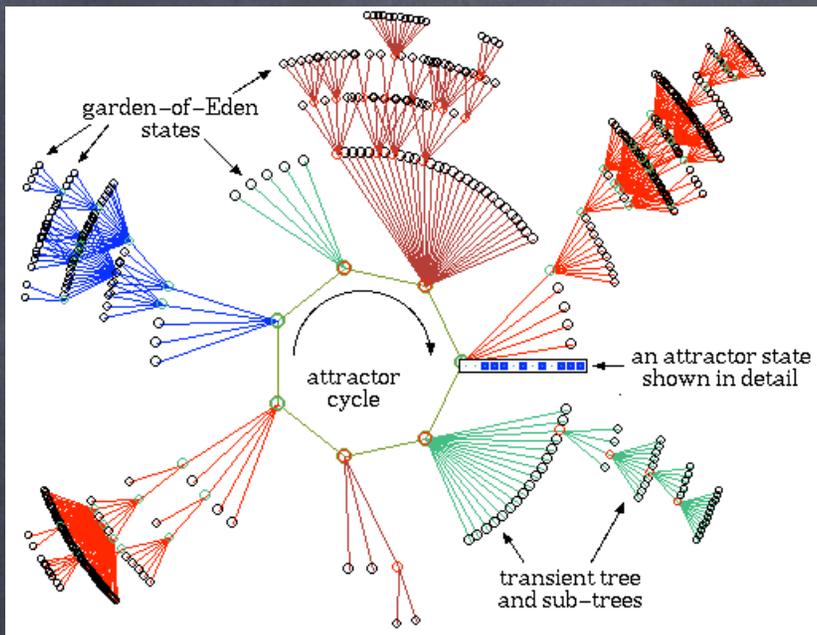


6. Regulatory Models that Mimic Phenotype and Dynamics, Part II

Warning: Statistical Physics.
It only works on average.

<http://regan.med.edu/CVRB-course.php>

Rhythm and its robustness



- Limit cycle attractors
 - ➔ no single steady state
 - ➔ potentially sensitive to fluctuating time delays

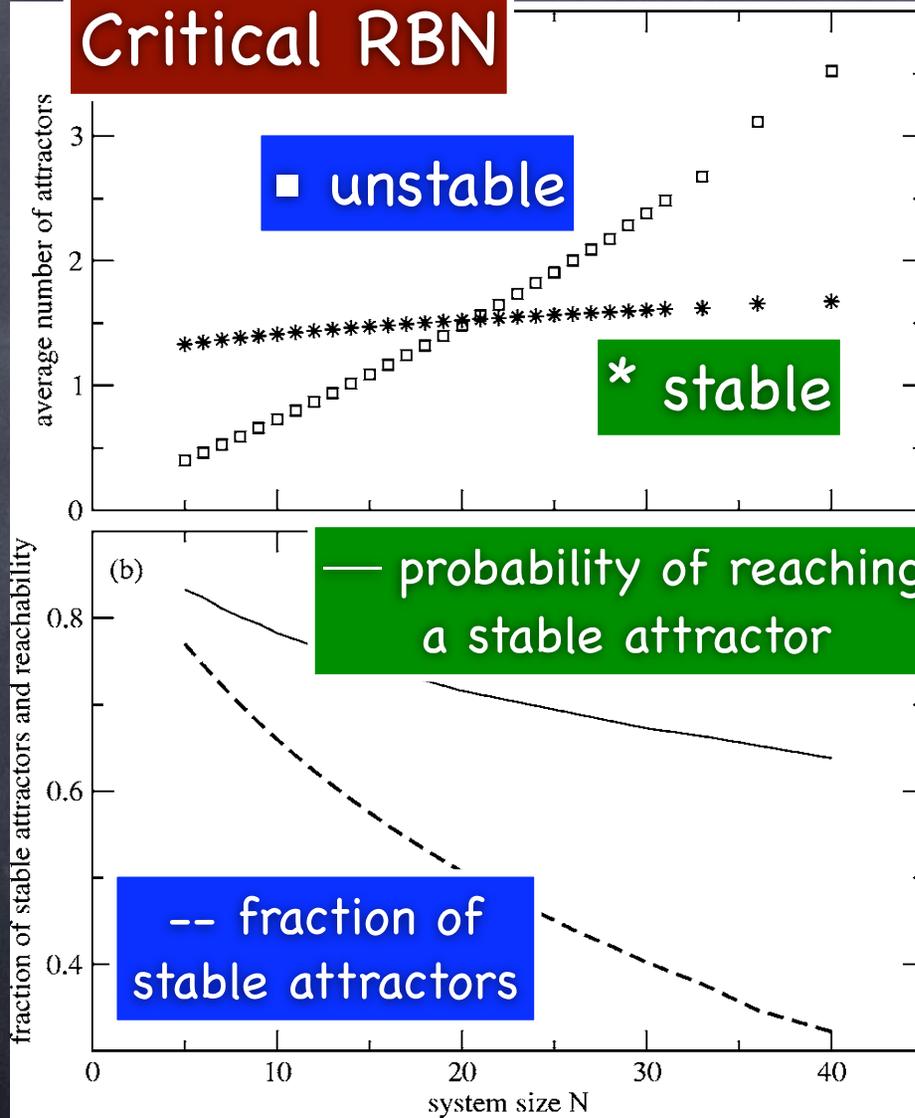
Are cyclic attractors artifacts of synchronous update?

