Local-to-Global Algorithms in Biology: Drosophila and Beyond (maybe ...)

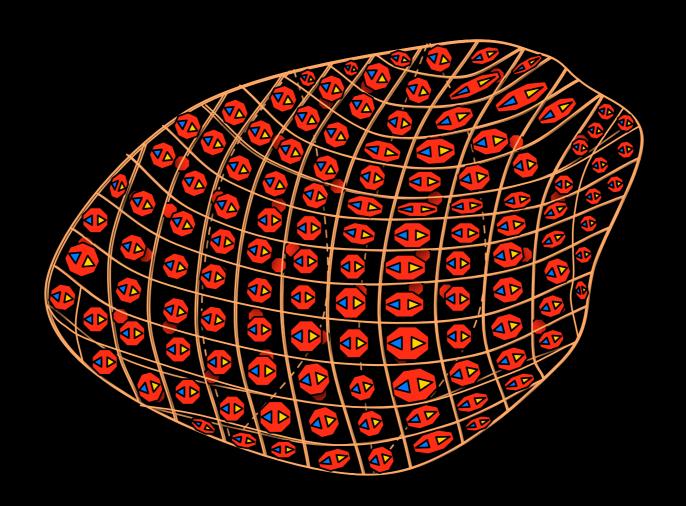
MIT Synthetic Biology Working Group

Dan Yamins

2007.12.05

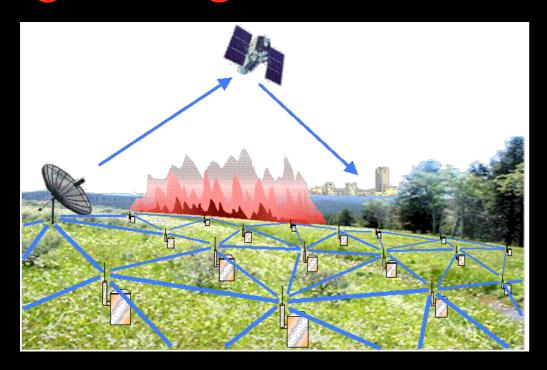
spatial multi-agent systems

a space with agents embedded in the space



local information and processing globally defined tasks

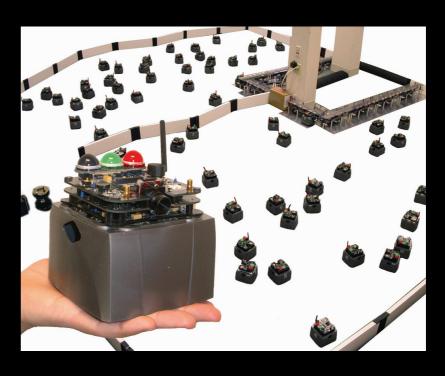
engineering:



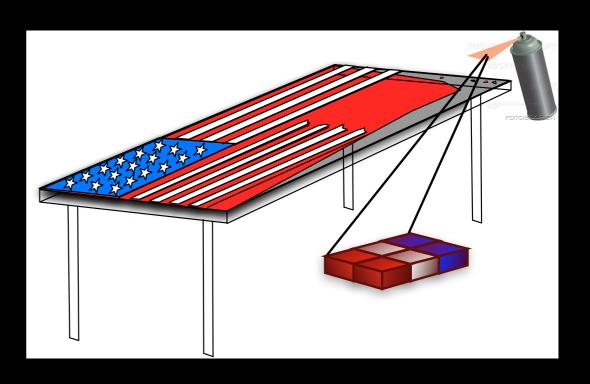
Sensor Networks



Saul Griffith's selffolding structures



McLurkin iRobot Swarm



Butera's "paintable computer" concept

biology:



drosophila embryo

- What does thinking of natural spatial multi-agent systems qua computers tell us scientifically?
- What agent resource capacities are required to solve a given task?
- Can we then turn around this knowledge to build a better bug?

outline

- I. Drosophila background
- II. Multi-agent systems Model of Drosophila
- III. An information bound
- IV. The Radius/State Tradeoff

The first four sections study a known system, showing how multi-agent systems reasoning can help us understand features of local-to-global systems design.

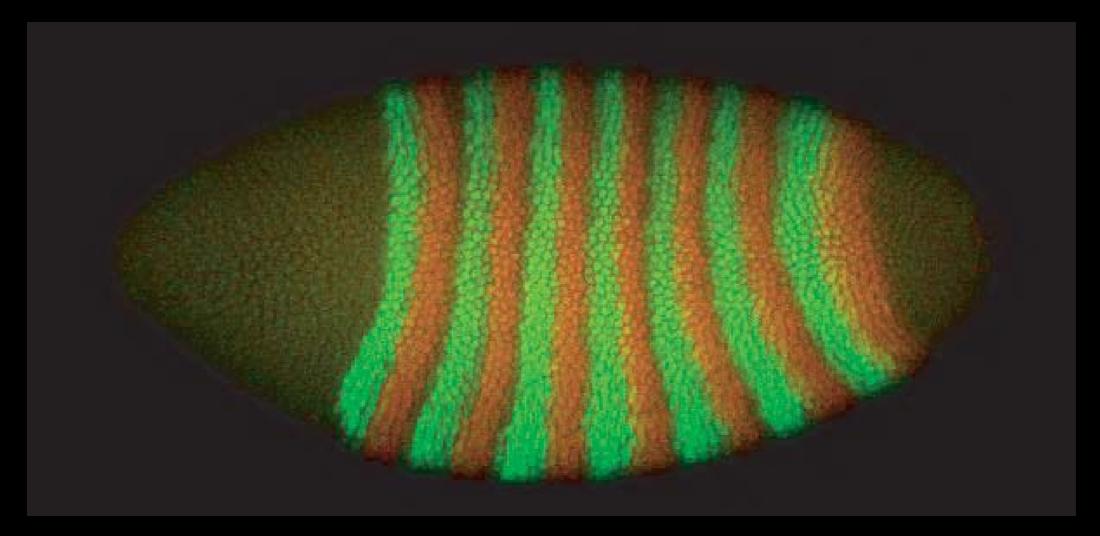
V. Designing Local Rules

The fifth section takes this knowledge, and applies it to develop the beginnings of a protocol for designing novel systems.

Drosophila Background

(Actually ... one stage in early Drosophila, utterly simplified)

Drosophila



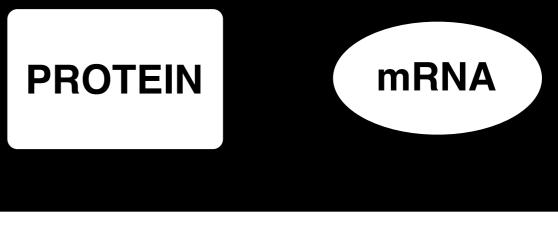
Drosophila melanogaster embryo about 100 minutes post-fertilization

