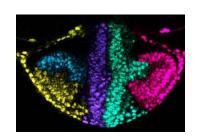


Heterogeneous collective motion or moving pattern formation? The two sides of embryogenesis combined by multi-agent modeling into a

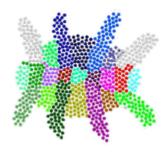


René Doursat

http://www.iscpif.fr/~doursat









Systems that are self-organized and architectured



free self-organization

the scientific challenge of complex systems: how can they integrate a true architecture?

the engineering challenge of complicated systems: how can they integrate selforganization?



deliberate design





decompose the system

self-organized architecture / architectured self-organization



Toward programmed self-organization

Self-organized (complex) systems

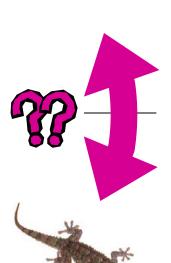
- ✓ a myriad of self-positioning, self-assembling agents
- ✓ collective order is not imposed from outside (only influenced)
- ✓ comes from purely *local* information & interaction around each agent
- ✓ no agent possesses the global map or goal of the system.
- ✓ but every agent may contain all the *rules* that contribute to it

Structured systems

eradicate any illusion of ID by <u>explaining</u> the <u>developmental</u> mechanisms of spontaneous architecture formation (not just evolutionary)

- ✓ true architecture: non-trivial, complicated morphology
 - hierarchical, multi-scale: regions, parts, details, agents
 - modular: reuse, quasi-repetition
 - *heterogeneous*: differentiation & divergence in the repetition
- ✓ random at the microscopic level, but reproducible (quasi deterministic) at the mesoscopic and macroscopic levels









Quick preview of multi-agent embryogenesis

An *abstract* (computational), *top-down* approach



- development as a fundamentally *spatiotemporal* phenomenon
- JAG: "start generically, treat cells as black boxes, add details gradually"
- highlighting the broad principles necessary to integrate and make sense of the data; proposing a *computational* model of these principles

And the broad principles are...



- *biomechanics* → collective motion → "sculpture" of the embryo
- *gene regulation* \rightarrow gene expression patterns \rightarrow "painting" of the embryo
- coupling between shapes and colors

Multi-agent models



- best positioned to integrate both
- account for heterogeneity, modularity, hierarchy
- each agent carries a combined set of biomechanical and regulatory rules



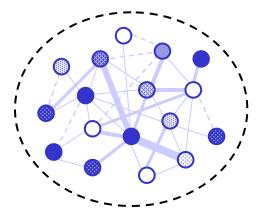
The self-made puzzle of embryogenesis

- 1. Self-organized <u>and</u> structured systems
- 2. A two-side challenge: heterogeneous motion / moving patterns
- 3. A multi-agent model of embryogenesis
- 4. Evolutionary development (evo-devo)



Self-organized <u>and</u> structured systems

Natural and human-made complex systems everywhere



- large number of elementary agents interacting locally
- simple individual behaviors creating a complex emergent collective behavior
- decentralized dynamics: no master blueprint or grand architect
- physical, biological, technical, social systems (natural or artificial)



pattern formation O = matter



biological development ○ = cell

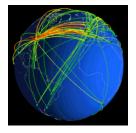


the brain & cognition \bigcirc = neuron





Internet & Web ○ = host/page



social networks
O = person





"Statistical" vs. "morphological" complex systems

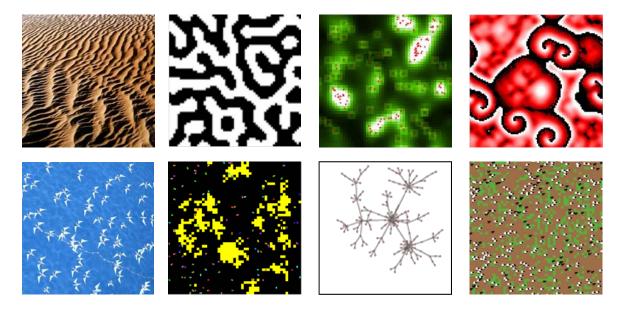
A brief taxonomy of systems

	Category	Agents / Parts	Local Rules	Emergent Behavior	A "Complex System"?
	2-body problem	few	simple	simple	NO
	3-body problem, low-D chaos	few	simple	complex	NO – too small
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	crystal, gas	many	simple	simple	NO – few params suffice to describe it
	patterns, swarms, complex networks	many	simple	"complex"	YES – but mostly random and uniform
	embryogenesis	many	sophisticated	complex	YES – reproducible and heterogeneous
	machines, crowds with leaders	many	sophisticated	"simple"	COMPLICATED - not self-organized



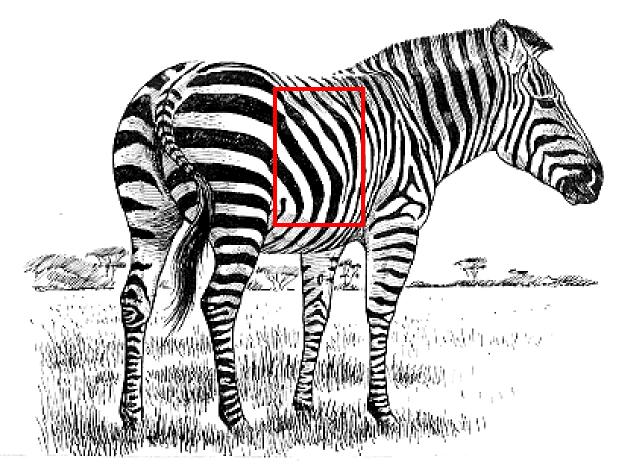
Statistical (self-similar) systems

- Many agents, simple rules, "complex" emergent behavior
 - → the "clichés" of complex systems: diversity of pattern formation (spots, stripes), swarms (clusters, flocks), complex networks, etc.



