

# Introduction to Random Boolean Networks

**Carlos Gershenson**

Centrum Leo Apostel, Vrije Universiteit Brussel.

Krijgskundestraat 33 B-1160 Brussel, Belgium

[cgershen@vub.ac.be](mailto:cgershen@vub.ac.be)

<http://homepages.vub.ac.be/~cgershen/rbn/tut>

<http://rbn.sourceforge.net>

# Topics (1)

- Introduction
- Classical Model (Kauffman)
  - Order, Chaos, and the Edge
    - Phase transitions in RBNs
  - Explorations
    - Attractor lengths
    - Convergence
  - Multi-valued Networks
  - Topologies
  - RBN Control

# Topics (2)

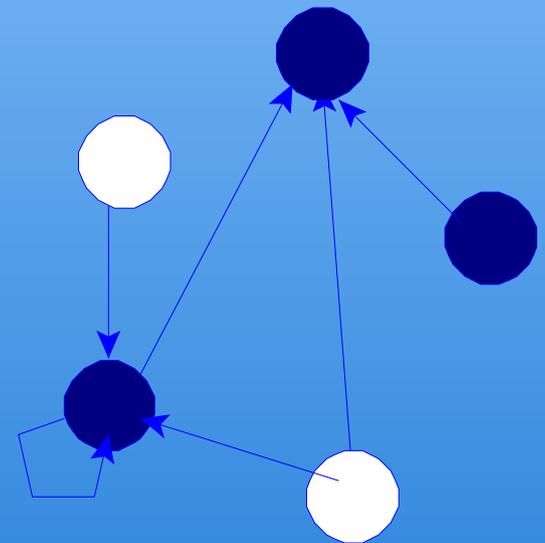
- Different Updating Schemes
  - Asynchronous RBNs
    - Rhythmic ARBNs
  - Deterministic Asynchronous RBNs
    - Thomas' ARBNs
  - Mixed-context RBNs
  - Classification of RBNs
- Applications
- Tools
- Future Lines of Research
- Conclusions

# Introduction

- RBNs originally models of genetic regulatory networks (Kauffman, 1969; Kauffman, 1993)
- Random connectivity and functionality
  - Useful with very complex systems
- Possibility to understand holistically living processes (e.g. for disease treatment)
- Possibility to explore *possibilities* of living and computational systems.

# Classical Model

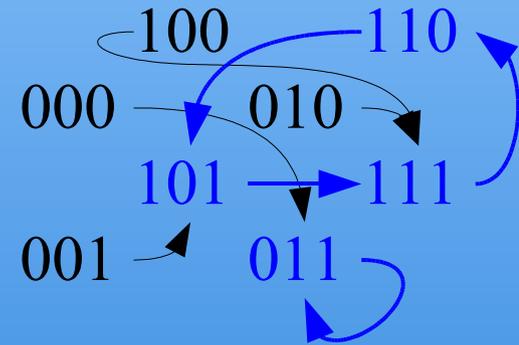
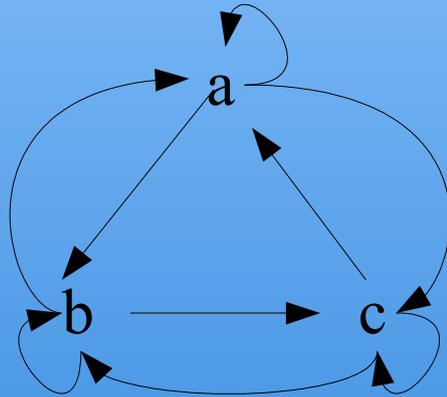
- Generalization of boolean CA
- $N$  boolean nodes,  $K$  connections per node
- Connectivity and logical functions generated randomly
- Synchronous updating



# Example

abc(t)	abc(t+1)
000	011
001	101
010	111
011	011
100	111
101	111
110	101
111	110

$N=3, K=3$



- Finite states ( $2^N$ )  $\Rightarrow$  **attractors** (dissipative system)
  - Point and cycle attractors

# Computational “Difficulties”

- Practically infinite possible networks
  - $2^{2^K}$  possible logic functions per node
  - $N!/(N-K)!$  possible ordered combinations per node

$$\text{possible nets} = \left( \frac{2^{2^K} N!}{(N-K)!} \right)^N$$

- though many equivalent (Harvey and Bossomaier, 1997)
- Can extract general properties with statistical samples
  - All possible initial states but small networks OR
  - Large networks but few initial states

# Order, Chaos, and the Edge (1)

- Neighbouring nodes in lattice
  - If changing, green; if static, red
  - Order: few green “islands”, surrounded by a red “frozen sea”
  - Chaos: green sea of changes, typically with red stable islands
  - Edge: green sea breaks into green islands, and the red islands join and percolate through the lattice (Kauffman, 2000, pp. 166-167)

# Order, Chaos, and the Edge (2)

- Network stability
  - “sensitivity to initial conditions”
  - “damage spreading”
  - “robustness to perturbations”
- Make a change in a state, connection, or rule, and see how changes affect the “normal” behaviour
  - Order: “Perturbed” network goes to the same dynamical path as “normal” net. Changes stay in green islands, damage does not spread

# Order, Chaos, and the Edge (2.5)

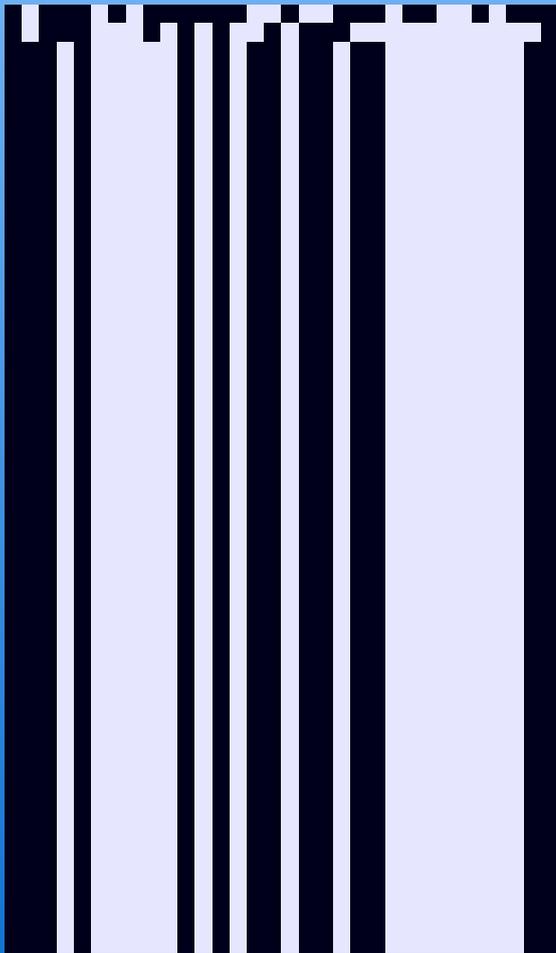
- Chaos: changes propagate, high sensitivity. Damage percolates through green sea
- Edge: changes can propagate, but not necessarily through all the network (Kauffman, 2000, pp. 168-170)