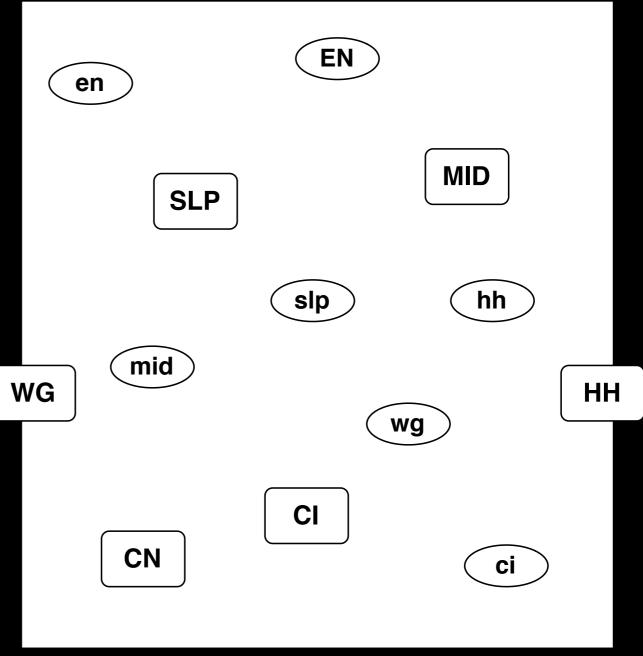
Drosophila

Let's think of the cell as a

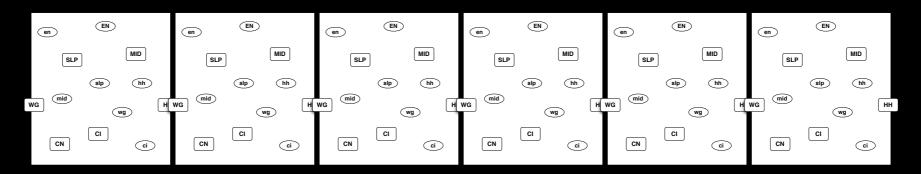
.... bag of biomolecules

Highlighted here are 13 substances that play are involved in the stripes in the picture on the previous slide

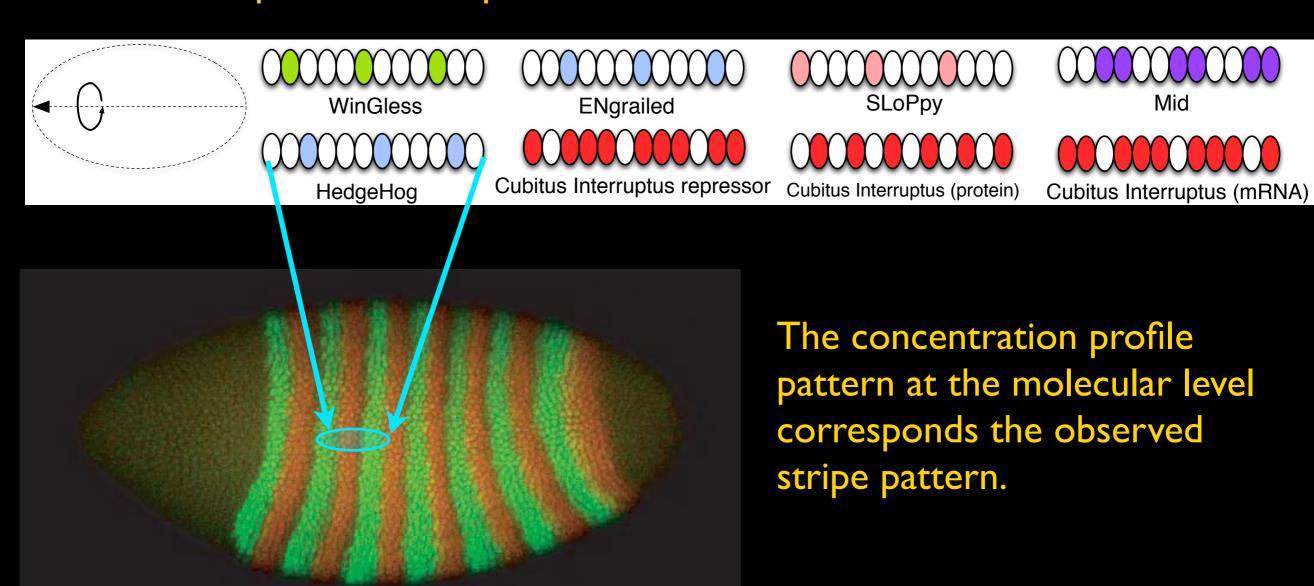




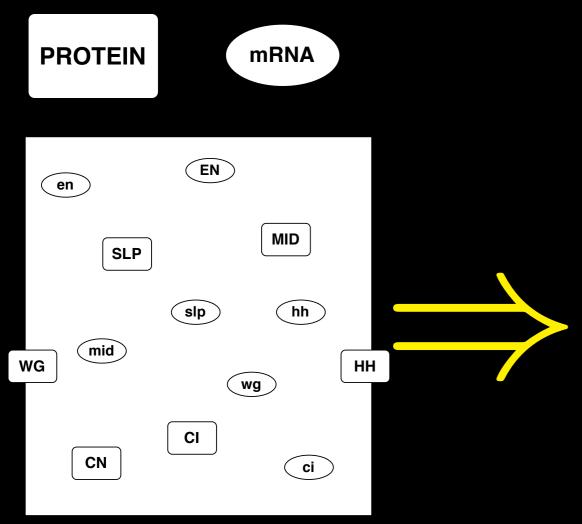
Considering cells along the A-P axis:



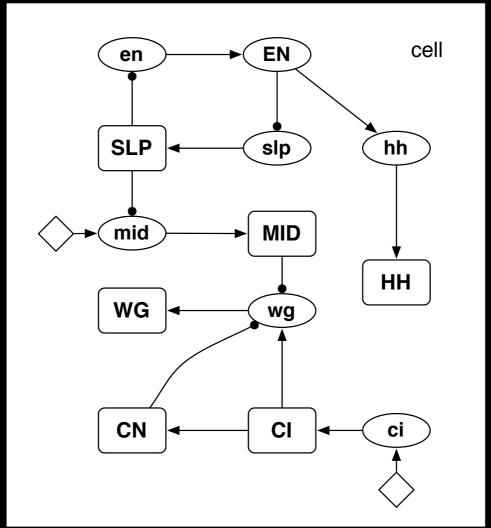
... there is a periodic pattern in the concentration of each of various substances depicted on the previous slide:



Drosophila



The bag of biomolecules ...

