



Logic-Based Modeling in Systems Biology

Alexander Bockmayr

LPNMR'09, Potsdam, 16 September 2009



- I. Systems biology
- II. Logic modeling of regulatory networks
 - A. Boolean logic
 - B. Multi-valued logic
- III. Logical analysis of network dynamics
- IV. Application: Bio-Logic



I. Systems biology



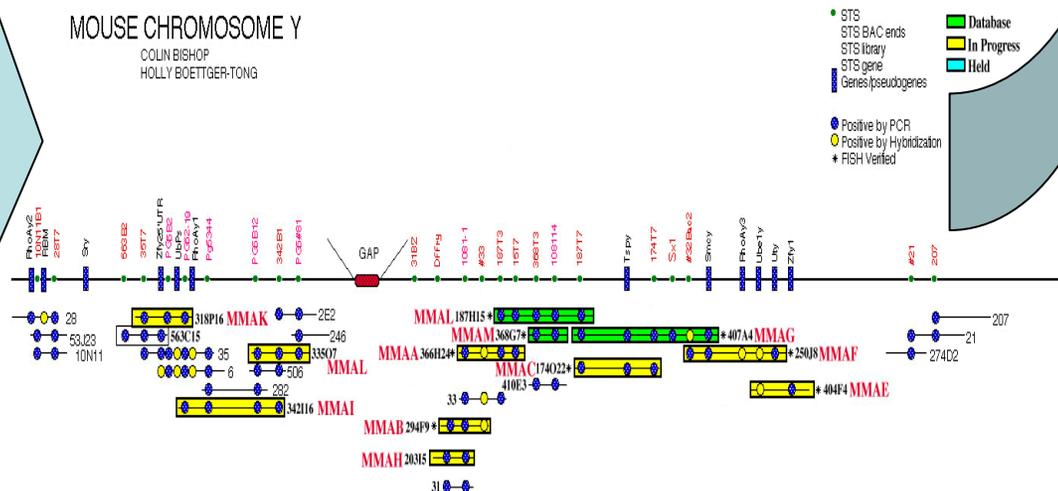
Molecular
biology



Systems
biology

MOUSE CHROMOSOME Y

COLIN BISHOP
HOLLY BOETTGER-TONG



Very active interdisciplinary research field



Various types of biological networks

- metabolic
- regulatory
- signaling, ...

Various modeling approaches

- continuous (ordinary/partial differential equations)
- stochastic (chemical master equation)
- discrete (logic, Petri nets, process calculi, ...)
- hybrid (continuous/stochastic, discrete/continuous)

Here: Logic-based discrete modeling of regulatory networks

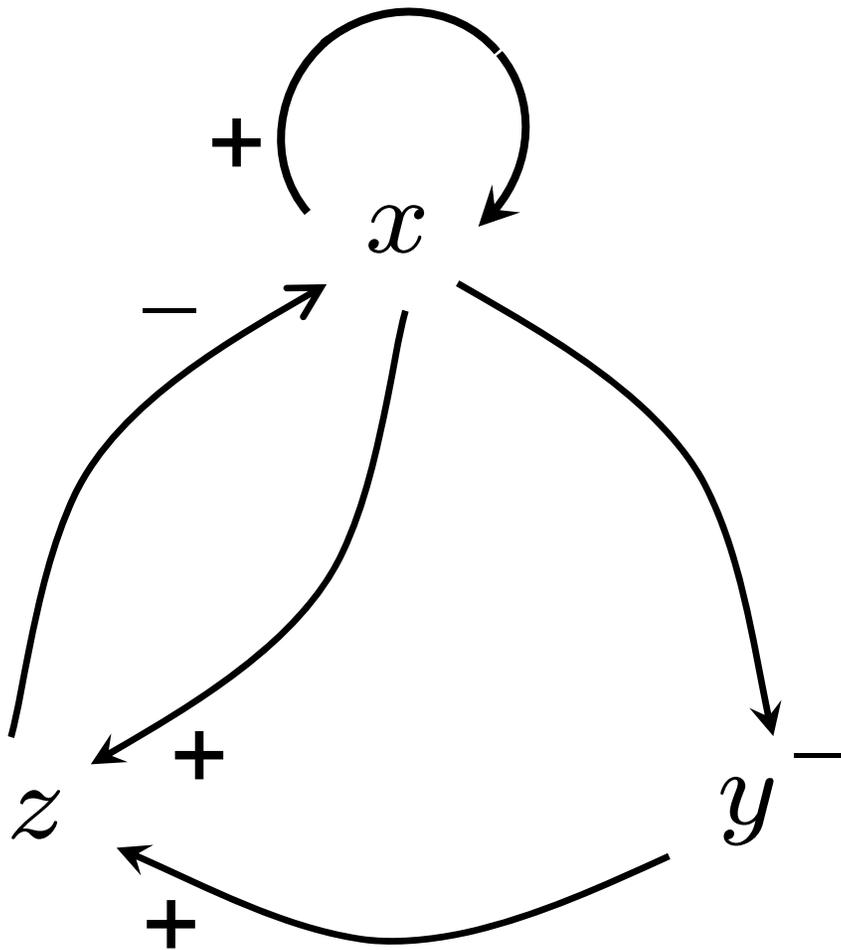


1. Logic modeling of the network **structure**
 - Boolean logic
 - Multi-valued logic
2. Logical analysis of the **dynamics**
 - Non-determinism
 - Temporal logic
 - Model checking



II. Logic modeling of regulatory networks

A) Boolean logic



Nodes

$$X_1, \dots, X_n \in \{0, 1\}$$

(component is active or not)

Arcs

Activation: $X_i \xleftarrow{+} X_j$

Inhibition: $X_i \xleftarrow{-} X_j$

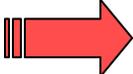


Sugita 61, Kauffman 69

1. Boolean variables $X_1, \dots, X_n \in \{0, 1\}$
2. Boolean mapping $F : \{0, 1\}^n \rightarrow \{0, 1\}^n$

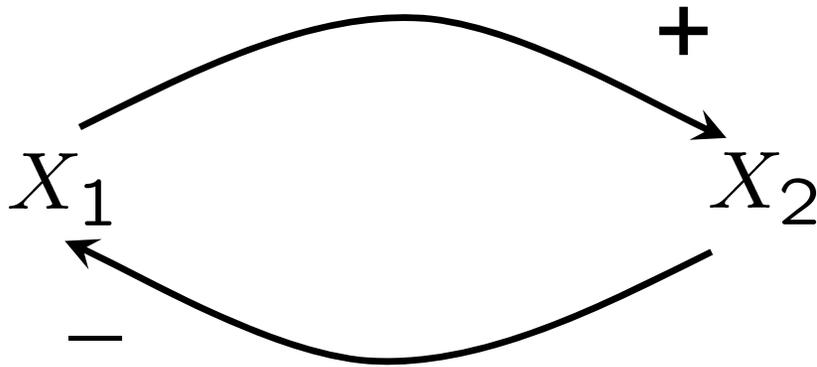
$F_i(X_1, \dots, X_n)$ describes how the next state of X_i depends on the current state of (X_1, \dots, X_n) .