- Noise in synchronous update
 - ➔ continuous time
 - ➔ low-pass filter on node switching
 - ➔ the "command" from the gate driving a node has to stay consistent for a time $s < 1$

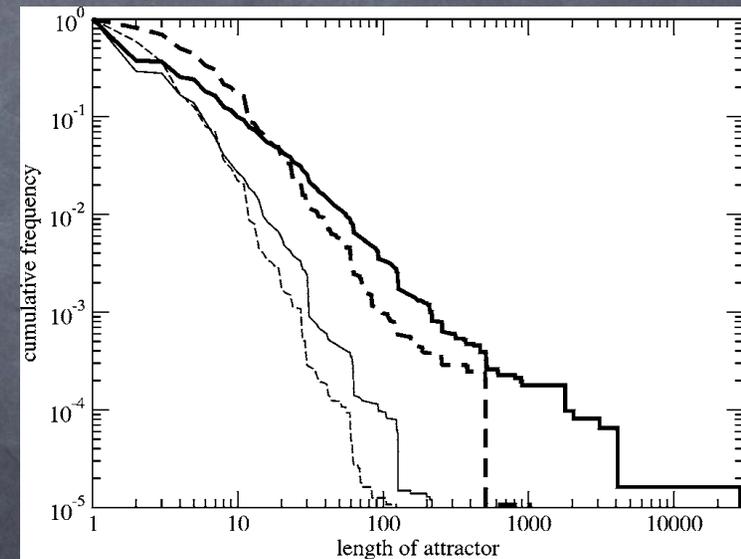
- **Reliable attractor:**
 - ➔ small random time delay on nodes: $\epsilon \ll s$
 - ➔ all possible perturbations in time delay lead to the same attractor

In random Boolean networks...

Critical RBN



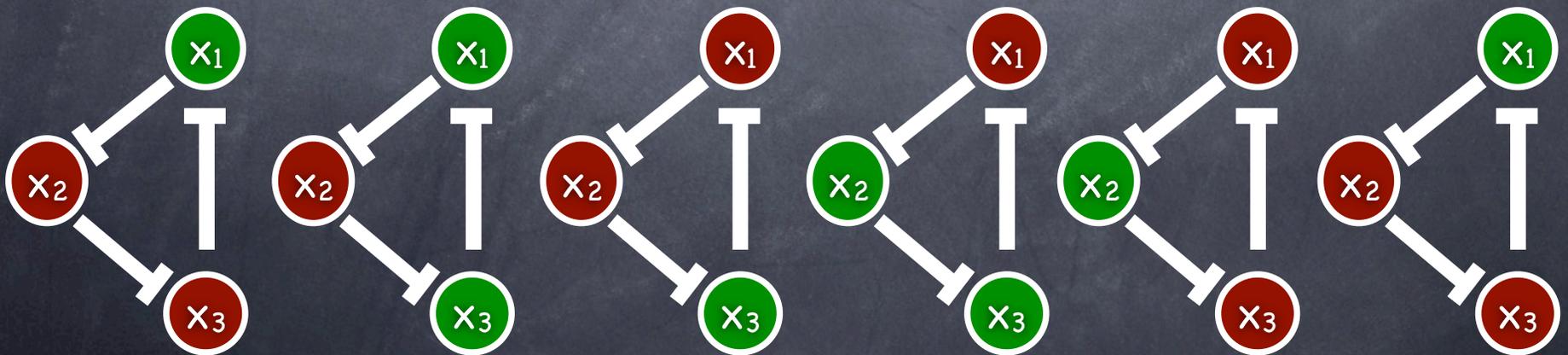
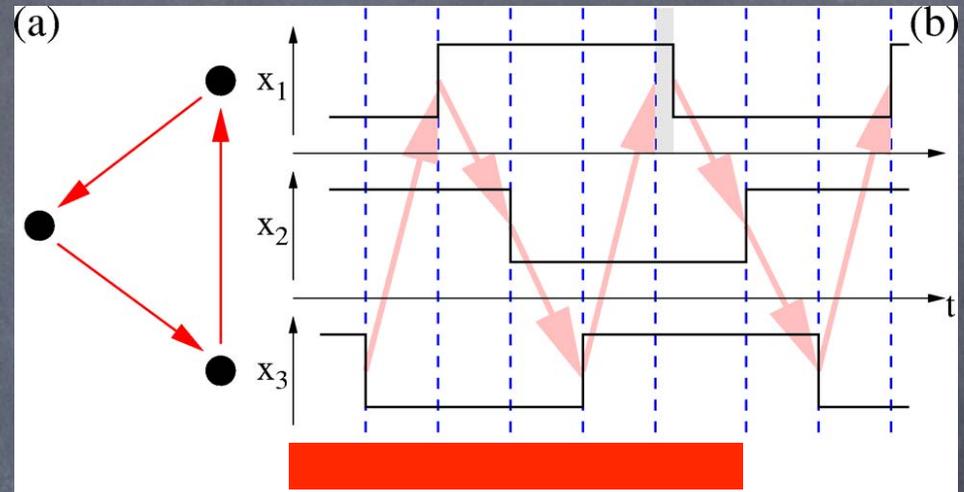
- Not so many stable attractors!
- They have larger basins of attraction



- They tend to have longer cycles!