- ✓ yet, often like "textures": repetitive, statistically *uniform*, information-poor
- ✓ spontaneous order arising from amplification of *random* fluctuations
- ✓ unpredictable number and position of mesoscopic entities (spots, groups)

Morphological (self-dissimilar) systems compositional systems: pattern formation ≠ morphogenesis

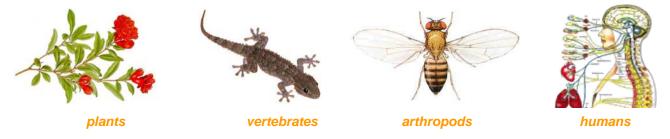


"I have the stripes, but where is the zebra?" or "The stripes are easy, it's the horse part that troubles me" —attributed to A. Turing, after his 1952 paper on morphogenesis



Morphological (self-dissimilar) systems

- Many agents, sophisticated rules, complex emergence
 - → natural ex: organisms (cells)



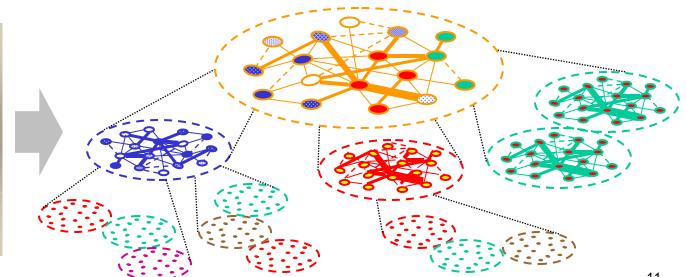
- ✓ mesoscopic organs and limbs have intricate, nonrandom morphologies.
- ✓ development is highly *reproducible* in number and position of body parts
- ✓ heterogeneous elements arise under information-rich genetic control
- Biological organisms are self-organized <u>and</u> structured
 - ✓ because the pieces of the puzzle (agent rules) are more "sophisticated" (than inert matter): depend on agent's type and/or position in the system
 - ✓ the outcome (development) is truly complex but, paradoxically, can also be more controllable and programmable



Beyond statistics: heterogeneity, modularity, reproducibility

- Complex systems can be much more than a "soup"
 - ✓ "complex" doesn't necessarily imply "homogeneous"...
 - → heterogeneous agents and diverse patterns, via positions
 - ✓ "complex" doesn't necessarily imply "flat" (or "scale-free")...
 - → modular, hierarchical, detailed architecture (at specific scales)
 - ✓ "complex" doesn't necessarily imply "random"...
 - → reproducible patterns relying on programmable agents







Statistical vs. morphological systems

Physical pattern formation is "free" – Biological (multicellular) pattern formation is "guided"

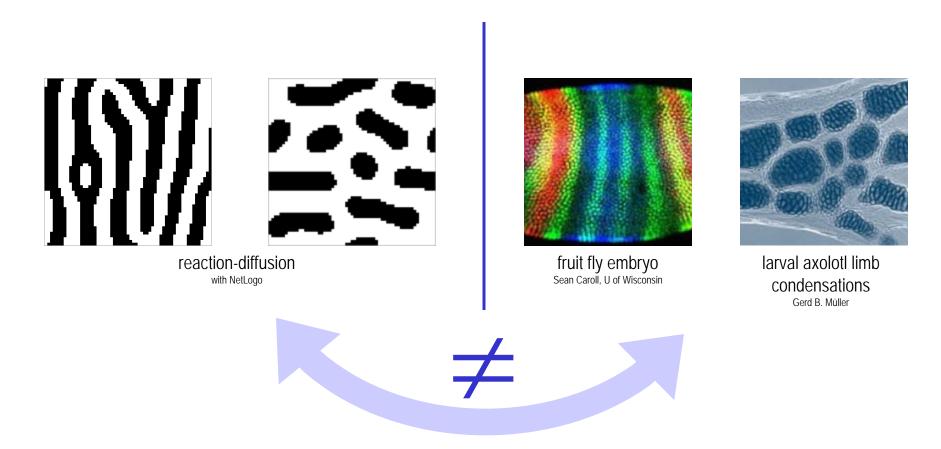


Fig. 8.2.



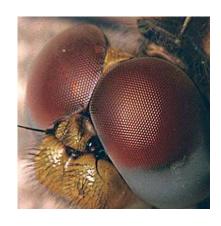
Statistical vs. morphological systems

- Multicellular forms = a bit of "free" + a lot of "guided"
 - ✓ domains of free patterning embedded in a guided morphology

unlike Drosophila's stripes, these pattern primitives are <u>not</u> regulated by different sets of genes depending on their position

spots, stripes in skin angelfish, www.sheddaquarium.org



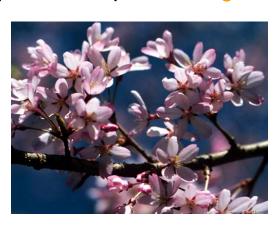


ommatidia in compound eye dragonfly, www.phy.duke.edu/-hsg/54

✓ repeated copies of a guided form, distributed in free patterns

entire structures (flowers, segments) can become modules showing up in random positions and/or numbers

flowers in tree cherry tree, www.phy.duke.edu/~fortney





segments in insect centipede, images.encarta.msn.com



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Embryogenesis couples assembly and patterning

➤ Sculpture → forms

Ádám Szabó, *The chicken or the egg* (2005) http://www.szaboadam.hu







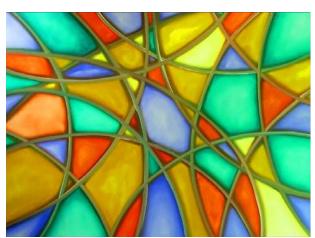


"shape from patterning"

✓ the forms are

"sculpted" by the selfassembly of the
elements, whose
behavior is triggered
by the colors

➤ Painting → colors



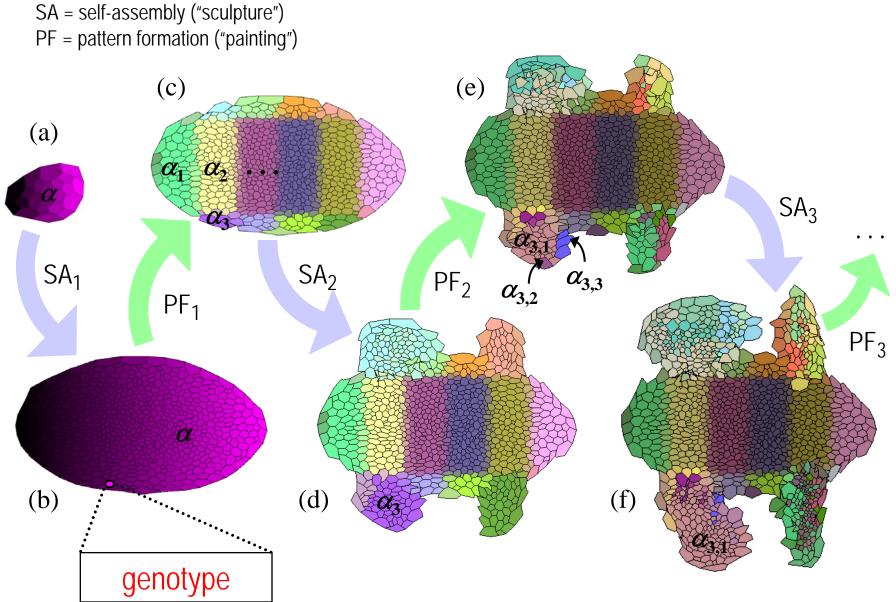
"patterns from shaping

✓ new color regions

 appear (domains of genetic expression)
 triggered by deformations



Embryogenesis couples assembly and patterning



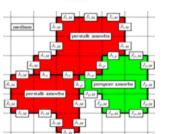


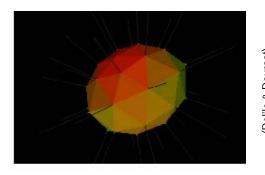
Embryogenesis couples mechanics and regulation