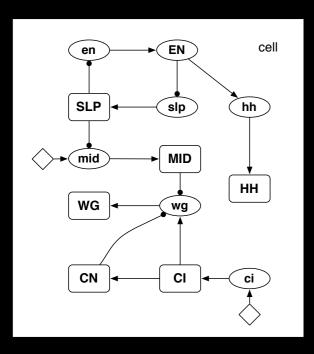


Positive Induction

Negative Inhibition

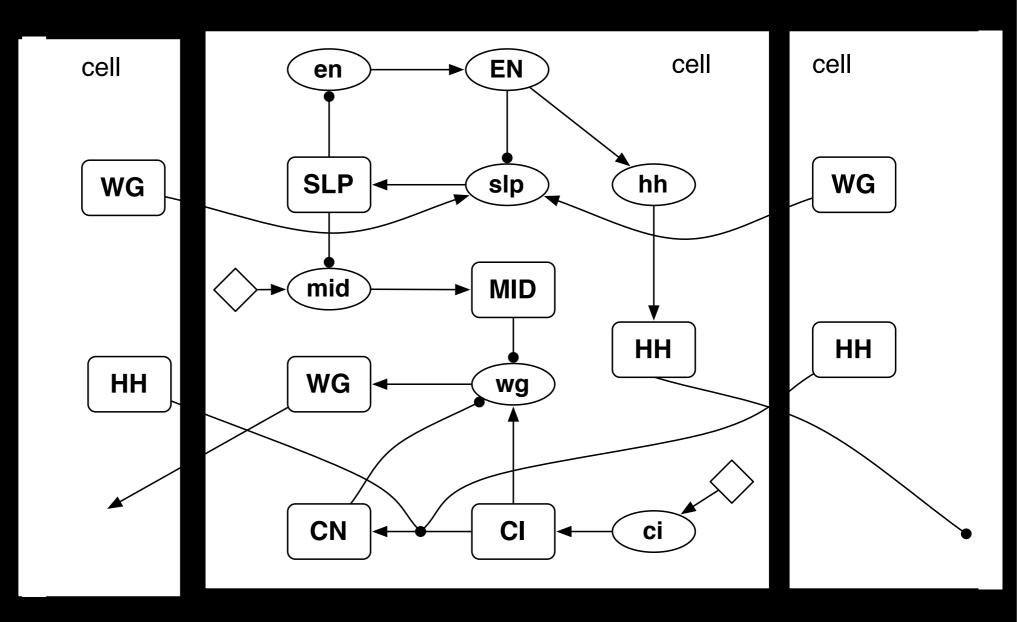


... is actually a Gene Regulatory Network

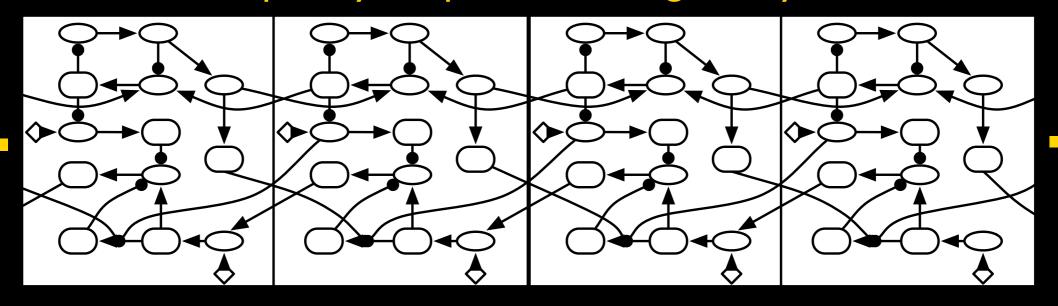


Some components of the Gene Regulatory Network in one cell ...

... influence those in neighboring cells.



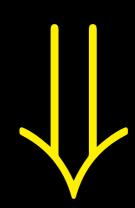
It's a spatially coupled Gene Regulatory Network.

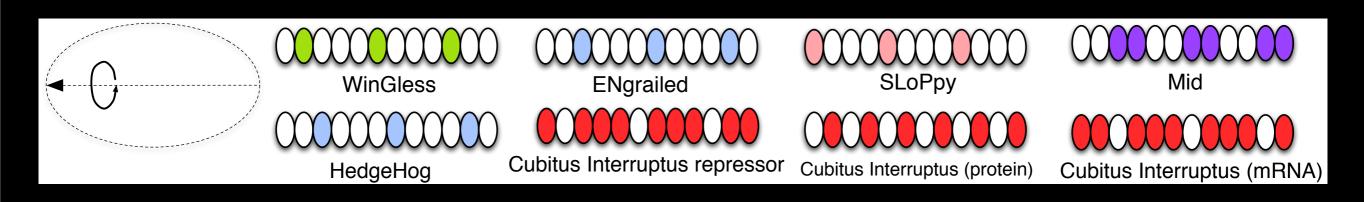


Such a network can be modeled as a coupled ODE:

$$\frac{dS_j^i}{dt} = f_j\left(S_1^i, S_2^i, \dots, S_N^i, S_1^{i-1}, S_2^{i-1}, \dots, S_N^{i-1}, S_1^{i+1}, S_2^{i+1}, \dots, S_N^{i+1}\right)$$

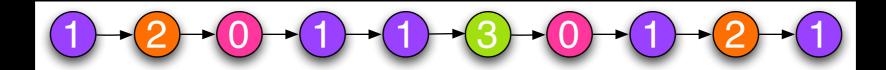
And the ODE will (a la Garrett Odell) have the pattern as a stable nondegenerate steady state:



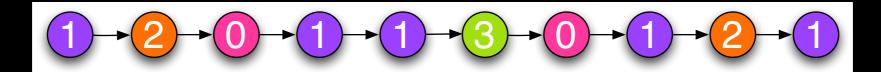


a multi-agent model of drosophila

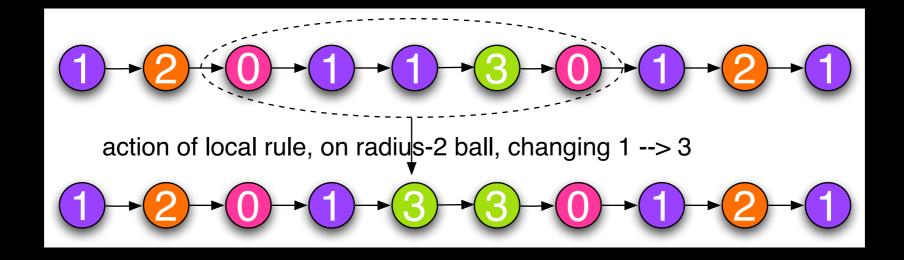
(a sketchy multi-agent model of drosophila)



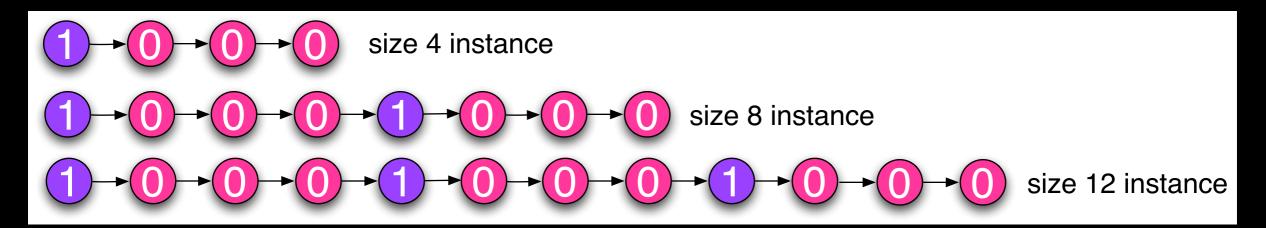
... a simple "I-D" organism, each cell of which has some internal state (0,1,2,3, etc...)

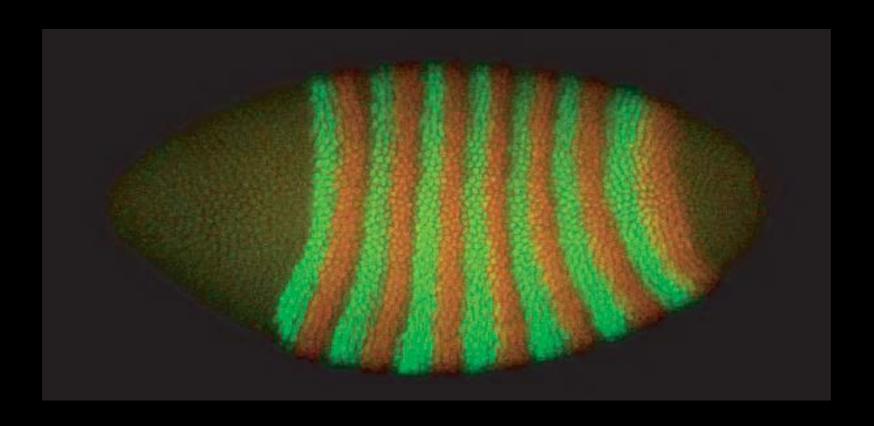


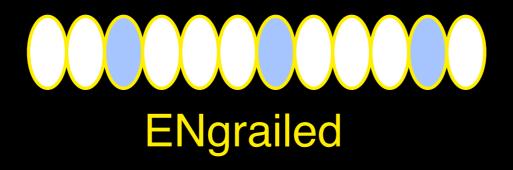
local rules by which the agent integrates information about states of nearby agents and takes a differentiation action



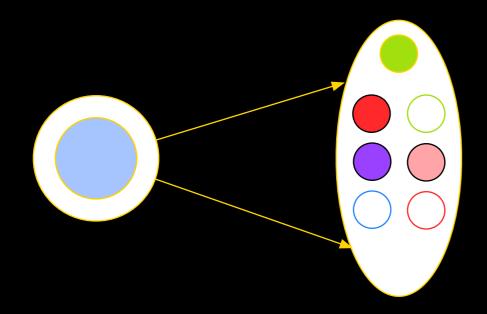
any radius-I (nearest neighbor) local rule is allowed.