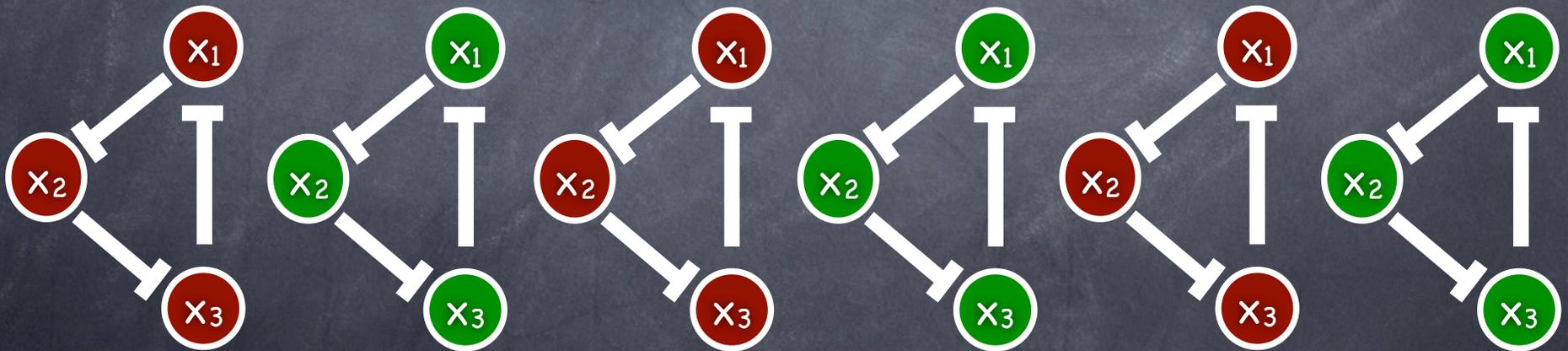
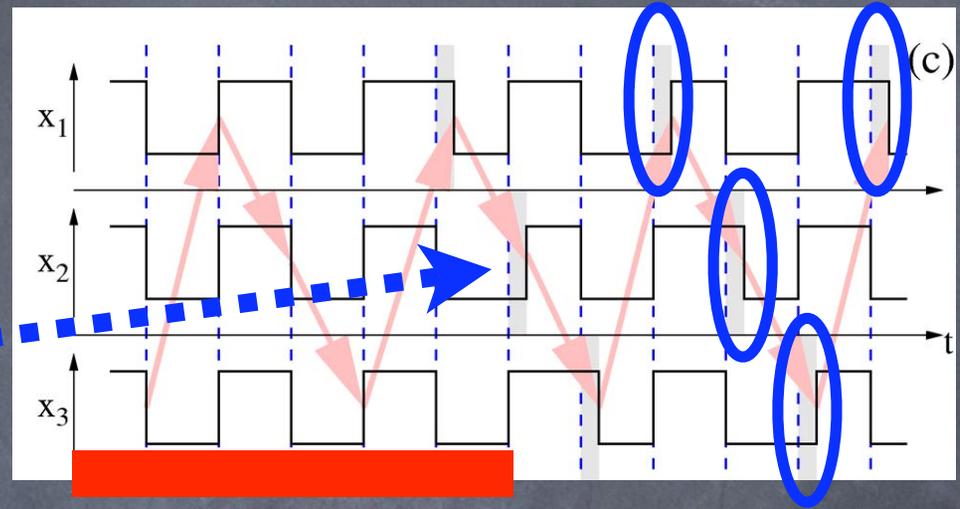
What type of wiring makes a non-reliable attractor?

- Extreme modularity (causally disconnected network components)
- More than one cascade of switching events!



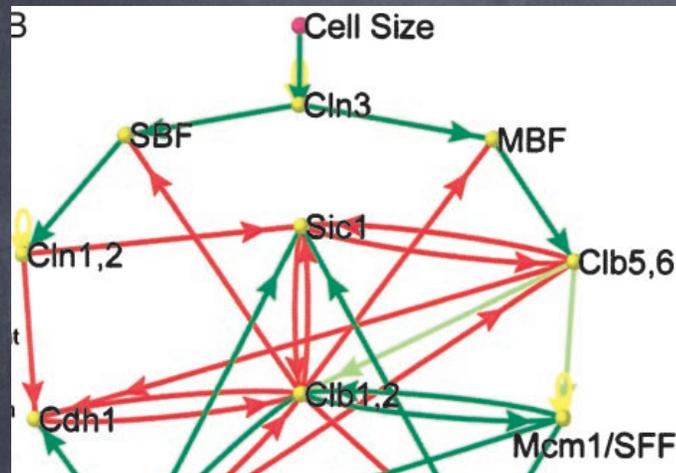
Juggling is unstable

- More than one cascade of switching events!
- Causal cascades can accumulate phase shift and loose synchrony



Causal switching chains need to stay in lock-step!

