well known*
a generative process enumerates the elements but
membership-test can be very hard
- *various kind of time evolution* (for the same set of rules)
- *demonstration by induction*
on the rules or on the derivation (e.g. growth function in L system)

- Topological collections
 - Structure
 - A collection of topological cells
 - An ***incidence relationship***



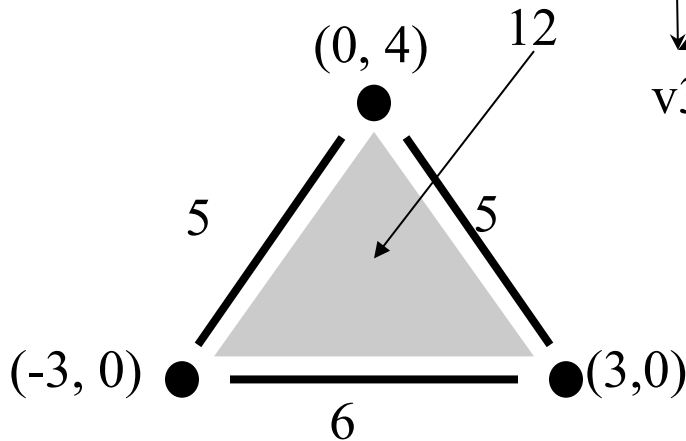
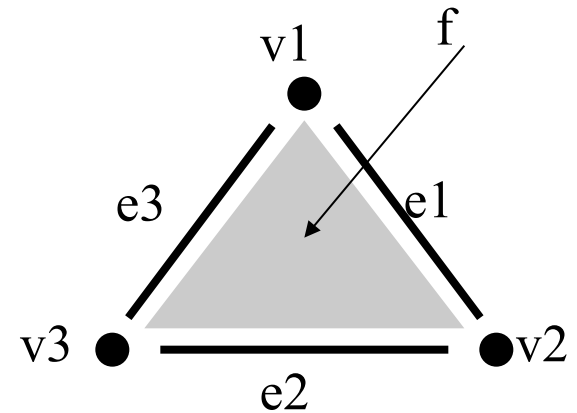
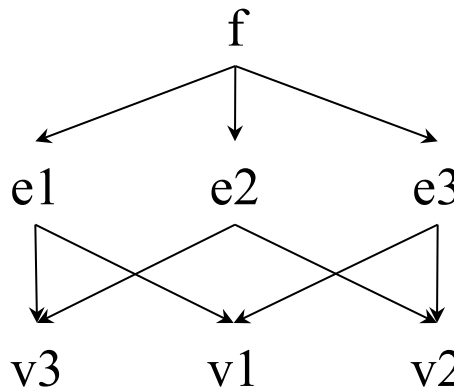
- Topological collections
 - Structure
 - A collection of topological cells
 - An incidence relationship

– Data: **association of a value with each cell**



Incidence relationship and lattice of incidence:

- $\text{boundary}(f) = \{v_1, v_2, v_3, e_1, e_2, e_3\}$
- $\text{faces}(f) = \{e_1, e_2, e_3\}$
- $\text{cofaces}(v_1) = \{e_1, e_3\}$



Topological chain

- coordinates with vertices
- lengths with edges
- area with f

$$\begin{pmatrix} 0 \\ 4 \end{pmatrix} \cdot v_1 + \begin{pmatrix} 3 \\ 0 \end{pmatrix} \cdot v_2 + \begin{pmatrix} -3 \\ 0 \end{pmatrix} \cdot v_3 + 5 \cdot e_1 + 6 \cdot e_2 + 5 \cdot e_3 + 12 \cdot f$$

Topology (open and closed sets)



simplices
=
Closed sets



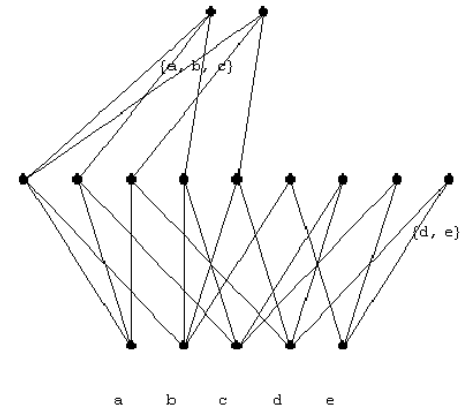
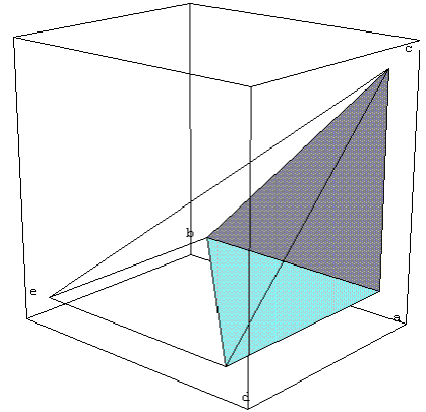
Simplicial Complex (set of sets closed by inclusion/intersection)



simplices
=
Lattice cones



Lattice (order relation: \wedge , \vee)



- Concise reformulation of classical approaches
- Extension

- Transformations

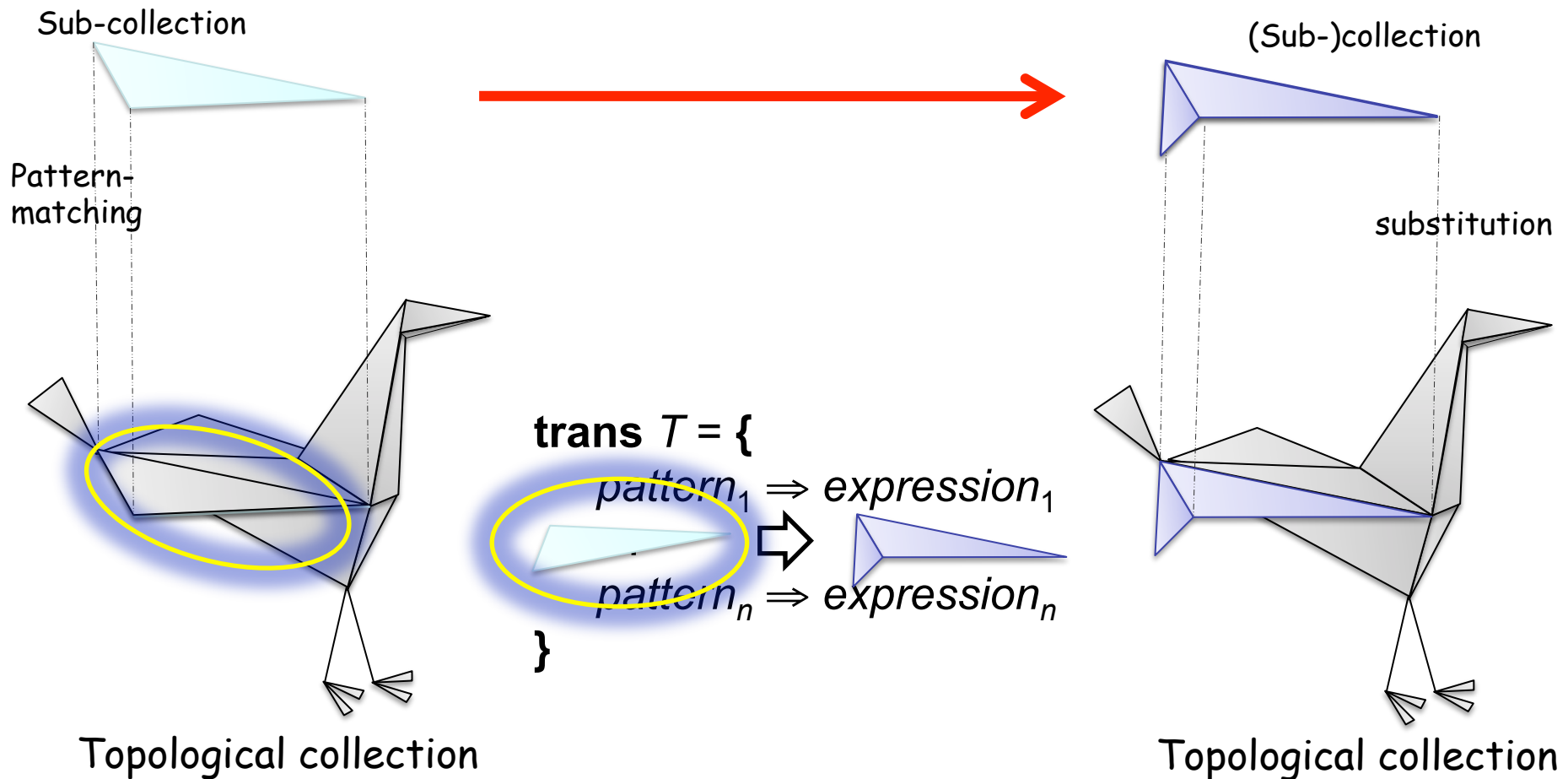
- Functions defined by case on collections

