

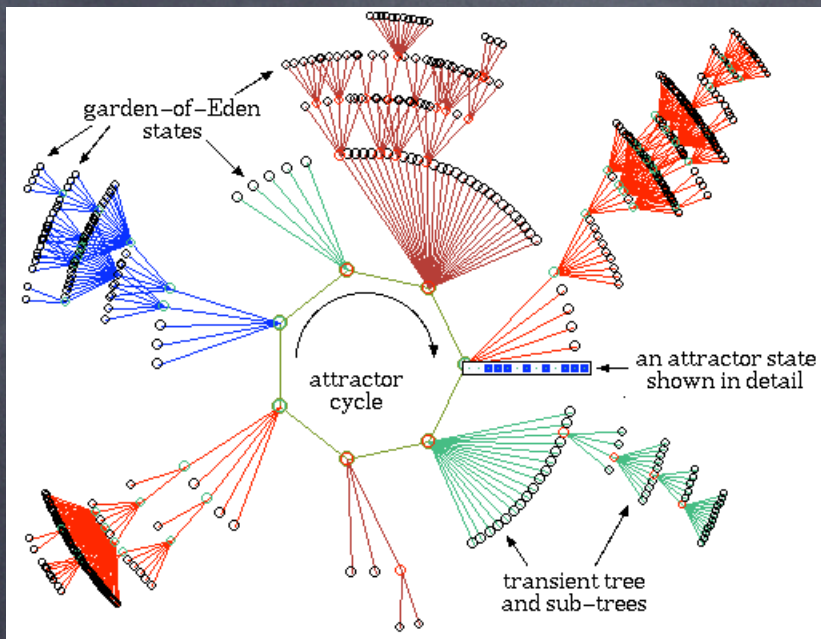
# 6. Regulatory Models that Mimic Phenotype and Dynamics, Part II

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

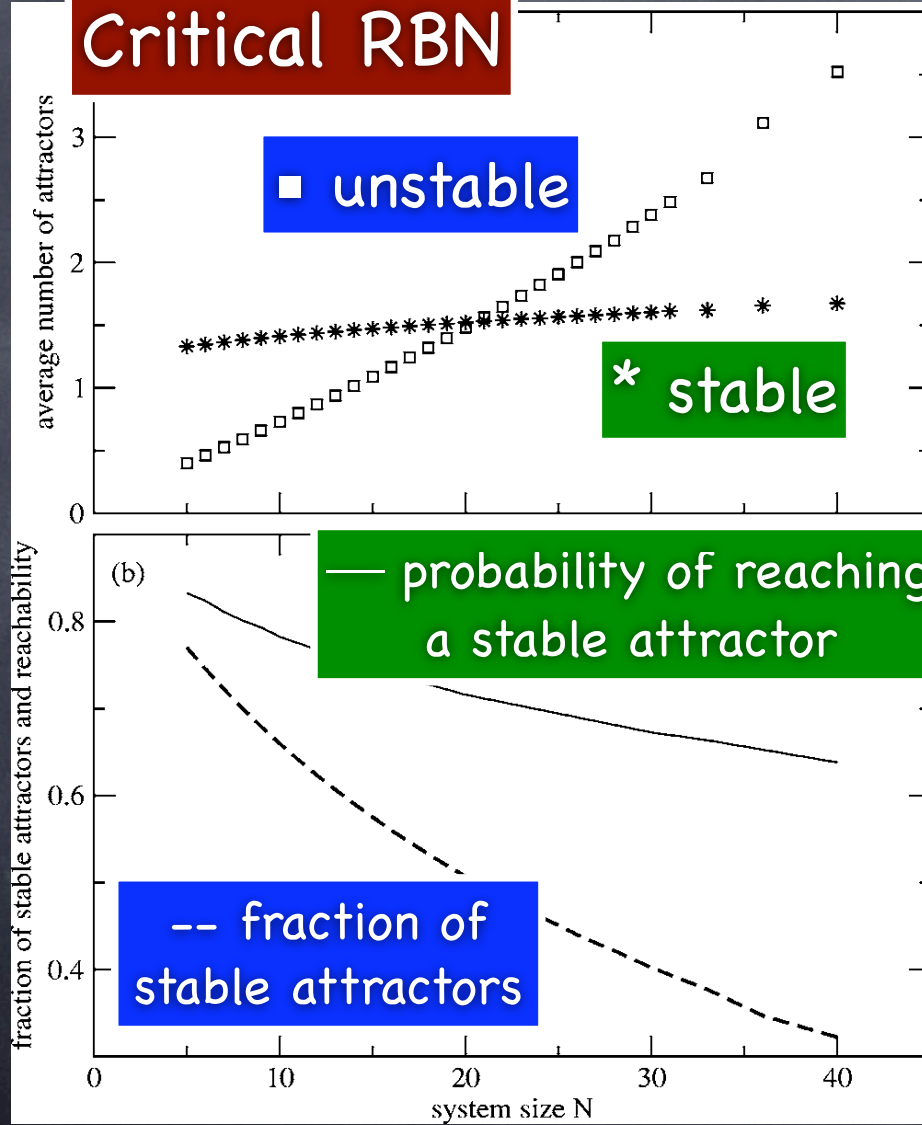
Are cyclic attractors artifacts of synchronous update?