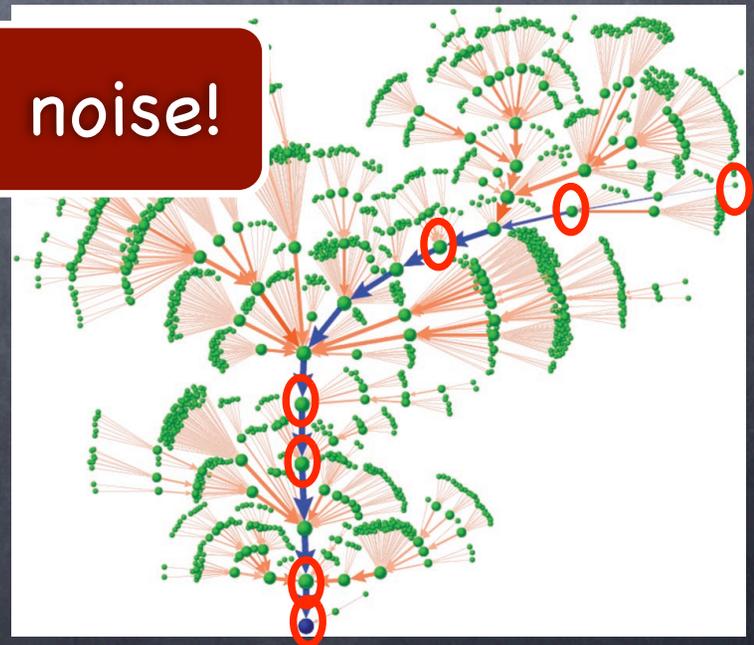
Back to biology: cell cycle revisited



- Same model of yeast CC
- Same update gates
- Noise in signal propagation time
 - continuous time
 - low-pass filter for switching
- Cycle: **state S1 triggers Cln3 activation**

Completely stable for small noise!

Buffer steps with change in just 1 node!

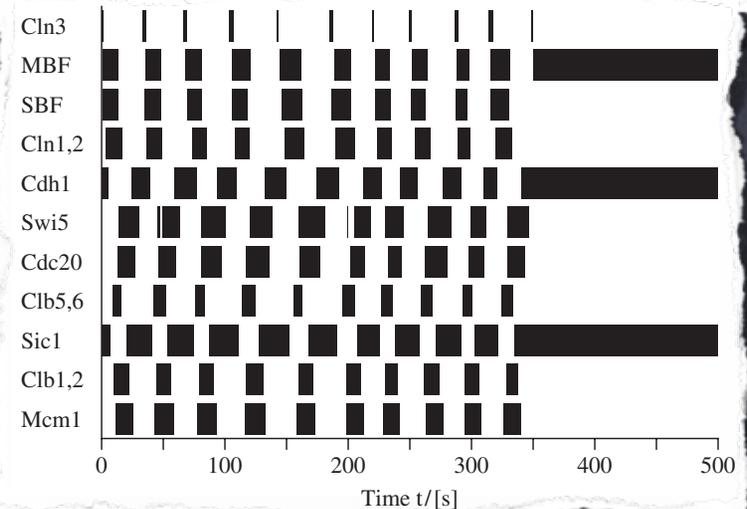
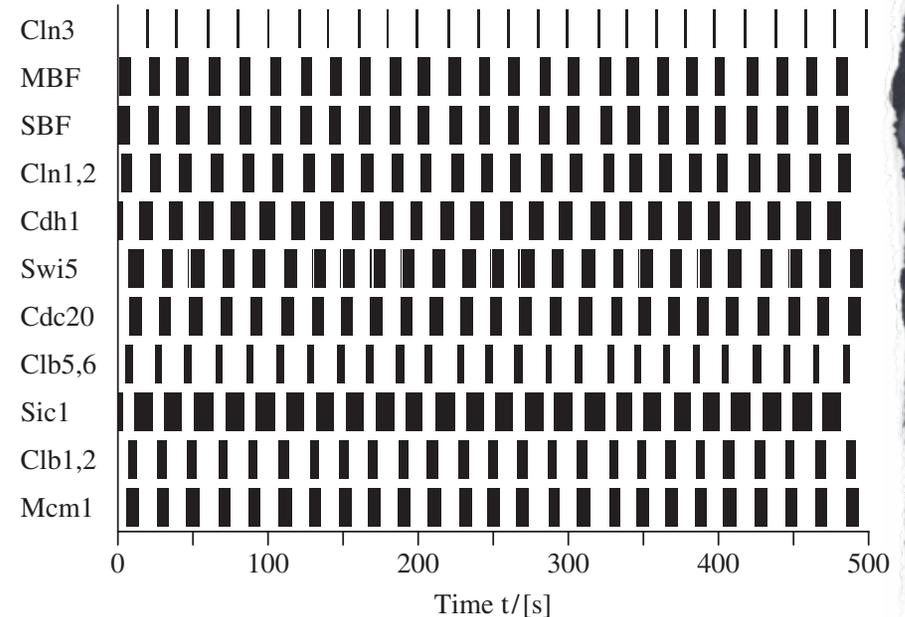


Does the cell cycle juggle?

- Stronger noise
 - allow delays past 1/2 the propagation time unit
- Looser stability measure
 - G1 is regularly assumed for a time period long enough to trigger CC restart

The system stays within the same attractor!