# Order, Chaos, and the Edge (3)

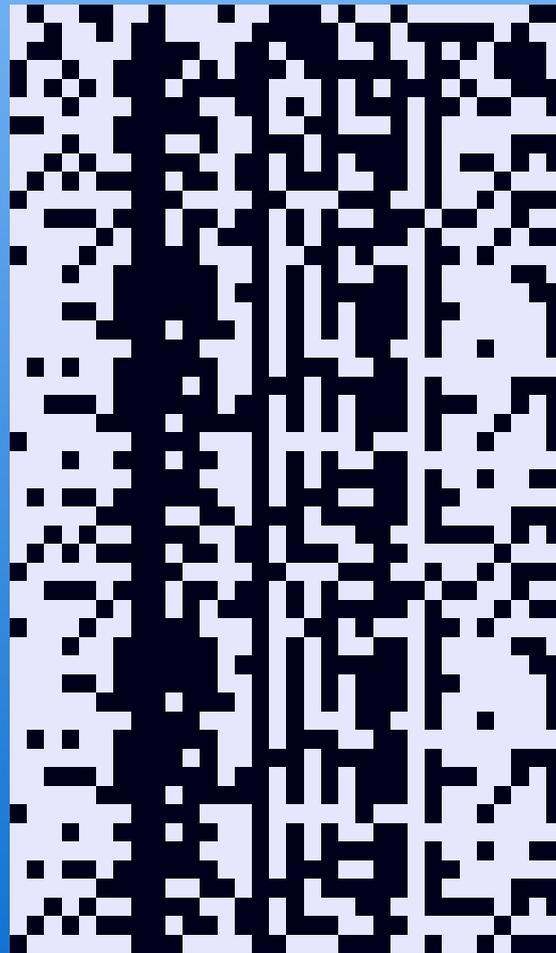
- Convergence vs. Divergence of Trajectories
  - Order: Similar similar states tend to converge to the same state
  - Chaos: similar states tend to diverge
  - Edge: nearby states tend to lie on trajectories that neither converge nor diverge in state space  
(Kauffman, 2000, p. 171)

**Life and Computation at the Edge of Chaos  
(Langton, 1990; Kauffman, 1993; 2000)**

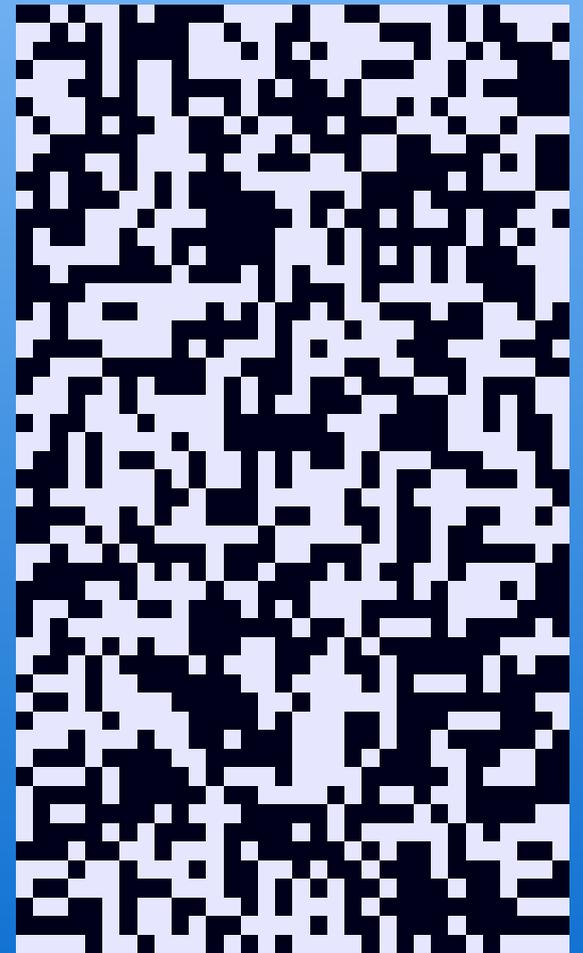
# Phase Transitions in RBNs



Ordered



Edge



Chaos

# Derrida's Annealed Approximation (1)

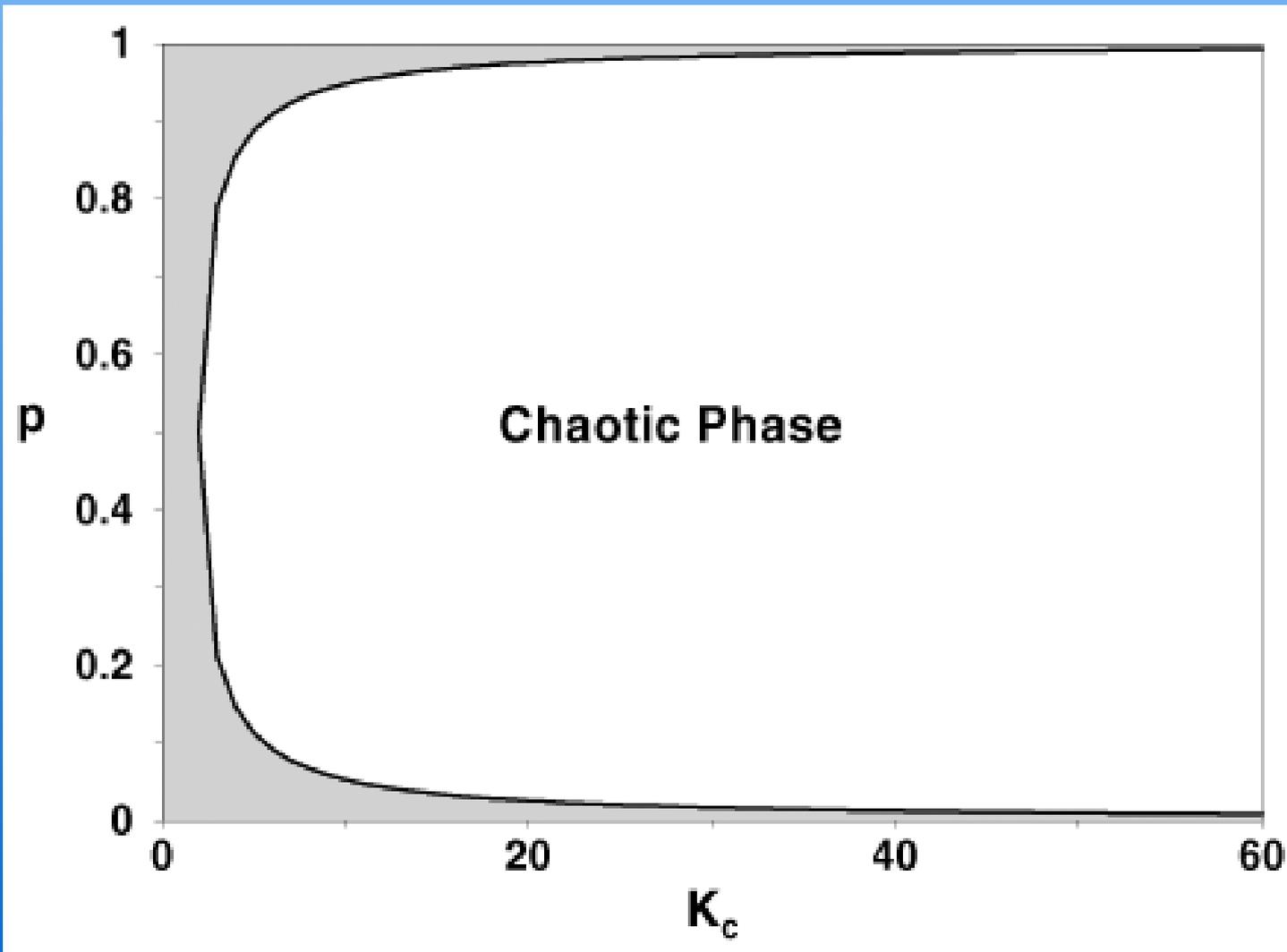
- Phase transition when  $K=2$  (Derrida and Pomeau, 1986)
- Also generalized for mean  $K$  and probability  $p$
- Measure overlap of state at  $t$  with state at  $t+1$  using normalized Hamming distance:

$$H(A, B) = \frac{1}{n} \sum_i^n |a_i - b_i|$$

- Then choose new rules and connections
- One dimensional map
  - At  $p=0.5$ , converges to 0 when  $K < 2$  (ordered)
  - when  $K > 2$ , divergence (chaos), critical phase  $K=2$