- Each case (pattern) matches a sub-collection

- Defining a rewriting relationship: ***topological rewriting***

$$\text{trans } T = \left\{ \begin{array}{l} \text{pattern}_1 \Rightarrow \text{expression}_1 \\ \dots \\ \text{pattern}_n \Rightarrow \text{expression}_n \end{array} \right\}$$

- Transformations



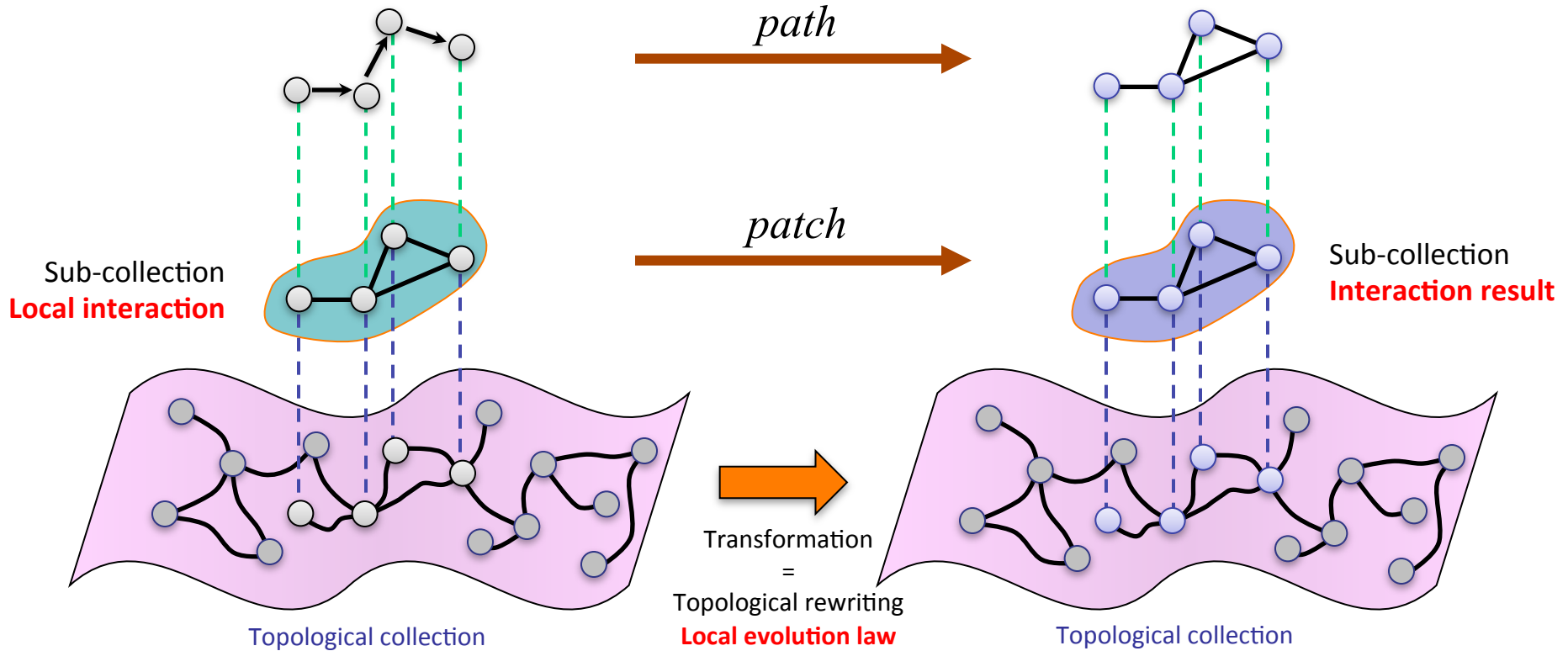
Topological rewriting = transformation

$1 + 2 \rightarrow \dots$ (arithmetic) term rewriting
↙
arithmetic operation

$a . b \rightarrow \dots$ string rewriting (\sim L systems)
↙
string concatenation: « . » is a formal associative operation

$2H + O \rightarrow H_2O$ multiset rewriting (\sim chemistry)
↙
multiset concatenation (= the chemical soup): « . » is AC

$v_1 \cdot \sigma_1 + v_2 \cdot \sigma_2 \rightarrow \dots$ **topological rewriting** (MGS)
↙
gluing cell in a cell complex: ... (AC and algebraic machinery)



Pattern matching : specifying a sub-collection of elements in interaction

- *Path transformation* (path = sequence of neighbor elements)
 - Concise but limited expressiveness
- *Patch transformation* (arbitrary shape)
 - Longer but higher expressiveness


Example: Diffusion Limited Aggregation (DLA)

- Diffusion: some particles are randomly diffusing; others are **fixed**
- Aggregation: if a **mobile** particle meets a **fixed** one, it stays fixed

```

trans dla = {
    `mobile , `fixed => `fixed, `fixed ;
    `mobile , <undef> => <undef>, `mobile
}

```


NEIGHBOR OF

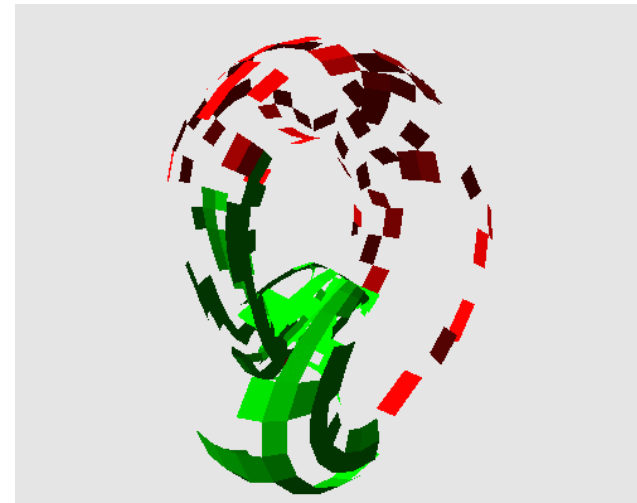
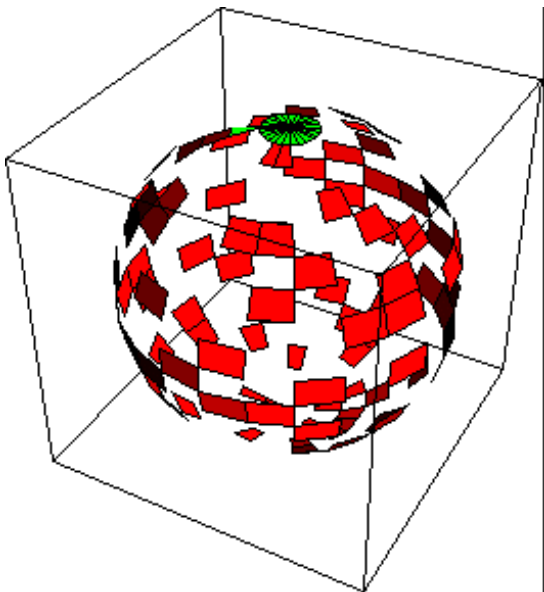