- Although attractor switch is possible



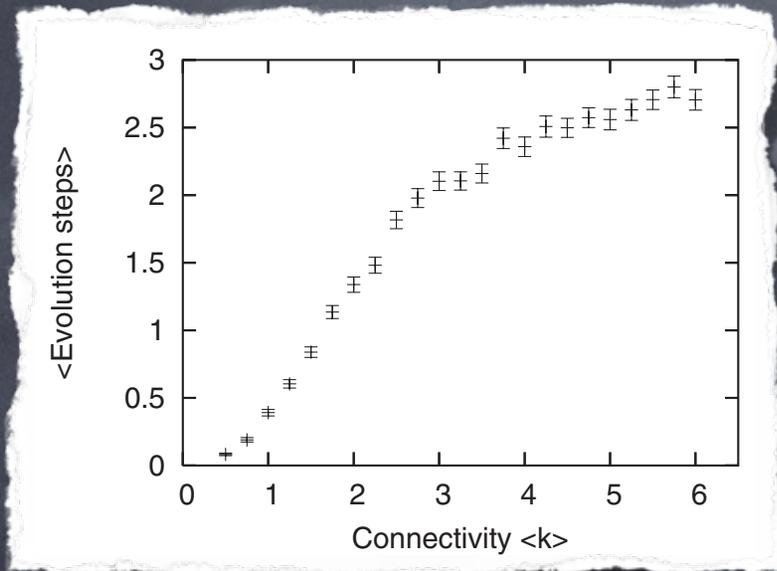
Is reliability evolvable?

- Random threshold networks
 - ➔ map attractor landscape
 - ➔ record stable and unstable attractors
 - ➔ **fitness score**
- Evolutionary dynamics
 - ➔ rewire 1 link
 - ➔ measure new fitness
 - ➔ if higher than original, keep new network

A) Full attractor landscape
sum of stable basin sizes

B) Functional attractor
largest stable attractor basin size

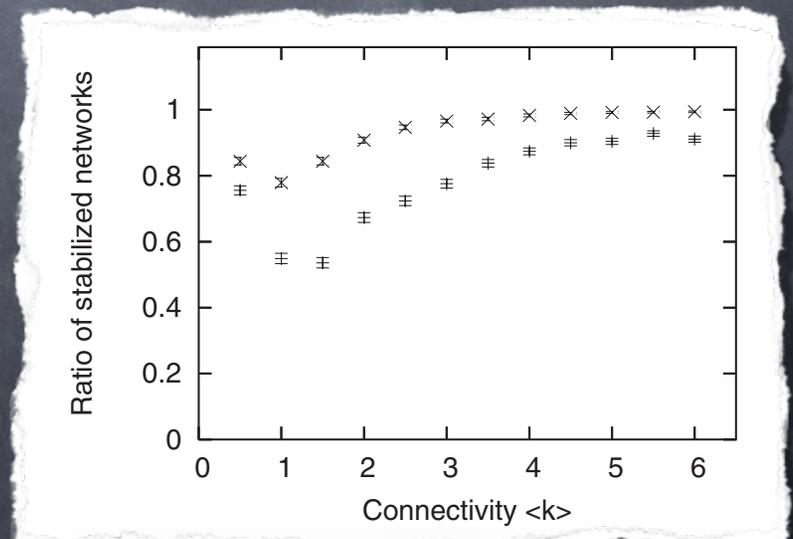
Evolving a stable landscape is easy!



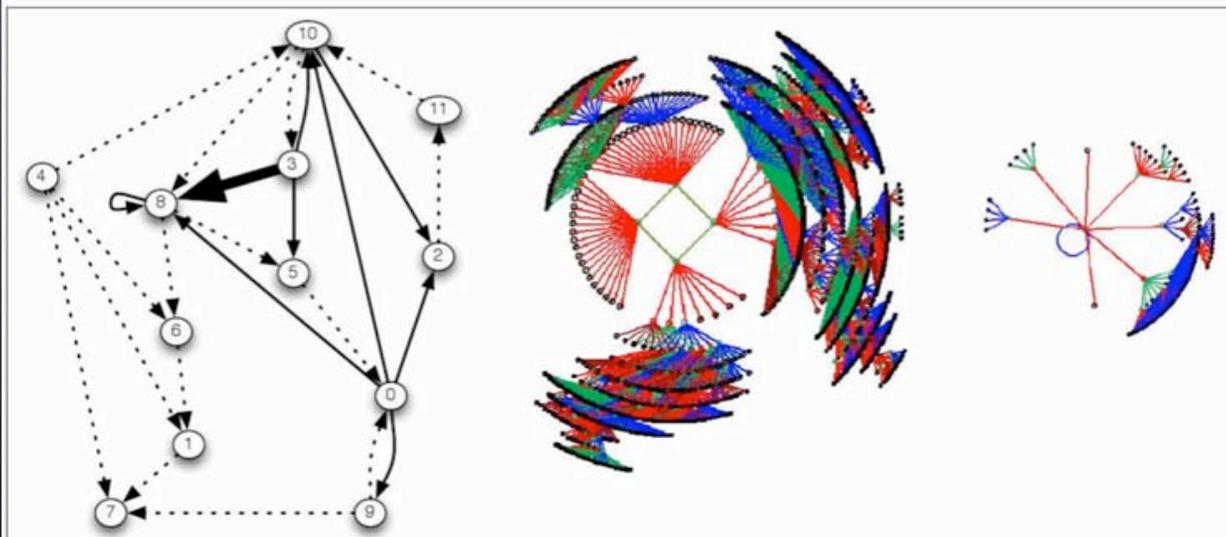
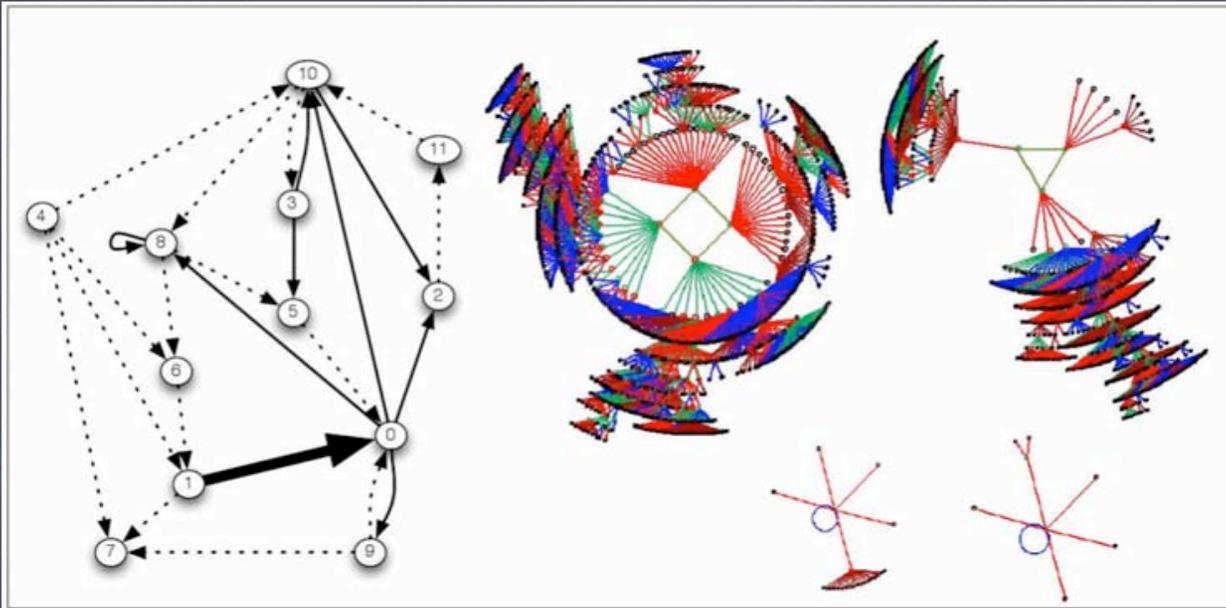
- How about one "functional" attractor basin?