# Derrida's Annealed Approximation (2)

Critical  $K$



$$\langle K \rangle = \frac{1}{2p(1-p)}$$

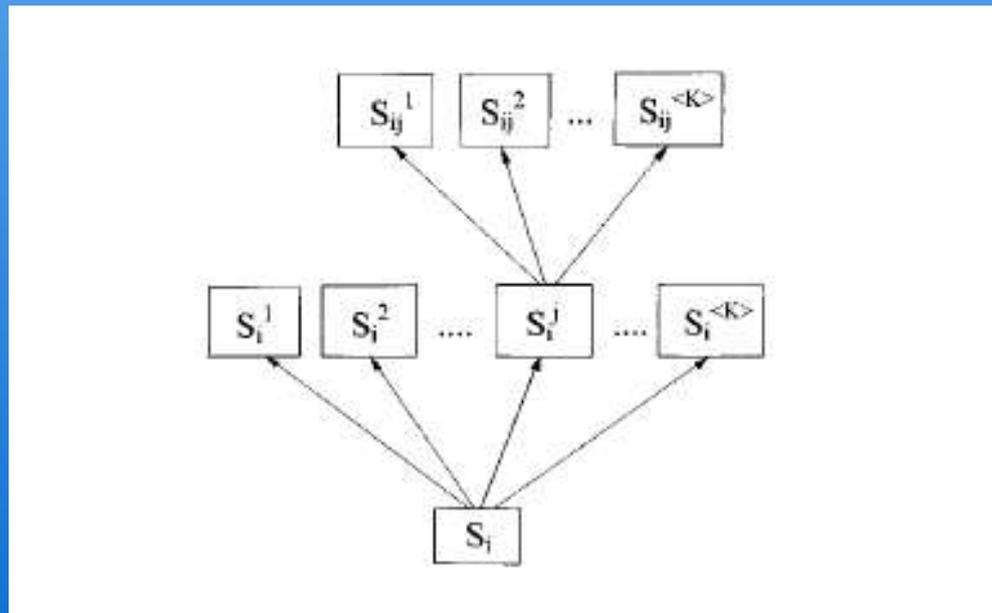
(Aldana, 2003)  
© Elsevier

# A Simpler Analytical Determination (1)

- Damage spreading when single nodes are perturbed (Luque and Solé, 1997b)
- Consider trees of nodes that can affect the state of other nodes in time
- As a node has more connections, there will be an increase in the probability that a damage in a single node ( $0 \rightarrow 1$  or  $1 \rightarrow 0$ ) will percolate through the network.

# A Simpler Analytical Determination (2)

- Let us focus only in one node  $i$  at time  $t$ , and a node  $j$  of the several  $i$  can affect at time  $t + 1$ . There is a probability  $p$  that  $j$  will be one, and a damage in  $i$  will modify  $j$  towards one with probability  $1 - p$ . The complementary case is the same. Now, for  $K$  nodes, we could expect that at least one change will occur if  $\langle K \rangle 2p(1 - p) \geq 1$ , which leads to Derrida's result



- This method can be also used for other types of networks

# Lyapunov exponents in RBNs

- Using the concept of boolean derivative (Luque and Solé, 2000)

$$\lambda = \log(2 p(1-p) K)$$

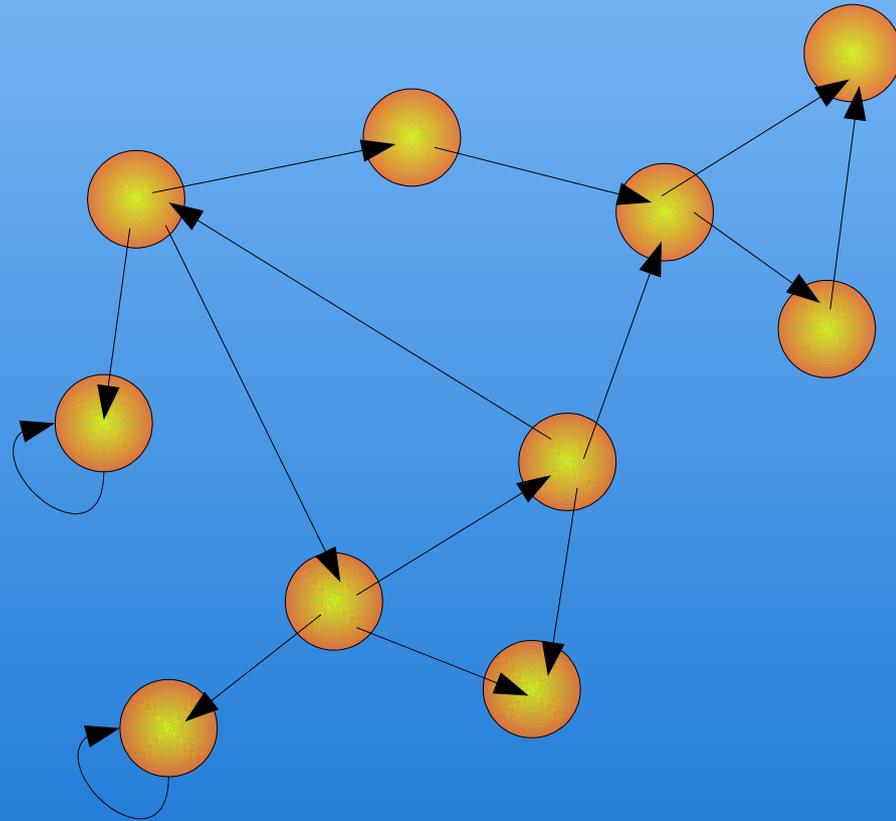
- where  $\lambda < 0$  represents the ordered phase,  $\lambda > 0$  the chaotic phase, and  $\lambda = 0$  the critical phase.
- **Beware:** Very high standard deviations!
- Theory can differ from practice...

# Explorations of the Classical Model

- E.g. number and length of attractors, sizes and distributions of their basins, and their dependence on different parameters ( $N$ ,  $K$ ,  $p$ , or topology) (Wuensche, 1997; Aldana et al., 2003)
- Analytical or statistical?

