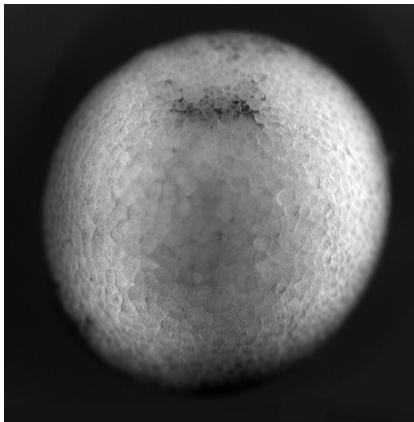
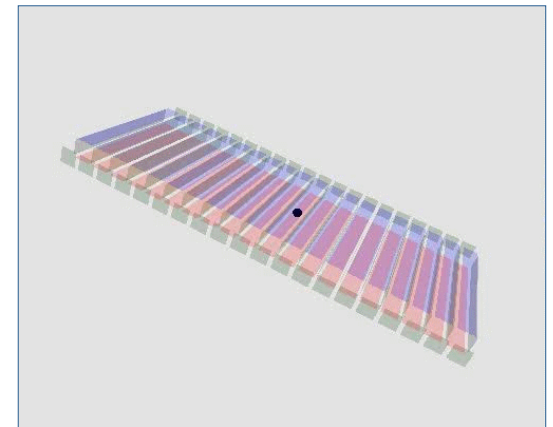


Interaction based simulation of dynamical systems with a dynamical structure $(DS)^2$ in MGS



mgs.spatial-computing.org



Jean-Louis Giavitto¹ & Antoine Spicher² & Olivier Michel²

¹IRCAM, CNRS & UPMC

²LACL, University Paris Est Créteil

SCSC – Jun. 2011

Outline



- Motivations : modeling morphogenesis
 - $(DS)^2$ and their modeling
 - The topological structures of interactions
- Topological collection and their transformation in MGS
 - Topological collections
 - Transformation
 - Examples
- Applications
 - The flock of birds
 - The growth of an epithelial tissue
 - Modeling a synthetic bacteria

Motivations



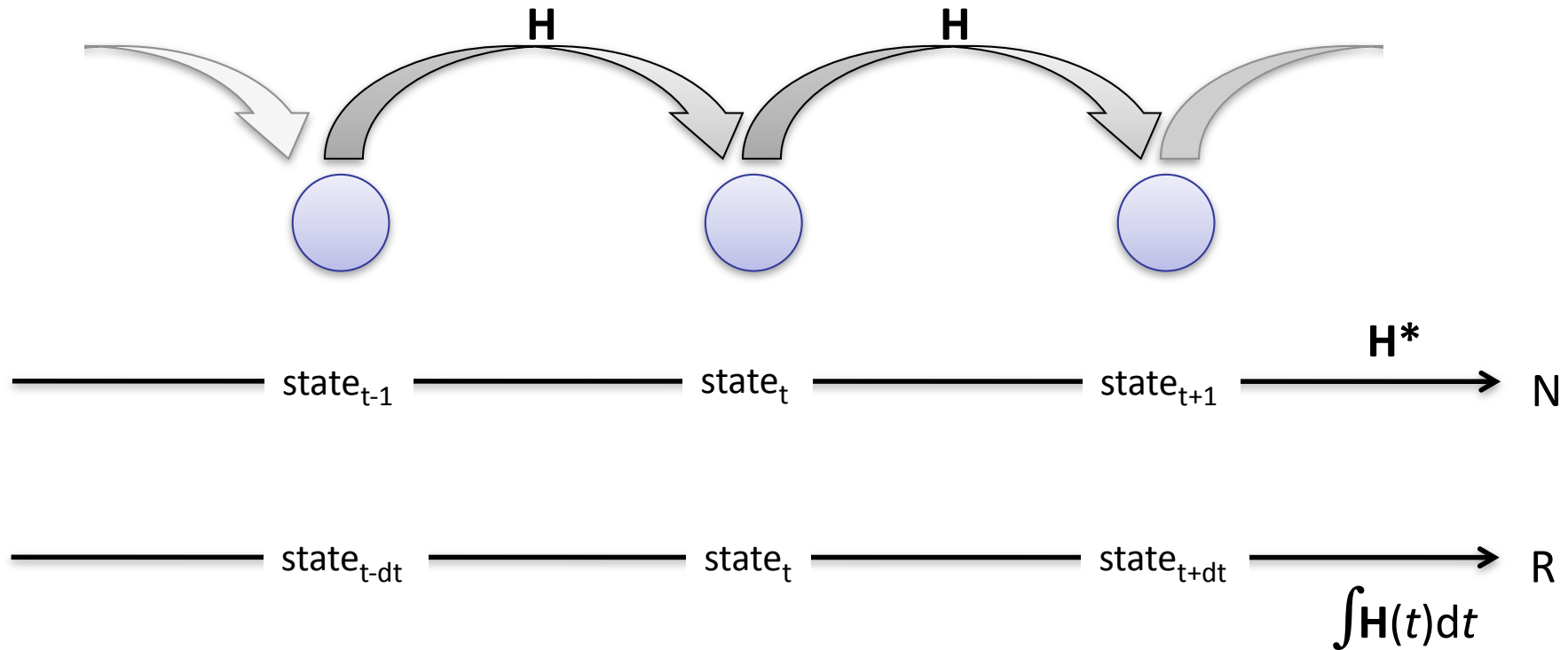
- Dynamical Systems *and* dynamical structure
 - Morphogenesis = patterning formation + growth
- $(DS)^2$
- The modeling of $(DS)^2$
 - Locality
- The topological structure of interaction
- A general picture



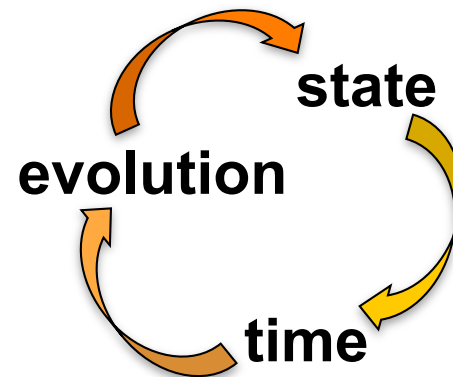
Dynamical systems and Dynamical Structures

(a problem outlined by A. Turing)

Specifying a dynamical system (for simulation)



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



Formalism for Dynamical System

- 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 medium/process problem

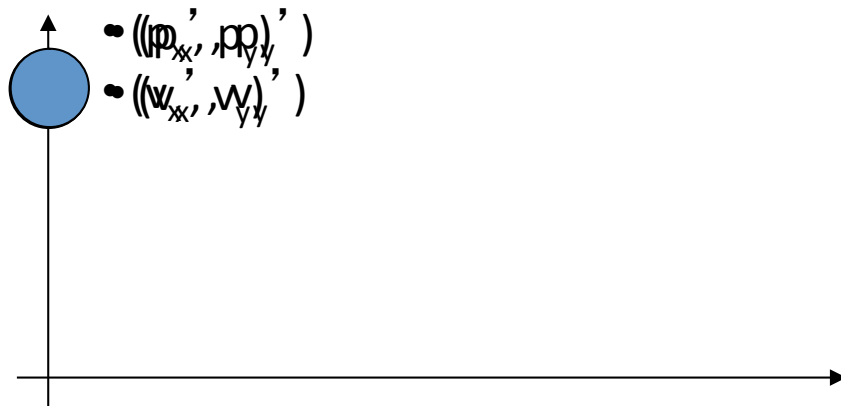


THE CHEMICAL BASIS OF MORPHOGENESIS

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)

The medium/process problem



THE CHEMICAL BASIS OF MORPHOGENESIS

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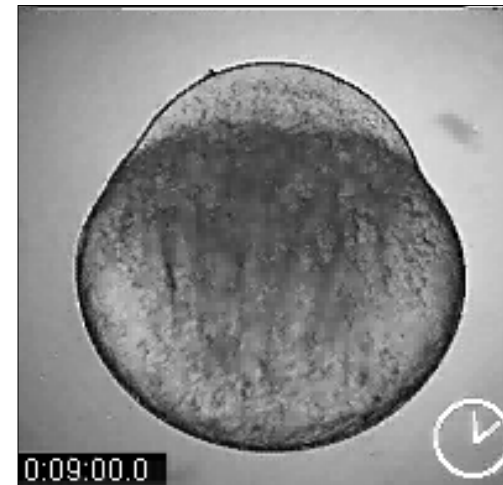
a falling ball



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**

Modelling morphogenesis: the predefined medium

The interdependence of the chemical and mechanical data adds enormously to the difficulty, and attention will therefore be confined, so far as is possible, to cases where these can be separated.

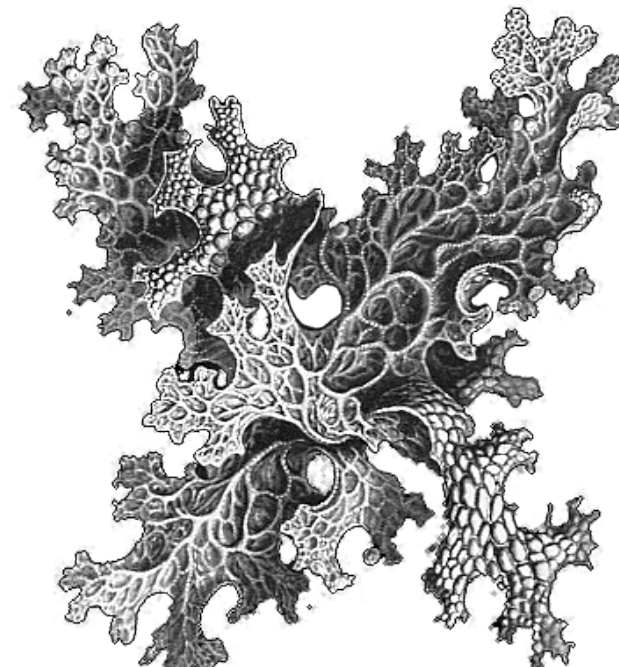
Suppose, for instance, that a 'leg-evocator' morphogen were being produced in a certain region of an embryo, or perhaps diffusing into it, and that an attempt was being made to explain the mechanism by which the leg was formed in the presence of the evocator. It would then be reasonable to take the distribution of the evocator in space and time as given in advance and to consider the chemical reactions set in train by it.