➔ evolution stops when half of the configuration space belongs to the stable, functional attractor

Larger $\langle k \rangle$ is better!

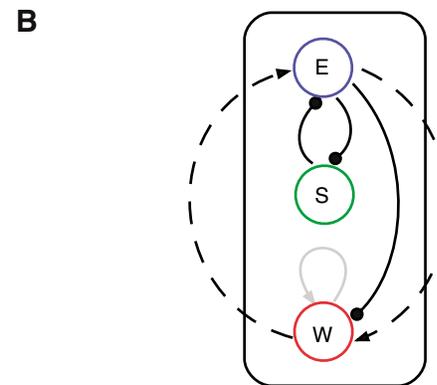
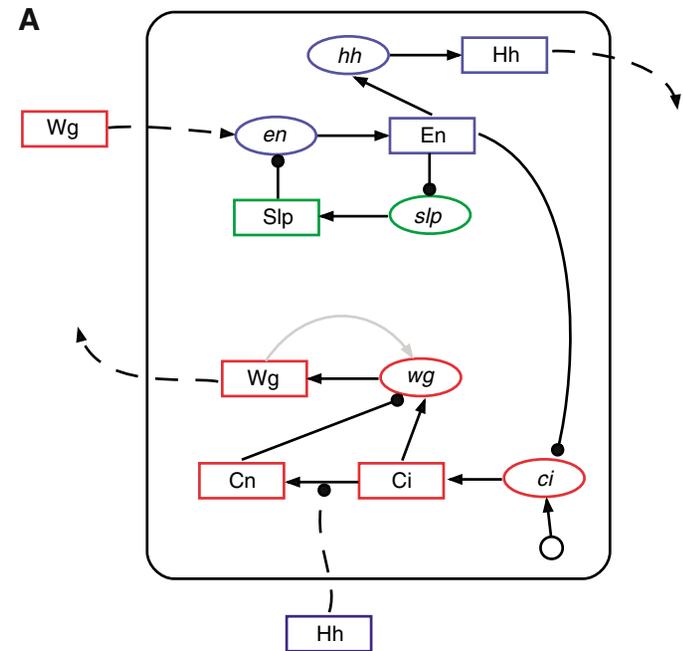


An example



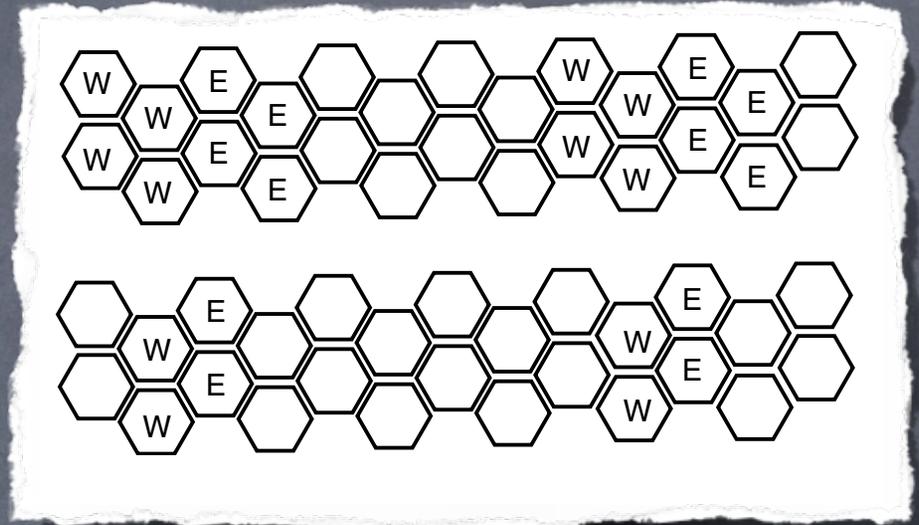
Does function dictate structure?