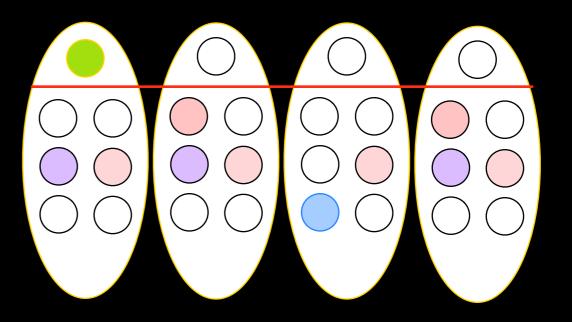






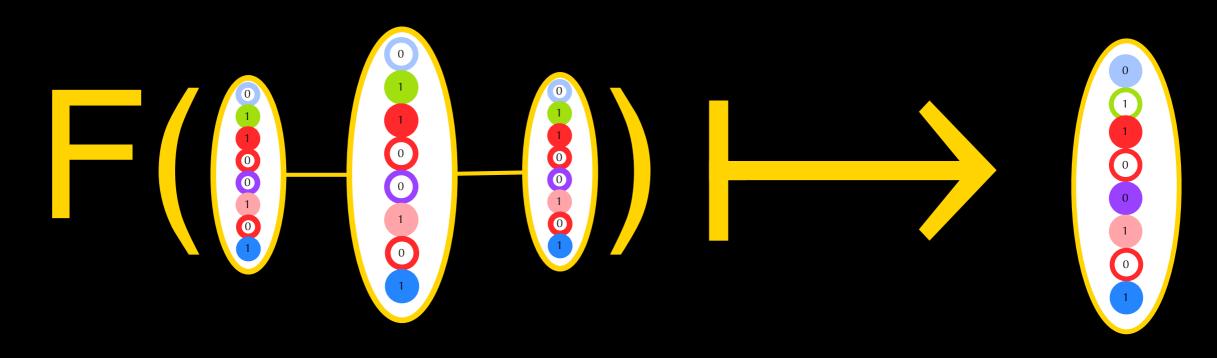
Add "slots."



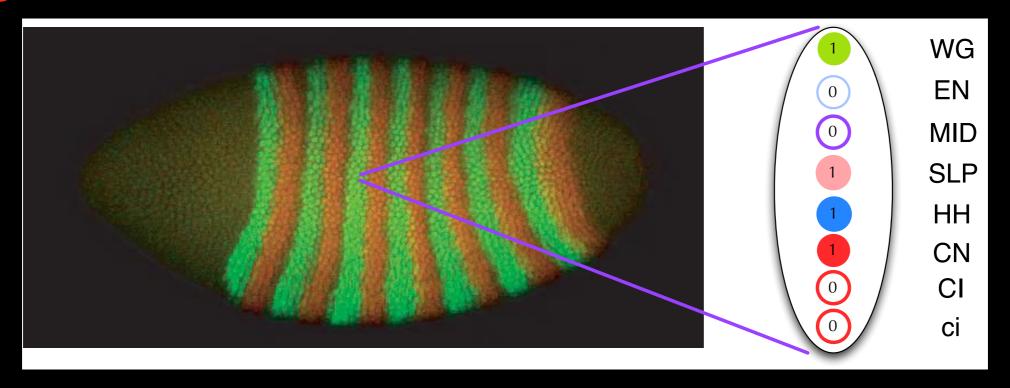


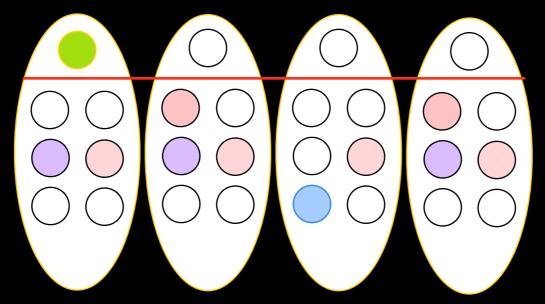
Agent with unstructured internal state.

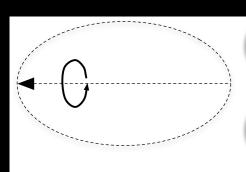
Structured "multi-register" internal states.

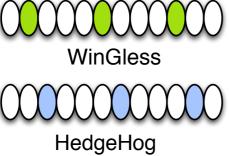


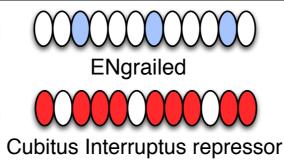
Multislot dynamics

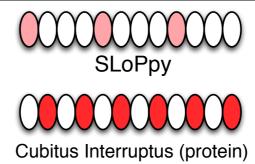


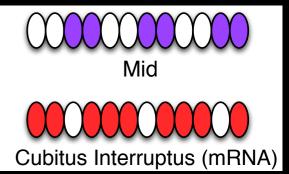


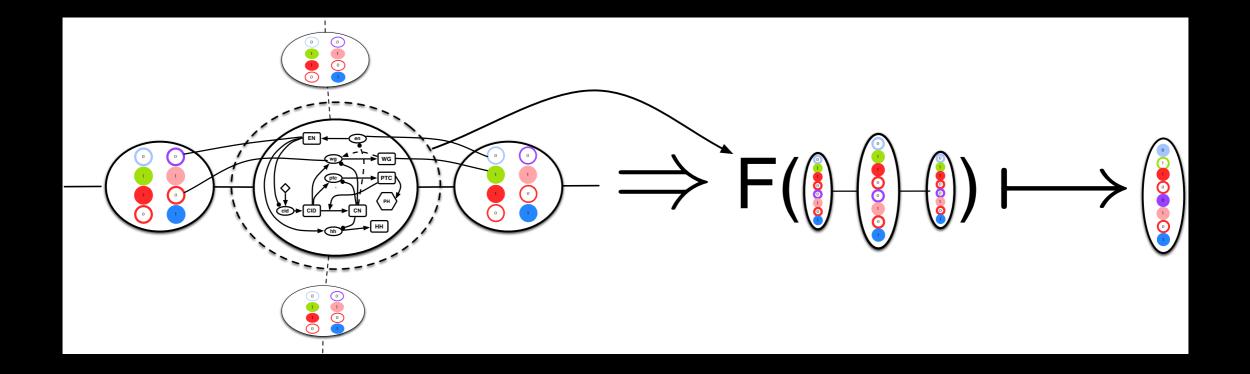












the gene regulatory network as a nearest-neighbor local rule

the "brains" of the cellular-agent

An informational bound

Local Checkability

Consider the repeat pattern:

Can this pattern be solved robustly with a nearest-neighbor rule?

Answer: No. Because the with a radius I rule, 000 would have to be a fixed state.