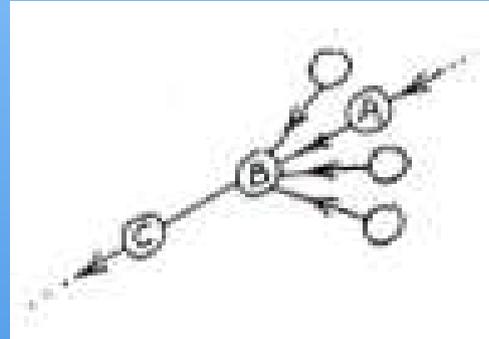
# Node Structure

- Descendants
  - Ancestors
  - Linkage loops
  - Linkage trees
- More connections,  
more loops, less  
stability



# State Space Structure

- *A predecessor* of B
- *C successor* of B



(Wuensche, 1998)

- Only one successor  $\Rightarrow$  CRBNs dissipative systems
- *In-degree*: number of predecessors
- *Garden-of-Eden* states: in-degree=0
- *Transient*: trajectory towards attractor

# Attractor Lengths (1)

- Analytic solutions of RBNs for  $K = 1$  (Flyvbjerg and Kjaer, 1988), and for  $K = N$  (Derrida and Flyvbjerg, 1987), but not for general case
- Statistical studies ( $p=0.5$ ) (Kauffman, 1969; 1993; Bastolla and Parisi, 1998; Aldana et al., 2003; ...)
  - $K=1$  probability of having long attractors decreases exponentially with  $N$ . Avg. number of cycles seems to be independent of  $N$ . The median lengths of state cycles are of order  $\sqrt{N/2}$ .

# Attractor Lengths (2)

- $K \geq N$ , average length of attractors and the transient times required to reach them grow exponentially (numerical investigations only of small networks). Typical cycle length grows proportional to  $2^{N/2}$ .
- $K = 2$ , (critical phase), both typical attractor lengths and average number of attractors grow algebraically with  $N$ .
  - $\sqrt{N}$ ? - undersampling (Kauffman, 1969; Kauffman, 1993; Bastolla and Parisi, 1998)
  - $N$ ? - small networks (Bilke and Sjunnesson, 2002; Gershenson, 2002)
  - Needs more research

# Convergence (1)

- Measured with  $G$ -density, in-degree frequency distribution (histogram), etc. (Wuensche, 1998).
- **ordered** phase, very high  $G$ -density, high in-degree frequency  $\Rightarrow$  high convergence, very short transient times. The basins of attraction are very compact, with many states flowing into few states.
- **critical** phase, in-degree distribution approximates a power-law. There is medium convergence.

# Convergence (2)

- **chaotic** phase, relatively lower  $G$ -density, and high frequency of low in-degrees. Basins of attraction are very elongated  $\Rightarrow$  very long average transient times. Low convergence.
- Other measures of convergence:
  - Walker’s “internal homogeneity” (Walker and Ashby, 1966)
  - Langton’s  $\lambda$  parameter (Langton, 1990)
  - Wuensche’s  $Z$  parameter (Wuensche, 1999).
    - Automatic classification of rule-space

# Multi-Valued Networks

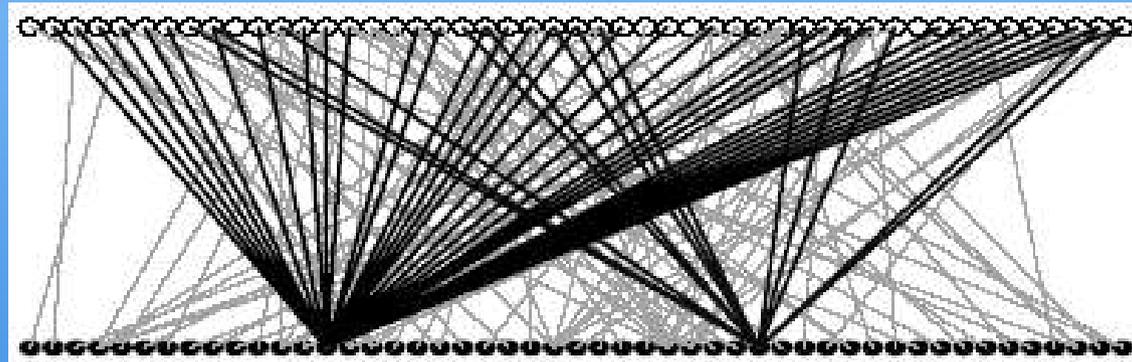
- More than 2 values per node (Solé et al., 2000; Luque and Ballesteros, 2004)
- results of Derrida are recovered for 2 values
- In nature, certain systems better described with more than two states. Particular models should go beyond the boolean idealization.
- However, for theoretical purposes, we could combine several boolean nodes to act as a multi-valued one
  - e.g. codifying in base two its state

# Topologies

- Topology can change drastically properties of RBNs (Oosawa and Savageau, 2002):
- more uniform rank distributions exhibit more and longer attractors and less entropy and mutual information (less correlation in their expression patterns)
- more skewed topologies exhibit less and shorter attractors and more entropy and mutual information
- A topology based on *E. coli* (scale-free), balances the parameters to avoid the disadvantages of the extreme topologies
- Most RBN studies use uniform rank distributions

# RBNs with scale-free topology (1)

- $P(k) = [\zeta(\gamma)k^\gamma]^{-1}, \gamma > 1$  (Aldana, 2003)

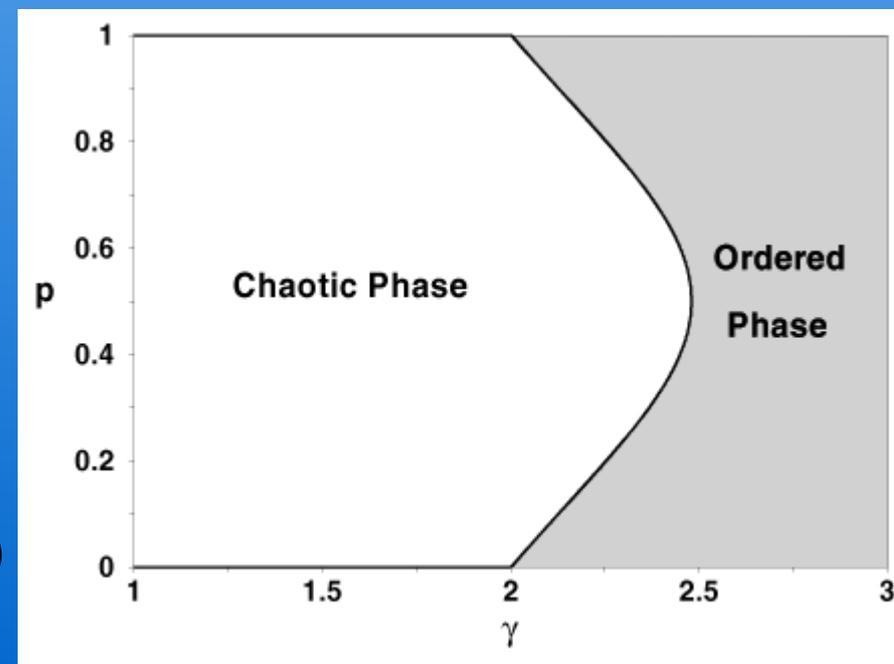


(Aldana, 2003)  
© Elsevier

- Using Derrida's method:

$$2p(1-p) \frac{\zeta(\gamma_c - 1)}{\zeta(\gamma_c)} = 1$$

(Aldana, 2003)  
© Elsevier



## RBNs with scale-free topology (2)

- The network properties at each phase (e.g. number and length of attractors, transient times) are analogous to homogeneous RBNs.
- **Evolvability** has more space in scale-free networks, since these can adapt even in the ordered regime, where changes in well-connected elements do propagate through the network.
- However, experimental evidence shows that most biological networks are scale free with exponent  $2 < \gamma < 2.5$ , i.e. near edge of chaos