- 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  $s < 1$

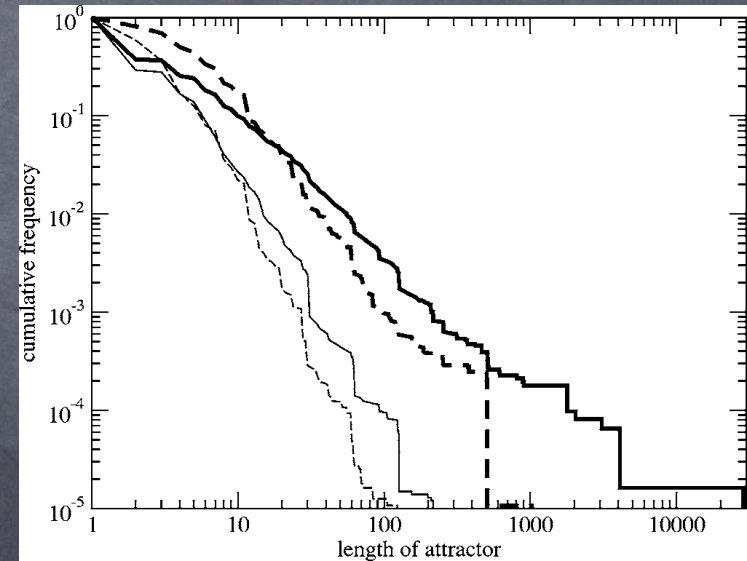
- **Reliable attractor:**
  - ➔ small random time delay on nodes:  $\epsilon \ll s$
  - ➔ all possible perturbations in time delay lead to the same attractor

# In random Boolean networks...

## Critical RBN



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

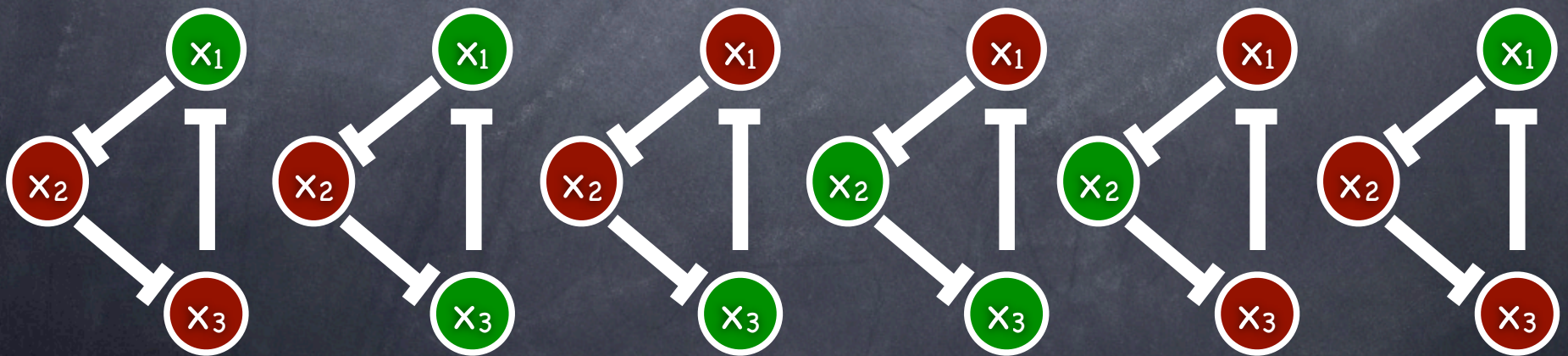
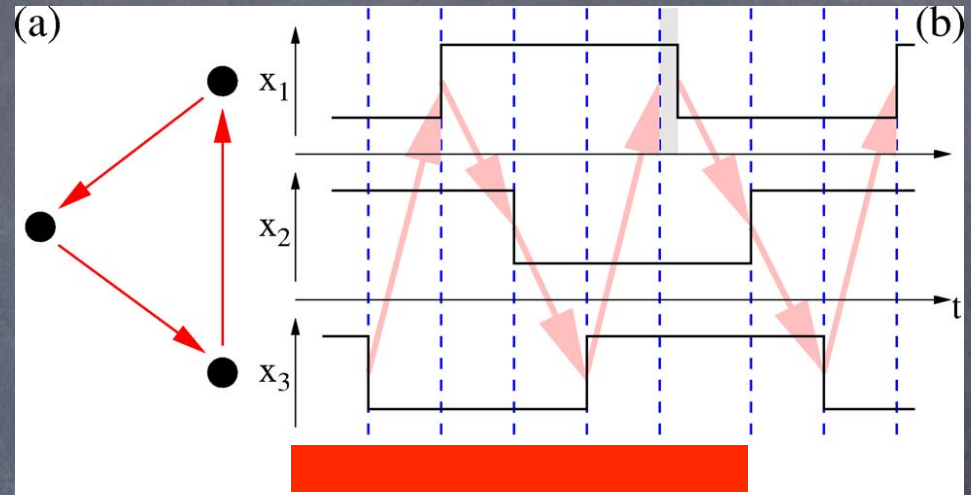