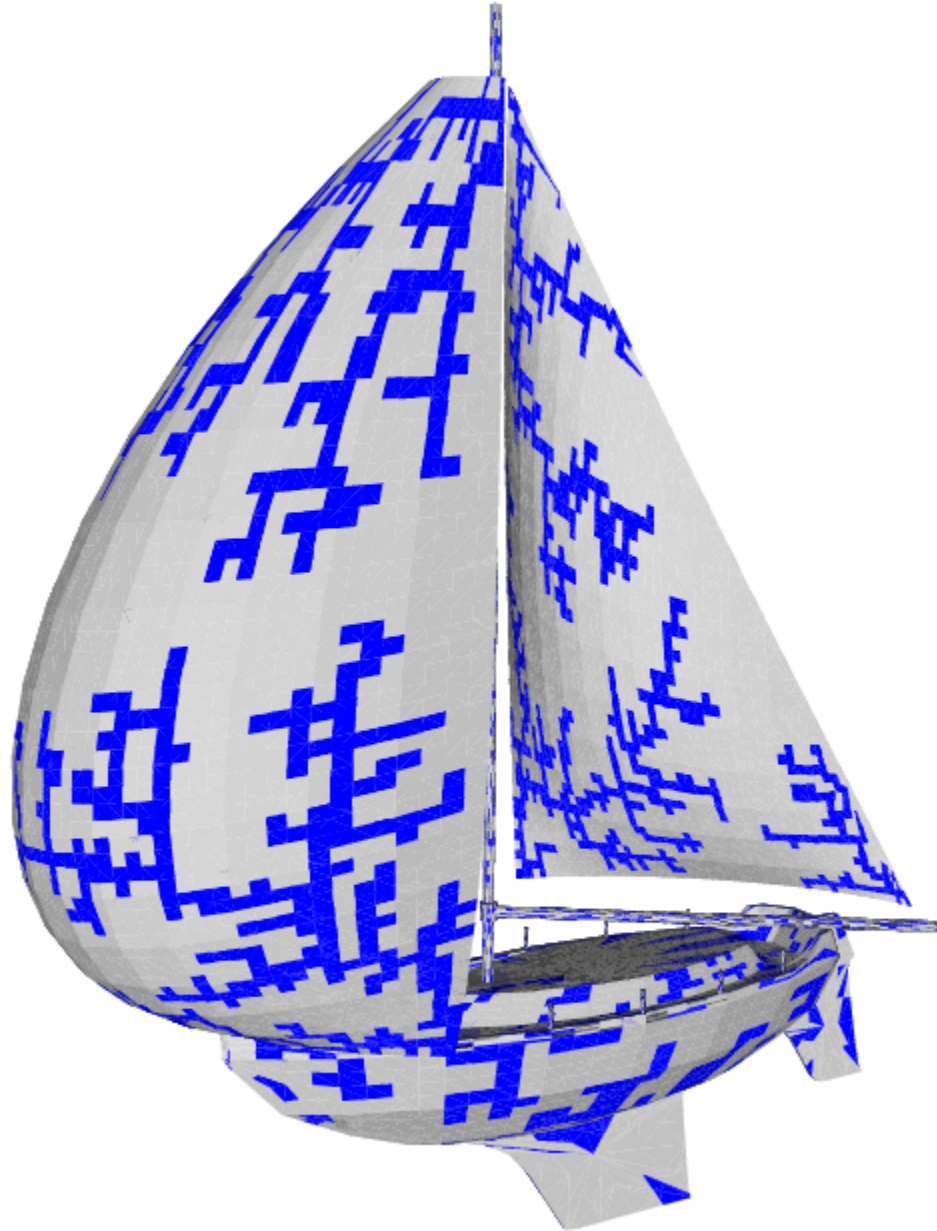
Example: Diffusion Limited Aggregation (DLA)

- Diffusion: some particles are randomly diffusing; others are **fixed**
- Aggregation: if a **mobile** particle meets a **fixed** one, it stays fixed

```
trans dla = {  
  `mobile , `fixed => `fixed, `fixed ;  
  `mobile , <undef> => <undef>, `mobile  
}
```

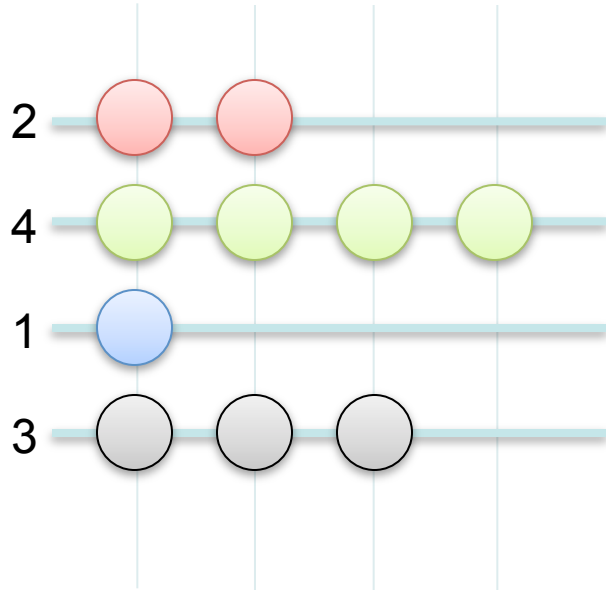
this transformation is an abstract process that can be applied to any kind of space



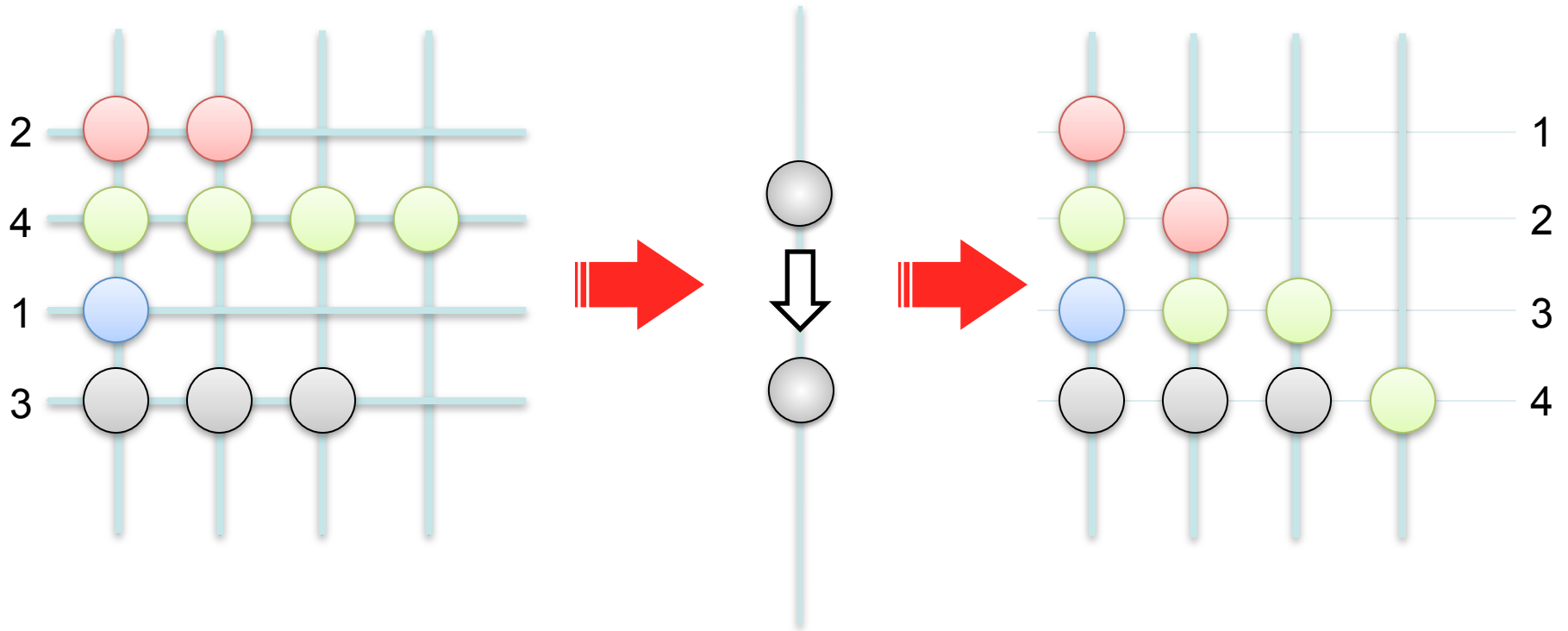


Algorithmic examples

Bead Sort

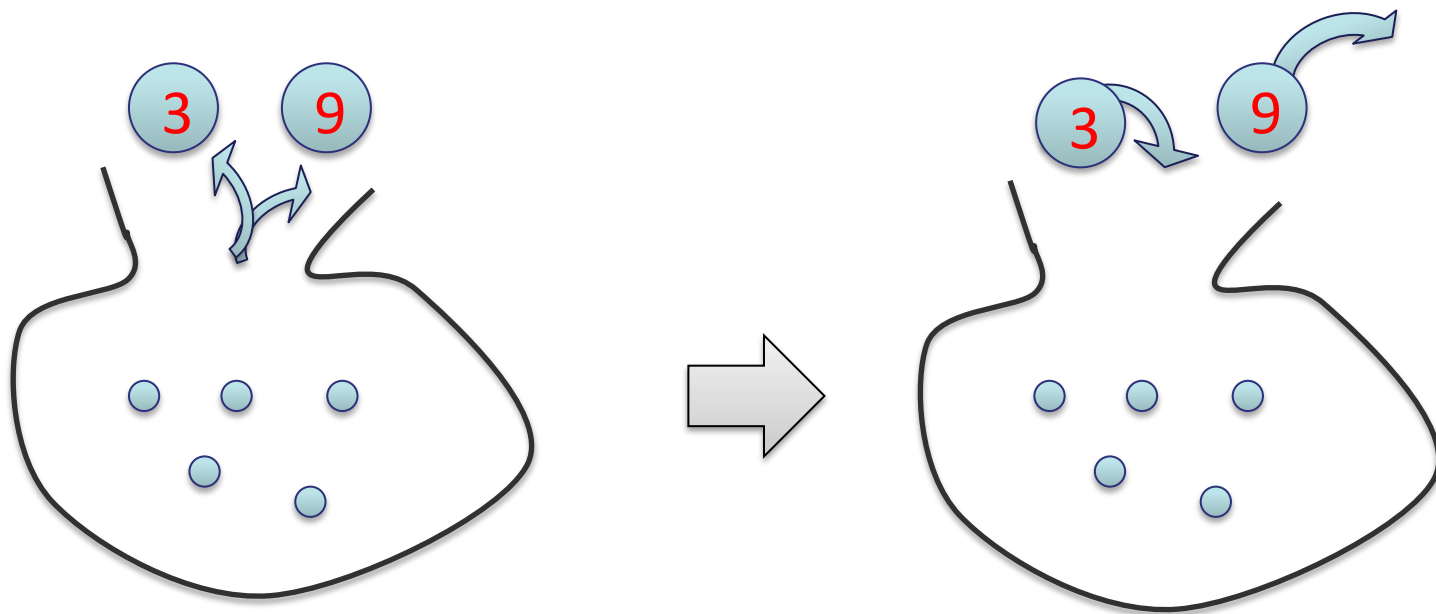


Bead Sort





```
Gbf NEWS = < North, South, East, West;  
          North+South=0, East+West=0>  
  
trans dla = {  
    `bead |south> `empty => `empty, `bead ;  
}
```