$$X_i^{\rightarrow} = F_i(X_1, \dots, X_n)$$

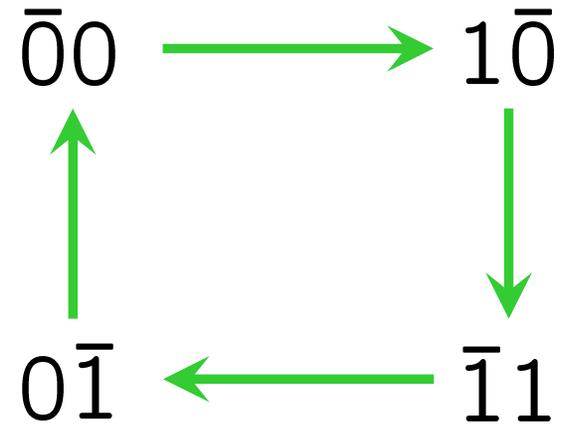
 discrete dynamics



Interaction graph



State transition graph



$$X_1, X_2 \in \{0, 1\}$$

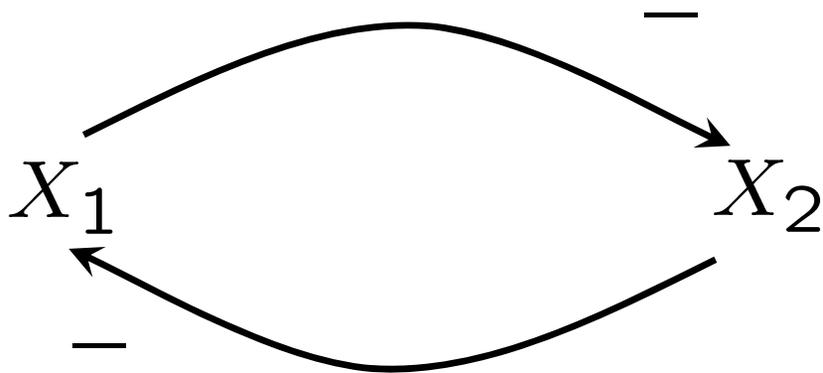
$$X_1^{\rightarrow} = F_1(X_1, X_2) = \neg X_2$$

$$X_2^{\rightarrow} = F_2(X_1, X_2) = X_1$$

X_1	X_2	X_1^{\rightarrow}	X_2^{\rightarrow}
0	0	1	0
0	1	0	0
1	0	1	1
1	1	0	1



Interaction graph



$$X_1, X_2 \in \{0, 1\}$$

$$X_1^{\rightarrow} = F_1(X_1, X_2) = \neg X_2$$

$$X_2^{\rightarrow} = F_2(X_1, X_2) = \neg X_1$$

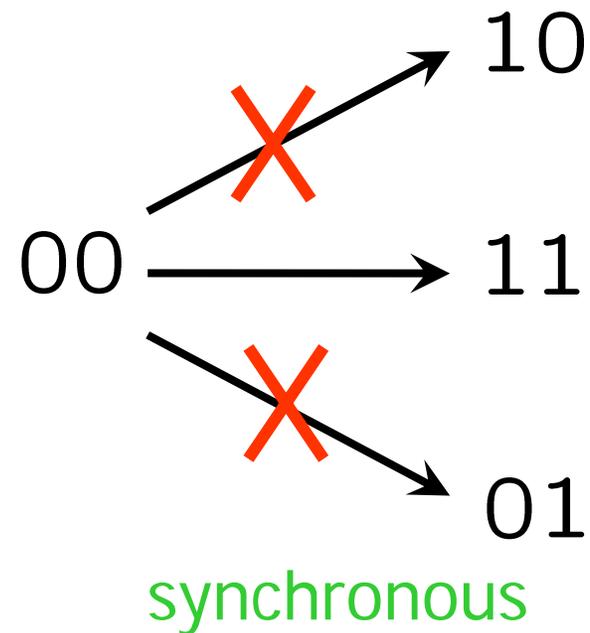
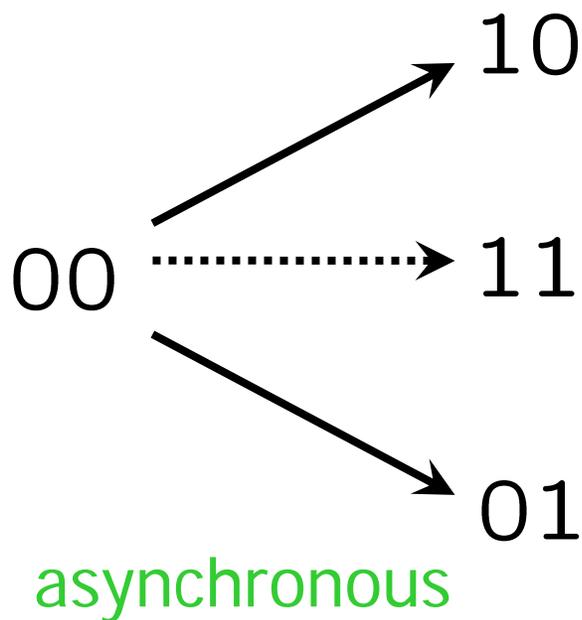
X_1	X_2	X_1^{\rightarrow}	X_2^{\rightarrow}
0	0	1	1
0	1	0	1
1	0	1	0
1	1	0	0



$$(X_1, X_2) = (0, 0), (X_1^{\rightarrow}, X_2^{\rightarrow}) = (1, 1)$$

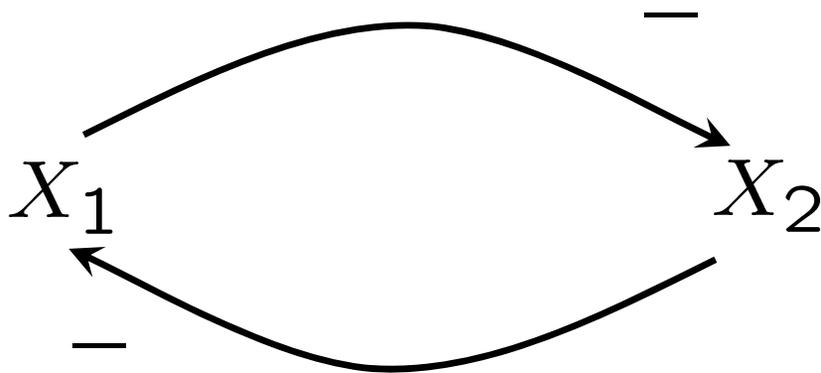
Thomas '73: Update only one variable at a time.