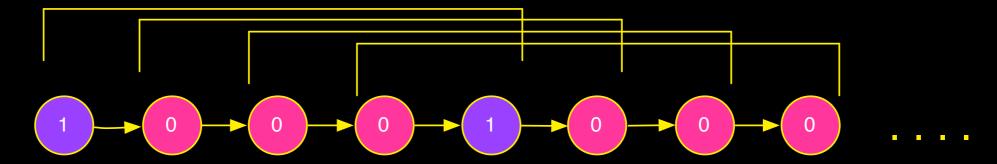
$$\begin{array}{c} 1 \\ \hline \\ 0 \\ \hline \\ \end{array}$$

local checkability; multi-agent reasoning about local equilibria

A function $\Theta: B_r \to \{0,1\}$ is a local check scheme for a pattern T if

$$\bigwedge_{i \le |X|} (\Theta(B_r(i, X)) = 1) \quad \Rightarrow \quad X \in T$$

This means: if all cells think "criterion Theta" is true ... then actually the organismal pattern is correct



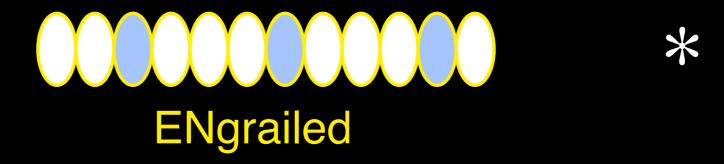
1000 has a radius-2 check scheme.

All repeated patterns are locally checkable.

Any pattern thus has a *minimal* check radius, a lower bound on how far local rules have to see to form the pattern robustly.

local checkability; multi-agent reasoning about local equilibria

Hey. Wait.



Local check radius of this pattern is = 2, and in fact to actually make a rule radius 5/2 is required.

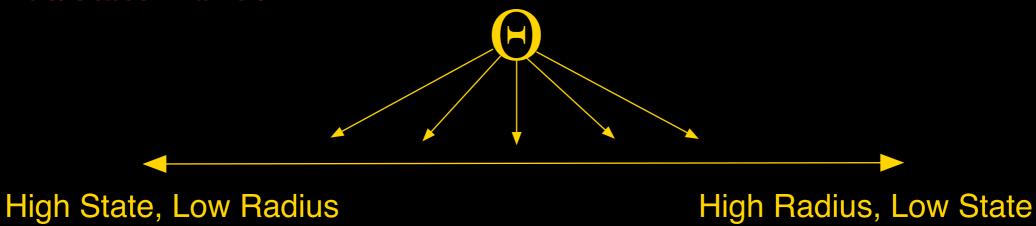
But the coupled gene network rules are nearest neighbor (i.e. r = 1).

Put another way, no equation of the form:

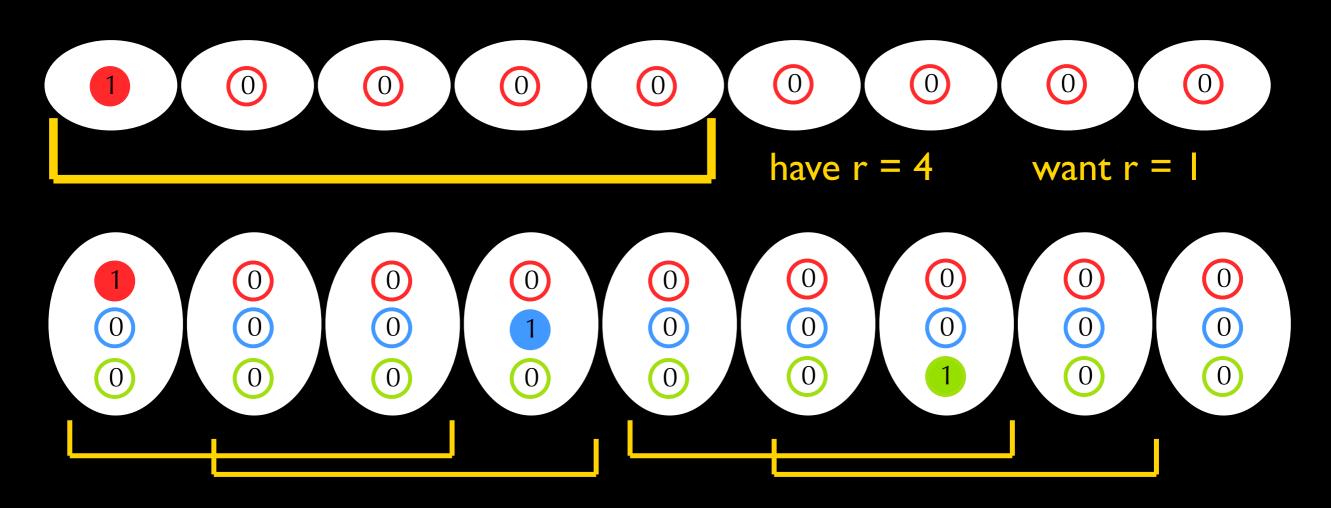
$$\frac{dS_j^i}{dt} = f_j\left(S_1^i, S_2^i, \dots, S_N^i, S_1^{i-1}, S_2^{i-1}, \dots, S_N^{i-1}, S_1^{i+1}, S_2^{i+1}, \dots, S_N^{i+1}\right)$$

can have a nondegenerate stable steady state of the form * if engrailed is decoupled.

The system is "trying" to make a 4-coordinate....

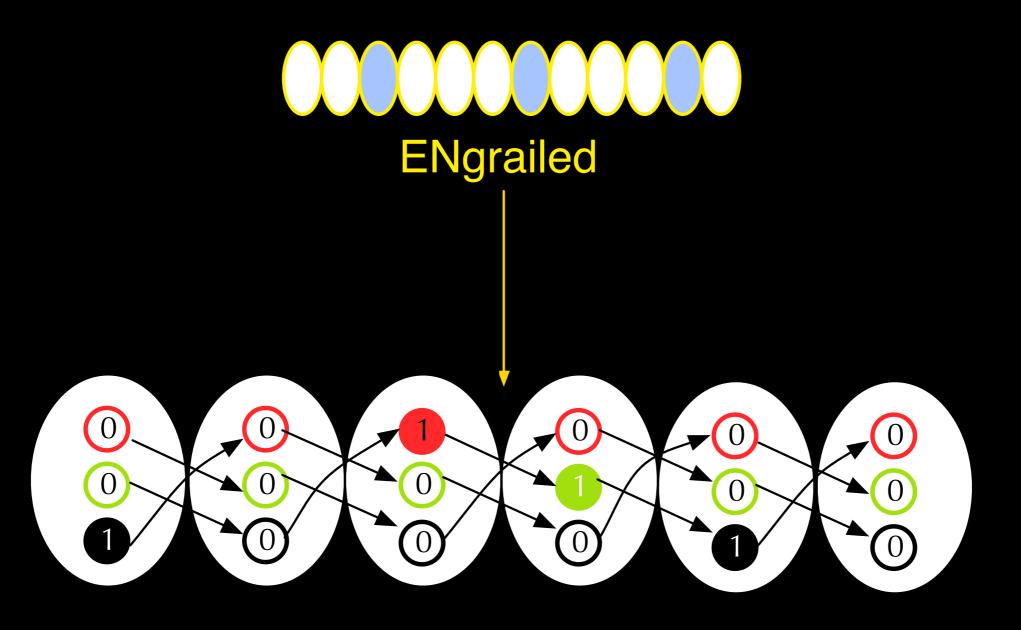


A generic "cut-and-shift" procedure for implementing the Radius -> tradeoff:

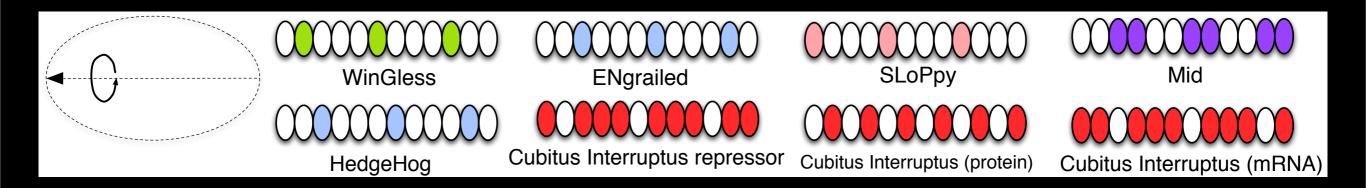


#slots = $\lceil (2r+1)/(2k+1) \rceil$ to go from radius r to radius k.