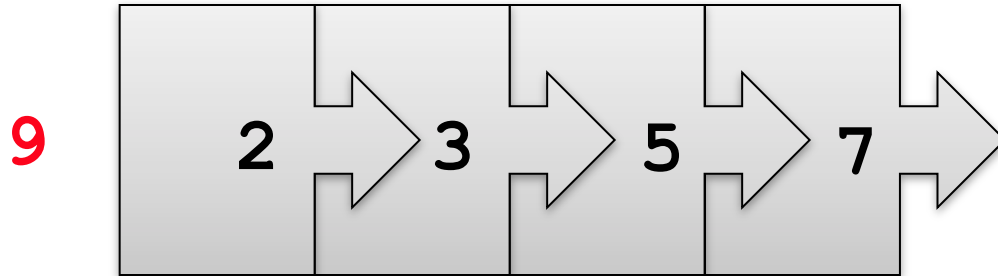


`trans Generate = {x, true} => x, {x + 1, true};`

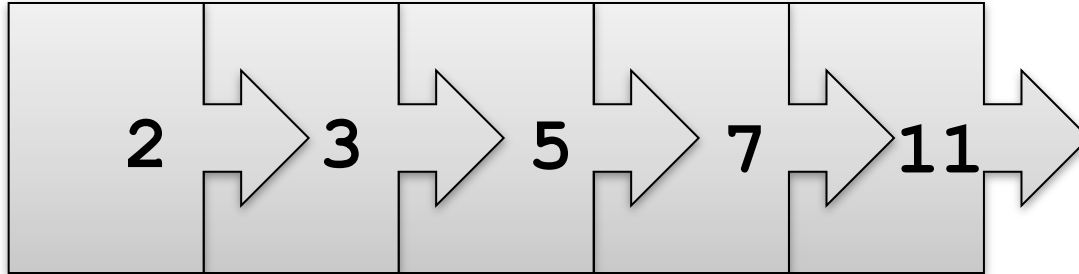
`trans Succeed = {x, true} => x;`

`trans Eliminate = (x, y / y mod x = 0) => x;`

`Eliminate[fixrule](Succeed(Generate[N]({2, true}, set : ())))`

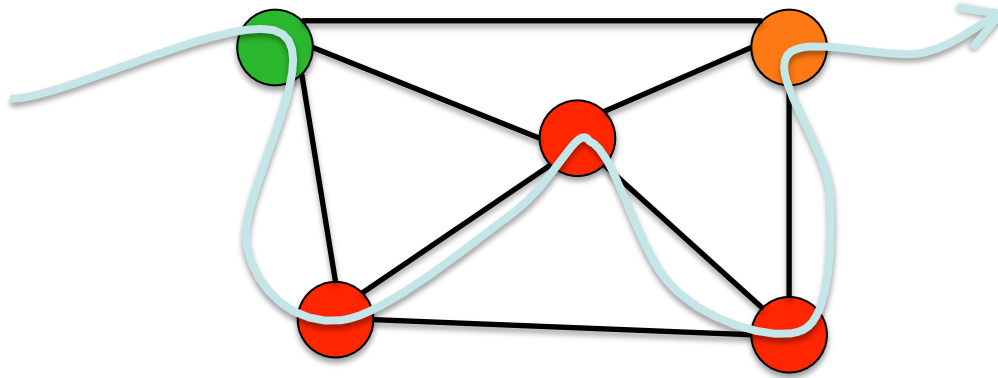


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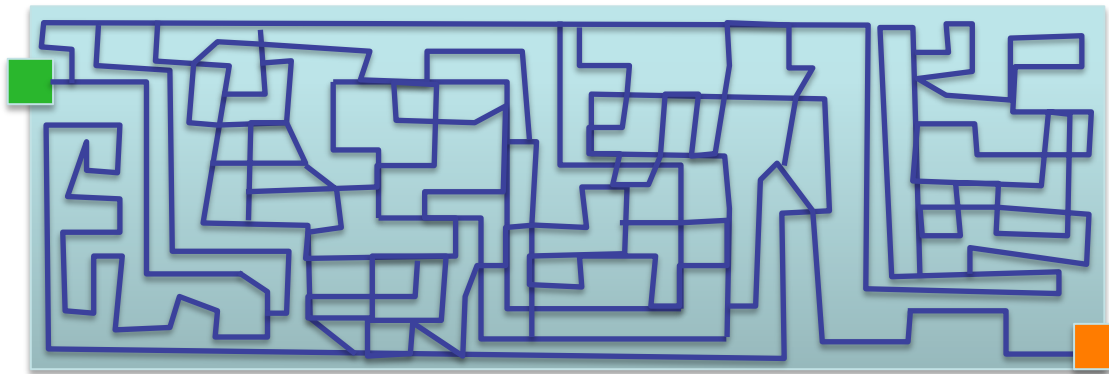


```
trans Eratos = {  
  Genere1 = n : integer / ~right n => n, {prime = n};  
  Genere2 = n : integer, {prime as x, ~candidate, ~ok}  
    => n + 1, {prime = x, candidate = n};  
  Test1 = {prime as x, candidate as y, ~ok} / y mod x = 0 => {prime = x};  
  Test2 = {prime as x, candidate as y, ~ok} / y mod x <> 0  
    => {prime = x, ok = y};  
  Next = {prime as x1, ok as y}, {prime as x2, ~ok, ~candidate}  
    => {prime = x1}, {prime = x2, candidate = y};  
  NextCreate = {prime as x, ok as y} as s / ~right s  
    => {prime = x}, {prime = y};  
}
```

Hamiltonian path

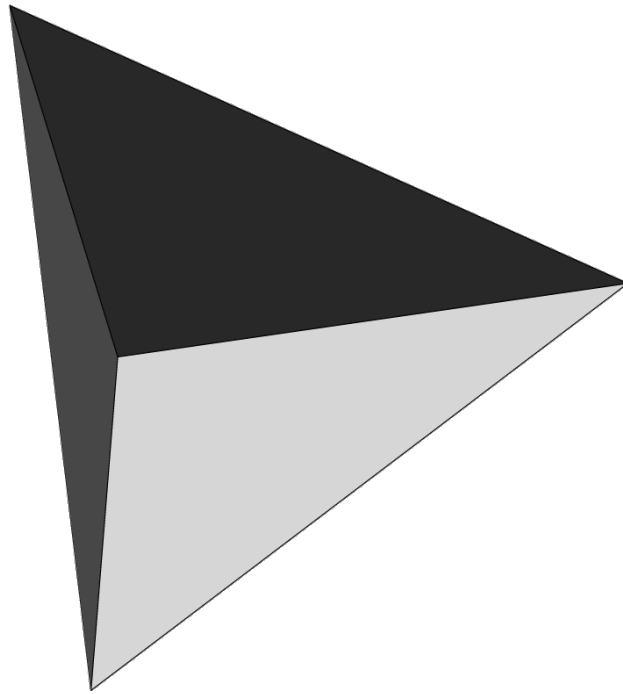


```
trans h_path = { `start , x* as p, `stop  
                / size(p) = n-2 => return p }
```