Compatible with

- the notion of morphogenetic field
- cell fate

But

- there is evidence for
**feedback loops between the shape
and the process inhabiting the shape**



from E. Haenkel (cited by C. Goodman-Strauss): example of a negative curvature surface. Curvature can be controlled while the surface is growing along a 'front'

Patterning vs. Growth

■ Dynamics ON form

patterning, motif formation in a predefined form

- Diffusion, reaction-diffusion, transport (continuous models)
- cellular automata (discrete models)

■ Dynamics OF form

growth, deformation of a shape

- Deformation of elastic bodies (continuous models)
- Lindenmayer systems (discrete models)

The interplay between state and form

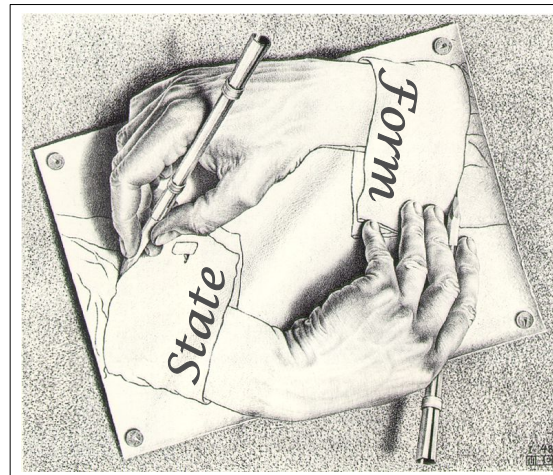
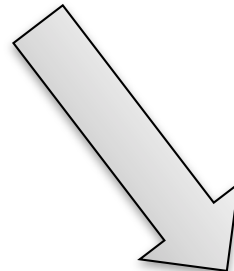


a developing embryo

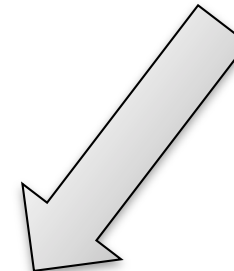
the state **as well as the structure of the state**
is changing in time

*(chemical and mechanical state of each cell **as well as the arrangement of the cells**)*

Dynamics ON form



Dynamics OF form



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

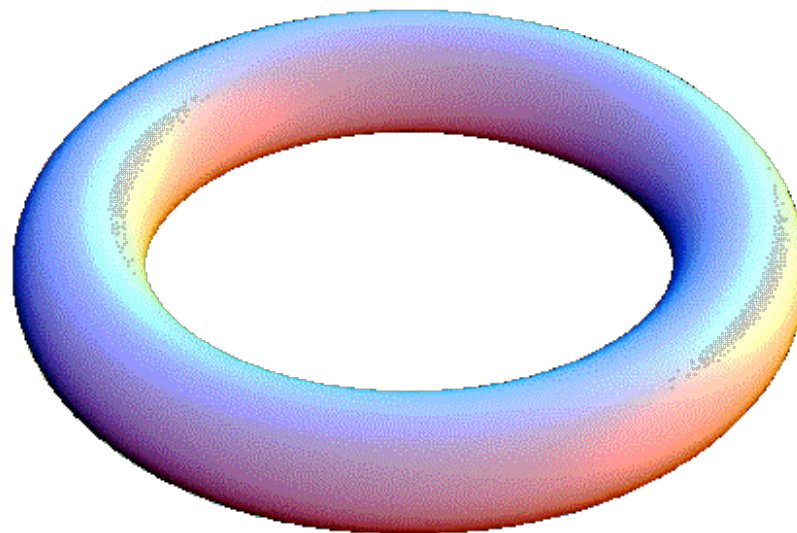
$(DS)^2$ versus DS

- Processes in the form are topologically (geometrically) meaningful
e.g. growth rate
- Topological (geometric) information is meaningful
e.g. domain of diffusion, information transfert

OK: the coupling is important.

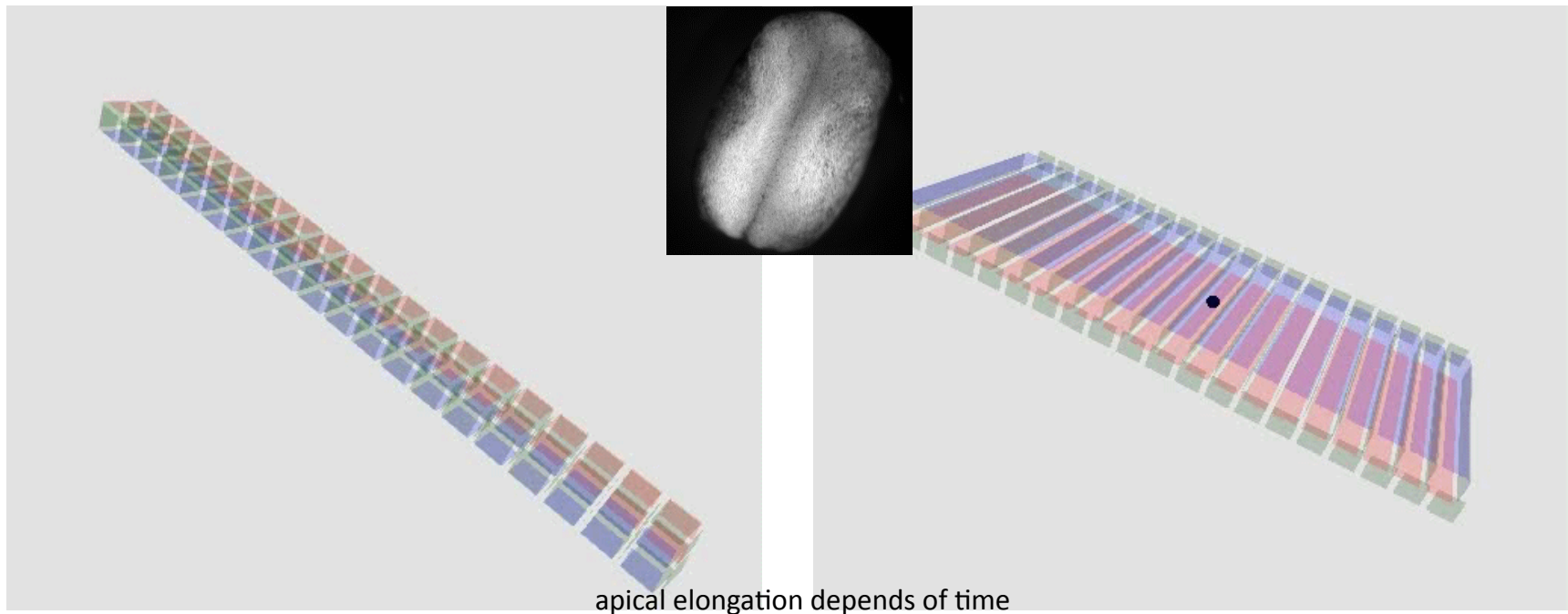
What can we do?

1. **“Dynamics ON form” toward DS^2**
 - parameterize (control) the shape by some quantities (e.g. curvature for a manifold or adjacency matrix for a graph)
 - link these quantities with processes in the shape
 - growth depending on concentration



Differential geometry is not enough

- Encoding a shape into continuous parameters is difficult
- It does not handle topological changes very well



- Ok, sometimes they are some tricks (e.g. level set) but they are very smart tricks

OK: the coupling is important. What can we do?

1. “Dynamics ON form” toward DS^2

- parameterize (control) the shape by some quantities (e.g. curvature for a manifold or adjacency matrix for a graph)
- link these quantities with processes in the shape
 - growth depending on concentration



2. “Dynamics OF form” toward DS^2

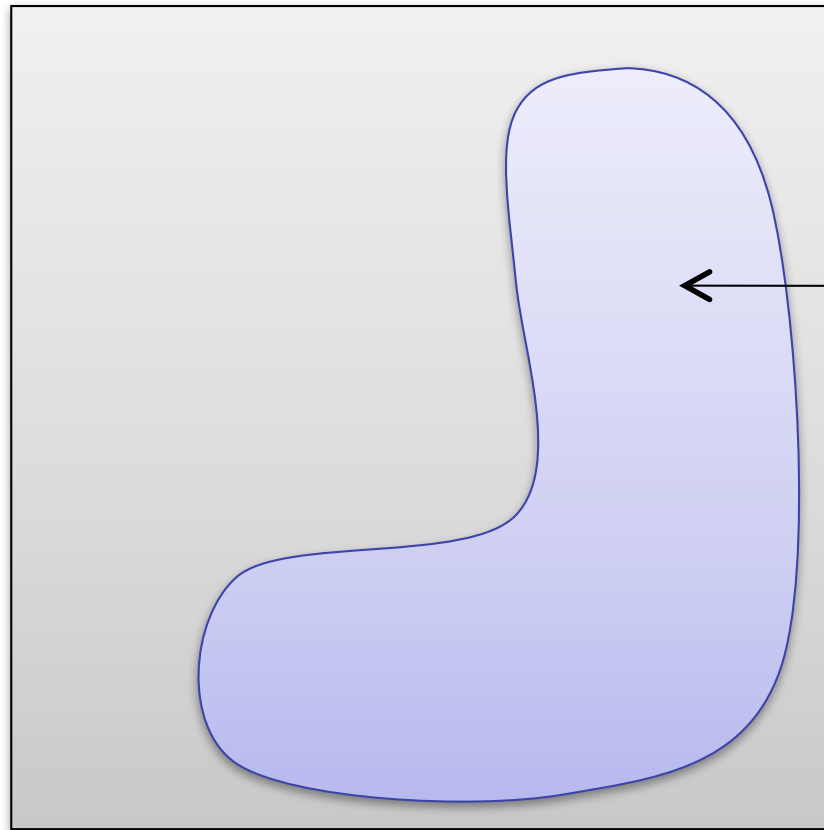
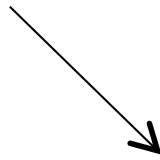
- enhance the form by parameters
- put a dynamic on these parameters
example: module in Lindenmayer systems