➡ **Nondeterminism:** Several successor states possible

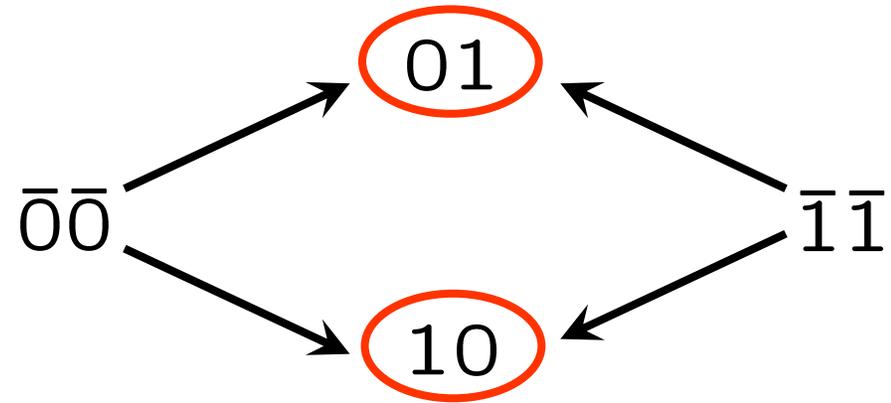




Interaction graph



State transition graph



$$X_1, X_2 \in \{0, 1\}$$

$$X_1^{\rightarrow} = F_1(X_1, X_2) = \neg X_2$$

$$X_2^{\rightarrow} = F_2(X_1, X_2) = \neg X_1$$

X_1	X_2	X_1^{\rightarrow}	X_2^{\rightarrow}
0	0	1	1
0	1	0	1
1	0	1	0
1	1	0	0



II. Logic modeling of regulatory networks

B) Multi-valued logic



If component j acts on n_j other components

⇒ (up to) n_j thresholds: $X_j \in D_j = \{0, \dots, n_j\}$

$X_j = k$: activity level of component j is above the k -th threshold and below the $(k+1)$ -th.

⇒ discrete update function

$$F : D_1 \times \dots \times D_n \rightarrow D_1 \times \dots \times D_n$$

$$X_i^{\rightarrow} = F_i(X_1, \dots, X_n) = F_i(X, \mathbf{K})$$

with discrete parameter vector \mathbf{K} .