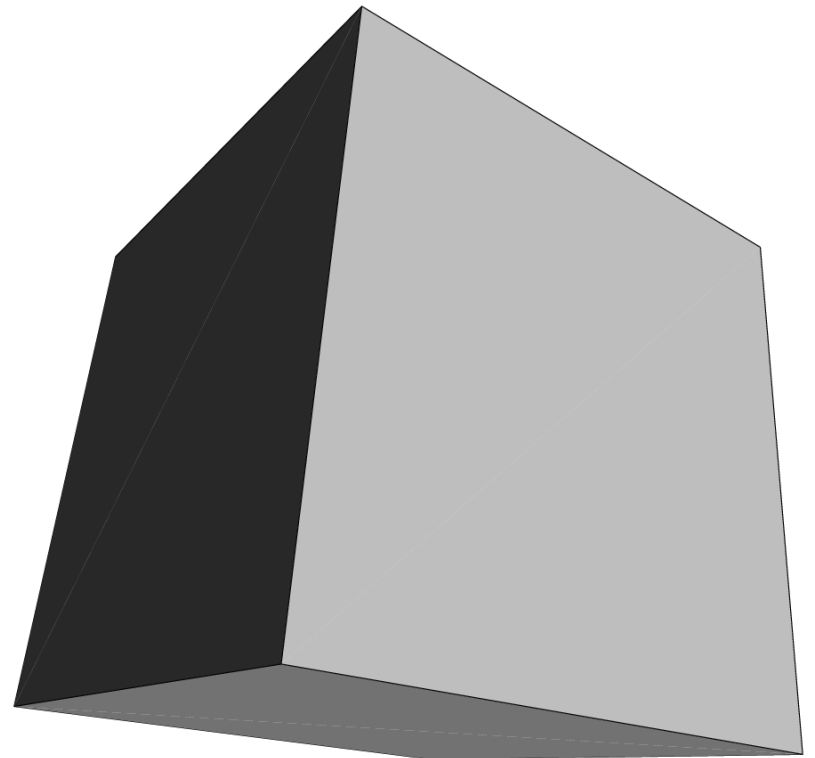


```
trans maze = { `input, c* as p, `output => return p }
```

Fractal construction by carving



Sierpinsky sponge (4 steps)



Menger sponge (2 steps)



- ...
- Various models of Phage λ
- Sperm crawling
- Neurulation
- Prototyping a
« synthetic multicellular bacteria »
- ...

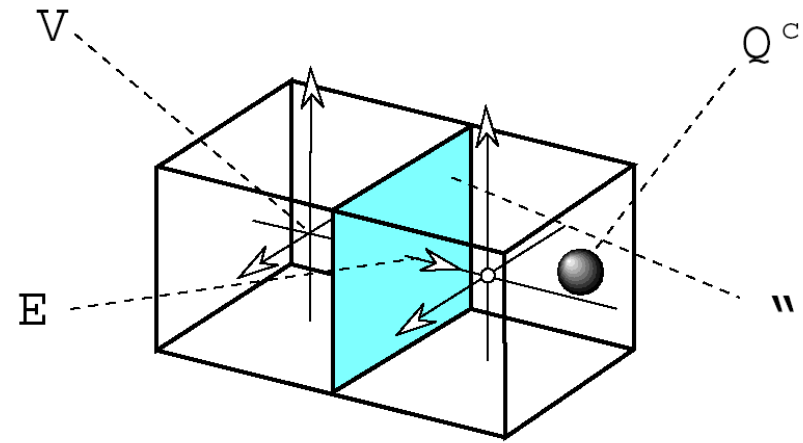
Example of electrostatic Gauss law [Tonti 74]

- Electric charge content ρ : **dimension 3**
- Electric flux Φ : **dimension 2**
- Law available on a arbitrary complex domain

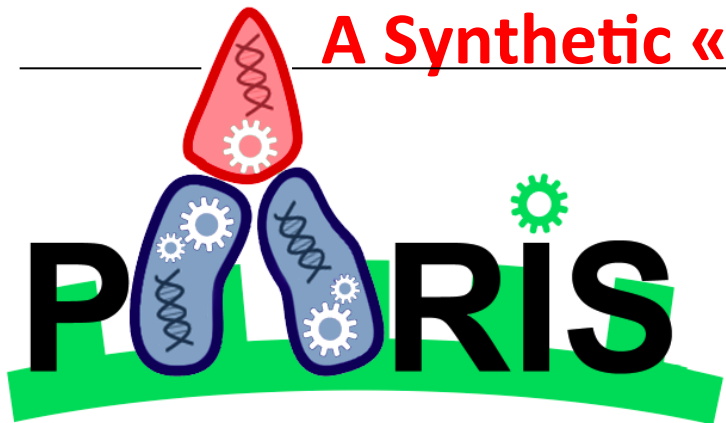
$$\phi = \iint w \cdot dS = \frac{Q^c}{\epsilon_0} = \iiint_{(V)} \frac{\rho}{\epsilon_0} d\tau$$

electric field in space:

- V: electric potential (dim 0)
- E: voltage (dim 1)
- w: electric flux (dim 2)
- Qc: electric charge (dim 3)



A Synthetic « Multicellular Bacterium »



Synthetic Biology is

- A) the design and construction of new biological parts, devices, and systems, and
- B) the re-design of existing, natural biological systems for useful purposes.

(Español)

Synthetic Biology Logo

- Home
- About
- Conferences
- Labs
- Courses
- Resources
- FAQ

Community news

- IET Synthetic Biology first issue includes iGEM 2006
- Synthetic Biology 3.0 Zurich proceedings. Download [here](#).
- BioBricks Foundation first [membership drive](#).
- [Synthetic Biology: Caught between Property Rights, the Public Domain, and the Commons](#)
- US HSPD-18. Guidance on openness and international transparency in biodefense work still needed.

Resources

- [Press articles](#)
- [Publications: citulike, connotes, PubMed](#)



David Bikard, Thomas Landrain, David Puyraimond, Eimad Shotar, Gilles Vieira, Aurélien Rizk, David Guegan, Nicolas Chiaruttini, Thomas Clozel, Thomas Landrain

Registry of Standard Biological Parts

http://parts.mit.edu/

Registry of Standard Biological Parts
Massachusetts Institute of Technology

About the Registry

- Using the Registry
- User Accounts

Parts, Devices & Systems

- About Parts
- Adding Parts
- Measuring Parts

Assembly

- Standard Assembly
- Assembly Tool
- DNA Synthesis
- DNA Repository

Educational Program

- IAP 2003/2004
- SBC 2004
- iGEM 2005

References

- Glossary
- FAQ
- Links
- Search

View Part

BBa_

Parts Catalog Click on the icons below to see parts by category. [more...](#)

Regulatory

Reporter

Inverter

RNA

Protein Generator

Tag

Parts List

Deleted

Cell Strain

RBS

CDS

Terminator

Composite

Cell-Cell Signalling

Measurement

Primer

Other

Plasmid

T7

Web Site Update

Registry web site changes in support of iGEM 2005 are under way.

- The new account manager is in place with better support for groups, group leaders, and editing.
- Part categories are becoming more detailed, see the signalling category for an example.
- The new part viewer and editor is on the way soon.
- New Rolling Assembly tool under development.