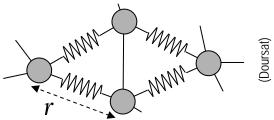
Cellular mechanics

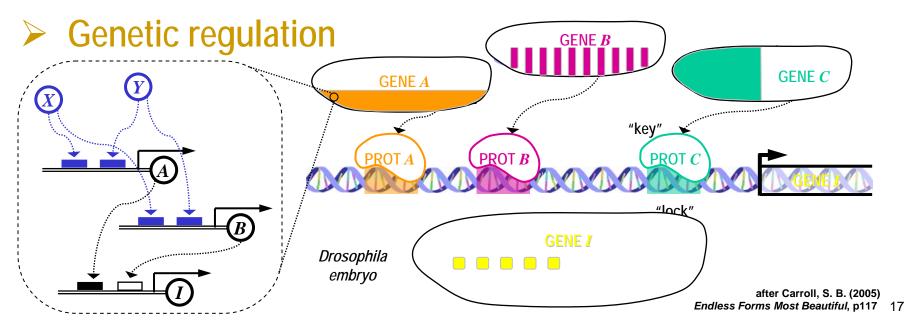
- adhesion
- deformation / reformation
- migration (motility)
- division / death

(Graner, Glazier, Hogeweg)









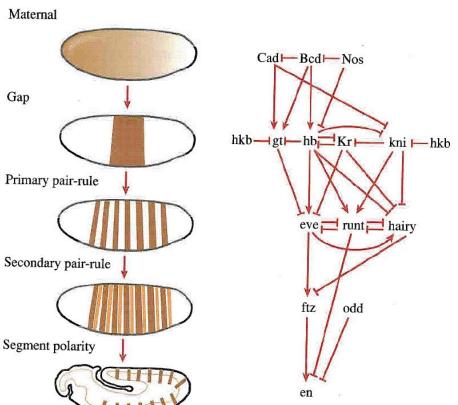
cellular Potts model



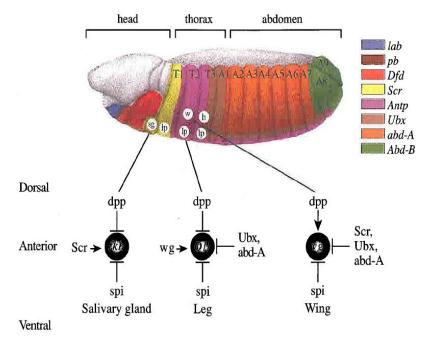
Gene regulatory pattern formation

Segmentation & identity domains in Drosophila

 periodic A/P band patterns are controlled by a 5-tier gene regulatory hierarchy



 ✓ intersection with other axes creates organ primordia and imaginal discs (identity domains of future legs, wings, antennae, etc.)



from Carroll, S. B., et al. (2001) From DNA to Diversity, p63



Embryogenesis couples mechanics and regulation

Cellular mechanics mechanical stress, modification of cell mechano-sensitivity size and shape growth, division, apoptosis differential adhesion Genetic regulation change of cell-to-cell contacts gene regulation change of signals, chemical messengers diffusion gradients ("morphogens")

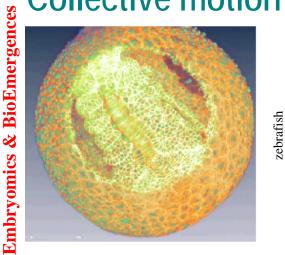


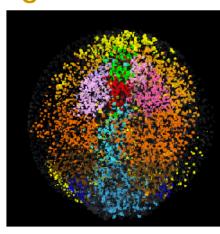
Nadine Peyriéras, Paul Bourgine, Thierry Savy,

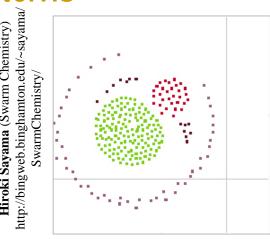
Benoît Lombardot, Emmanuel Faure et al.

Embryogenesis couples motion and patterns

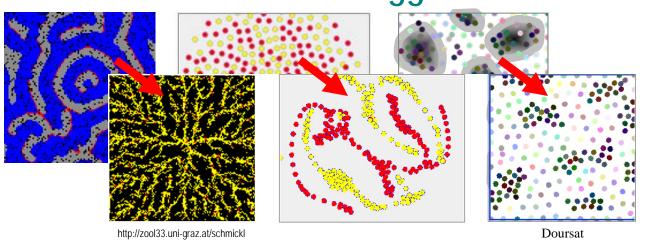
Collective motion regionalized into patterns

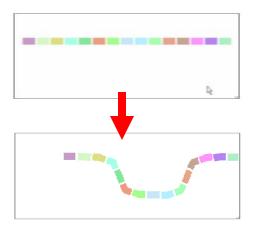






Pattern formation that triggers motion







The self-made puzzle of embryogenesis

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Why multi-agent modeling?

Equations and laws can be hard or impossible to find...

✓ "The study of non-linear physics is like the study of nonelephant biology." —Stanislaw Ulam

 the physical world is a fundamentally nonlinear and out-of-equilibrium process

 focusing on linear approximations and stable points is missing the big picture in most cases

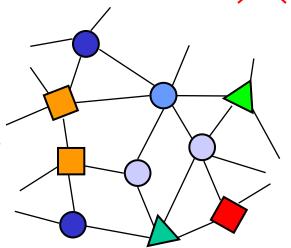
- ✓ let's push this quip: "The study of nonanalytical complex systems is like the study of non-elephant biology." —??
 - complex systems have their own "elephant" species, too: dynamical systems that can be described by diff. eqs or statistical laws
 - many real-world complex systems do not obey neat macroscopic laws





Why multi-agent modeling?