The Topological Structure of Interactions

Decompose a system into subsystems following the elements in interaction

A system in some state

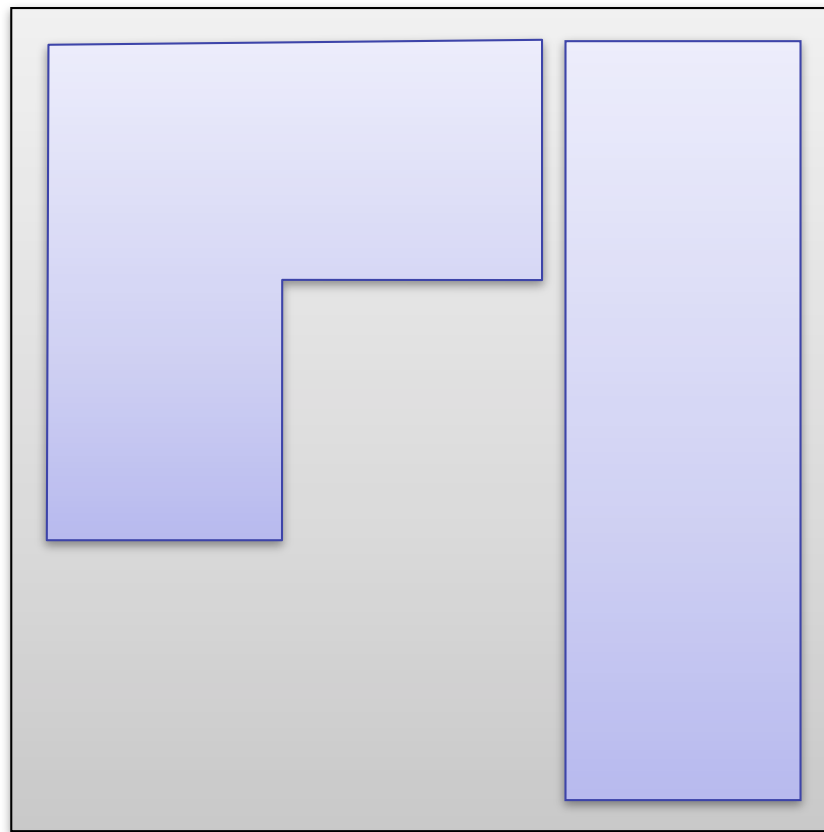


*Part of a system
that evolves.*

*Can be identified
by comparison
with the previous
global state*

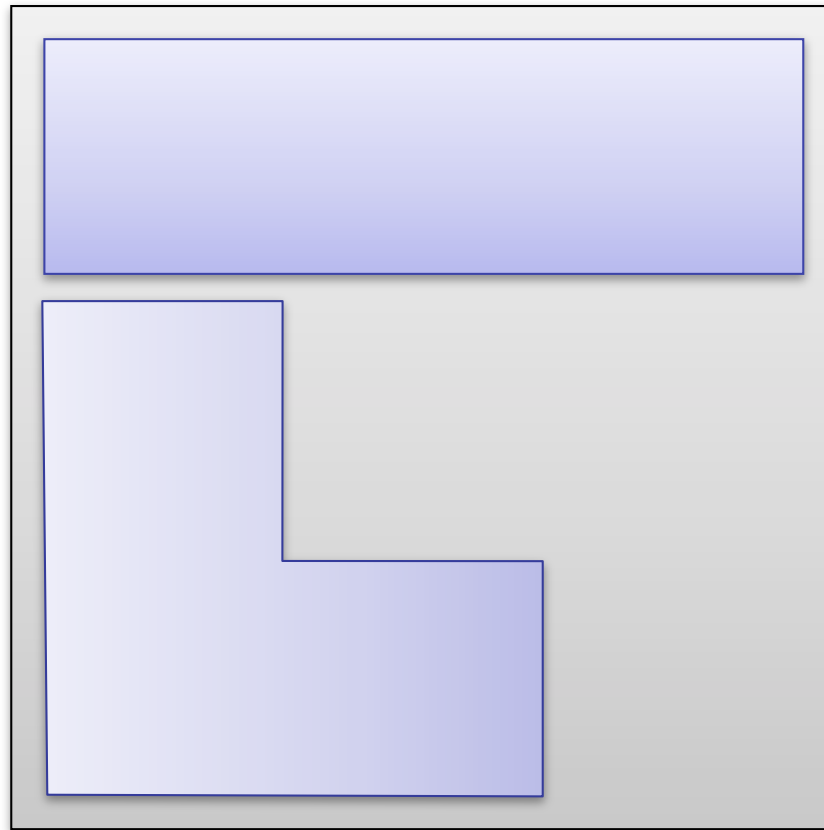
Decompose a system into subsystems following the elements in interaction

$t = 1$



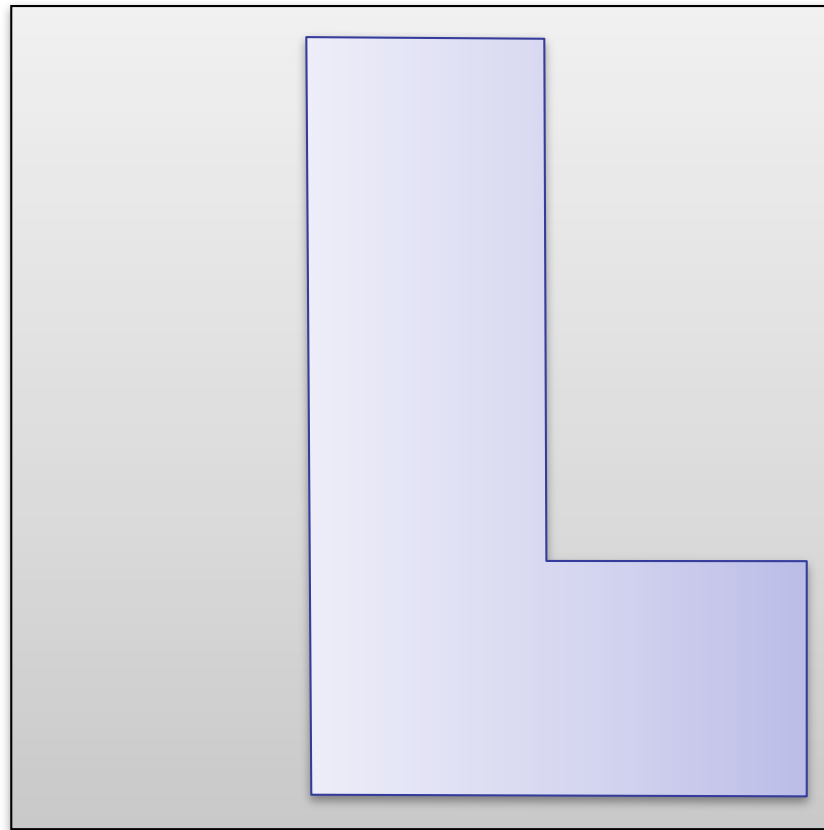
Decompose a system into subsystems following the elements in interaction

$t = 2$



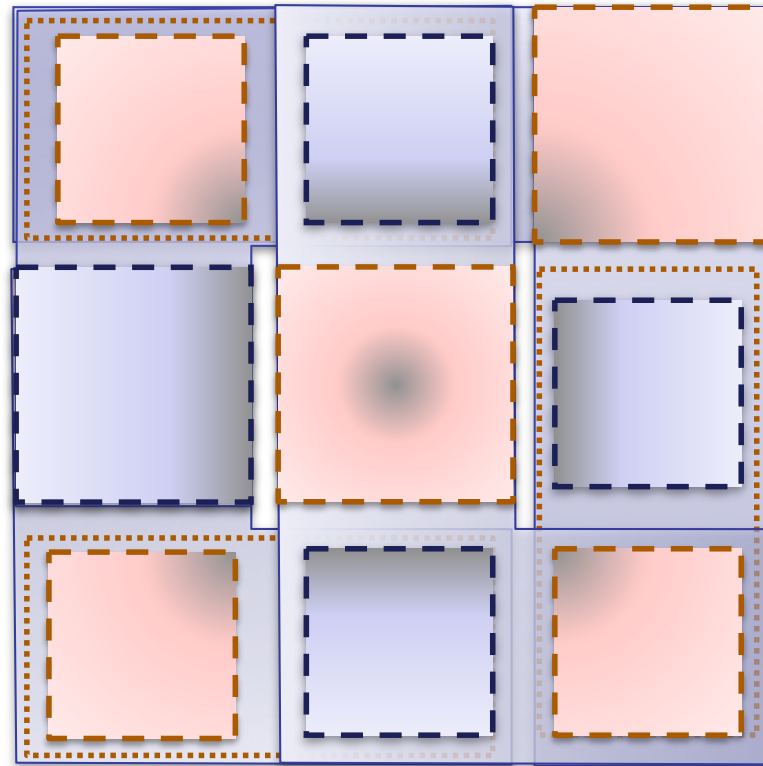
Decompose a system into subsystems following the elements in interaction

$t = 3$

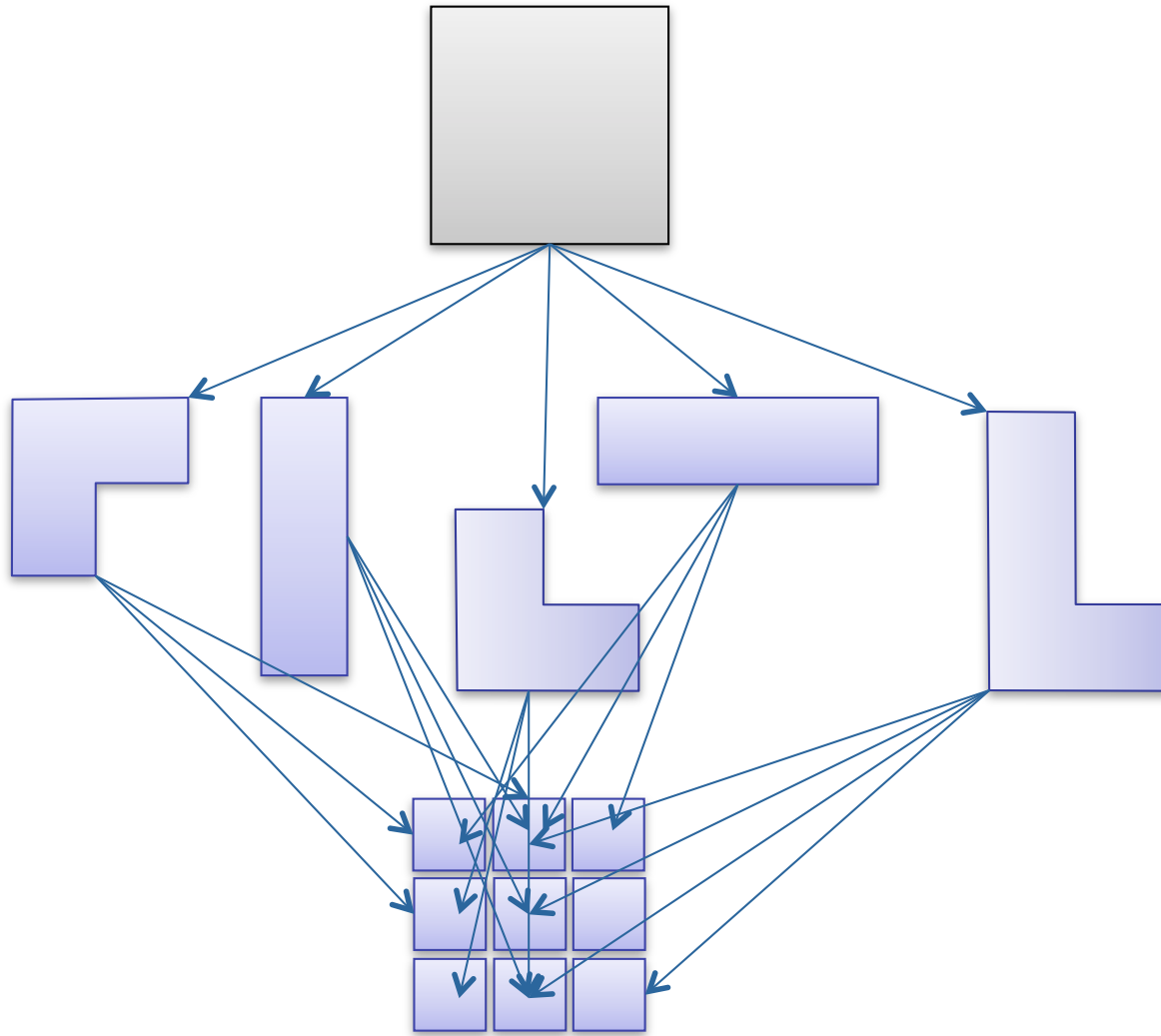


Decompose a system into subsystems following the elements in interaction

**The interactions decomposes the systems into elementary parts.
An interaction implies one or several elementary parts.**