Educational Programs

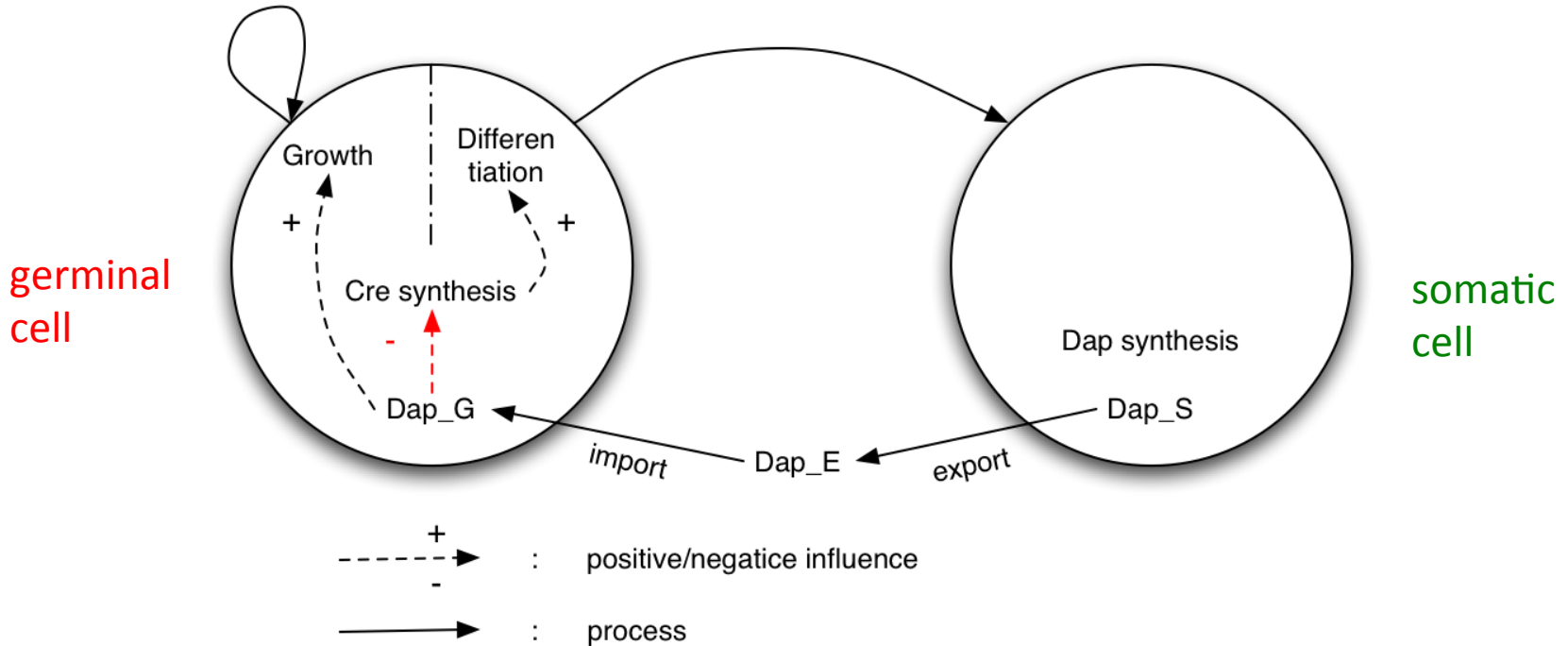
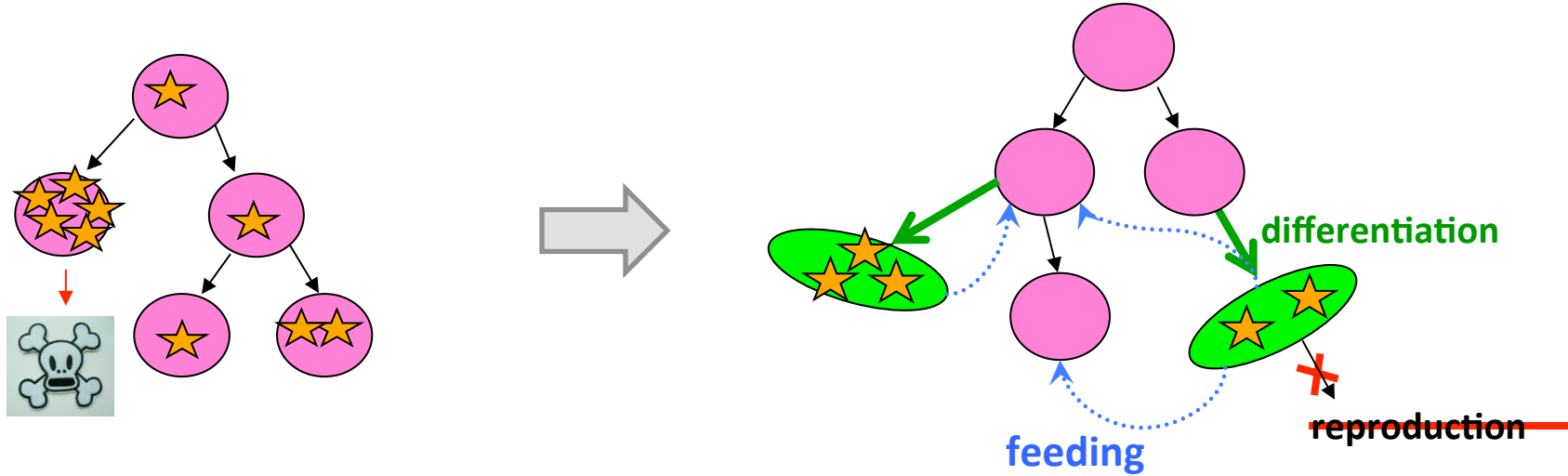
The Registry supports design classes where students make simple systems from standard, interchangeable biological parts and operate them in living cells.

Thirteen schools are participating in the 2005 Intercollegiate Genetically Engineered Machine competition (iGEM 2005). The schools are: Berkeley, Caltech, Cambridge, Davidson, ETH Zurich, Harvard, MIT, Oklahoma, Penn State, Princeton, Toronto, UCSF, and UT Austin.

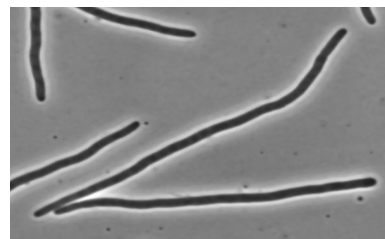
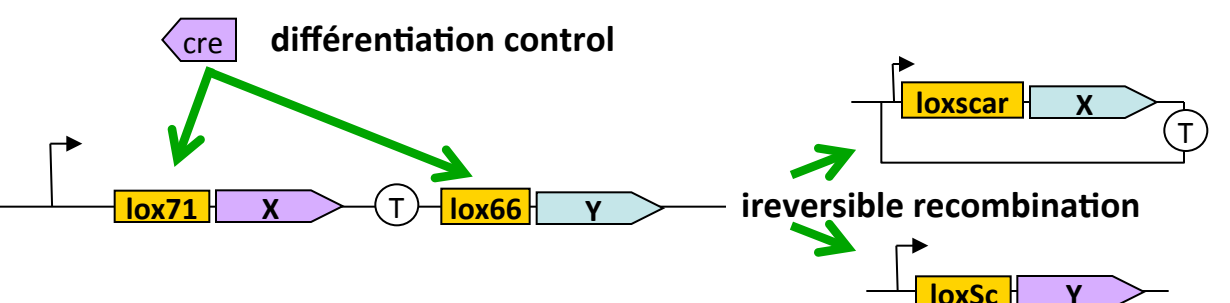
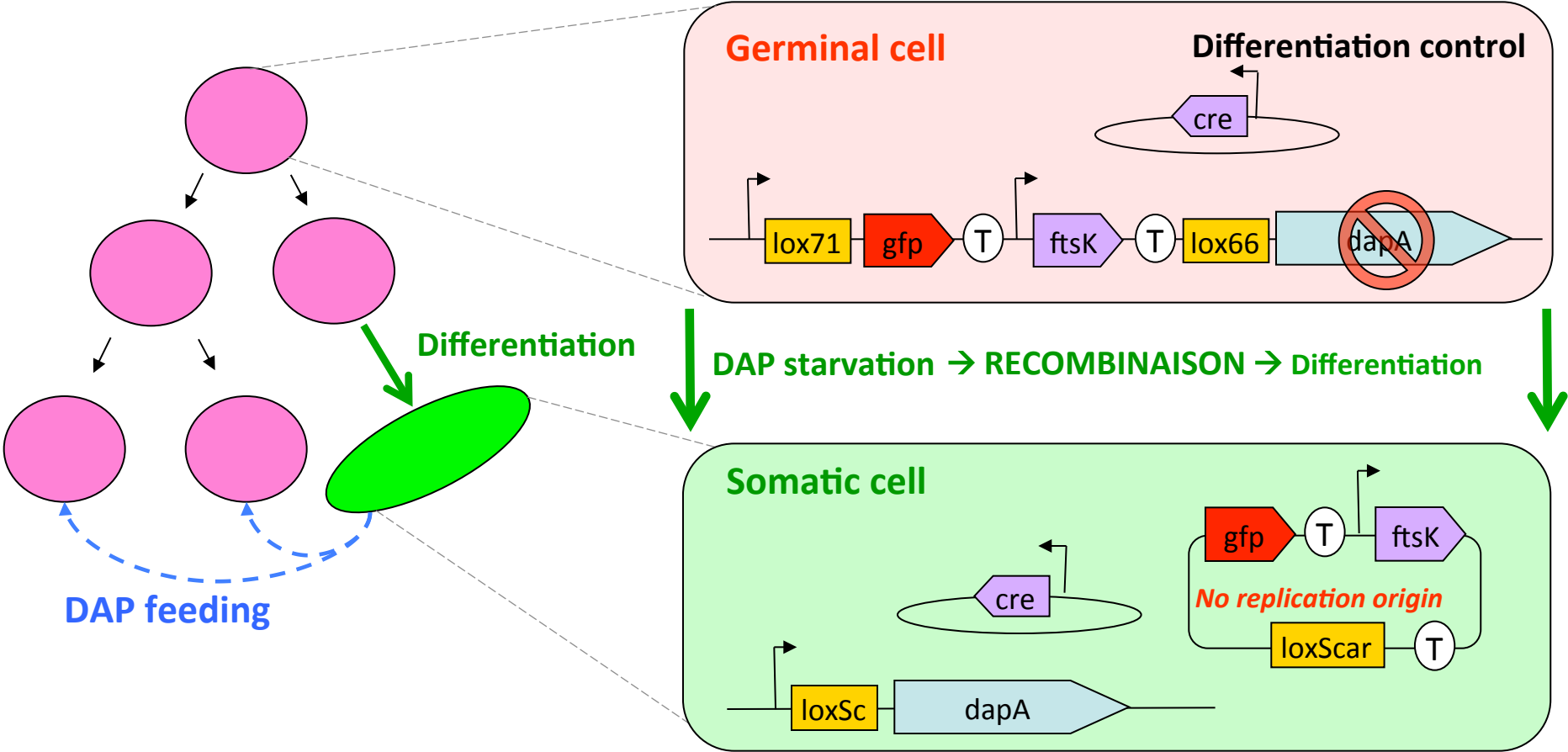
Employment

The Registry is looking for full-time Technical Assistants and Web Programmers. Please contact Staffing Services at MIT for details: [Technical Assistant](#), [Web Programmer](#).