# RBN Control (1)

- External inputs? (e.g. molecular clocks related to sunlight)
- Methods of chaos control have been successfully applied to chaotic RBNs (Luque and Solé, 1997a; 1998; Ballesteros and Luque, 2002)
- Use a periodic function to drive a very chaotic network into a stable pattern. If a periodic function determines the states of some nodes at some time, these will have a regularity that can spread through the rest of the network, developing into a global periodic pattern

# RBN Control (2)

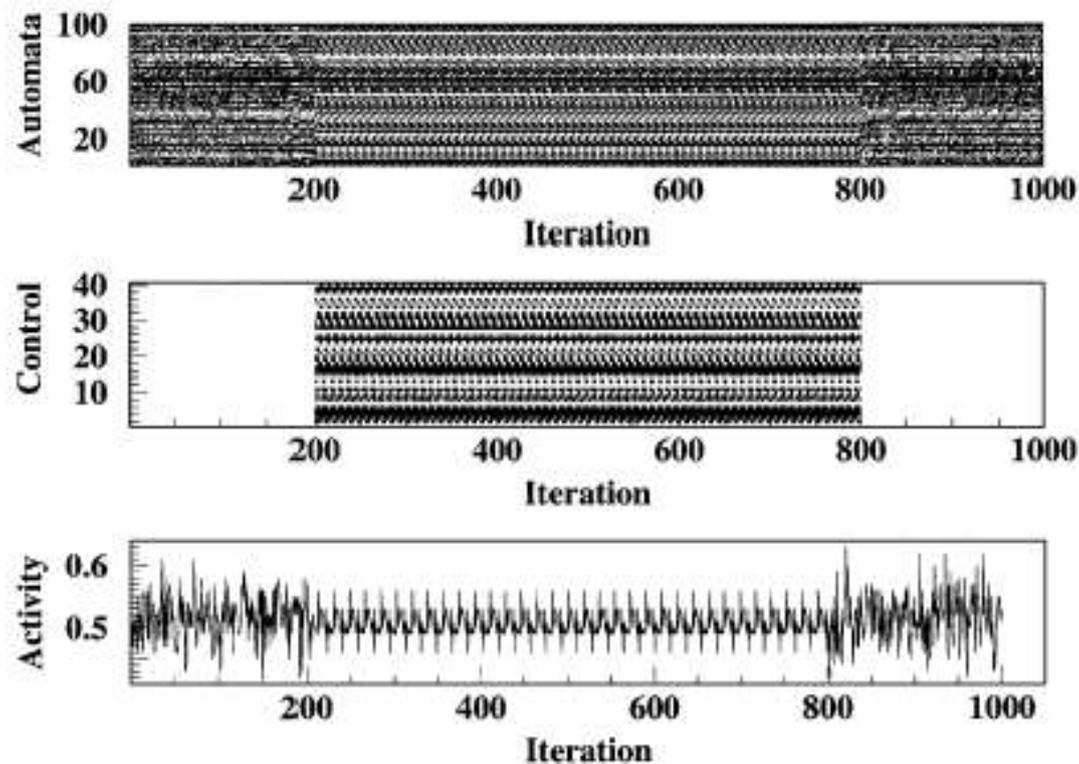
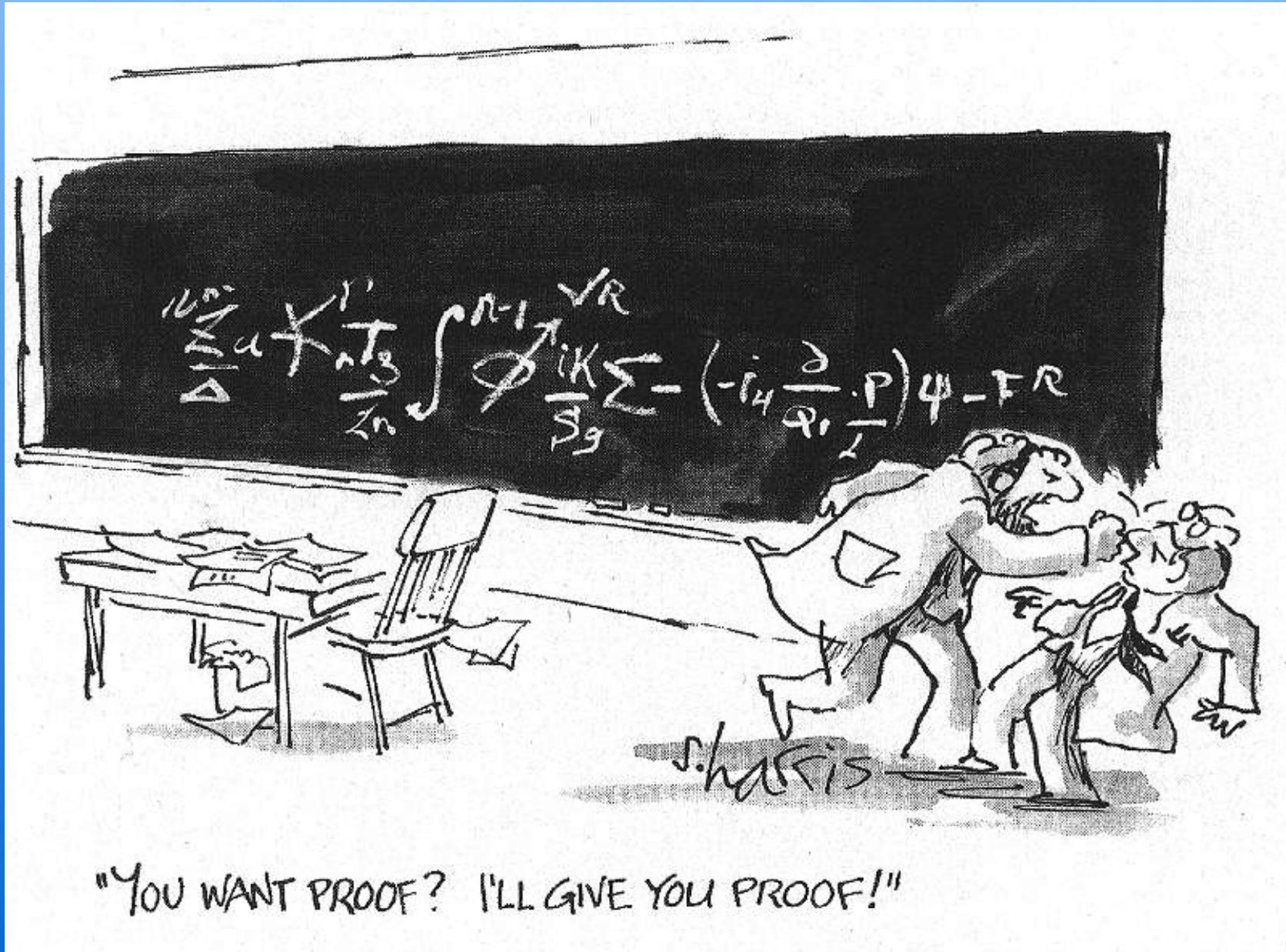


Fig. 6. RBN of size  $n = 40$  in ordered state ( $K = 2$ ,  $p = 0.5$ ) and period  $\tau = 9$  controlling the chaotic RBN described in Fig. 2. It can be seen how the first one induces an ordered behaviour of period 18 in the chaotic RBN.