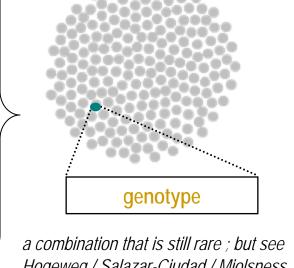
- Equations and laws can be hard or impossible to find in...
 - ✓ systems that *no macroscopic quantity* suffices to explain (DEE)
 - no law of "concentration", "pressure", or "gross domestic product"
 - even if global metrics can be designed to give an indication about the system's dynamical regimes, they rarely obey a given equation or law
 - ✓ systems that require a non-Cartesian decomposition of space (Pxx)
 - network of irregularly placed or mobile agents
 - ✓ systems that contain heterogeneity
 - segmentation into different types of agents
 - at a fine grain, this would require a "patchwork" of regional equations
 - ✓ systems that are dynamically adaptive
 - the topology and strength of the interactions depend on the short-term activity of the agents and long-term "fitness" of the system in its environment





Different approaches and families of models

- Biological, bio-inspired or artificial models
 - focused on spatial differentiation patterns (little or no motion)
 - reaction-diffusion (PDEs, cellular automata)
 - gene networks (Boolean or concentrations) on a fixed lattice
 - "amorphous computing"
 - focused on motion (little or no patterning)
 - Cellular Potts Model (on predefined cell types)
 - aggregation, self-assembly
 - collective motion, flocking, cellular sorting

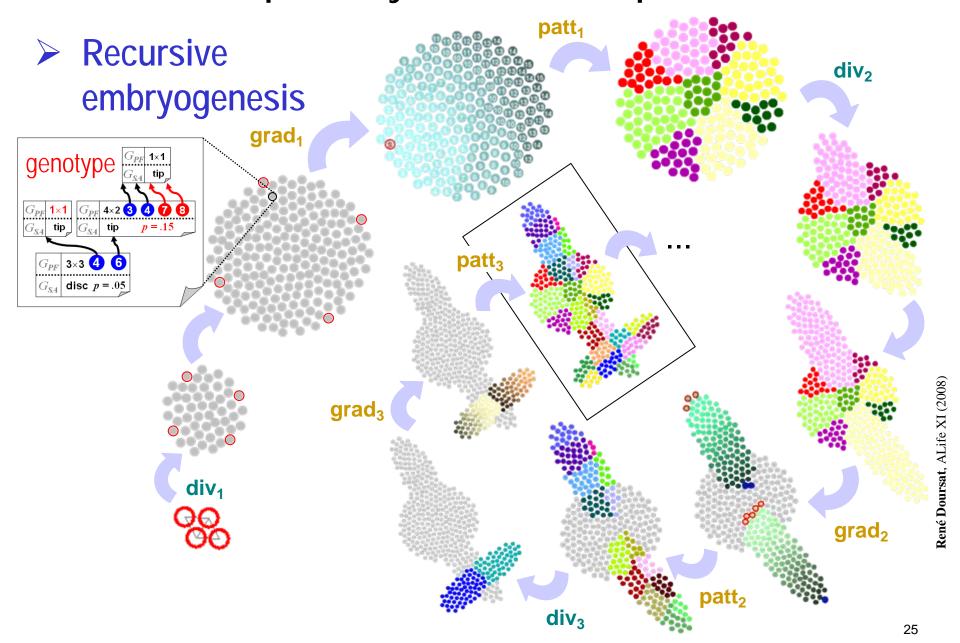


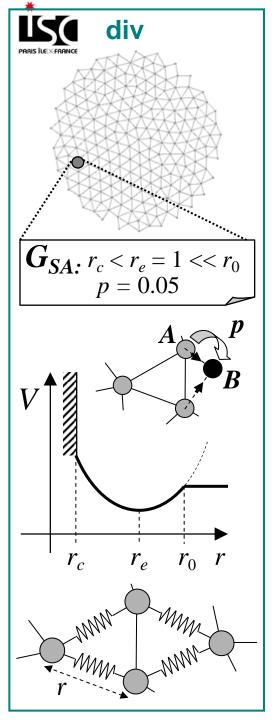
Hogeweg / Salazar-Ciudad / Miolsness..

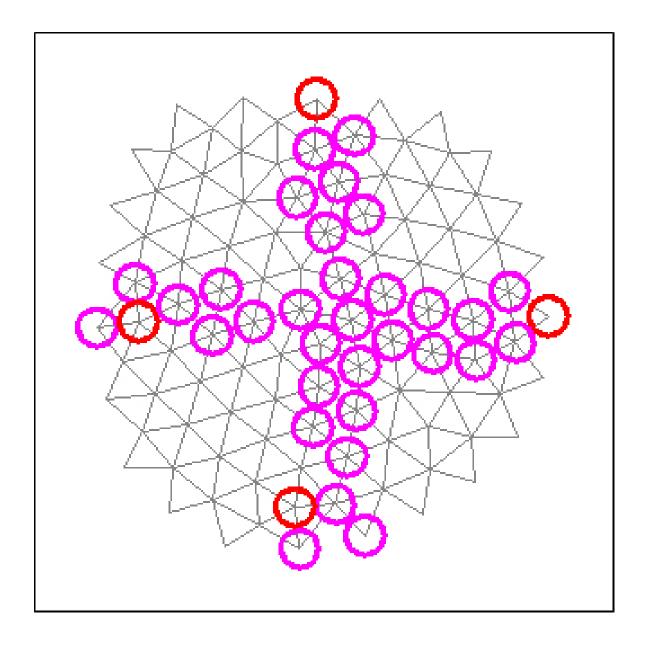
- at different scales
 - macroscopic models (densities, differential geometry) \rightarrow no individual information
 - mesoscopic models (cellular centers, Potts) \rightarrow no membrane geometry or nuclei
 - microscopic models (elastic polyedra, drop models) \rightarrow cellular deformations