- They tend to have longer cycles!



# What type of wiring makes a non-reliable attractor?

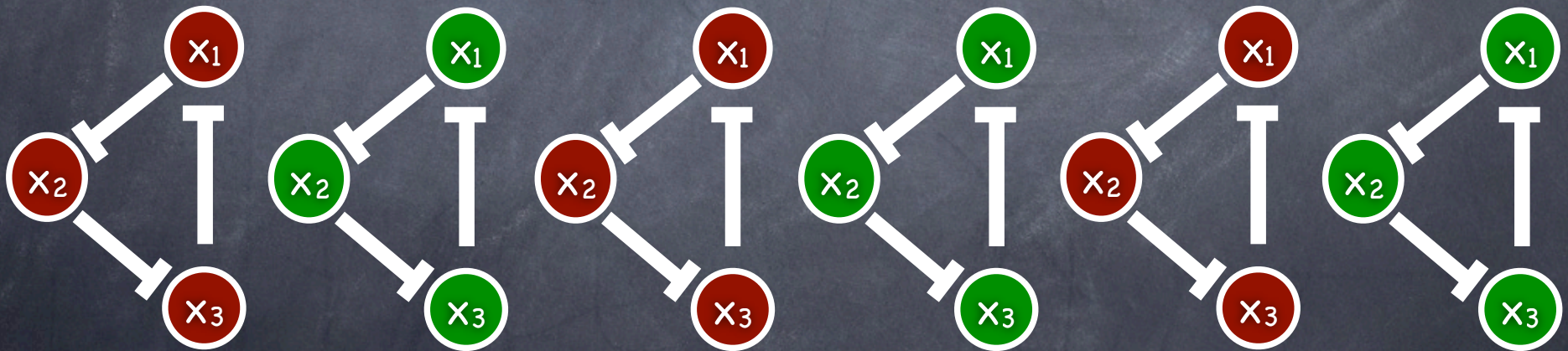
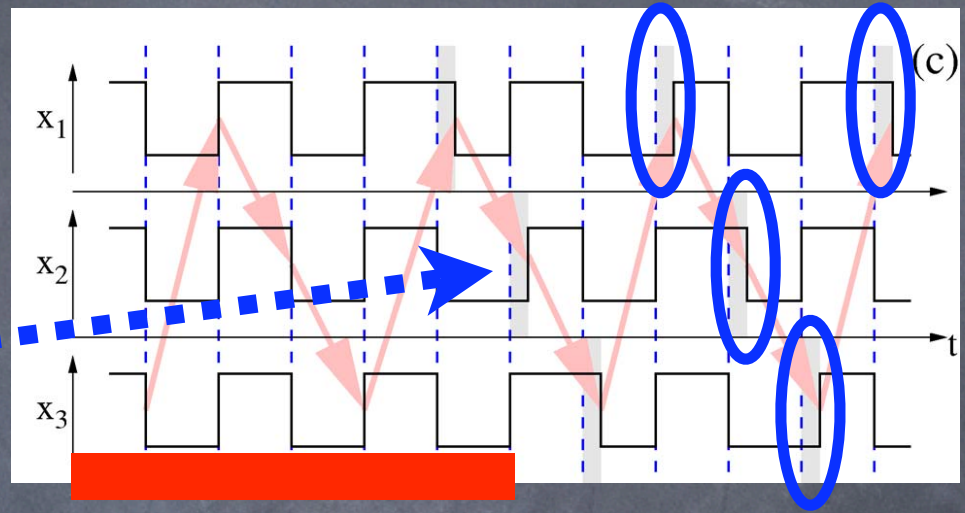
- Extreme modularity (causally disconnected network components)
- More than one cascade of switching events!