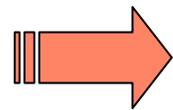
State

$$X = (X_1, \dots, X_n), \quad X_i \in \{0, \dots, n_i\}$$

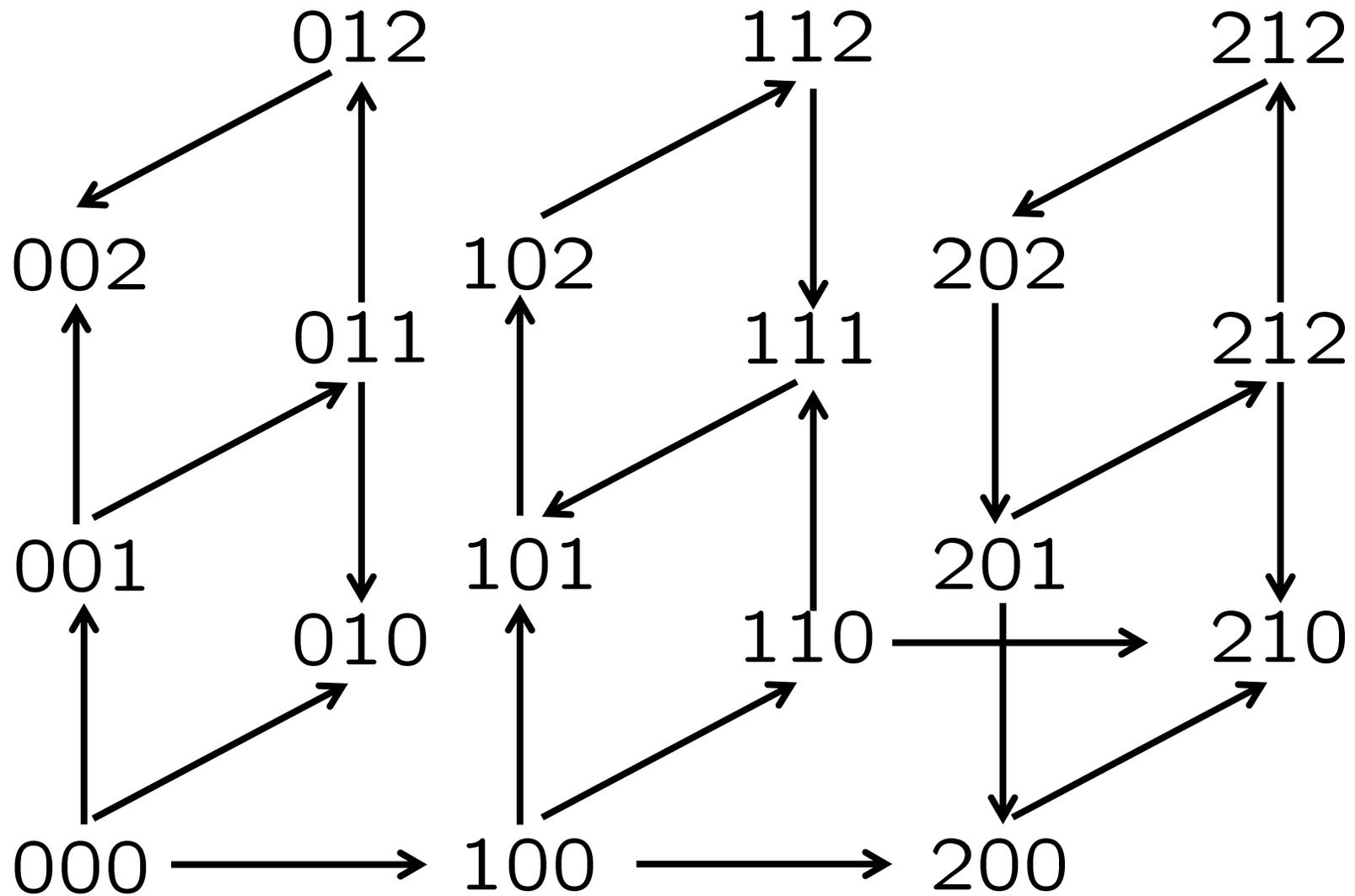
State transitions

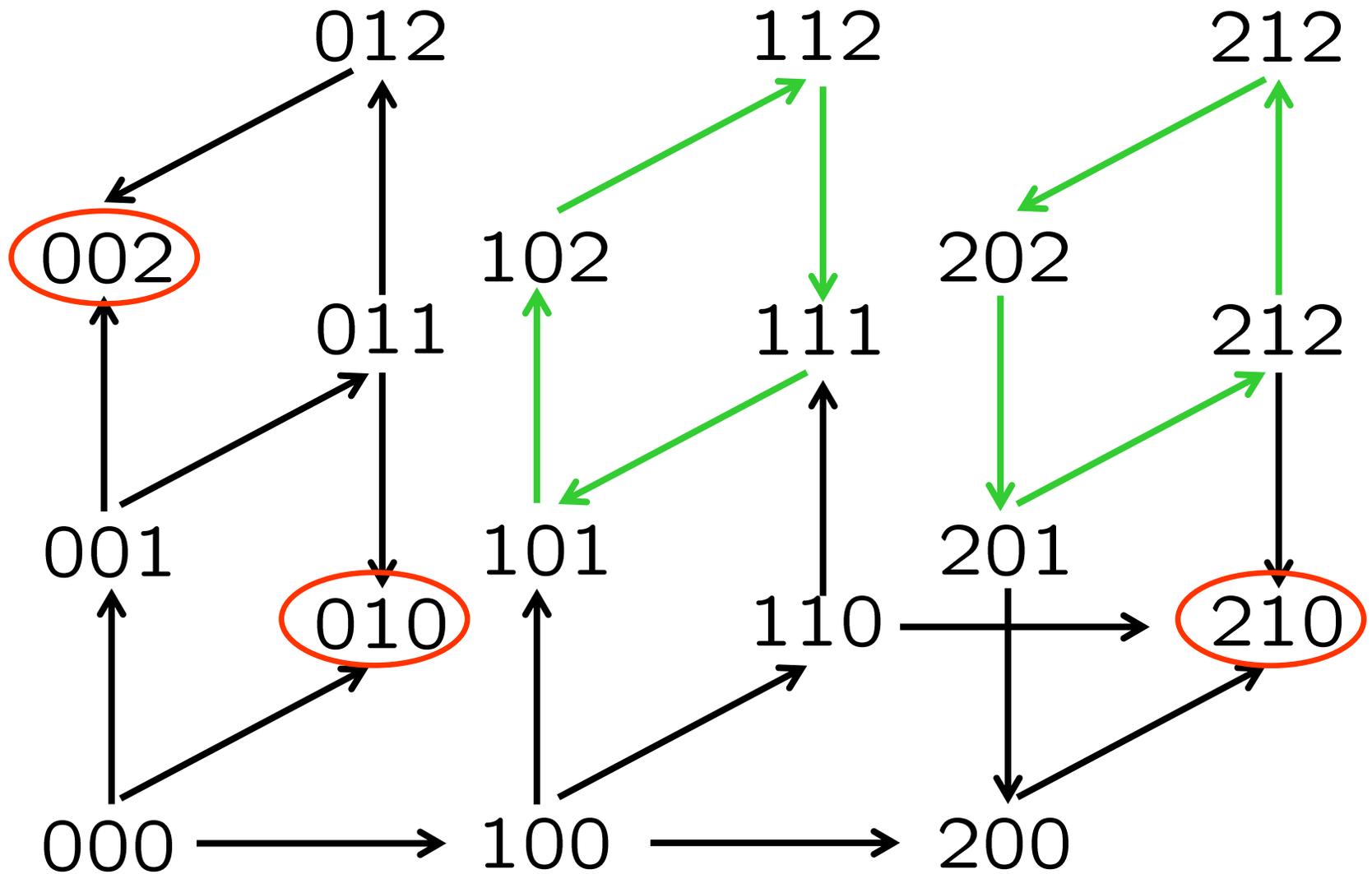
$$(X_1, \dots, X_n) \longrightarrow (X_1, \dots, X_i \pm 1, \dots, X_n)$$

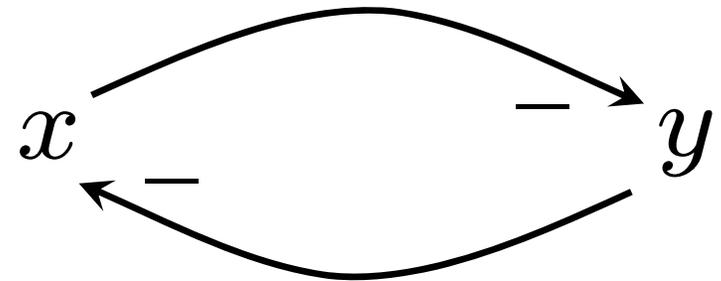
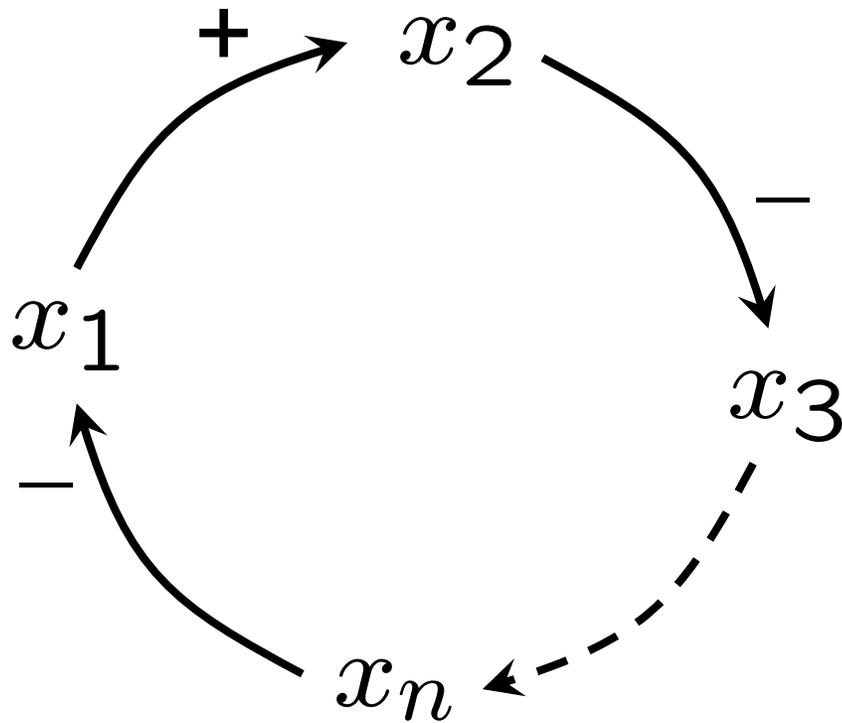
if $X_i^{\rightarrow} > X_i$ resp. $X_i^{\rightarrow} < X_i$