## RBN Control (3)

- A high percentage of nodes should be controlled to achieve periodic behaviour. However, once we control a small chaotic network, we can use this to control a larger chaotic network, and this one to control an even larger one, and so on
- This shows that it is possible to *design* chaotic networks controlled by few external signals to force them into regular behaviour
- And scale-free chaotic RBNs? Could control only high-ranking nodes?

# Intermission...



# RBN attractors as cell types, lengths as replication time?

- “order for free” (Kauffman, 1969; Kauffman, 1993)
- Drawbacks:
  - Precise number of genes, junk DNA, redundancy
  - Attractor number linear or sqrt dependence?
  - Scale-free topology
  - Biased functions
  - **Genes do not march in step!**

# Updating Schemes

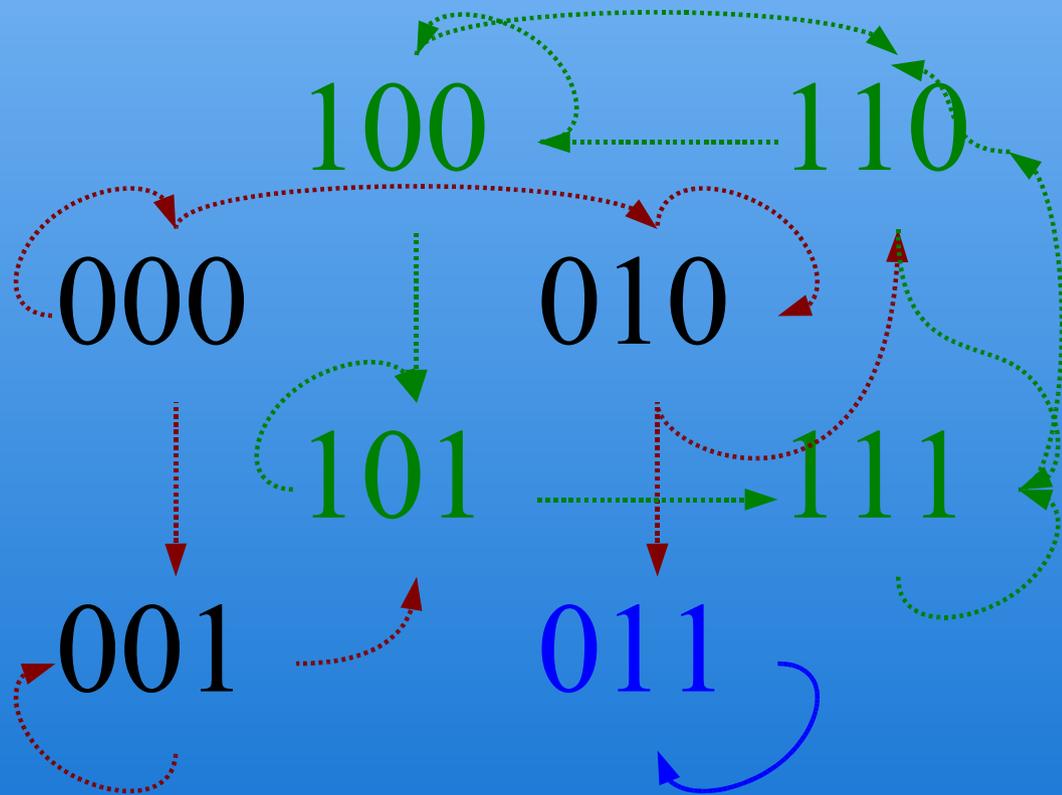
- Change of updating scheme can change drastically behaviour of a system
  - prisoner's dilemma (Huberman and Glance, 1993)
  - Conway's game of life (Bersini and Detours, 1994)

# Asynchronous RBNs (1)

- **ARBNs**: Pick up a node randomly, update network (Harvey and Bossomaier, 1997)
  - Asynchronous AND non-deterministic
- No cycle attractors
  - Point attractors (the same than CRBNs)
    - In theory, on average 1 per net. In practice, less.
  - “loose” attractors ( $K > 1$ )
- Different from **CRBN** behaviour
  - RBN useful genetic model???

# Example

$abc(t)$	$abc(t+1)$
000	011
001	101
010	111
011	011
100	111
101	111
110	101
111	110



# Rhythmic Asynchronous RBNs (1)

- If cells asynchronous, how could they achieve rhythm?
- Evolve RBNs and see... (Di Paolo, 2001)
- “Ring” topology (Rhofshagen and Di Paolo, 2004)
  - Only one linkage loop in pruned net
  - “Medusa” topologies found in yeast (Lee et al., 2002)
- What about more than one rhythmic attractor?



# Deterministic Asynchronous RBNs

- Cells not synchronous, but not purely stochastic
- **DARBNs**: Introduce parameters  $P_i$  and  $Q_i$  per node

$$P_i, Q_i \in \mathbb{N}, P_i > Q_i, P_{max} \geq P_i, Q_{max} \geq Q_i$$

- Update a node when  $\text{mod of time over } P_i == Q_i$ 
  - $P_i$  - period
  - $Q_i$  – translation
- If more than one node should be updated at a time step, do this in a sequential order
- Asynchronous, deterministic (Gershenson, 2002)

# Deterministic Generalized Asynchronous RBNs

- **DGARBNs:** Like DARBNs, but if more than one node should be updated, do this synchronously
- Semi-synchronous, deterministic (Gershenson, 2002)

# Generalized Asynchronous RBNs

- **GARBNs:** Like ARBNs, but select randomly nodes to update synchronously
- Semi-synchronous, non-deterministic (Gershenson, 2002)



# RBNs and Updating Schemes

- Many properties change drastically (Gershenson, 2002)
- All RBNs share point attractors, but basins can change
- More difference in attractor length due to determinism than synchronicity
- All have similar “edge of chaos” (Gershenson, 2004a,b)
- All perform *complexity reduction* (Gershenson, 2004b)
  - Including loose attractors
- Can map any deterministic RBN into a CRBN (Gershenson, 2002)
  - Encode in base two the periods, add new nodes