# Juggling is unstable

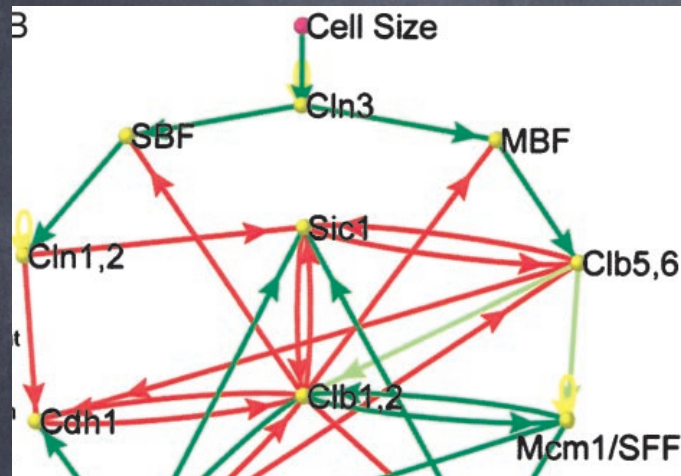
- More than one cascade of switching events!
- Causal cascades can accumulate phase shift and loose synchrony



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!

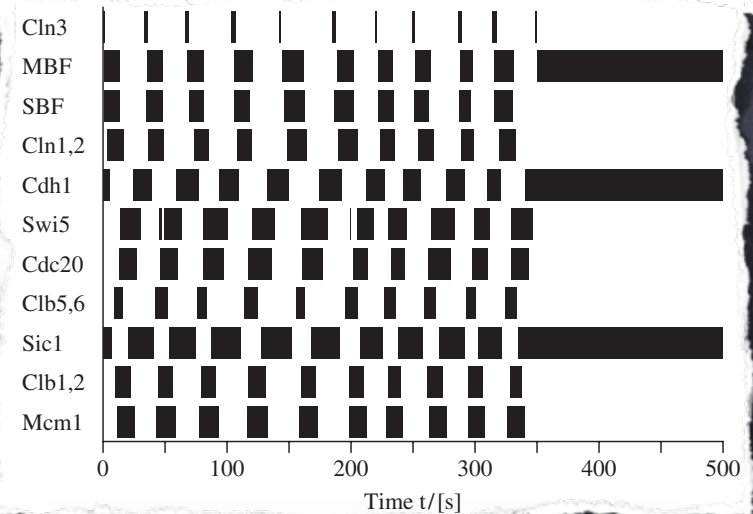
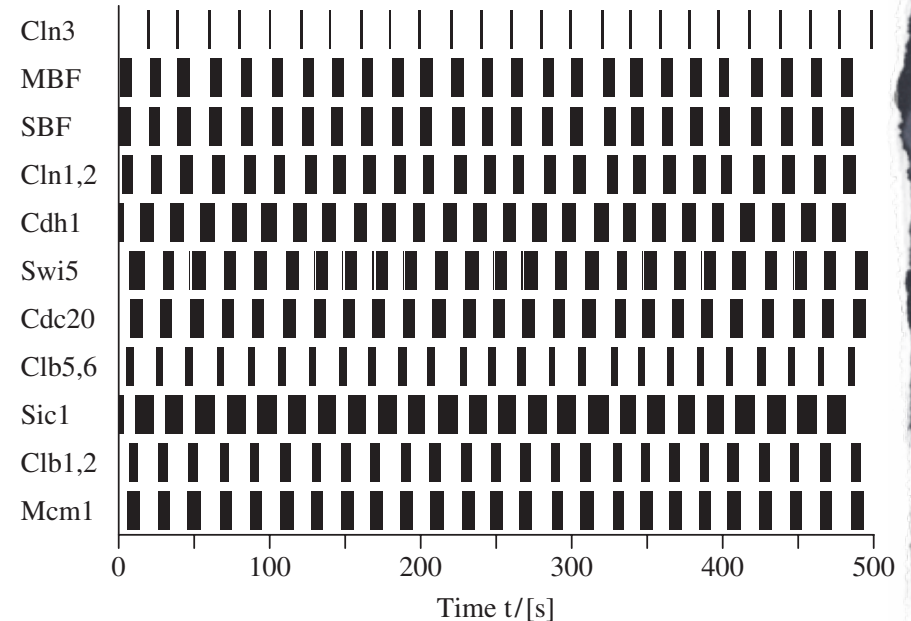


# Does the cell cycle juggle?

- 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?