The Paris iGEM project: a « multicellular bacteria » to decouple growth and transgene expression



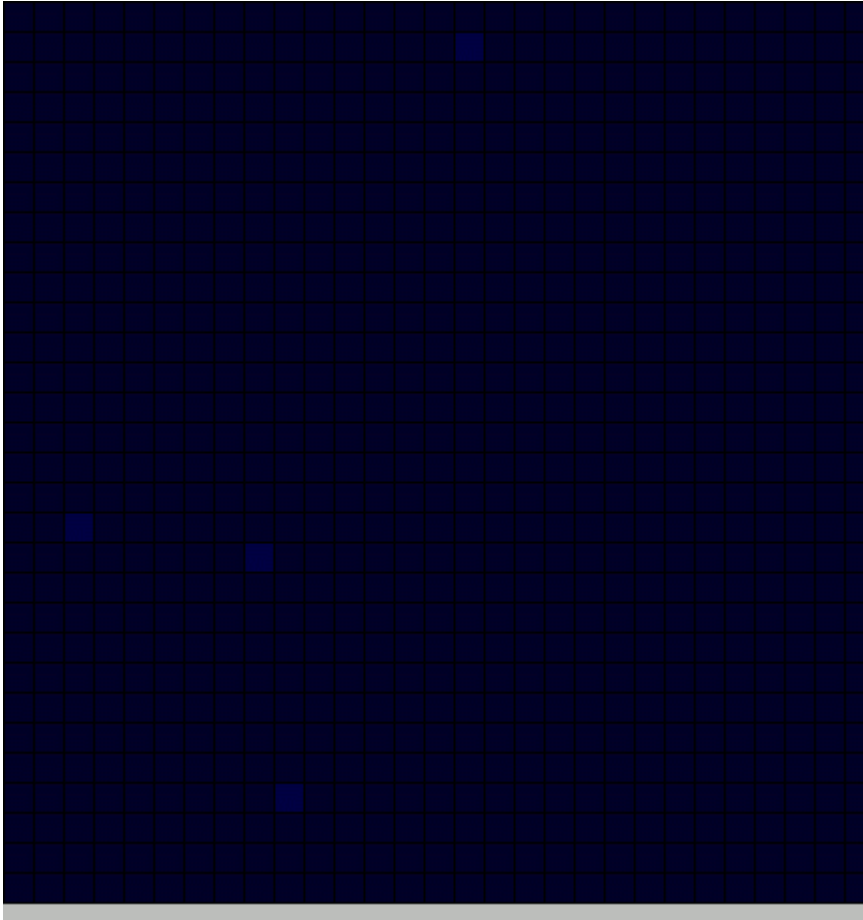
Implementation using BioBricks



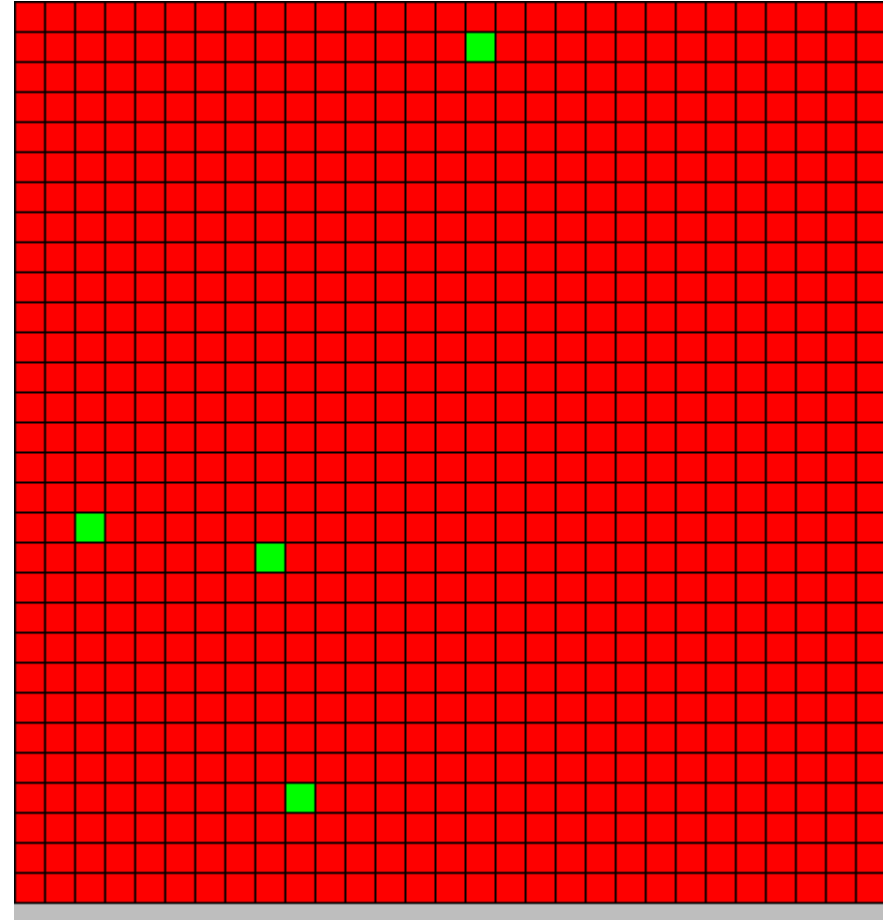
ftsK
needed for
cellular
division

Proof of Concept: Simulation to answer 4 questions

- How does differentiation induces feeding? (proof of concept)
cellular automaton (in MGS)



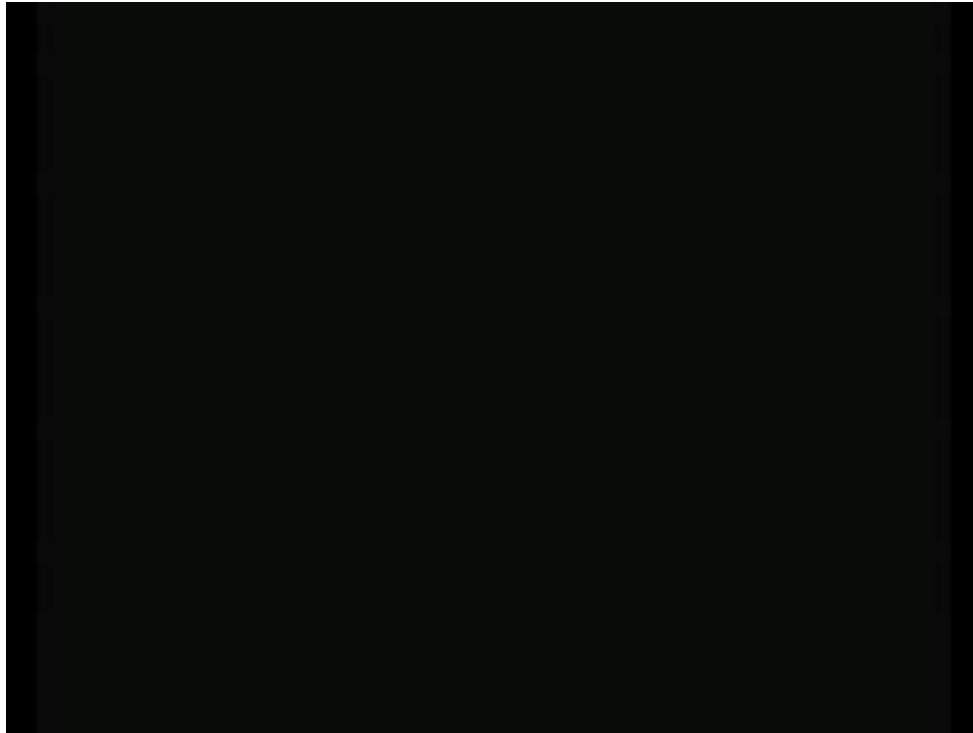
diffusion of DAP



somatic and germ cell

Proof of Concept: Simulation to answer 4 questions

- How does differentiation induces feeding? (proof of concept) cellular automaton (in MGS)
- How do spatial organization and distribution evolve?
agents based system (in MGS)