- Let's find ALL small networks that could perform the segment polarity patterning in *Drosophila*
 - ➔ how many of these are robust?
- Enumerated all possible 3-node networks
 - ➔ each can regulate itself and others
 - ➔ each link can be inter- and extracellular
 - ➔ restrict to 2 of 3
 - ➔ 14,348,907 topologies



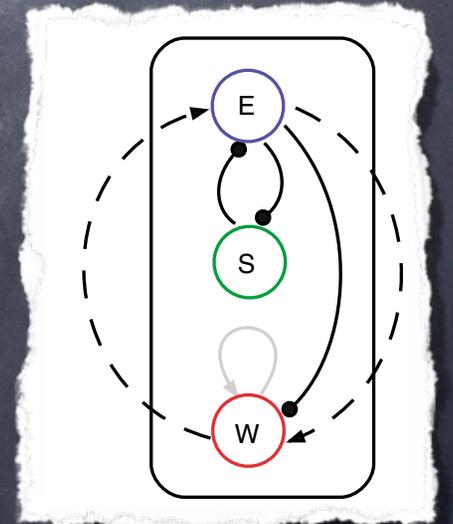
Does function dictate structure?

- A network is functional if:
 - ➔ has perform correct patterning
 - ➔ robustness: fraction of parameter space that can perform the function
 - ➔ (ODE's), parameters sampled at random

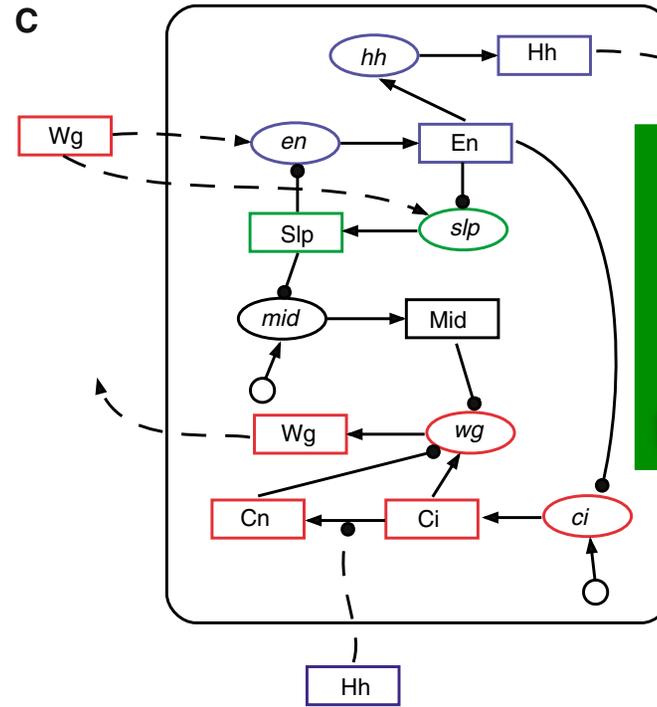
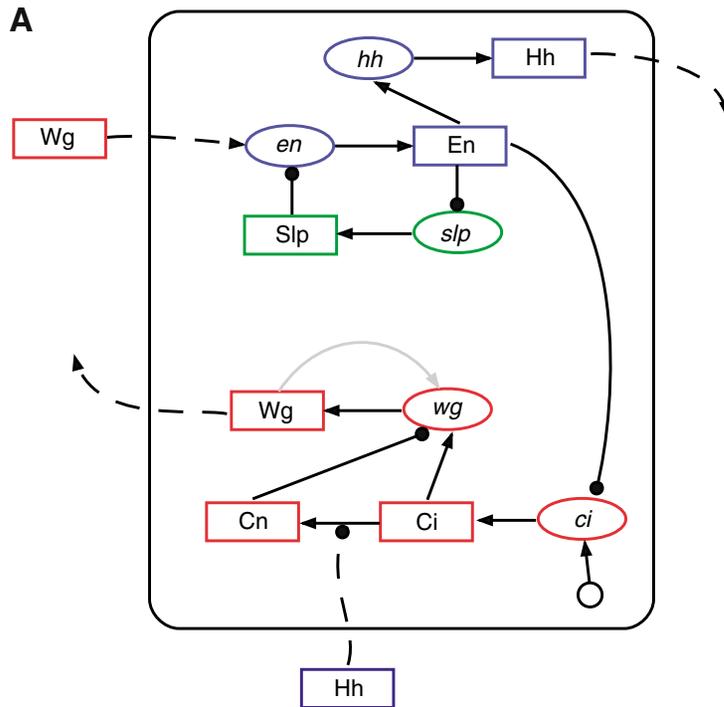


Biological topology: high score, not highest.