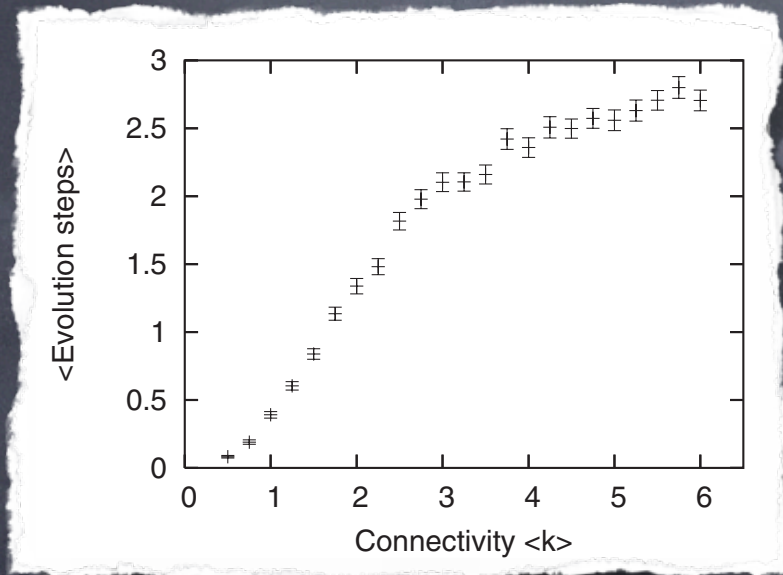
- Random threshold networks
  - ➔ map attractor landscape
  - ➔ record stable and unstable attractors
  - ➔ **fitness score**
- Evolutionary dynamics
  - ➔ rewire 1 link
  - ➔ measure new fitness
  - ➔ if higher than original, keep new network

A) Full attractor landscape  
sum of stable basin sizes

B) Functional attractor  
largest stable attractor basin size



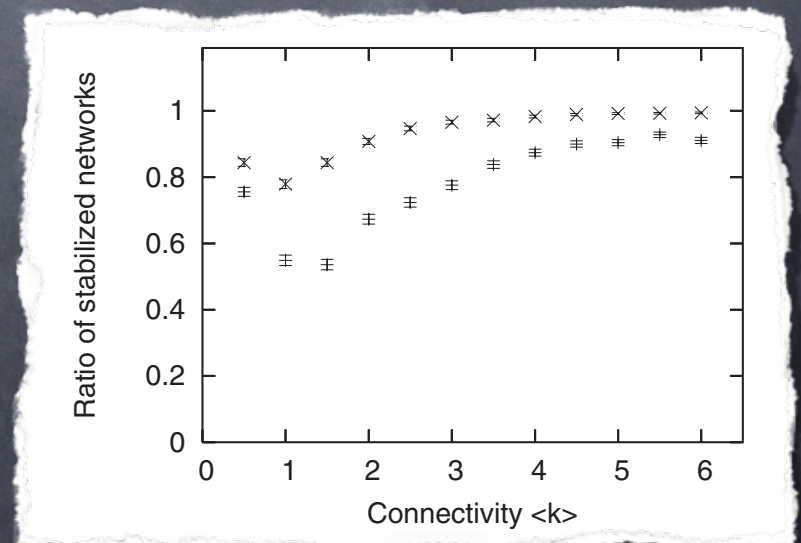
# Evolving a stable landscape is easy!