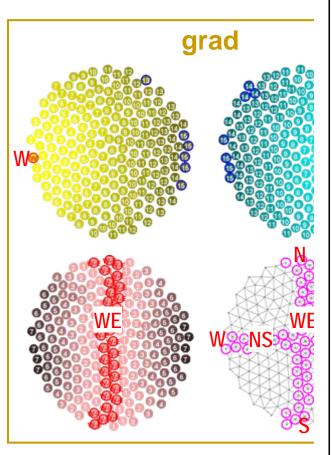
Example of hybrid mesoscopic model





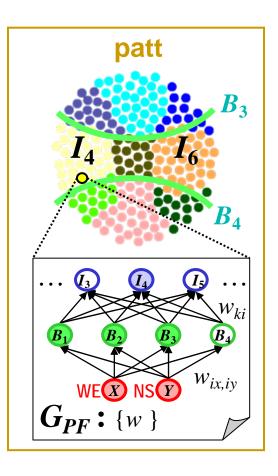


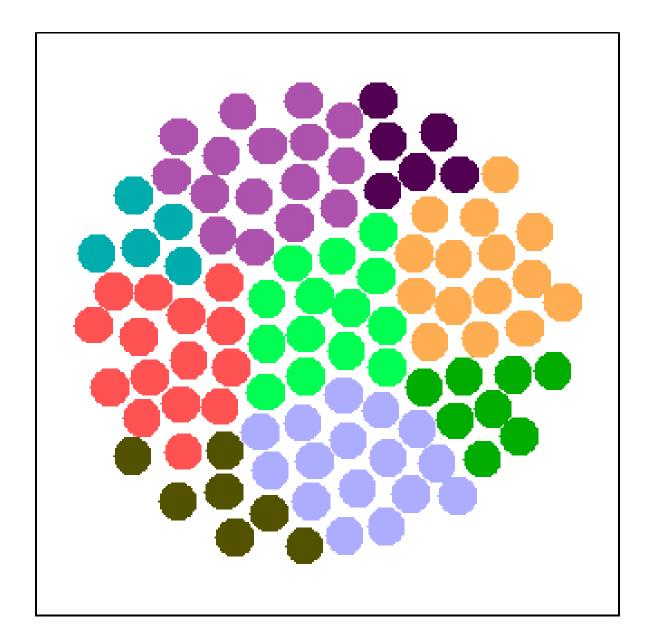


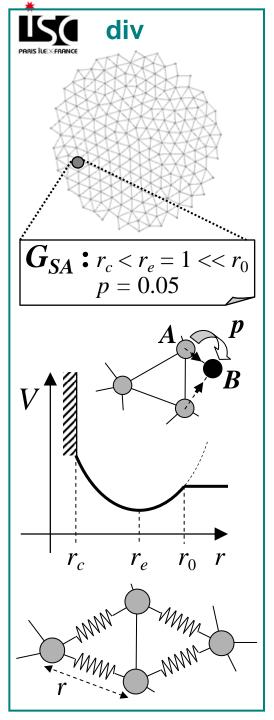


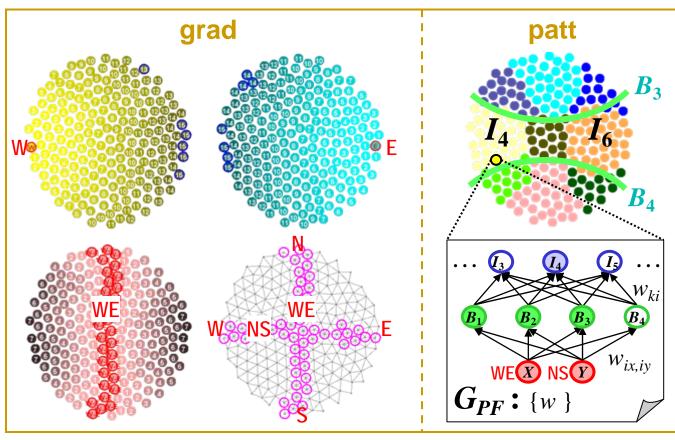


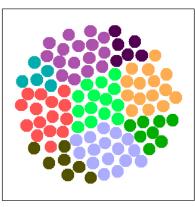


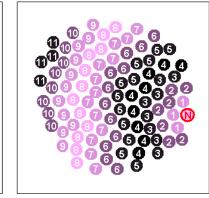


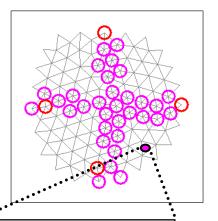










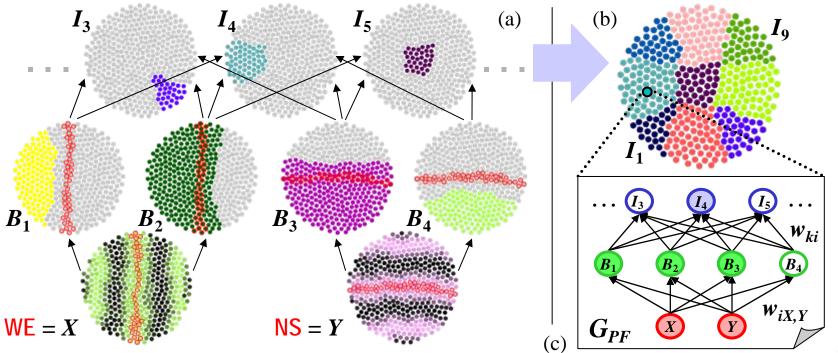




Virtual gene atlas

Programmed patterning (patt): the hidden embryo map

- a) same swarm in different colormaps to visualize the agents' internal patterning variables X, Y, B_i and I_k (virtual *in situ hybridization*)
- b) consolidated view of all identity regions I_k for k = 1...9
- c) gene regulatory network used by each agent to calculate its expression levels, here: $B_1 = \sigma(1/3 X)$, $B_3 = \sigma(2/3 Y)$, $I_4 = B_1B_3(1 B_4)$, etc.

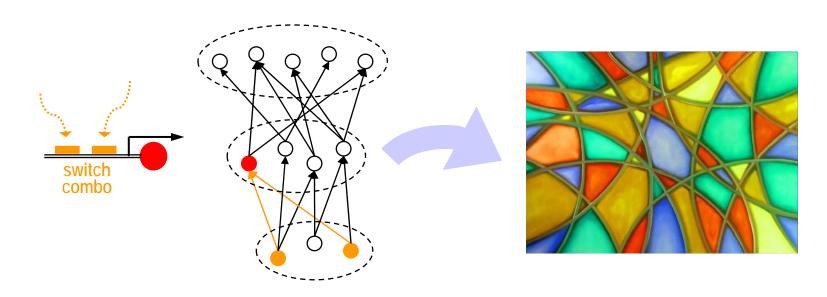




From feedforward to recurrent gene regulation

Summary: simple feedforward hypothesis

- ✓ developmental genes are broadly organized in tiers, or "generations": earlier genes map the way for later genes
- ✓ gene expression propagates in a directed fashion: first, positional morphogens create domains, then domains intersect

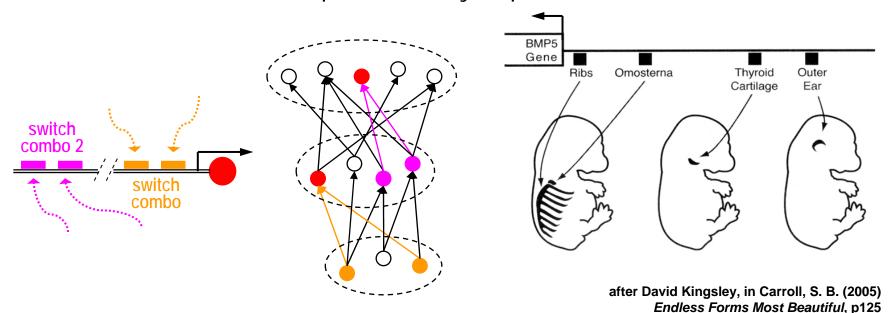




From feedforward to recurrent gene regulation

> Toolkit genes are often multivalent

- exception to the feedforward paradigm: "toolkit" genes that are reused at different stages and different places in the organism
- ✓ however, a toolkit gene is triggered by different switch combos, which can be represented by duplicate nodes in different tiers