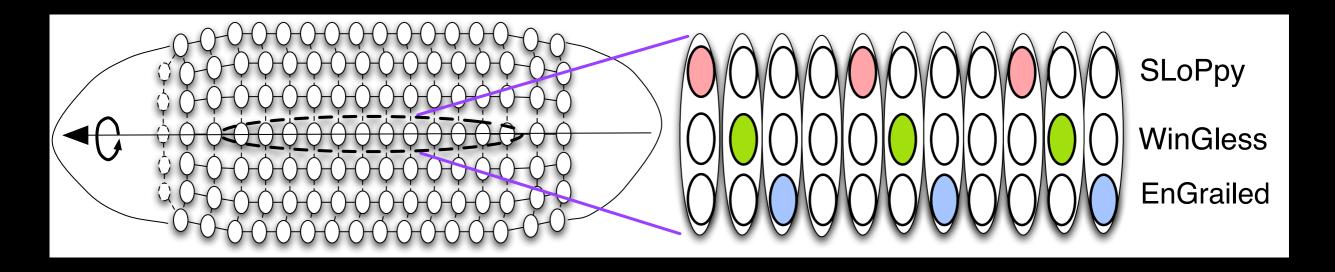
Let's apply radius/state tradeoff "cut-and-shift" procedure.



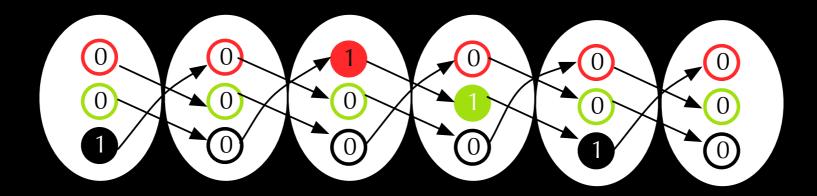
To get a make the trade, the algorithm adds two extra "slots."



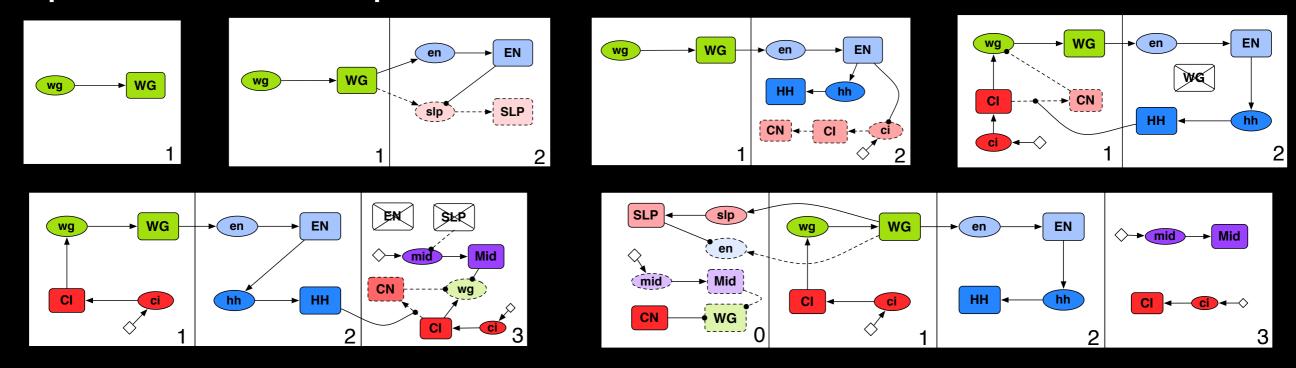
Focus on the three proteins: Sloppy, Wingless, Engrailed.



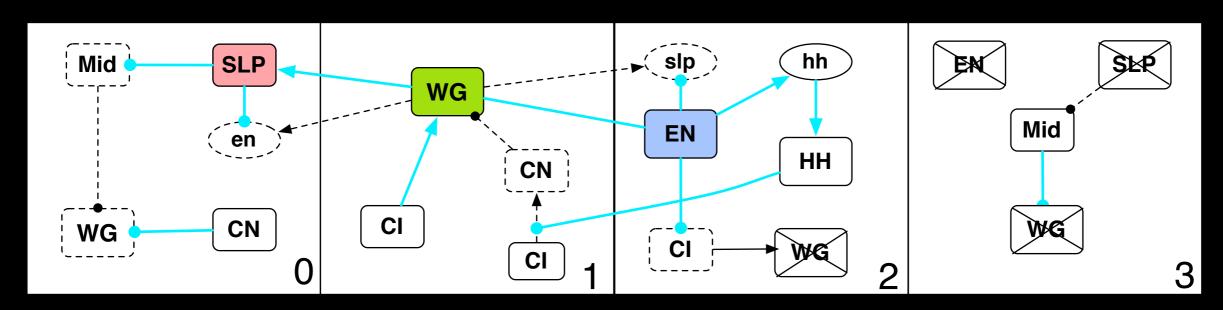
Interesting superficial resemblance:

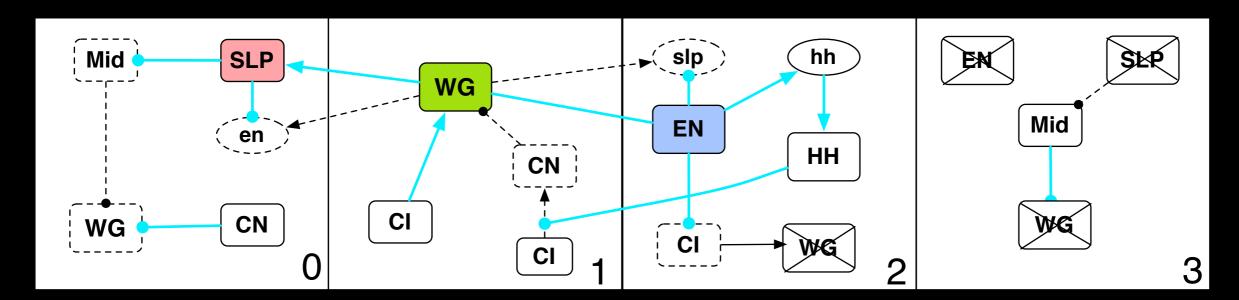


Much stronger form of validation comes by isolating a "path" within the gene regulatory network that stabilizes the three proteins in a spatial feedback sequence:

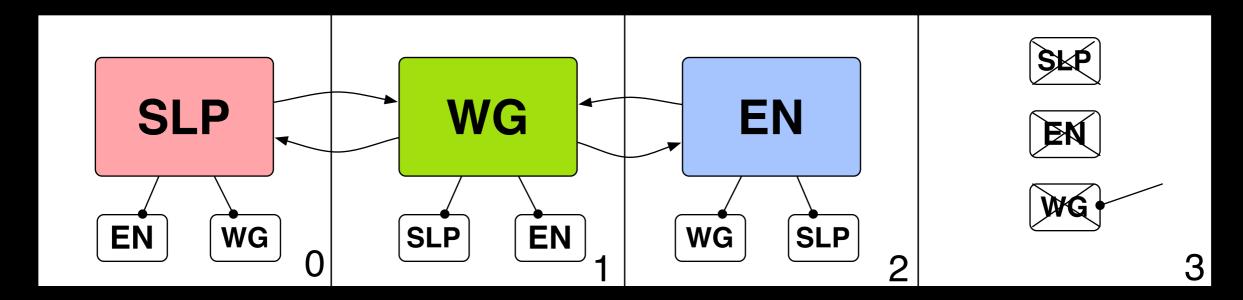


The dynamically activated subnetworks:





Consolidating the intermediates:



Conclusion: the coordination of multiple input-output state relations that unfold over space and time allow cellular agents with nearest-neighbor views to generate long-range coordinate patterns.