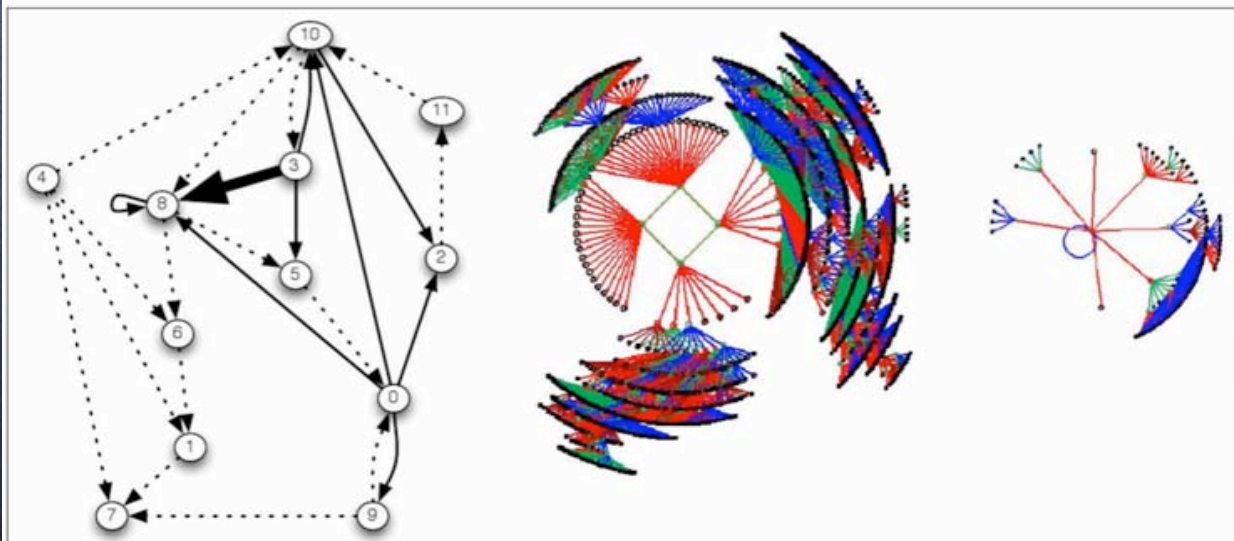
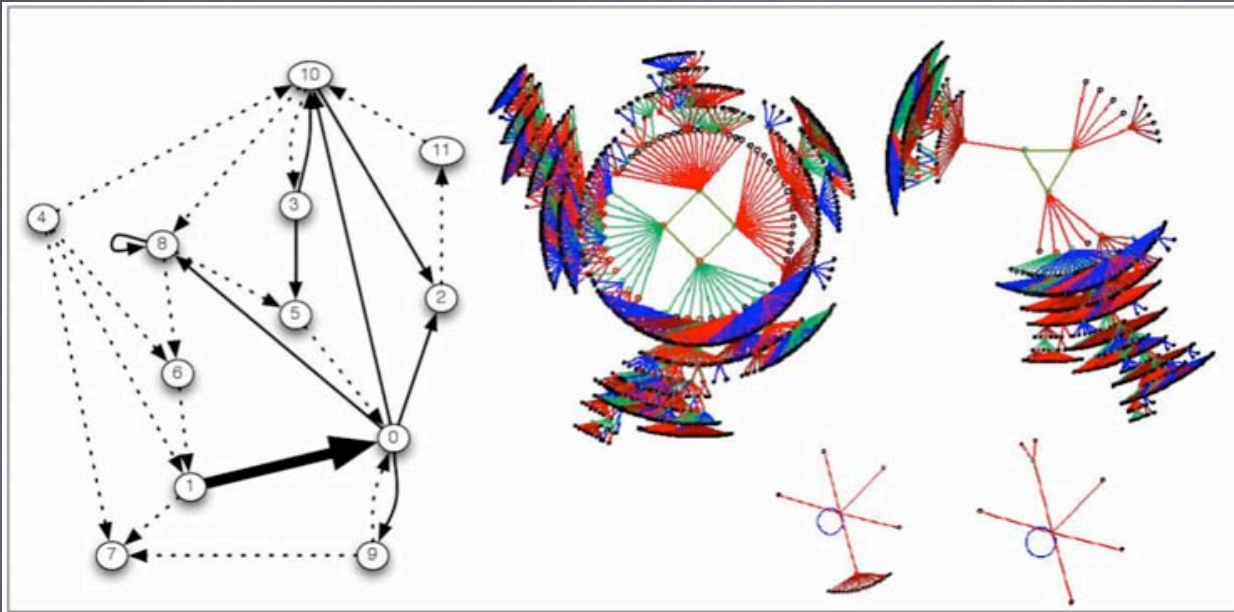
- How about one "functional" attractor basin?

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

Larger  $\langle k \rangle$  is better!