# Thomas' ARBNs

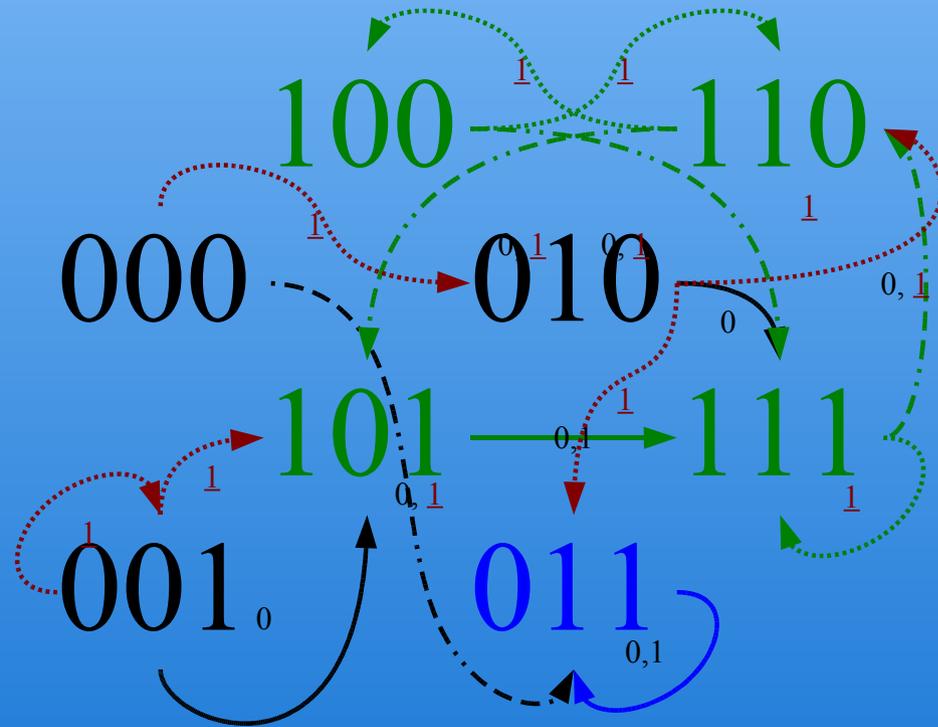
- ARBNs using delays (deterministic or stochastic) (Thomas, 1973; Thomas, 1978; Thomas, 1991)
- Used for analysis of precise networks, their circuits, and feedback loops. For ensembles???
- A positive feedback loop (direct or indirect autocatalysis) implies two point attractors
  - *Multistationarity*
- A negative feedback loop implies periodic behaviour, i.e. point or cycle attractors
  - *Homeostasis*

# Mixed-context RBNs (1)

- Sets  $\underline{P}$  and  $\underline{Q}$  ( $P_i$ 's and  $Q_i$ 's) as *context* of a network (Gershenson, Broekaert, and Aerts, 2003)
  - External factors can change precise updating periods
- Same DGARBN can have different behaviours with different contexts
- **MxRBNs**:  $M$  “pure” contexts, one chosen randomly at each  $R$  time steps
- Semi-synchronous, “quantum-like”
  - Non-determinism introduced in a very particular and controlled fashion

# Example

abc(t)	abc(t+1)
000	011
001	101
010	111
011	011
100	111
101	111
110	101
111	110

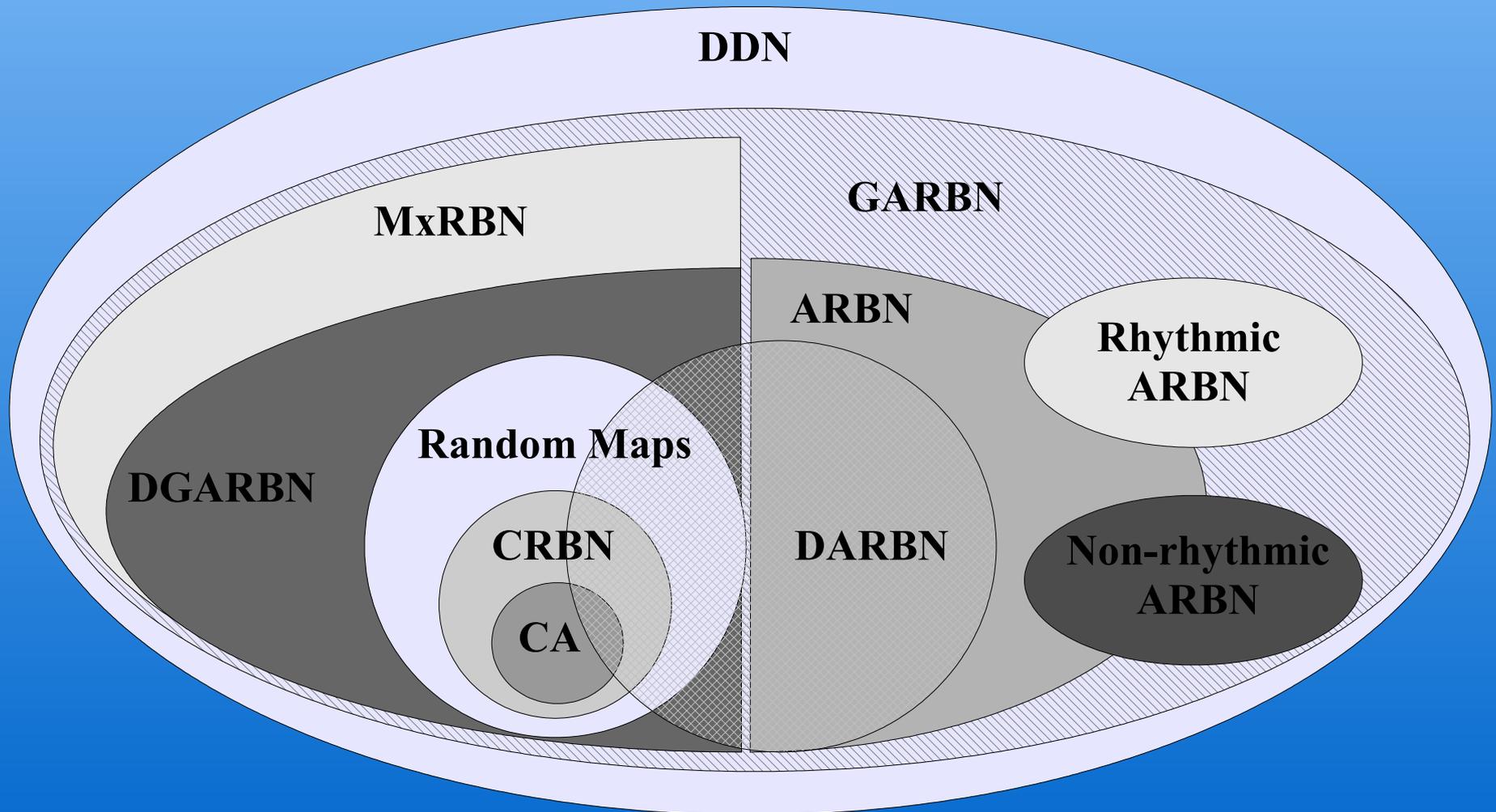


$P1's = \{1,1,2\}$ ,  $Q1's = \{0,0,0\}$   
 $P2's = \{2,1,1\}$ ,  $Q2's = \{0,0,0\}$