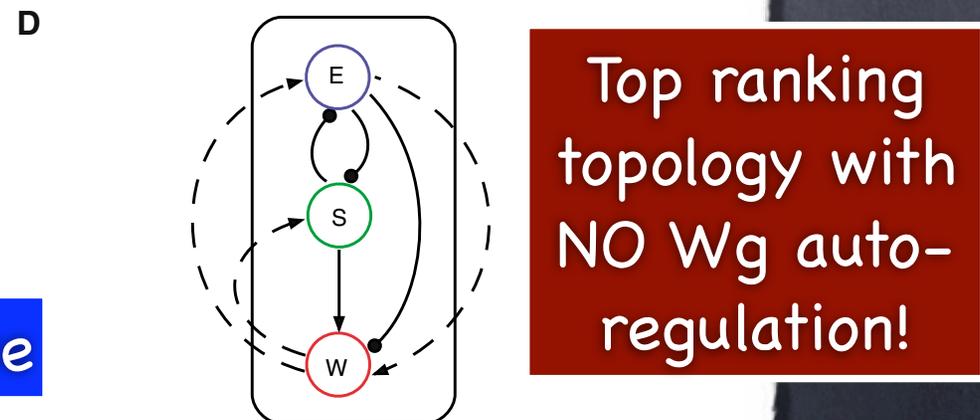
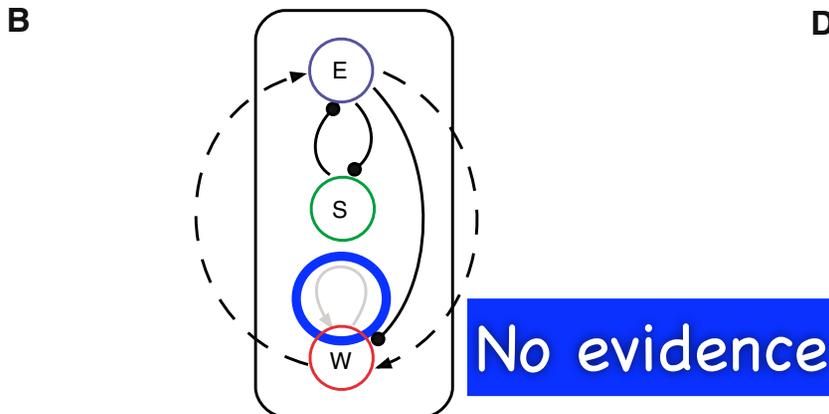
Excluding direct auto-regulation on E and S



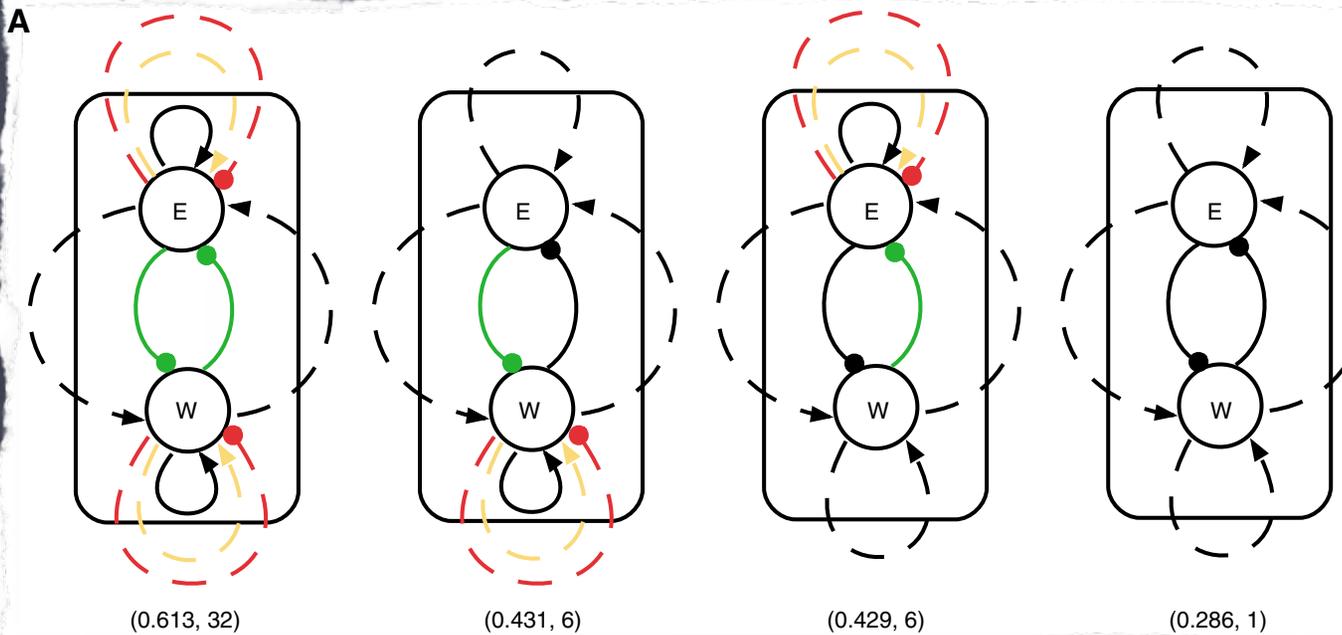
A twist on polarity models



Evidence
for *Slp* →
Wg
regulation!



Actually, 2 nodes are enough for patterning...



Positive feedback on E and W

E and W neighbors express W/nothing OR E/nothing! Bistable in E and W

Mutual intracellular E - W inhibition

Sharp boundary: E next to W only

4 core topologies: black links are required for robust patterning

- Green: neutral
- Orange: bad
- red: very bad

7. Transcriptional Regulation from Microarray Data

June 22
12 PM