# An example

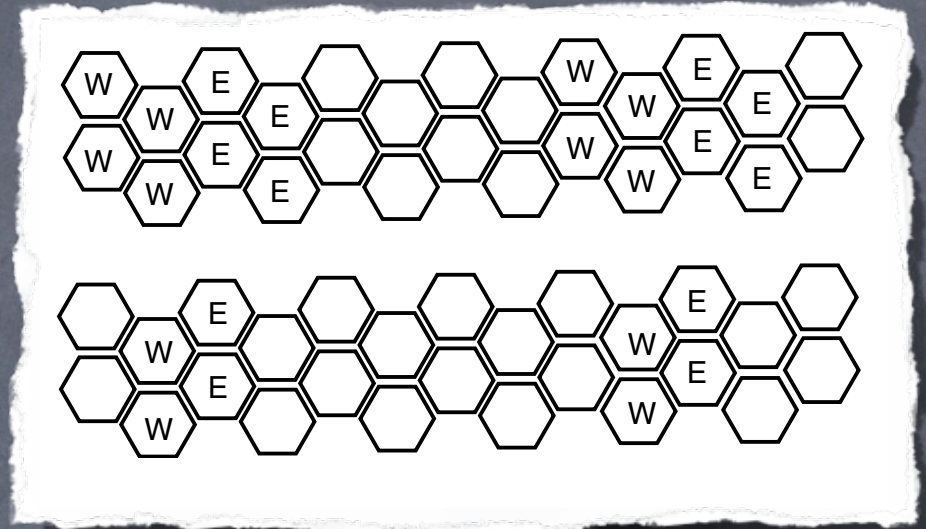






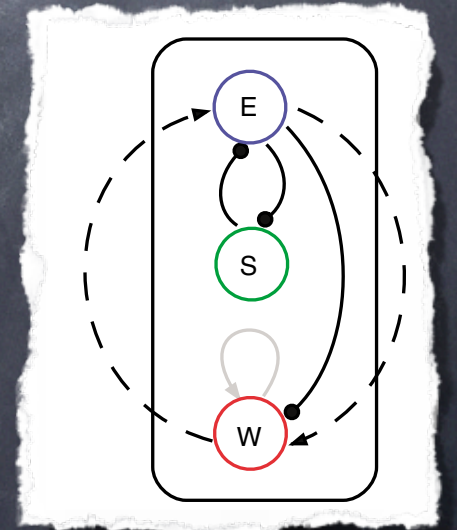
# Does function dictate structure?

- 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



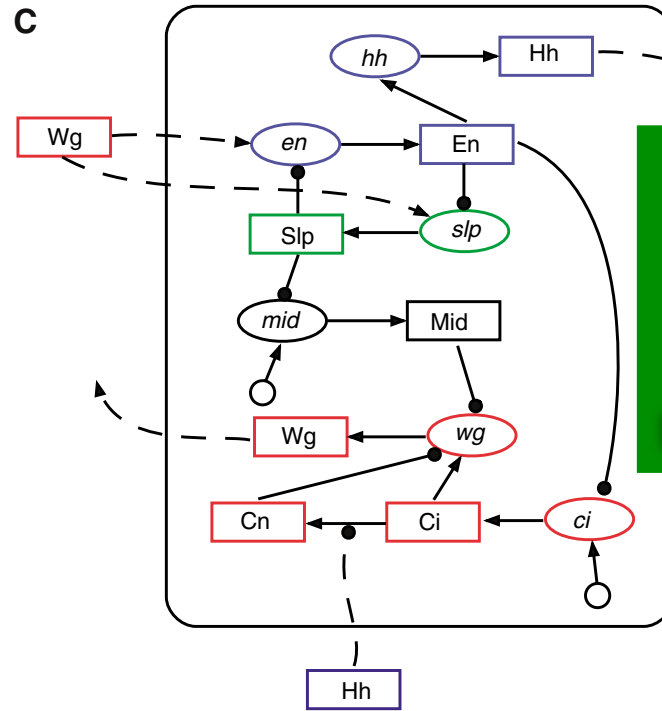
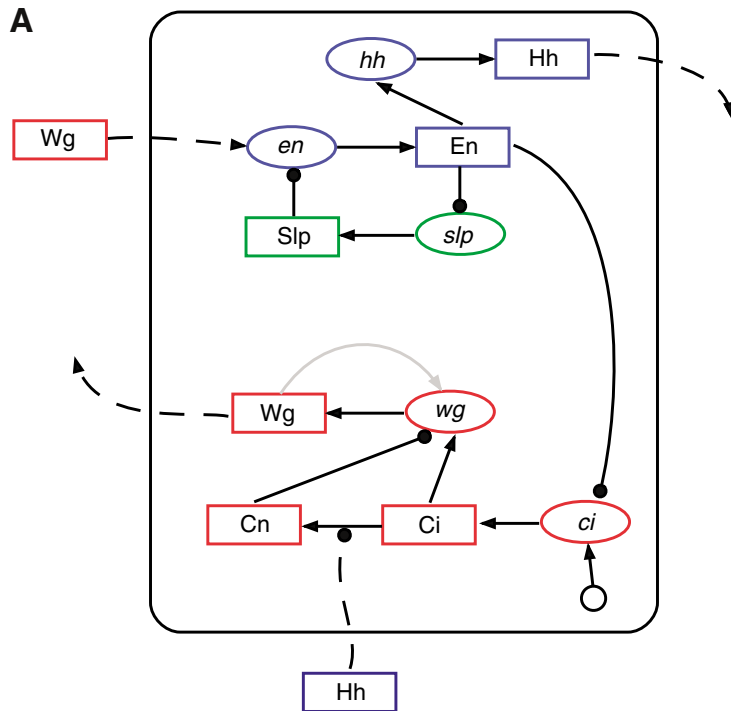
Biological topology: high score, not highest.