Decompose a system into subsystems following the elements in interaction



the inclusion structure
between the
elementary and
interacting parts is a
lattice

a (simplicial) complex
is a better (topological)
equivalent
representation

The grand picture



1. Describe a dynamical system following the interaction of its parts
2. Each part is characterized by a (local) state
3. The global state of the system is the “sum” of its local state and their topological organization
4. An interaction makes evolve a (small) subset of local states
5. An interaction potentially changes the topological organization of state

Topological Collections and their Transformation in MGS

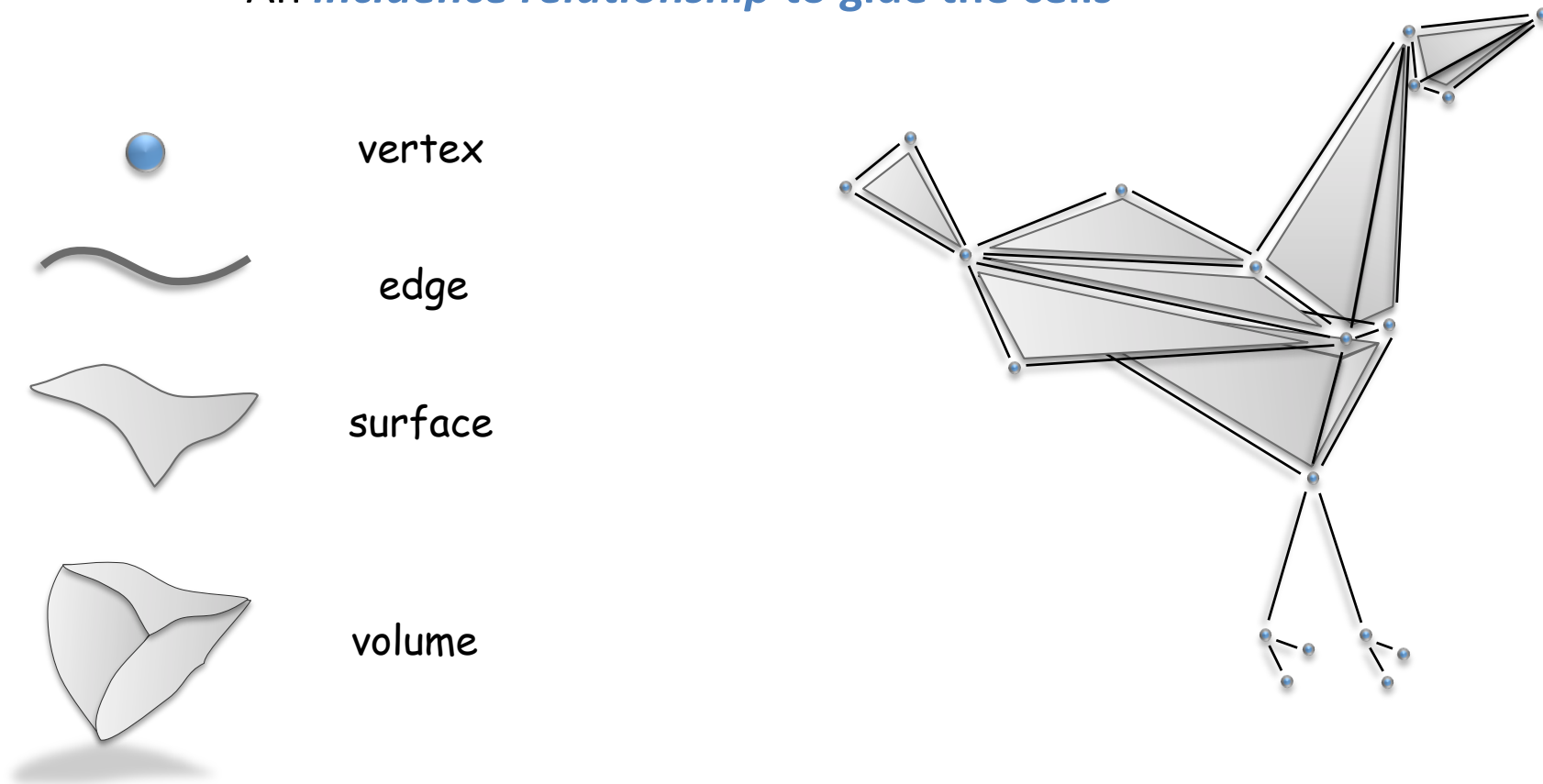
- Topological collections
- Topological Rewriting
- MGS examples

MGS Proposition

■ Topological collections

□ Structure

- A collection of topological cells
- An *incidence relationship to glue the cells*



MGS Proposition

■ Topological collections

□ Structure

- A collection of topological cells
- An incidence relationship

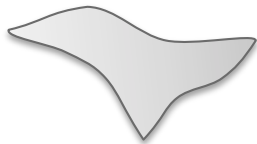
□ Data: **association of a value with each cell**



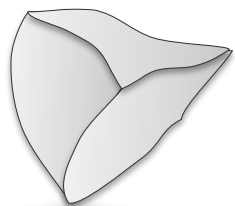
0-cell



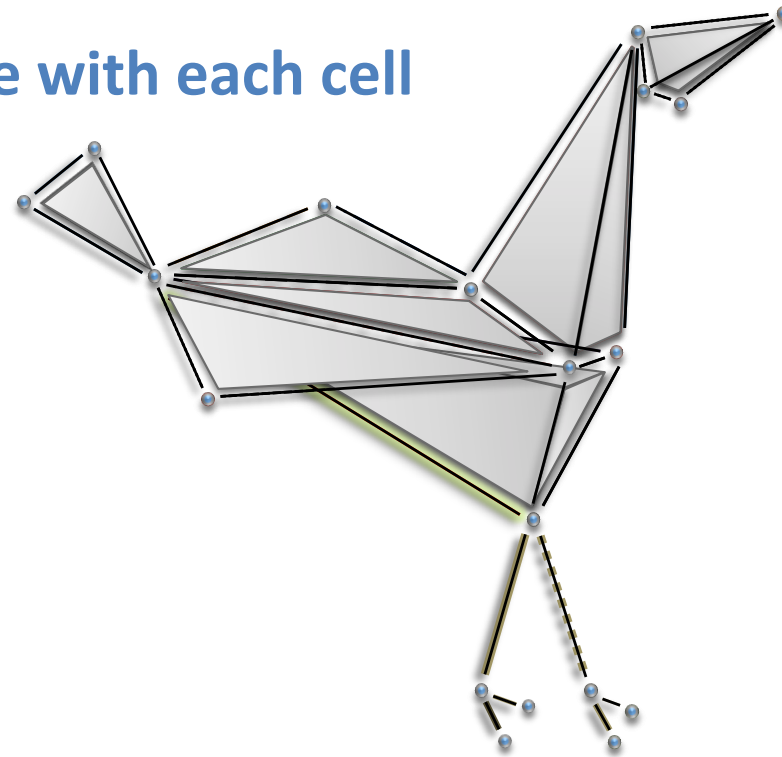
1-cell



2-cell



3-cell



MGS Proposition

■ Transformations

- Functions defined by case on collections

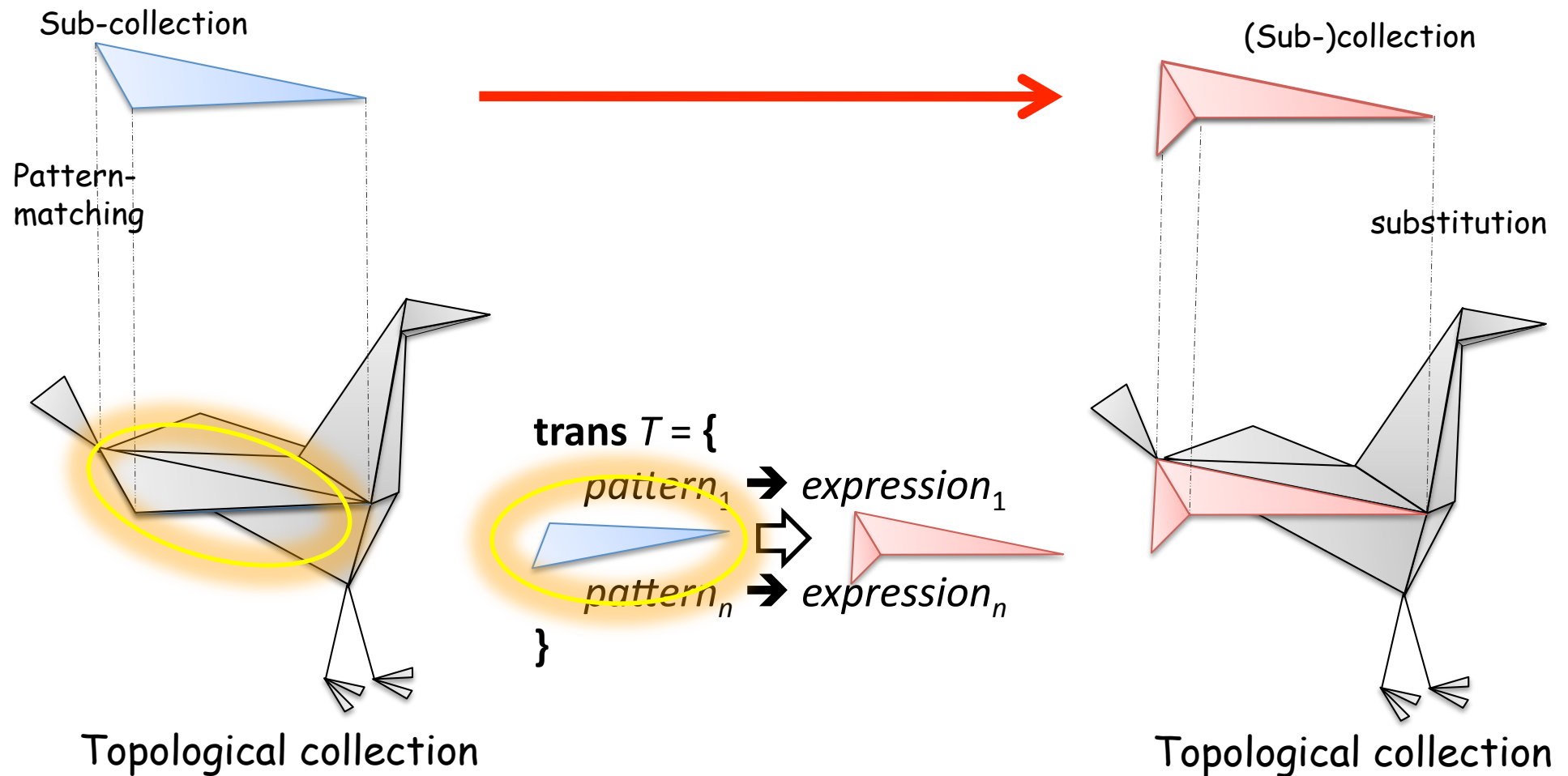
Each case (pattern) matches a sub-collection