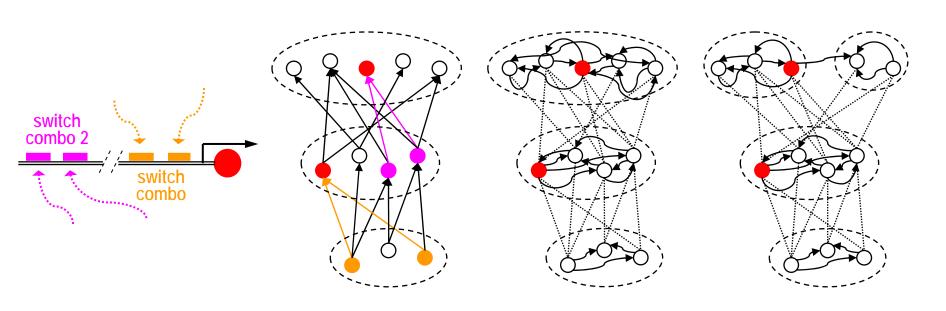




From feedforward to recurrent gene regulation

More realistic variants of GRNs

- ✓ add recurrent links within tiers → domains are not established independently but influence and sharpen each other
- ✓ subdivide tiers into subnetworks → this creates modules that
 can be reused and starts a hierarchical architecture





Multiscale embryogenesis

- Morphological refinement by iterative growth
 - ✓ details are not created in one shot, but gradually added. . .