Excluding direct auto-regulation on E and S

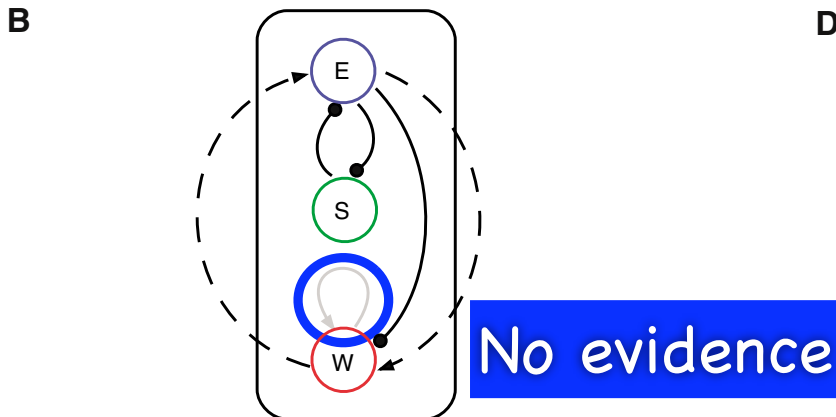




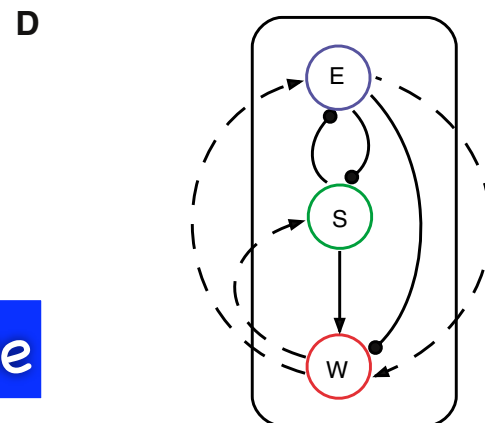
# A twist on polarity models



Evidence  
for *Slp* →  
*Wg*  
regulation!

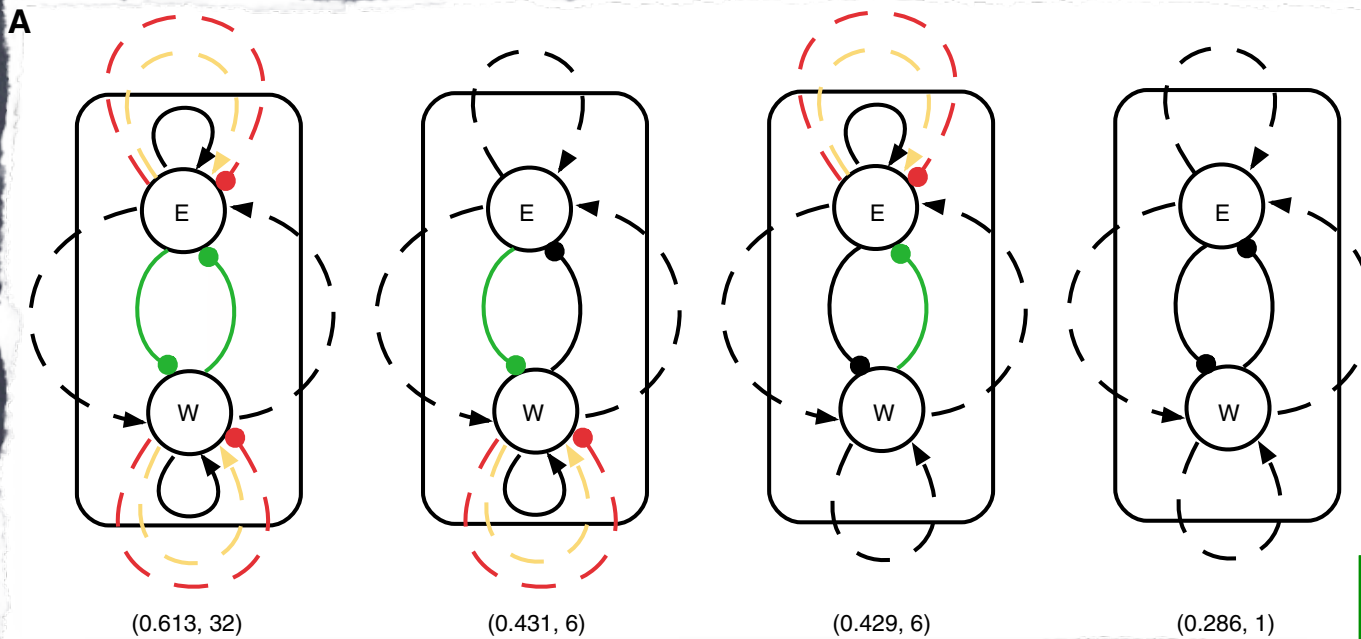


No evidence



Top ranking  
topology with  
NO *Wg* auto-  
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