- Defining a rewriting relationship: *topological rewriting*

```
trans T = {  
  pattern1 → expression1  
  ...  
  patternn → expressionn  
}
```

MGS Proposition

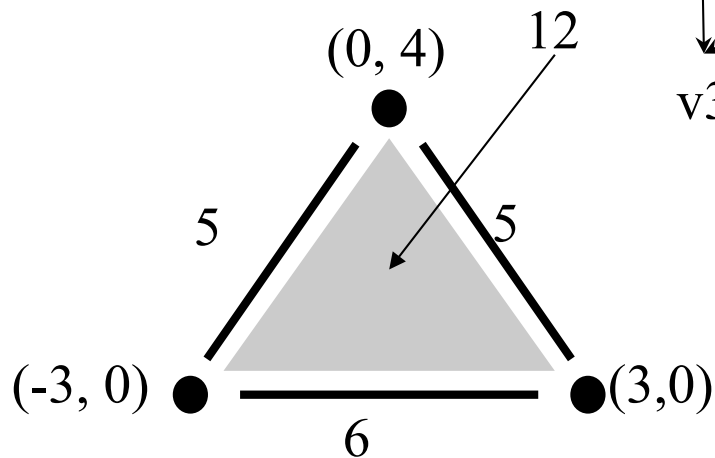
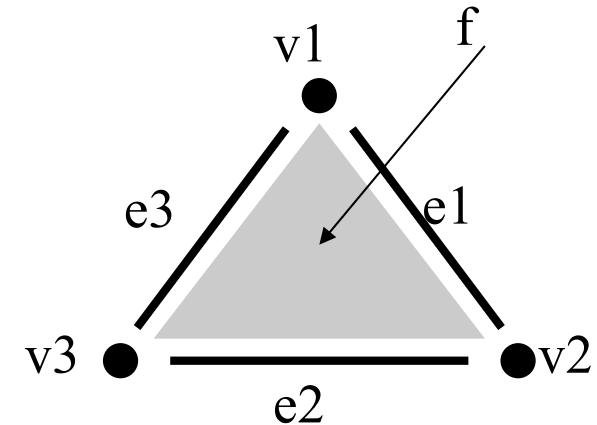
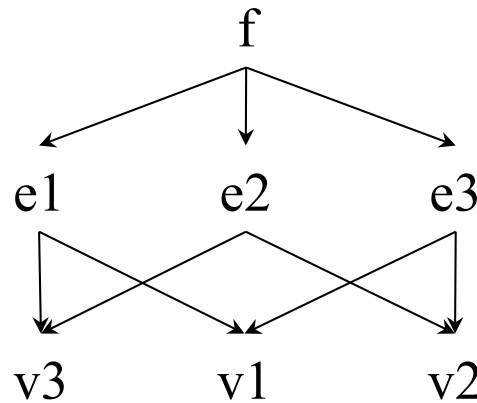
■ Transformations



Abstract Simplicial Complex and simplicial chains

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

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

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



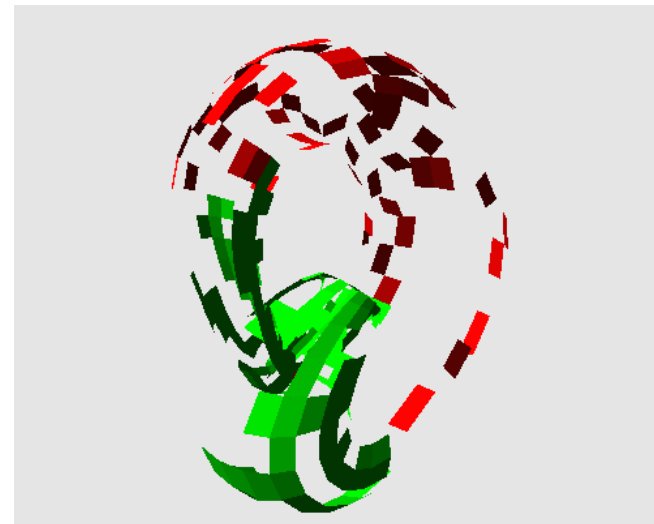
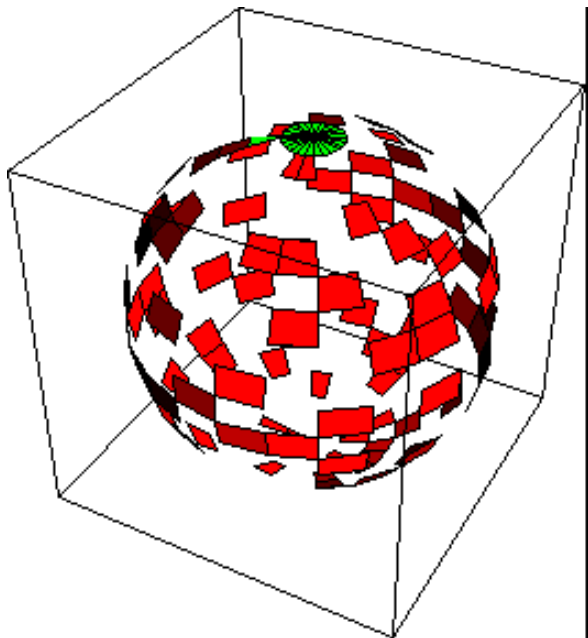
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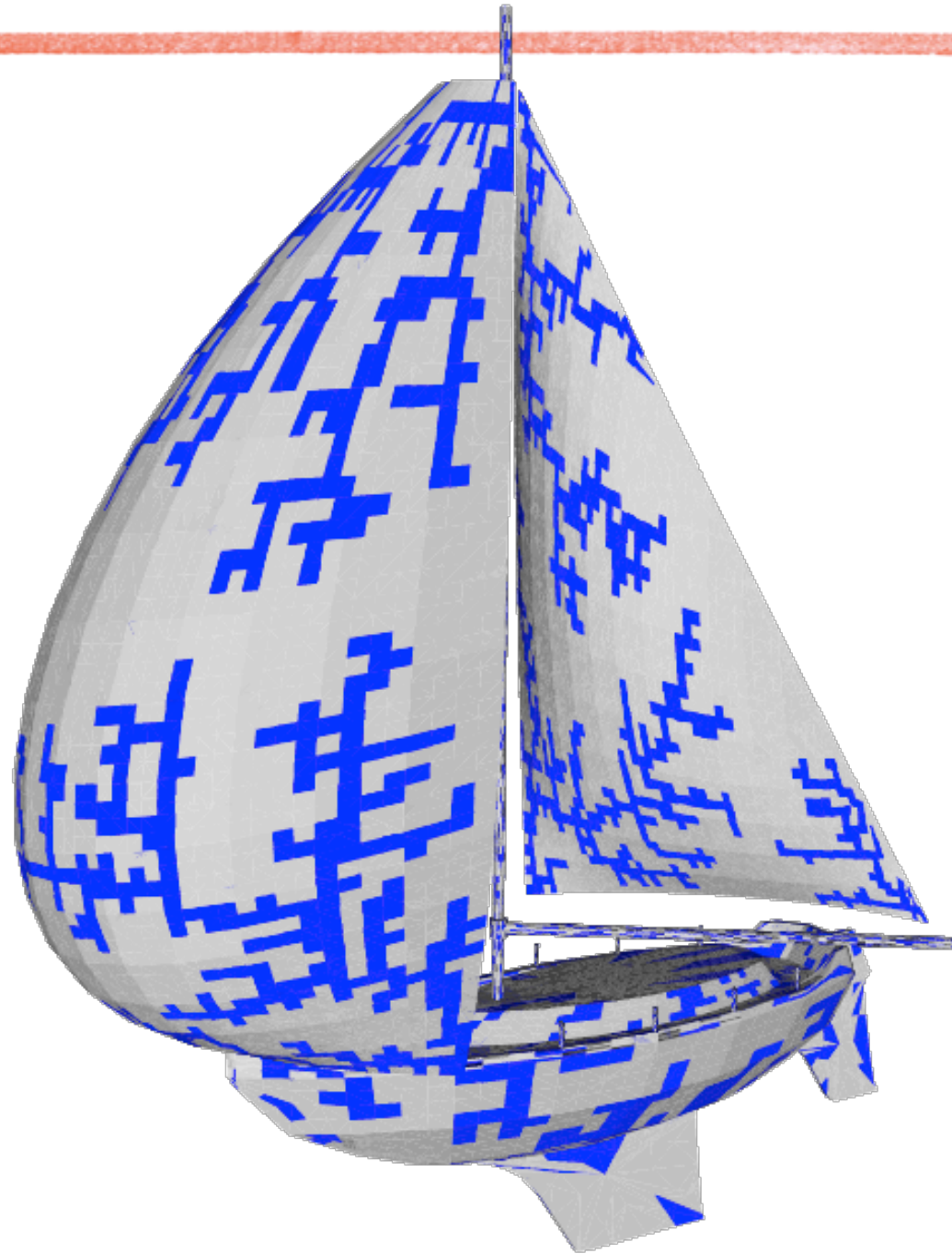
```
trans dla = {  
  `mobile , `fixed => `fixed, `fixed ;  
  `mobile , <undef> => <undef>, `mobile  
}
```