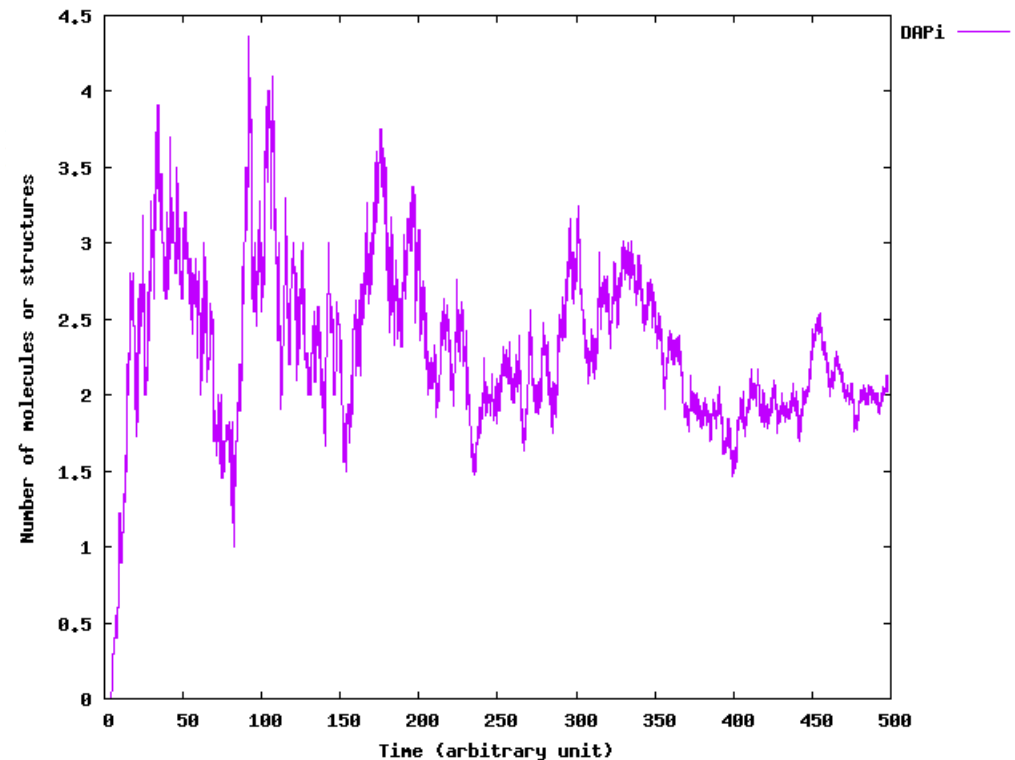
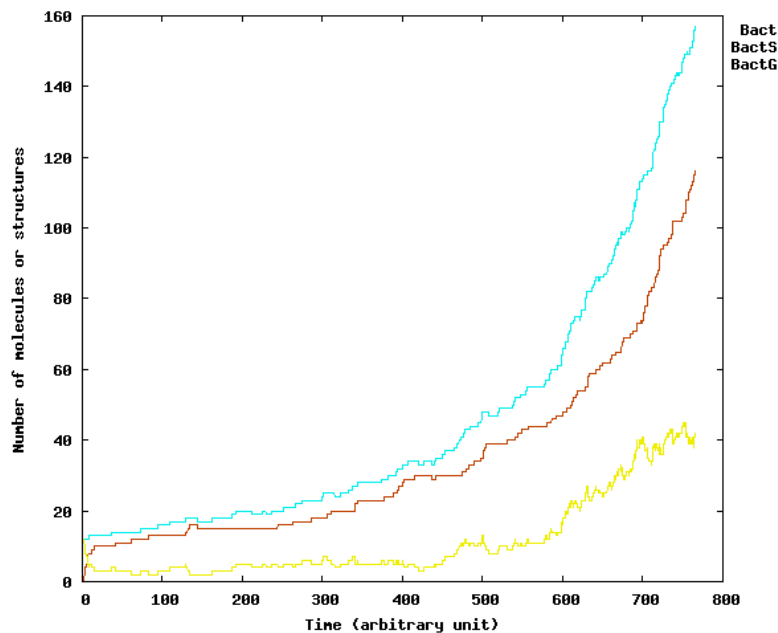
Proof of Concept: Simulation to answer 4 questions



- How does differentiation induces feeding? (proof of concept)
cellular automaton (in MGS)
- How do spatial organization and distribution evolve?
agents based system (in MGS)
- **How robust and tunable is the model?**
ODE kinetics (matlab)

Proof of Concept: Simulation to answer 4 questions

- How does differentiation induces feeding? (proof of concept)
cellular automaton (in MGS)
- How do spatial organization and distribution evolve?
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- How robust and tunable is the model?
ODE kinetics
- **How sensitive is the system to noise?**
Gillespie based simulation (in MGS)



Conclusions and Perspectives

Success

- Polytypisme is good
- Rule application strategies are good
- Patterns/rules are expressive and usually concise
- Clean semantics

Shortcomings

- Rules may be heavy (e.g. 100 variables for the fractal sponge)
graphical drawing of rules
look for better notations (e.g. path pattern)
- Efficiency
well...
- Implicit methods (solvers) are hairy
use explicit ones

- An intrinsic complexity theory
e.g., w.r.t. interactions
- A logic of spatial interactions
- Relationships to physics
a discrete differential calculus (cf. PhysicaD 08)
- Internalizing time
- Implementation
pattern-matching compilation and optimization, specific
abstract combinatorial complex, parallelism
- Non standard applications
e.g., in knowledge representation
or in music analysis (Louis Bigo talk)

A topological manifesto



Spatial computing proposes to celebrate corporeality of data rather than trying to deny it.

Simon Greenworld (MIT medialab)

- The logical approach in computer science
computation = deduction
(the Curry-Howard isomorphism)
- Other paradigms can be fruitfull : topology
computation = moving in a space
- Try to perceive space (and time) in programs
(rather than logical operations)
purposes: technical, heuristic, didactical

Call for Papers: Spatial Computing Workshop 2011

at 5th IEEE International Conference on Self-Adaptive and Self-Organizing Systems
Ann Arbor, Michigan, USA, October 3, 2011

Organizers: Jacob Beal (BBN Technologies, USA) Stefan Dulman (Delft University, the Netherlands) Olivier Michel (University Paris-Est Créteil, France) Antoine Spicher (University Paris-Est Créteil, France)

Submission Deadline: July 4th, 2011

Many self-organizing or self-adaptive systems are “spatial computers” – collections of local computational devices distributed through physical space, in which:

- the difficulty of moving information between any two devices is strongly dependent on the distance between them, and
- the “functional goals” of the system are generally defined in terms of the system’s spatial structure.

Systems that can be viewed as spatial computers are abundant, both natural and man-made. For example, in wireless sensor networks and animal or robot swarms, inter-agent communication network topologies are determined by the distance between devices, while the agent collectives as a whole solve spatially-defined problems like “analyze and react to spatial temperature variance” or “surround and destroy an enemy.”

Similarly, in reconfigurable microchip platforms, moving data between adjacent logic blocks is much faster than moving it across the chip, which in turn favors problems with spatial structure like stream processing. In biological embryos, each developing cell’s behavior is controlled only by its local chemical and physical environment, but the eventual structure of the organism is a global property of the

Thanks



- Antoine Spicher

- Olivier Michel

<http://mgs.spatial-computing.org>

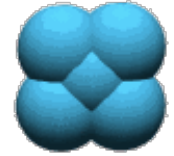
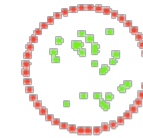
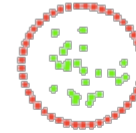
- PhD and other students

Louis Bigo

J. Cohen, P. Barbier de Reuille,

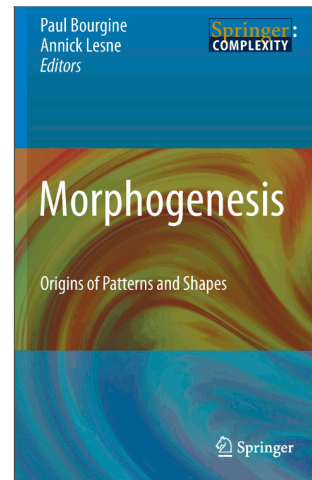
E. Delsinne, V. Larue, F. Letierce, B. Calvez,

F. Thonerieux, D. Boussié *and the others...*



- Past and presents Collaborations

- A. Lesne (IHES, stochastic simulation)
- P. Prusinkiewicz (UoC, declarative modeling)
- P. Barbier de Reuille (meristeme model)
- C. Godin (CIRAD, biological modeling)
- H. Berry (INRIA, stochastic simulation)
- G. Malcolm (Liverpool, rewriting)
- J.-P. Banâtre (IRISA, programming)
- P. Fradet (Inria Alpes, programming)
- F. Delaplace (IBISC, synthetic biology)
- P. Dittrich (Jena, chemical organization)
- F. Gruau (LRI, language and hardware)
- P. Liehnard (Poitiers, CAD, Gmap and quasi-manifold)



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