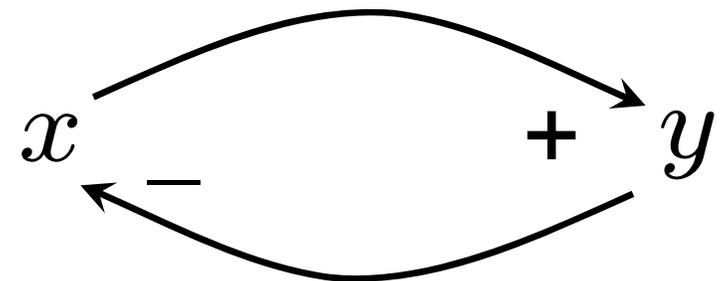
discrete non-deterministic dynamics







Positive 2-circuit



Negative 2-circuit

Sign of circuit =
Product of signs of arcs

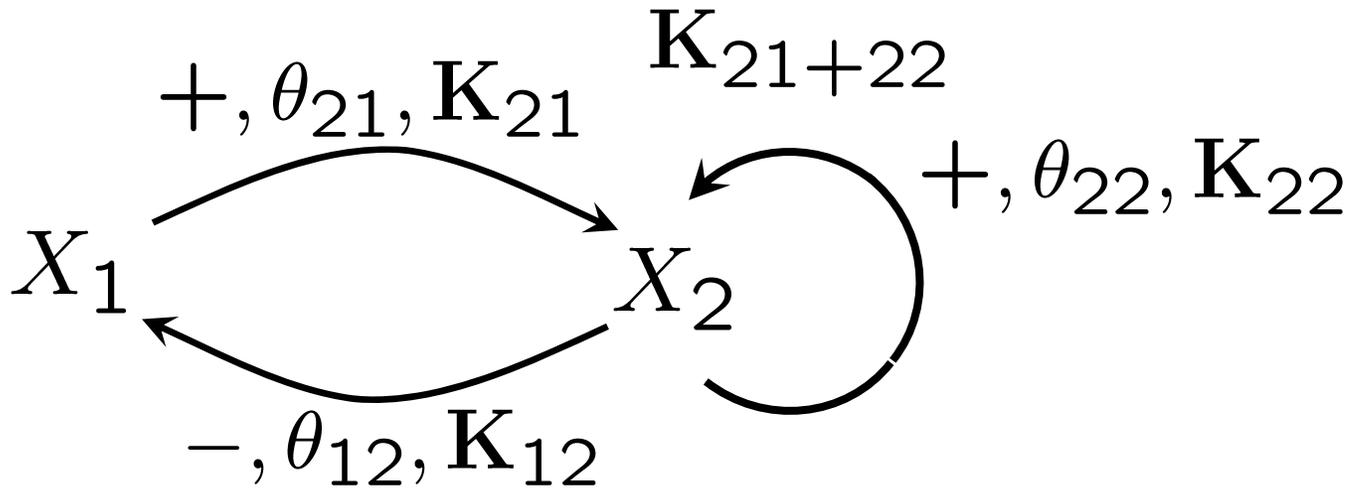


Thomas ´81

A positive circuit in the interaction graph is a necessary condition for multistationarity.

A negative circuit in the interaction graph is a necessary condition for stable periodic behavior.

[Proofs exist in various scenarios.]