NEIGHBOR OF

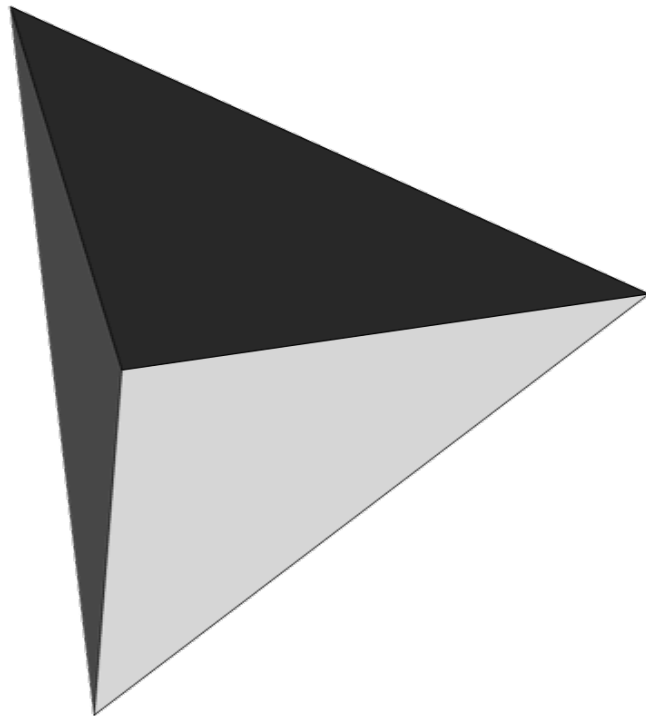
this transformation is an **abstract** process that can be applied to any kind of space



Polytypisme

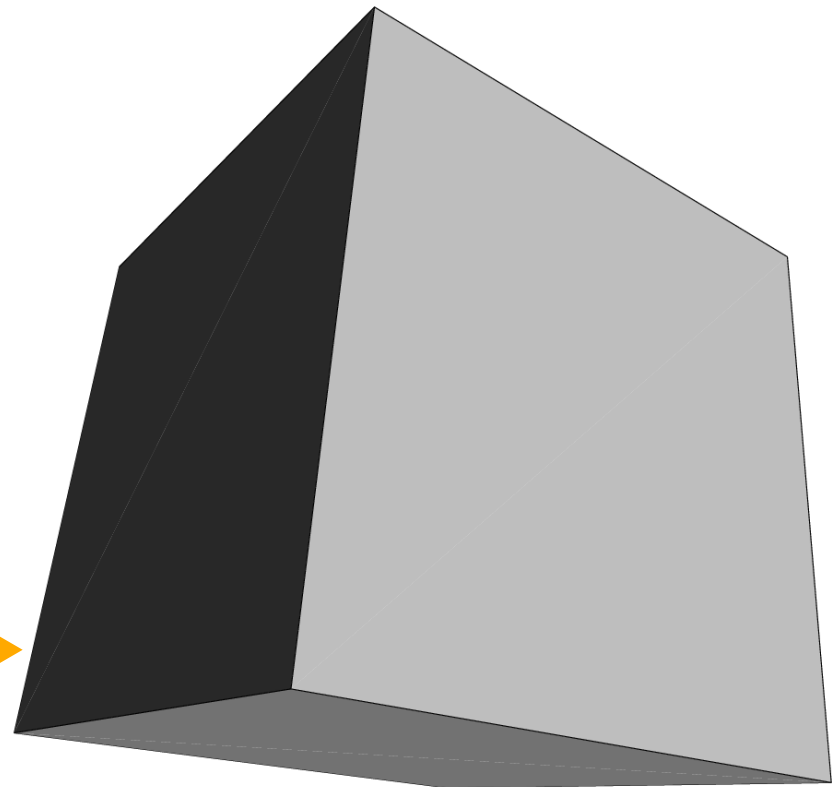


Fractal construction by carving



Sierpinsky sponge (4 steps)

Menger sponge (2 steps)



The Spatial Approach

- 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

Why higher dimensional objects and not just graphs?

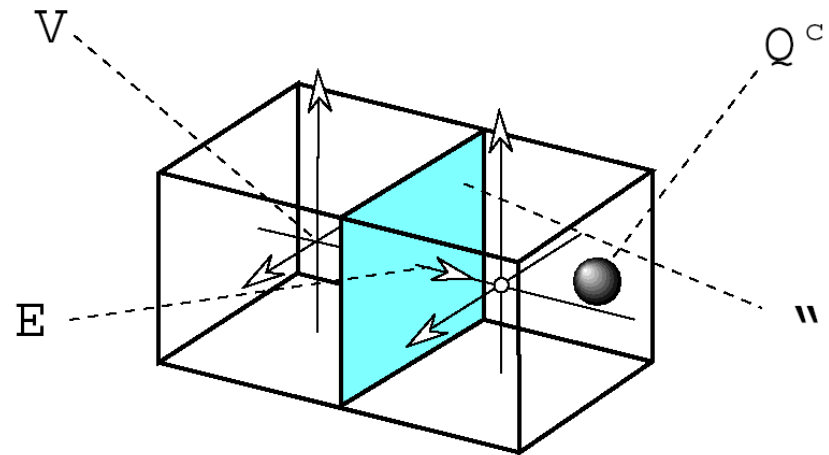
Example of electrostatic Gauss law [Tonti 74]

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

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

electric field in space:

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



Properties w.r.t. (DS)²

- *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

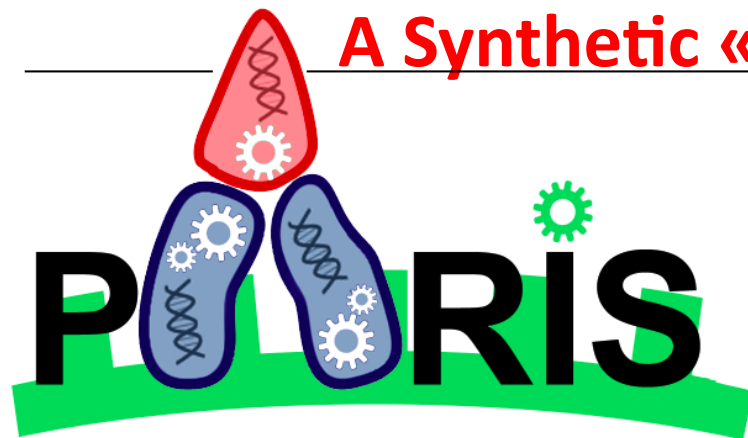
- *the phase space is well defined but not well known*

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)



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: citaulike, connotes, PubMed](#)

Registry of Standard Biological Parts

http://parts.mit.edu/

Parts

Registry of Standard Biological Parts

MIT
Massachusetts Institute of Technology

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 T 7

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

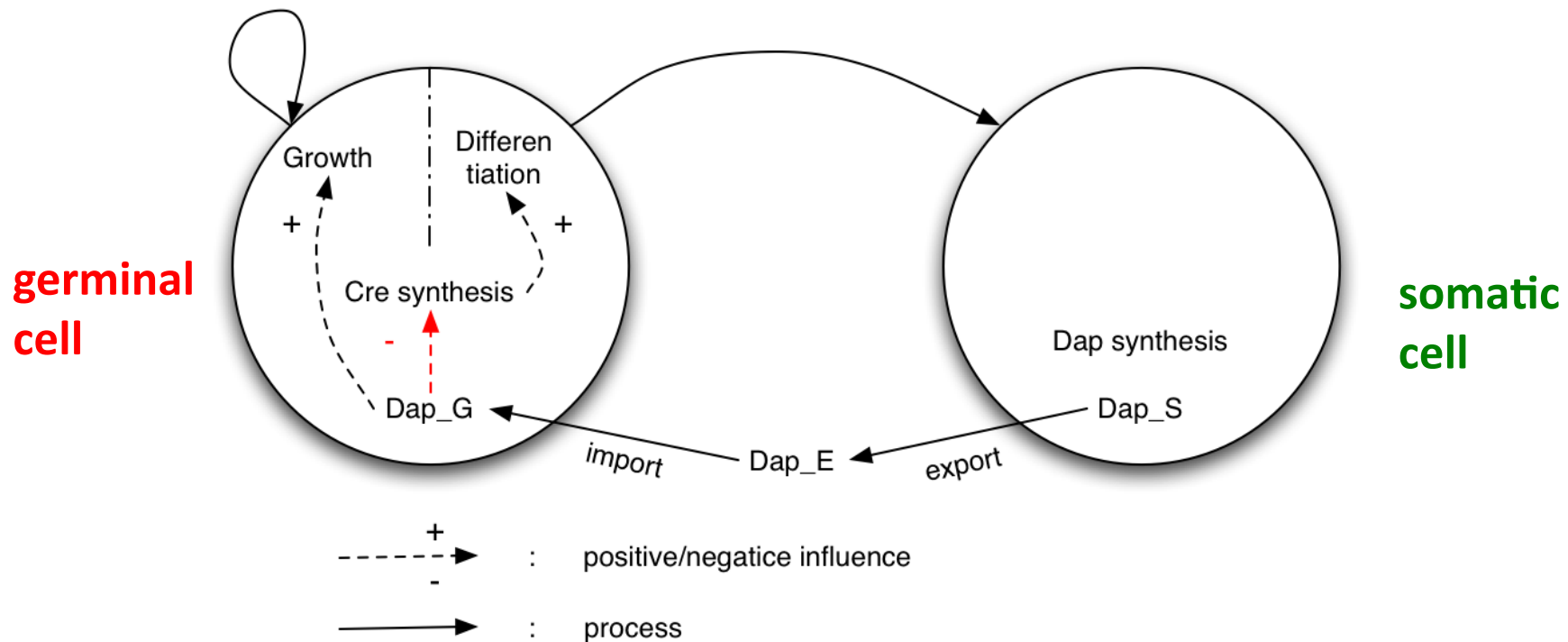
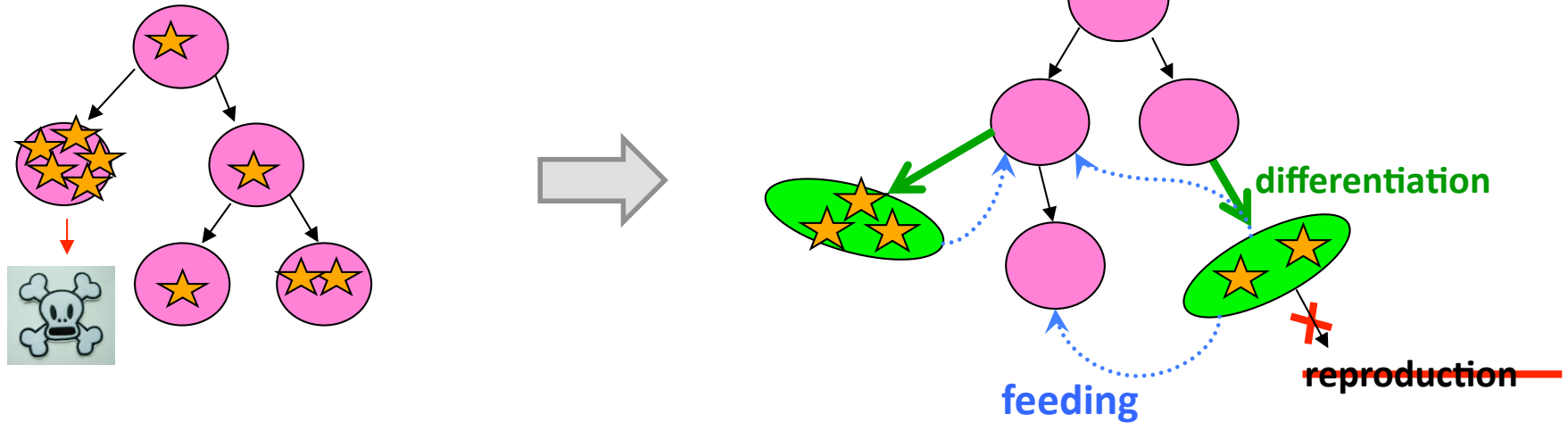
The Registry is looking for full-time Technical Assistants and Web Programmers. Please contact Staffing Services at MIT for details: [Technical Assistant](#), [Web Programmer](#).