Experimental: look in related drosophilids for modifications (DePace lab)

Beyond Drosophila

(kinda, sorta, maybe)

 What does thinking of natural spatial multi-agent systems qua computers tell us scientically?

It tells about local information requres, helping understand:

 What agent resource capacities are required to solve a given task.

But the things we learned (i.e. cut and shift) were general.

 Can we then turn around this knowledge to build a better bug?

so that we could, for example, make general patterns?

Goal: Design local rules to make given patterns.

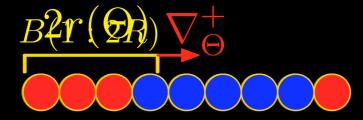
Repeat patterns have a well-defined gradient.

$$\nabla_{\Theta}(B)^{+} = \begin{cases} i, & \text{if } B \circ i \text{ is consistent with } \Theta \\ B(2R), & \text{otherwise} \end{cases}$$

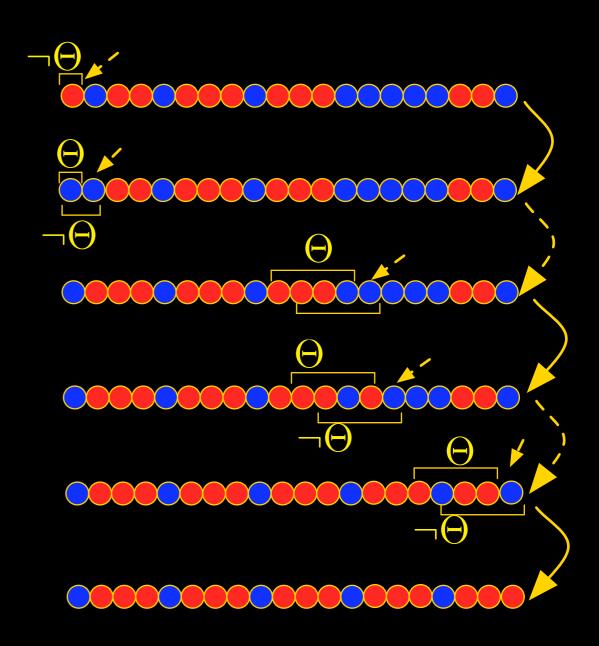
For example: 1000-repeat pattern (which has an r = 2 check scheme),

Now simply define:

$$F(B) = \nabla_{\Theta}^{+}(B(1:2r))$$

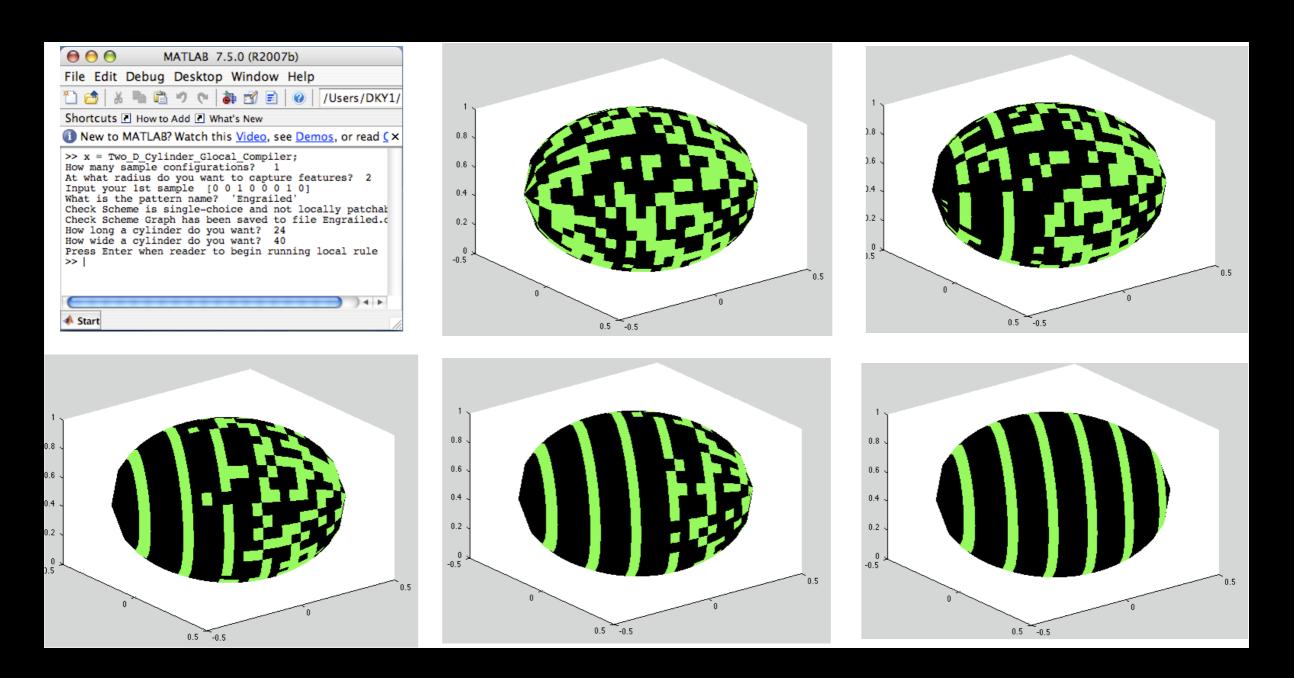


Gradient "spreads correctness."



Gradient waves ... Multiple "waves" all at once in actuality

Not hard to implement in higher dimensions.



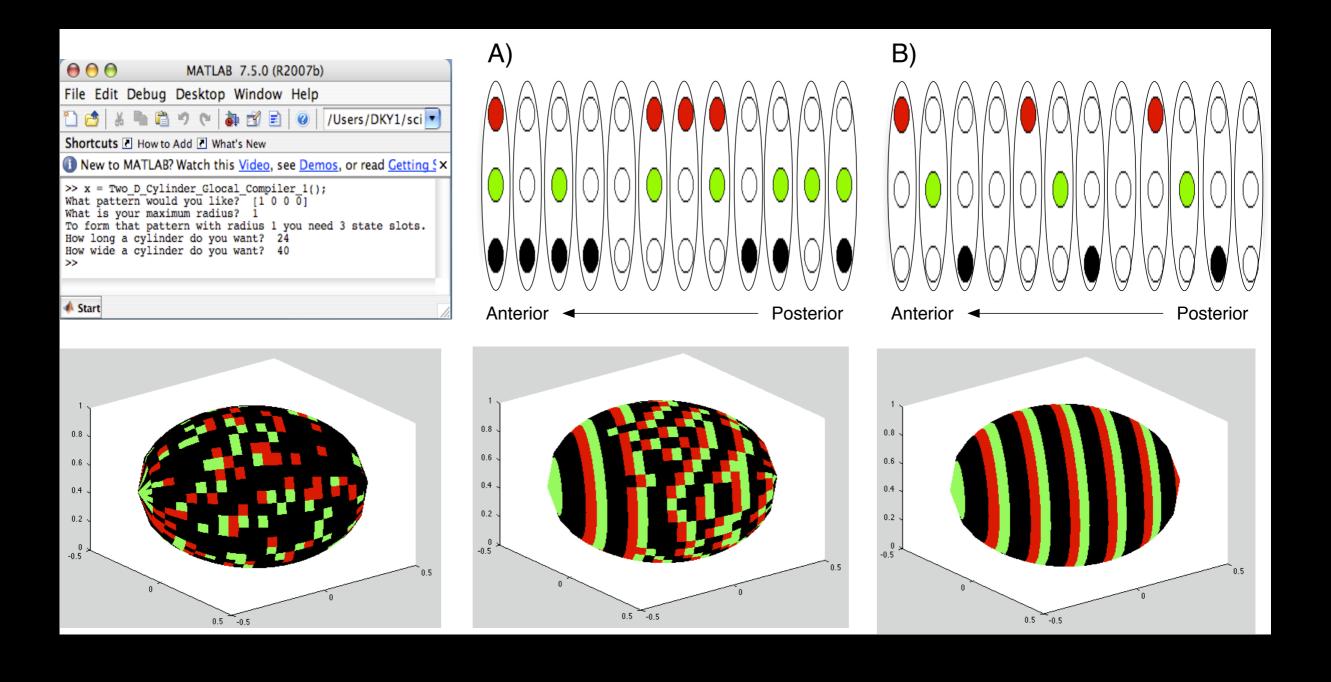
Resulting rule is 100% robust to:

- Initial condition perturbations.
- Timing issues.

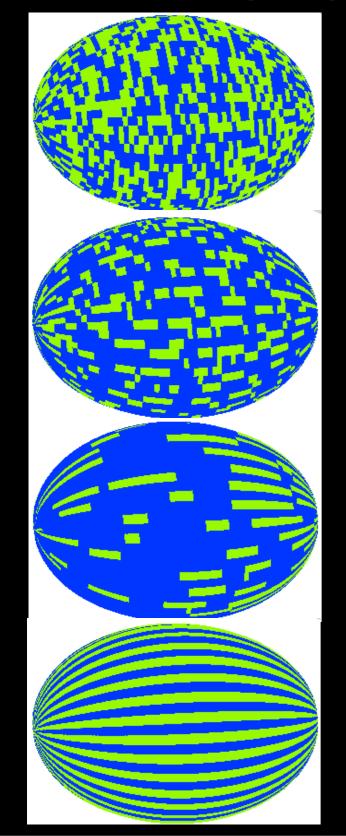
But: the rule to solve a repeat pattern of size |q| takes a radius of |q|.

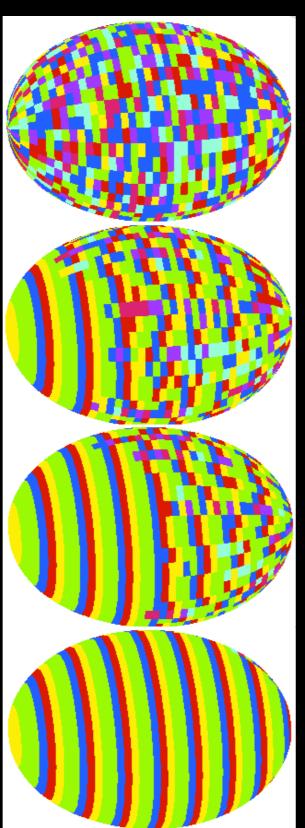
Can run a radius-minimization algorithm.

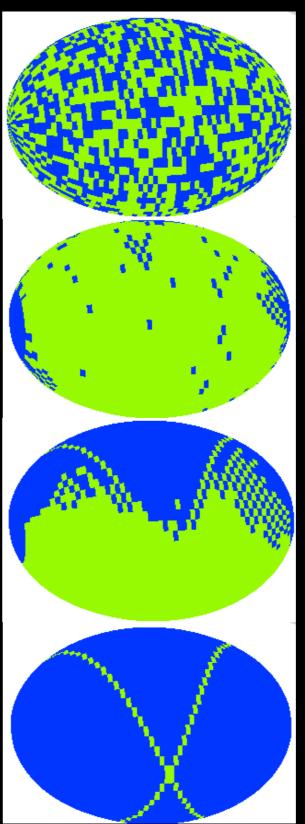
And then, couple in the radius/state tradeoff.

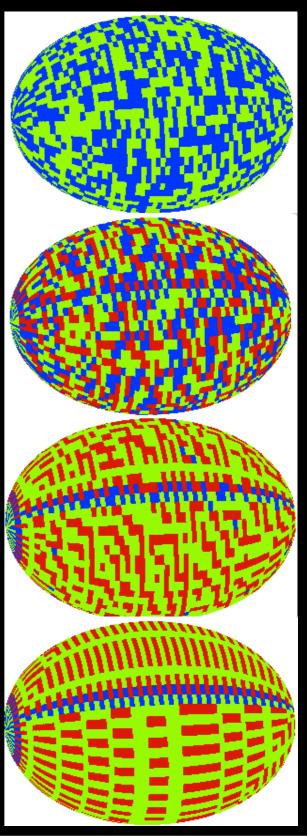


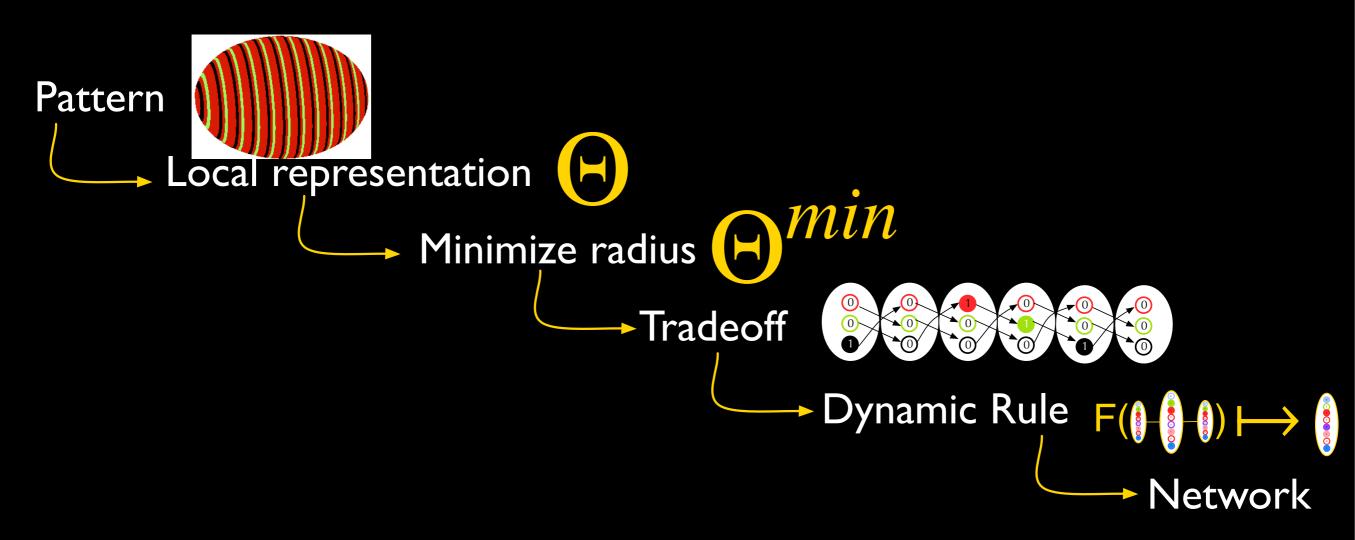
Similar principles can be applied to develop a language for building a diverse array of patterns.













 Some VERY simple ideas in multi-cellular synthetic biology might be "understandable" (or at any rate, imagineable) and connectable to cellular programs

Is anything here doable? even remotely?

 What is the simplest system that could possibly exploit some of these ideas?

What I'd really like to explore is evolution.