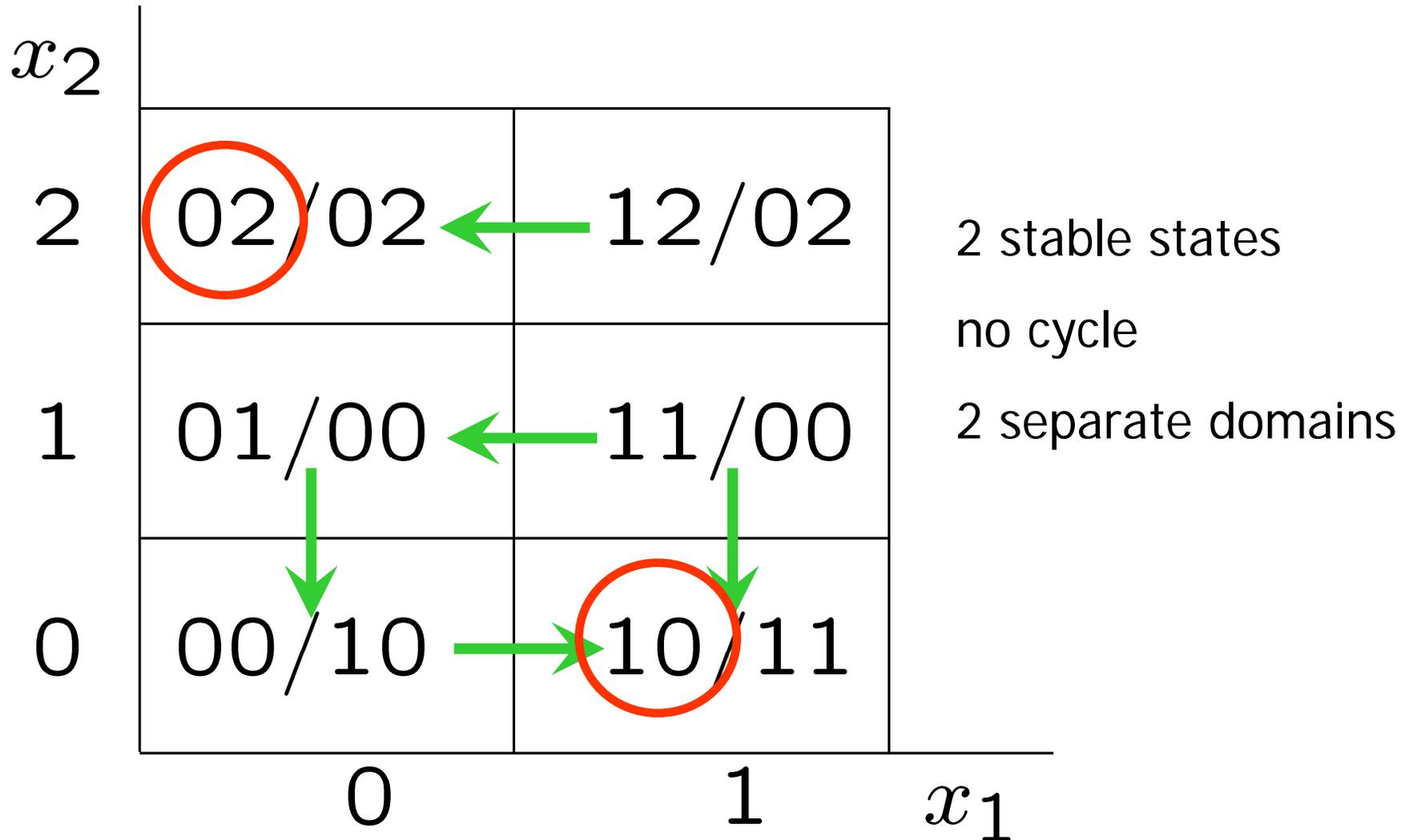


- $X_1 \in \{0, 1\}$
- $X_2 \in \{0, 1, 2\}$
- Assume $\theta_{12} < \theta_{22}$, i.e., when activated, X_2 acts first on X_1 , then on itself.

X_1	X_2	X_1^{\rightarrow}	X_2^{\rightarrow}
0	0	\mathbf{K}_{12}	0
0	1	0	0
0	2	0	\mathbf{K}_{22}
1	0	\mathbf{K}_{12}	\mathbf{K}_{21}
1	1	0	\mathbf{K}_{21}
1	2	0	\mathbf{K}_{21+22}