Production at rosalind - 4.4.05

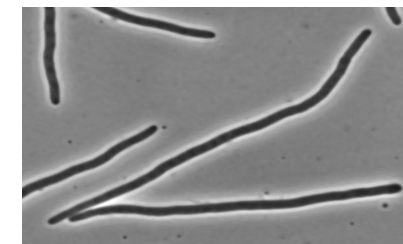
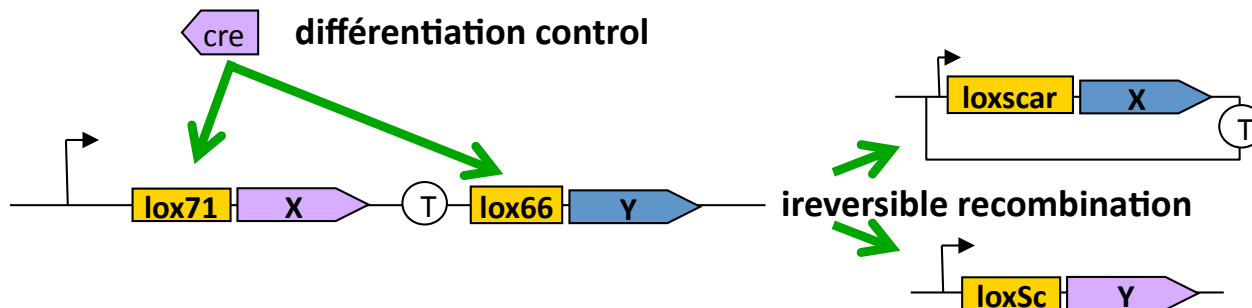
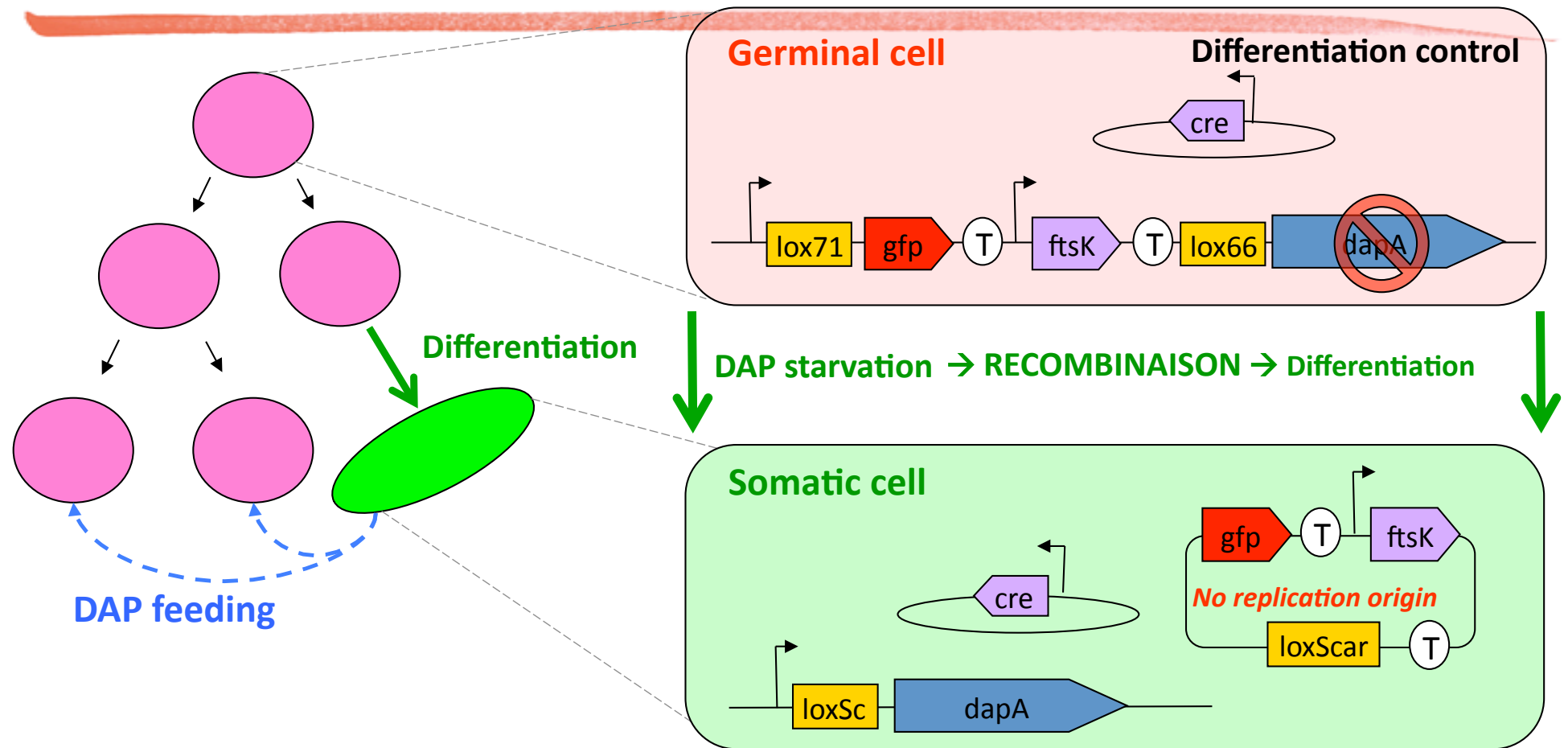


David Bikard, Thomas Landrain, David Puyraimond, Eimad Shotar, Gilles Vieira, Aurélien Rizk, David Guegan, Nicolas Chiaruttini, Thomas Clozel, Thomas Landrain