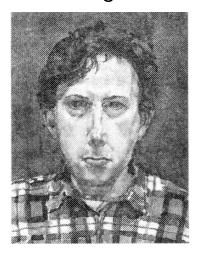




✓ ... while, at the same time, the canvas grows



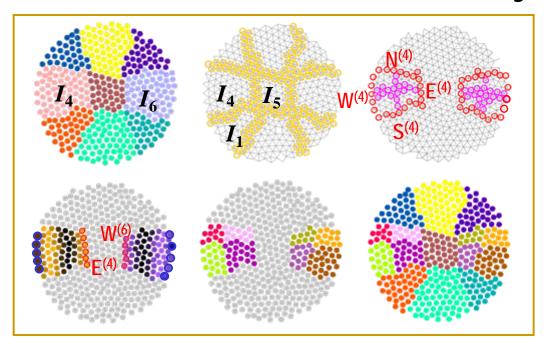




from Coen, E. (2000) The Art of Genes, pp131-135

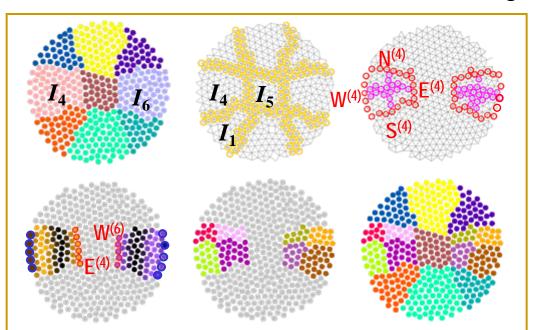


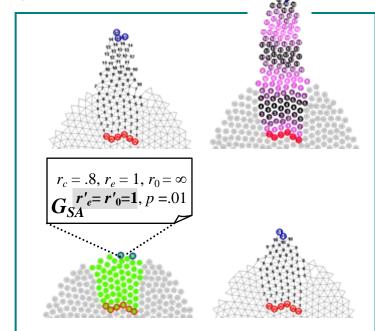
Multiscale embryogenesis





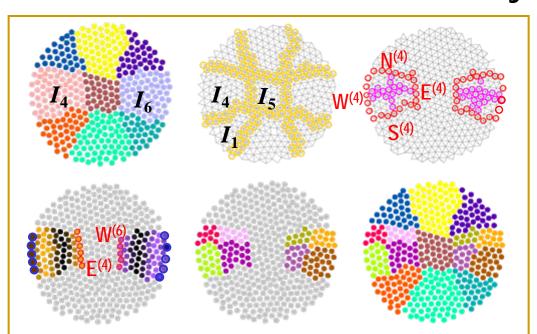
Multiscale embryogenesis

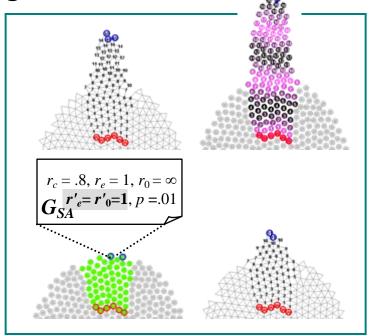


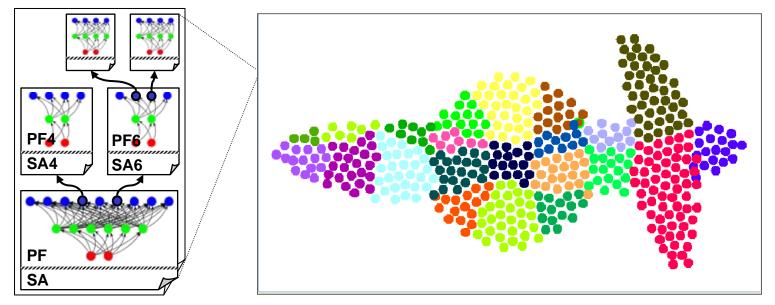




Multiscale embryogenesis

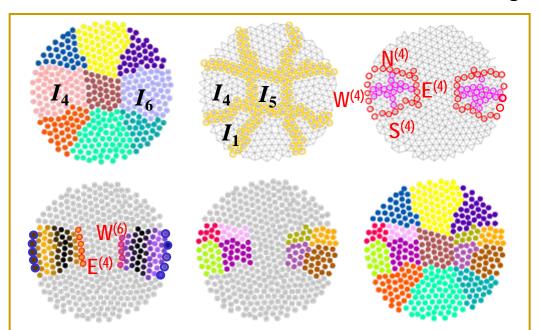


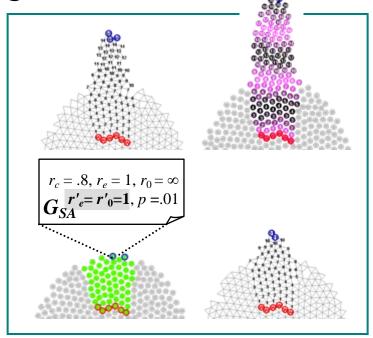


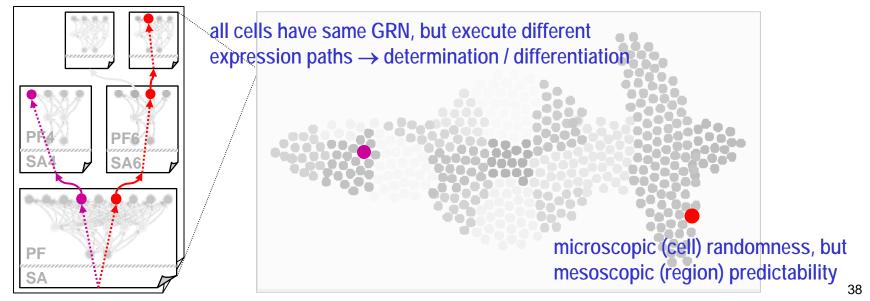




Multiscale embryogenesis









The self-made puzzle of embryogenesis

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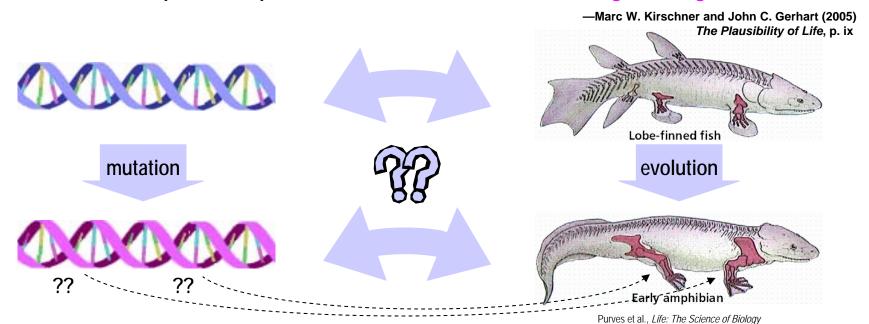


Evolutionary innovation by development

Development: the missing link of the Modern Synthesis

"When Charles Darwin proposed his theory of evolution by variation and selection, explaining selection was his great achievement. He could not explain <u>variation</u>. That was Darwin's dilemma."

"To understand novelty in evolution, we need to understand organisms down to their individual building blocks, down to their deepest components, for these are what undergo change."





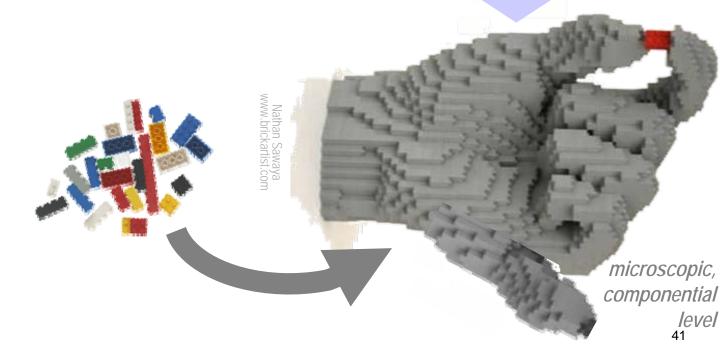
The self-made puzzle of "evo-devo" engineering

Development: the missing link of the Modern Synthesis



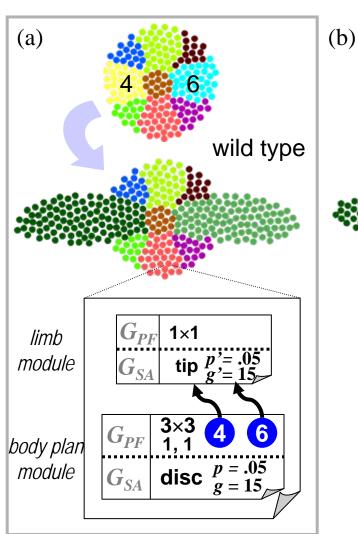
macroscopic, emergent level

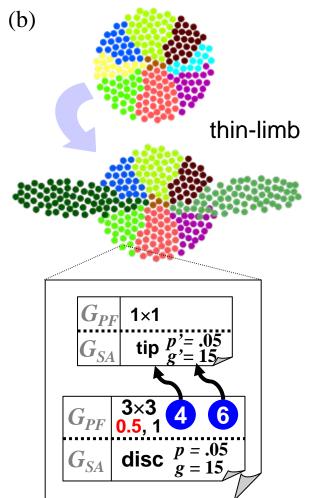
"To understand novelty in evolution, we need to understand organisms down to their individual building blocks, down to their deepest components, for these are what undergo change."