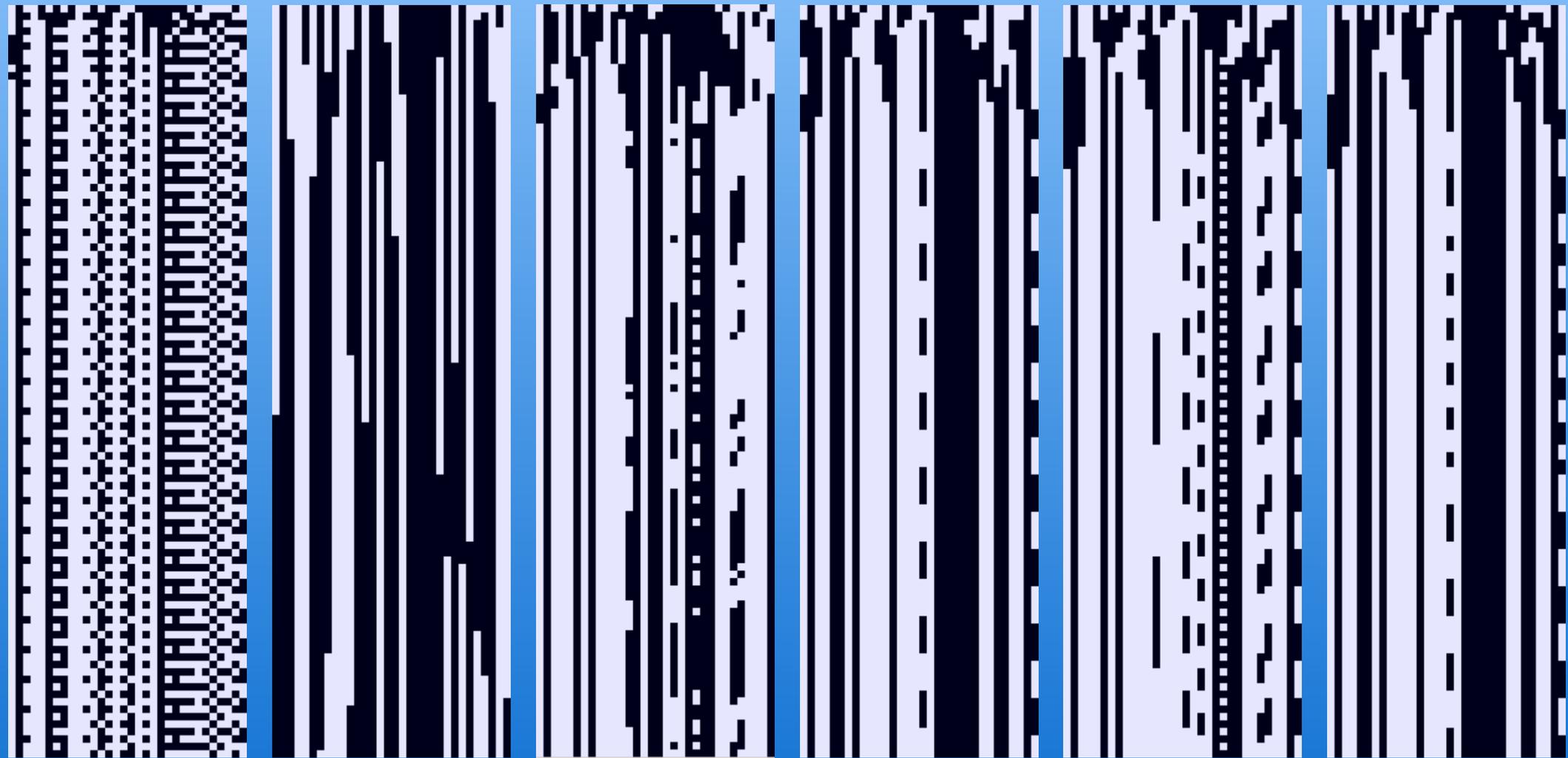
# How much non-determinism?

- GARBNs:  $N$  “coin flips” per time step
- ARBNs: one coin flip per time
- MxRBNs: one coin flip per  $R$  time steps
  - The higher the value of  $R$  and the lower number of  $M$  contexts, the less stochasticity there will be
- MxRBNs similar number of attractors than ARBNs and GARBNs, but much more complexity reduction (even more than CRBNs)
  - Information can be “thrown” into the context

# Classification of RBNs



# Dynamics Example



CRBN

ARBN

GARBN

DARBN

DGARBN

MxRBN

Same net & initial state.  $N = 32$ ,  $K = 2$ ,  $p = 0.5$ ,  $P_{max} = 5$ ,  $Q_{max} = 4$ ,  $M = 2$ ,  $R = 10$ .

# Applications

- Genetic regulatory networks
- Evolution and computation
- Neural networks (Huepe and Aldana, 2002)
- Social modelling (Shelling, 1971)
- Robotics (Quick et al., 2003)
- Music generation (Dorin, 2000).
- Mathematics
  - Cellular automata (von Neumann, 1966; Wolfram, 1986; Wuensche and Lesser, 1992)
  - Percolation theory (Stauffer, 1985)
- ...

# Genetic Regulatory Networks (1)

- Nodes as genes: “on-off” (activation), interaction via proteins (Kauffman, 1969)
- Generic properties in ensemble studies (Kauffman, 2004)
- Analysis and prediction of genomic interaction, data mining (Somogyi and Sniegowski, 1996; Somogyi et al., 1997; D’haeseleer et al., 1998)
- probabilistic boolean networks (PBNs): infer possible gene functionality from incomplete data (Shmulevich et al., 2002)

# Genetic Regulatory Networks (2)

- Experimental evidence of cell types as attractors of RBNs (Huang and Ingber, 2000)
  - Very strong correlation for some genes as a cell type is mechanically forced
  - Not all genes determine cell type (but e.g. metabolism)
- Continuous states GRN models (Glass and Kauffman, 1973; Kappler et al., 2002).
  - Use of differential equations in which gene interactions are incorporated as logical functions
  - no need for a clock to calculate the dynamics
  - Ensemble studies???

# Evolution and Computation (1)

- Evolvability is expected at the edge of chaos
- Network evolvability properties:
  - robustness, redundancy, degeneracy, modularity (Fernández and Solé, 2004)
- Life performs computations (Hopfield, 1994)
  - Understanding computation networks helps us to understand life and its possibilities
  - “How can computational networks be evolved?” close to “How could life evolve?”

# Evolution and Computation (2)

- Evolution of RBNs using genetic algorithms (Stern, 1999; Lemke et al., 2001)
- Evolvable hardware (Thompson, 1998)
  - Evolution of logical circuits in reconfigurable hardware
- Issues of evolvability also interesting for genetic algorithms, genetic programming, etc.
- ...

# Tools (1)

- **DDLab** (Andy Wuensche)
  - synchronous RBNs and CA, multi-valued networks
  - Dynamics and basins of attraction visualization
  - It includes a wide variety of measures, data, analysis and statistics
  - Very well documented, runs on most platforms.
  - <http://www.ddlab.com>
- **RBN Toolbox for Matlab** (Christian Schwarzer and Christof Teuscher)
  - Simulation and visualization of RBNs.
  - Different updating schemes, statistical functions, etc.
  - <http://www.teuscher.ch/rbntoolbox>

# Tools (2)

- **RBNLab** (Carlos Gershenson)
  - Simulation and visualization of RBNs with different updating schemes
  - Point, cycle, and loose attractors, other statistics...
  - Java, code and program at <http://rbn.sourceforge.net>
- **BN/PBN Toolbox for Matlab** (Harri Lähdesmäki and Ilya Shmulevich)
  - CRBNs and PBNs.
  - functions for simulating network dynamics, computing statistics (a lot), inferring networks from data, visualization and printing, intervention, membership testing of Boolean functions, etc.
  - <http://www2.mdanderson.org/app/ilya/PBN/PBN.htm>

# Future Lines of Research

- Ensemble approach (Kauffman, 2004)
- RBNs for data mining and GRN analysis
- Evolvability and adaptability at an abstract level
- Generalizations, combinations, and refinements of the different types of RBNs
  - e.g. scale-free multi-valued DGARBNs, etc
- General analytical solution for CRBNs
- ...

# Conclusions

- Tutorial only overview, but main topics covered
- RBNs interesting due to generality
  - Many conclusions with few assumptions
  - Illustrate order-complexity-chaos for non-physicists
- Which model is best? It depends...
- Theory vs. practice? Balance!
- An inviting research topic