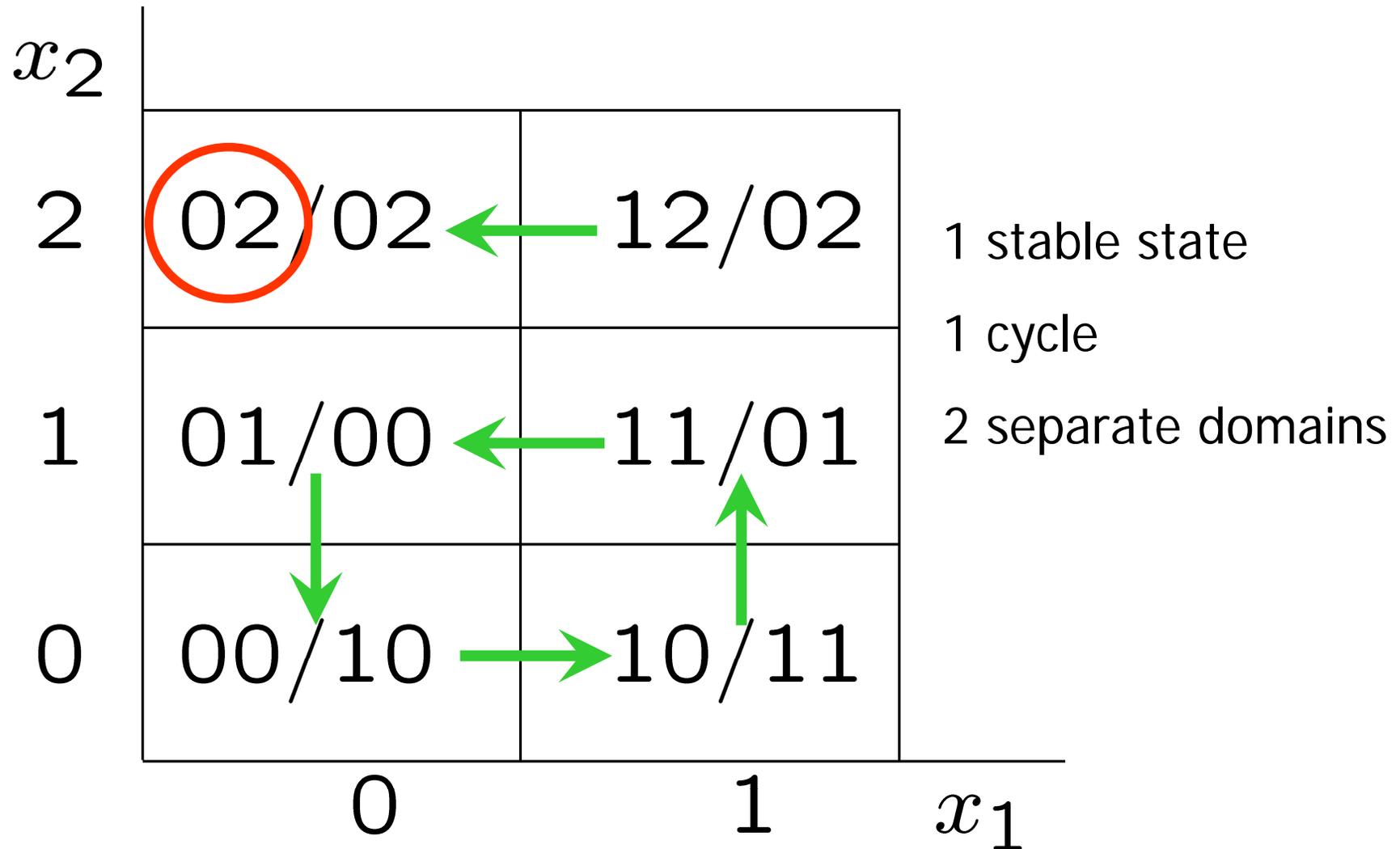


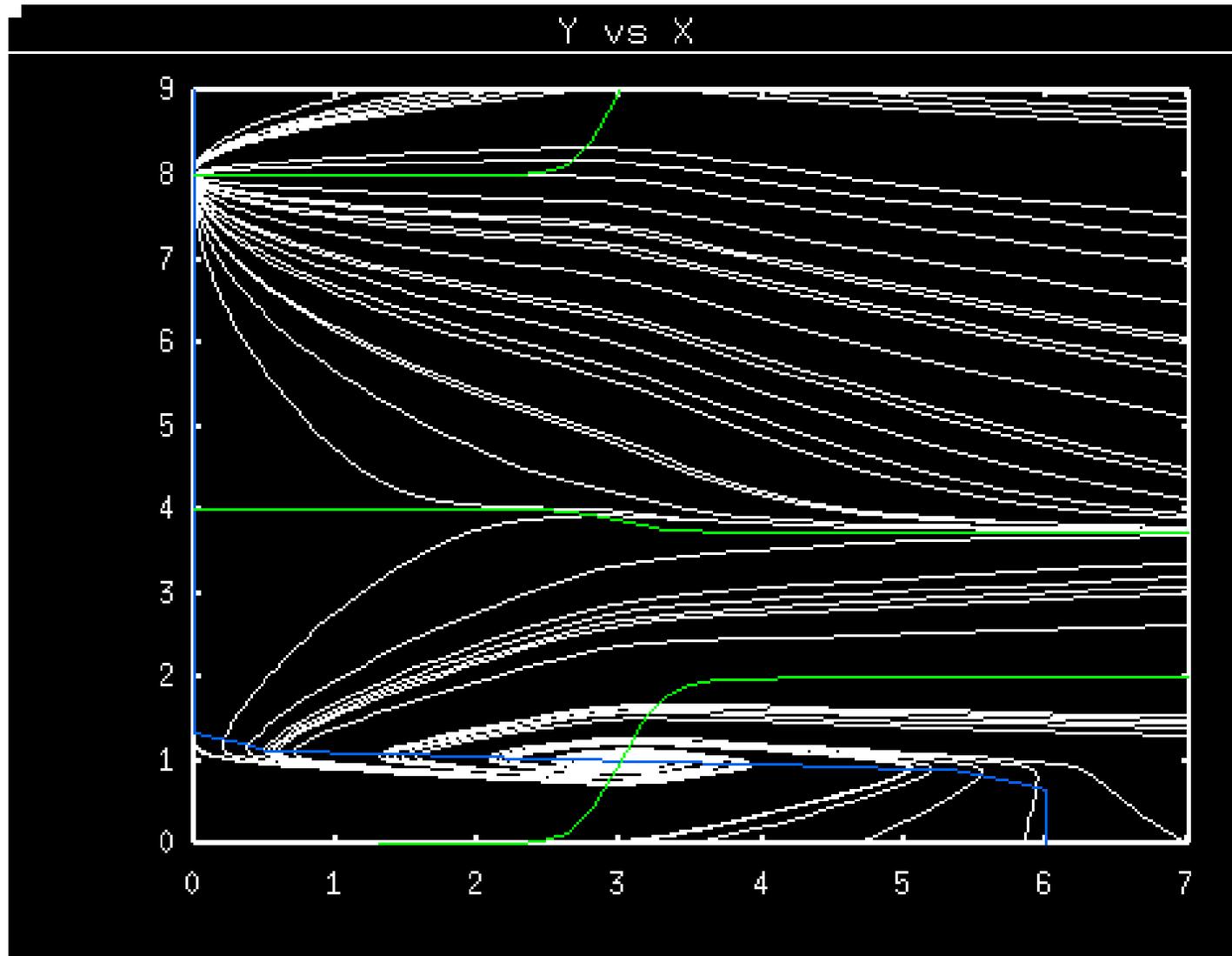
$$K_{12}=1, K_{21} = 0, K_{22} = K_{21+22} = 2$$





$$K_{12}=1, K_{21} = 1, K_{22} = K_{21+22} = 2$$





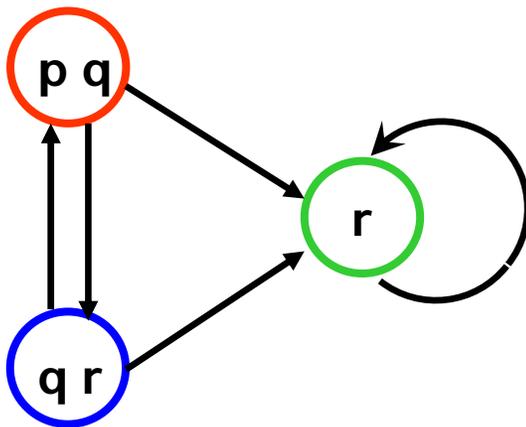


III. Logical analysis of the dynamics



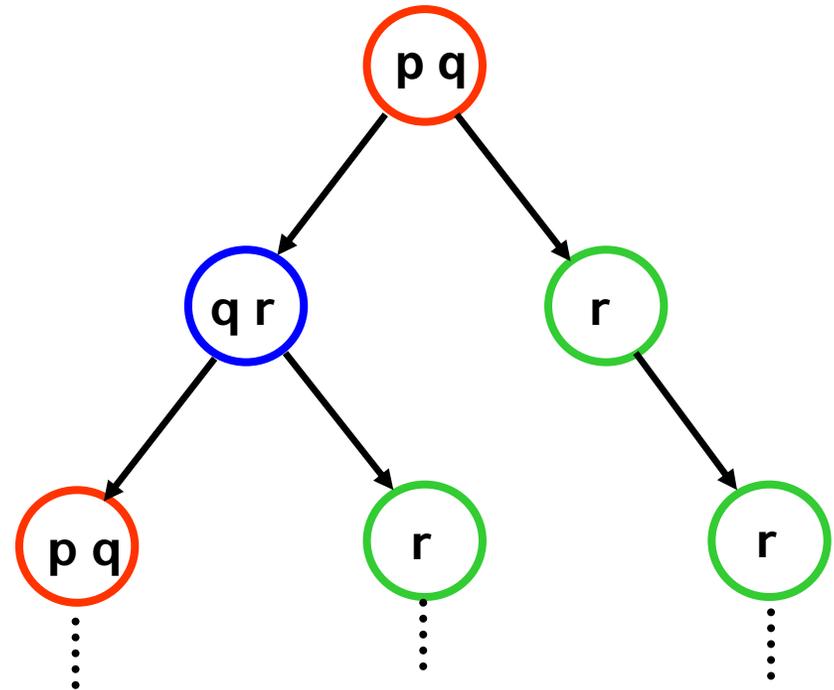
Clarke/Emerson and Sifakis 81

State transition graph
(Kripke model)



exponentially large

Infinite computation tree



 check properties expressed in some temporal logic.