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



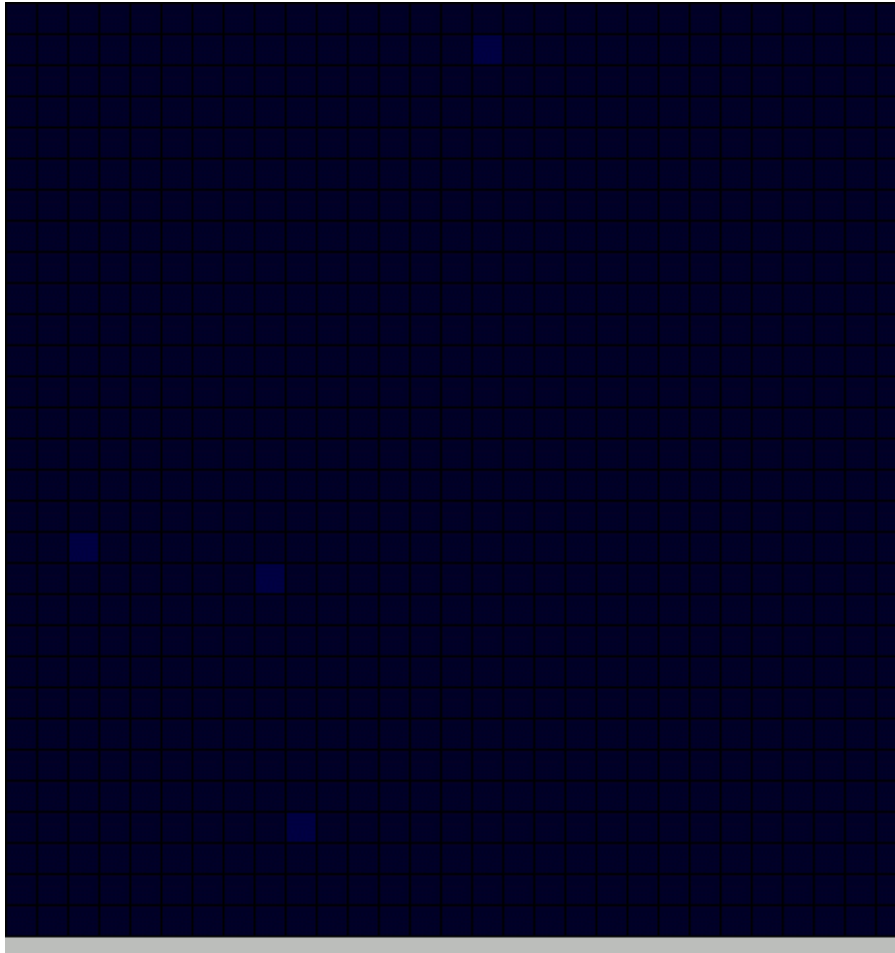
Implementation using BioBricks



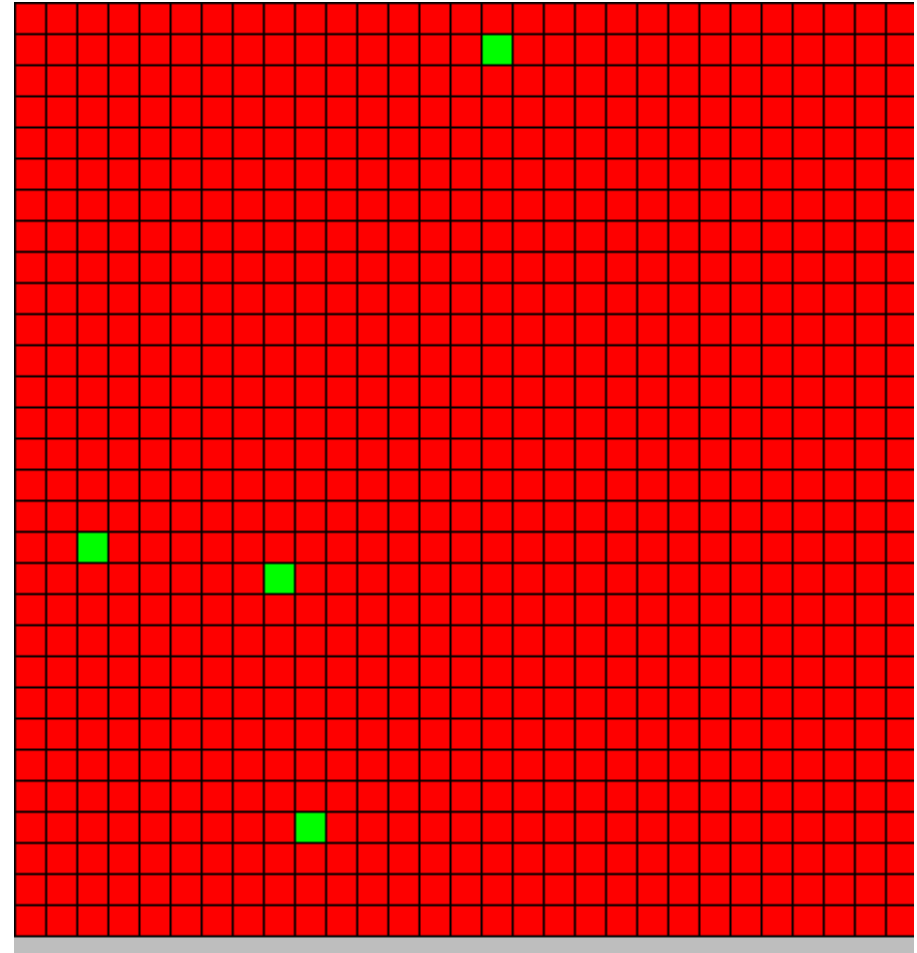
ftsK
needed for
cellular
division

Proof of Concept: Simulation to answer 4 questions

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



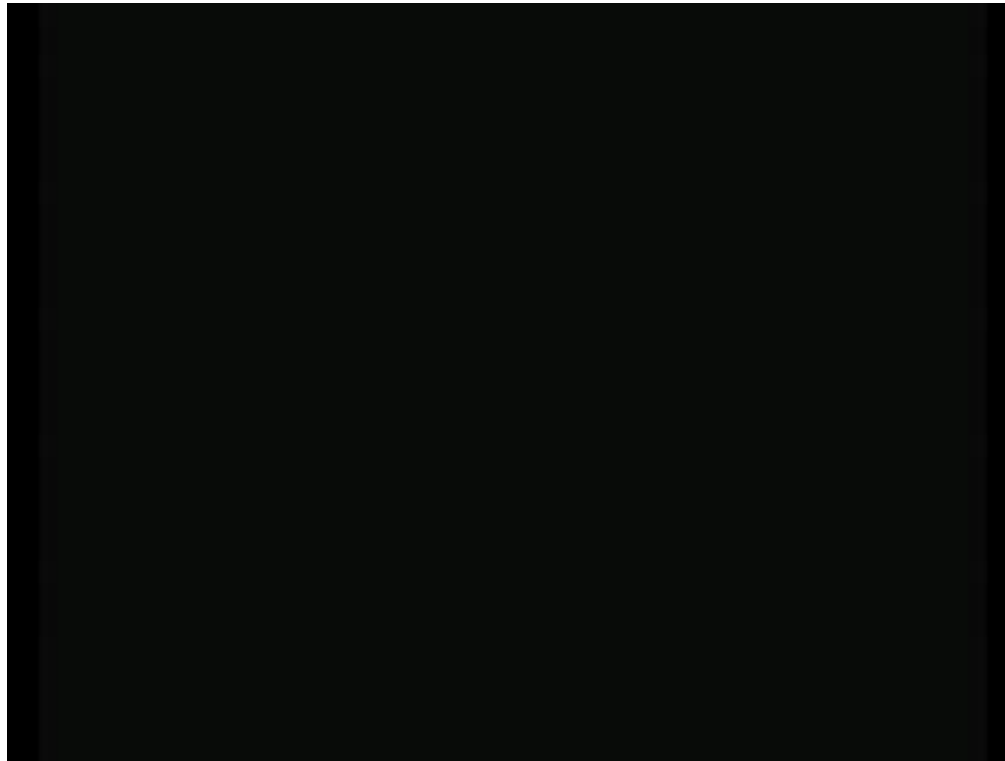
diffusion of DAP



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?
(diffusion in a growing medium) **agents based system (in MGS)**

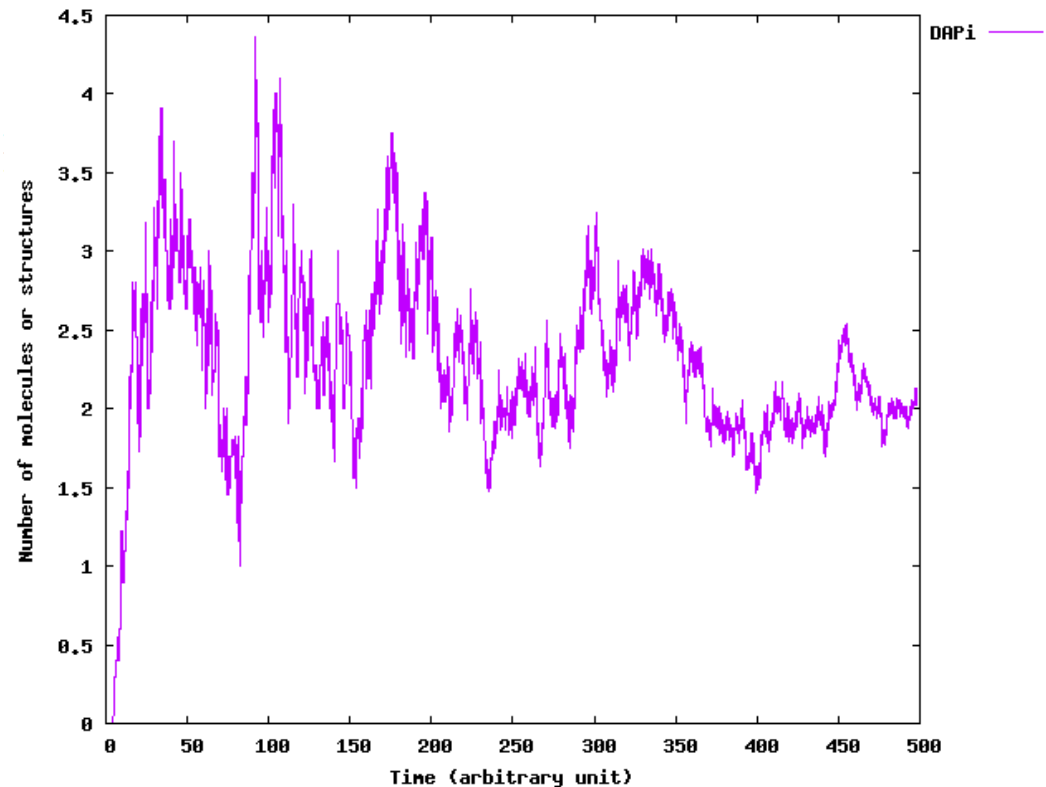
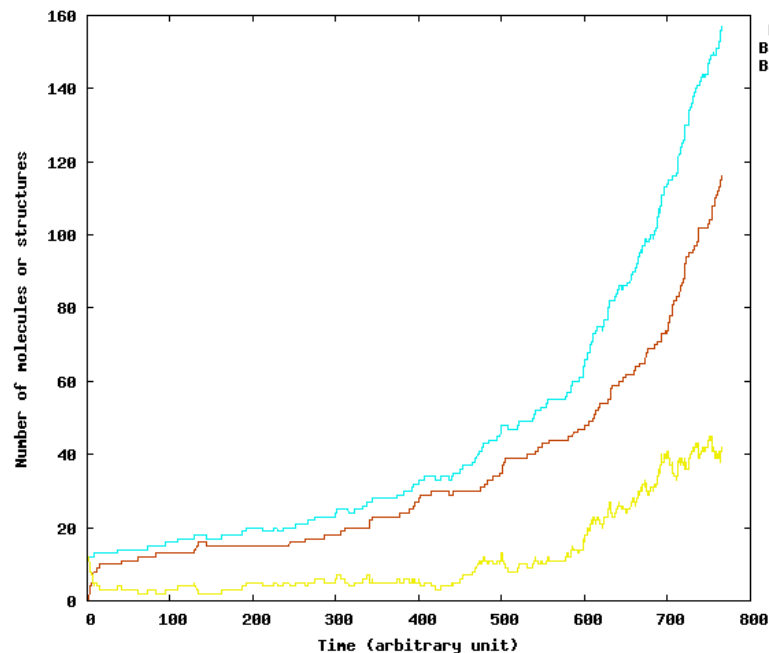


Proof of Concept: Simulation to answer 4 questions

- **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?
agents based system (in MGS)
- How robust and tunable is the model?
ODE kinetics
- How sensitive is the system to noise?
Gillespie based simulation (in MGS)



MGS drawbacks and successes

Success

- Polytypisme is 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

Conclusion & Perspectives

- A spatial & declarative approach to (DS)2 modeling & simulation
- Generic, formal, expressive & concise language
 - Implementation (mgs.spatial-computing.org)
 - Validation in many fields
 - Self-assembly processes
 - Systems biology
 - Protocol verification
 - ...
- Future work
 - Typing, optimizing, compiling transformations
 - Refinement of chemical programs (Gamma, P systems)
 - Music(ologic) representations & analysis

Thanks

ibisc

université
evry
val-d'essonne

genagole

cirs

UNIVERSITÉ
PARIS-EST
CRETEIL
VAL DE MARNE

UPEC

lcl

ircam
Centre
Pompidou

- Jean-Louis
- Antoine Spicher
- Olivier Michel

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

- **PhD and other students**

Julien, Antoine, P. Barbier de Reuille,
T. Louail, E. Delsinne, V. Larue, F. Letierce,
B. Calvez, F. Thonerieux, D. Boussié,
iGem'07 Paris team, *and the others...*

- **Some past and present collaborations**

- A. Lesne (IHES, stochastic simulation)
- P. Prusinkiewicz (Calgary, declarative modeling)
- C. Godin (CIRAD, biological modeling)
- H. Berry (LRI, stochastic simulation)
- G. Malcolm (Liverpool, rewriting)
- J.-P. Banâtre (IRISA, programming)
- P. Fradet (InriaAlpes, programming)
- F. Delaplace (IBISC, synthetic biology)
- D. Pumain (Géographie-Cité, city growth)
- R. Doursat (ISC, morphogenetic engineering)
- F. Gruau (U. PXI, language and hardware)
- P. Liehnard (Poitiers, CAD, Gmap and quasi-manifolds)

Jean-Louis



Antoine



Olivier