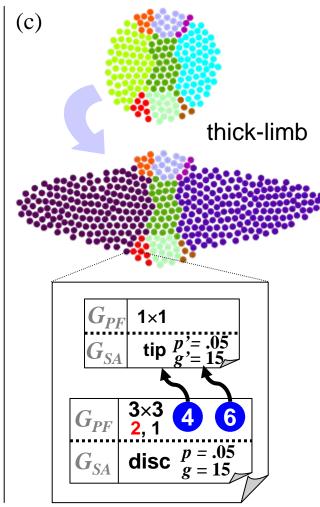




Quantitative mutations: limb thickness

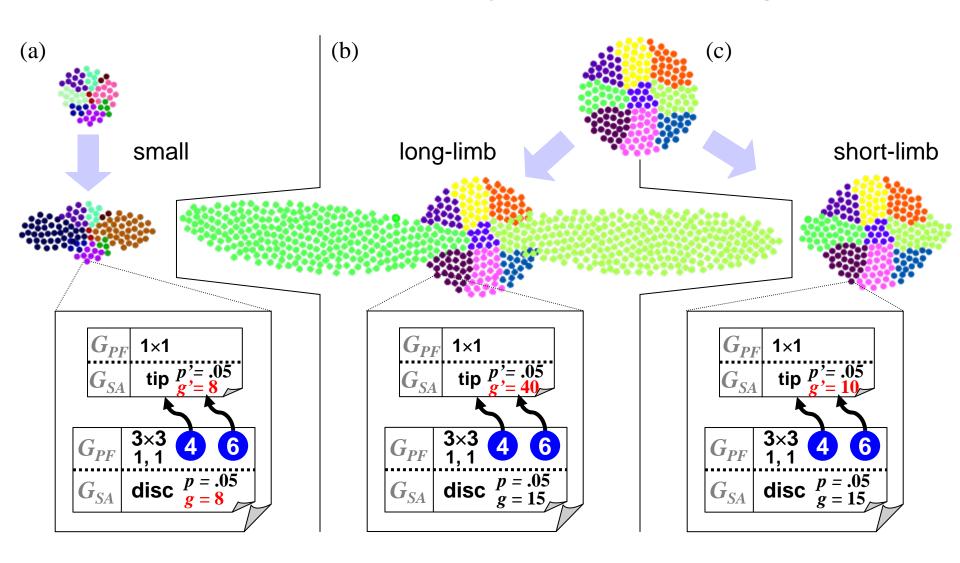






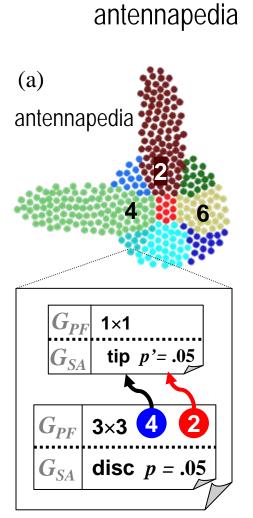


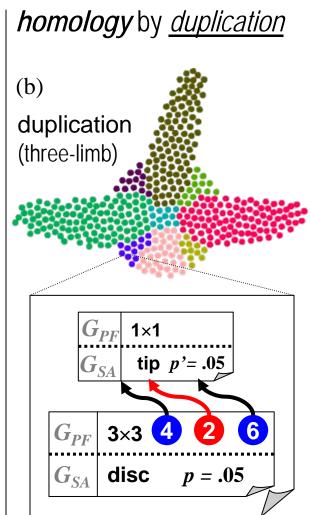
Quantitative mutations: body size and limb length

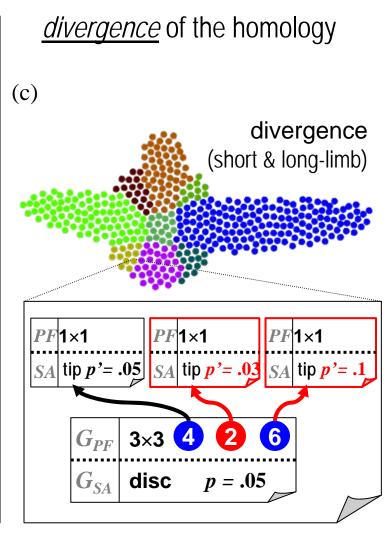




Qualitative mutations: limb position and differentiation

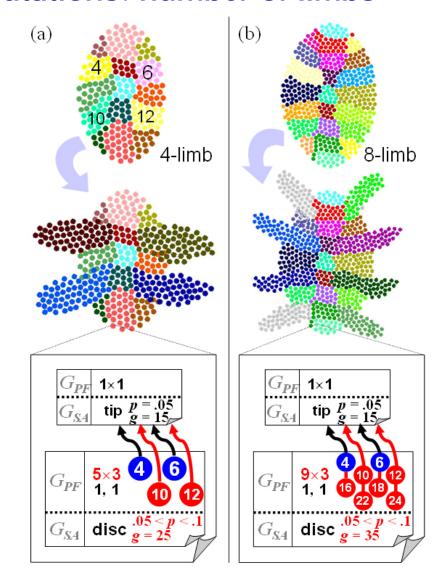




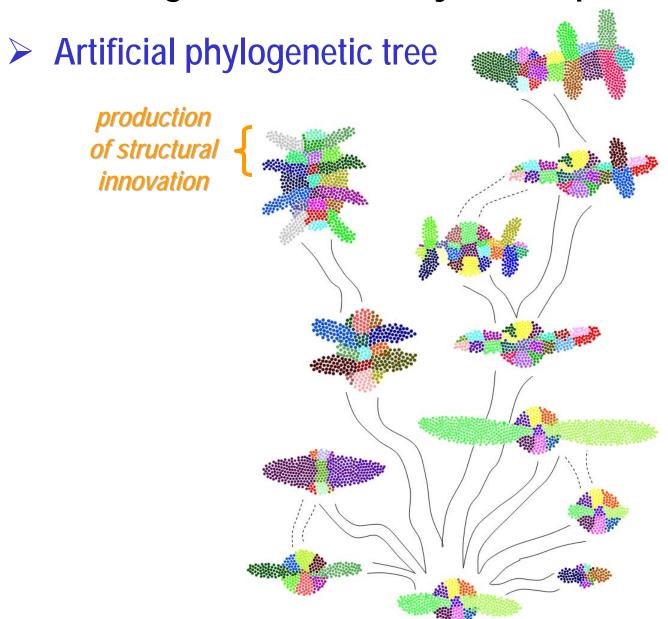




Qualitative mutations: number of limbs





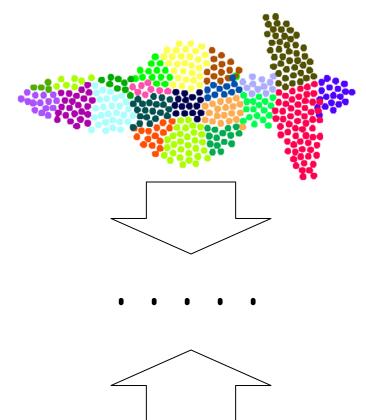


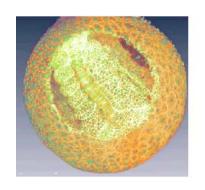


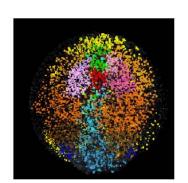
Work toward more accurate biological modeling

top-down, abstract, heuristic approach







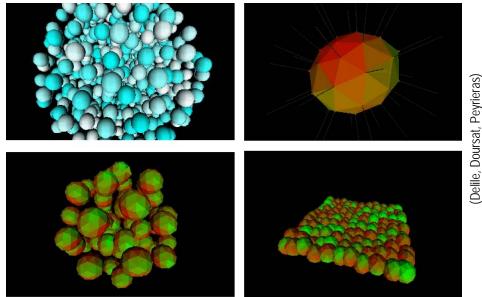




Work toward more accurate biological modeling

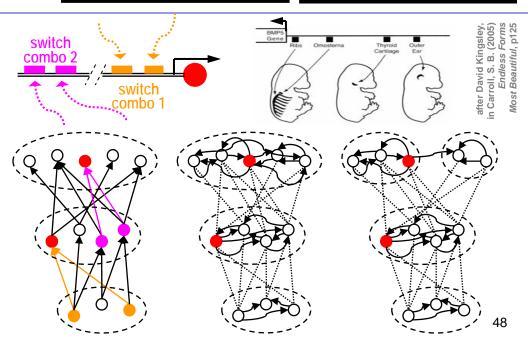
More accurate mechanics

- **√** 3-D
- ✓ individual cell shapes
- ✓ collective motion, migration
- ✓ adhesion



Better gene regulation

- ✓ recurrent links
- ✓ gene reuse
- ✓ kinetic reaction ODEs
- ✓ attractor dynamics





The self-made puzzle of embryogenesis

- 1. Self-organized and structured systems
- A two-side challenge: heterogeneous motion / moving patterns
- 3. A multi-agent model of embryogenesis
- 4. Evolutionary development (evo-devo)