Atomic formulae : p, q, r, \dots , e.g. $X_i = 1$

Linear time operators :

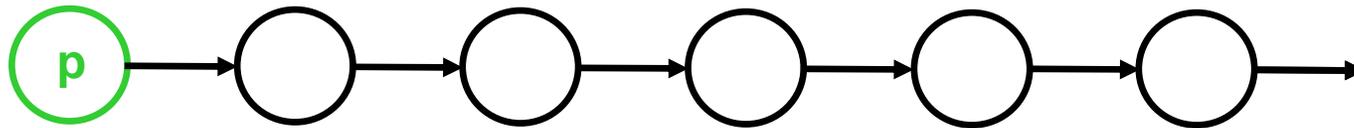
- X p : p holds next time
- F p : p holds sometimes in the future
- G p : p holds globally in the future
- p U q : p holds until q holds

Path quantifiers :

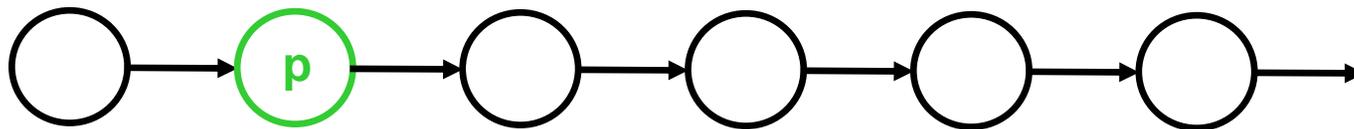
- A : for every path
- E : there exists a path



Now



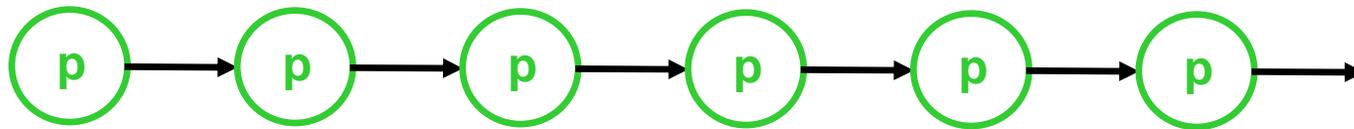
p



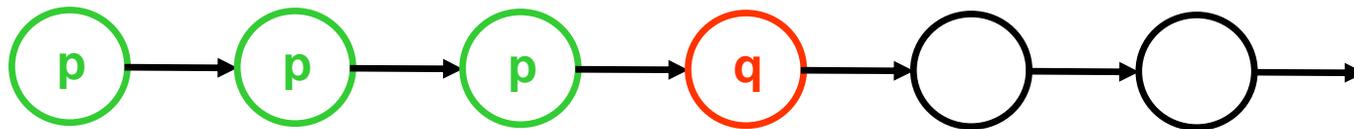
Xp



Fp



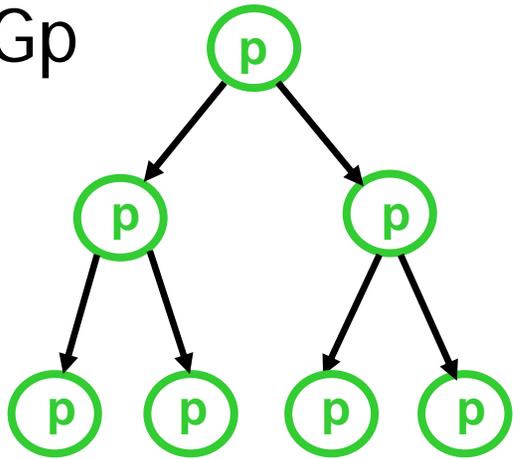
Gp



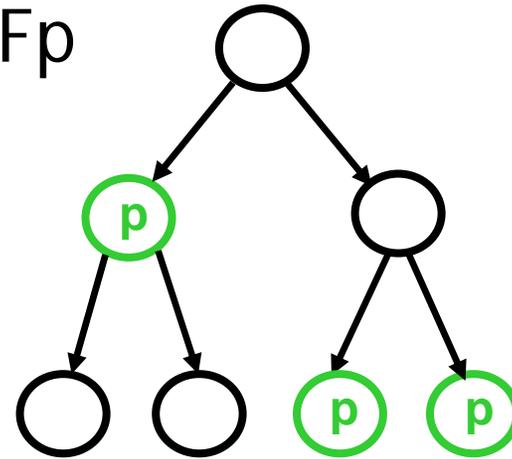
pUq



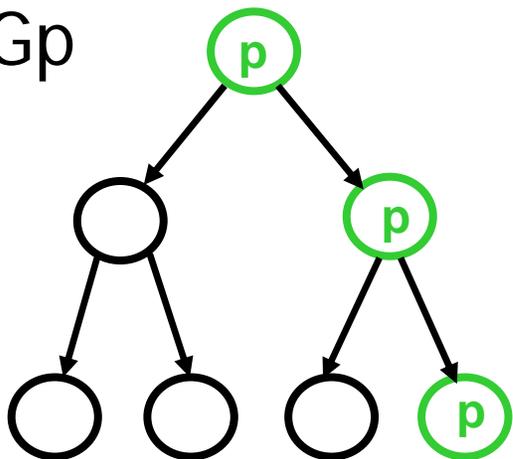
AGp



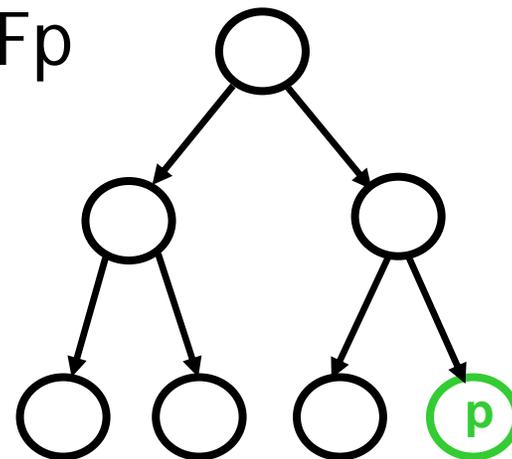
AFp



EGp



EFp





Input

- Interaction graph / state transition graph
- Temporal logic formula (CTL)

Output

Set of states in which the formula is true

Example

$$(x = 0) \Rightarrow AG(\neg(x = 2))$$

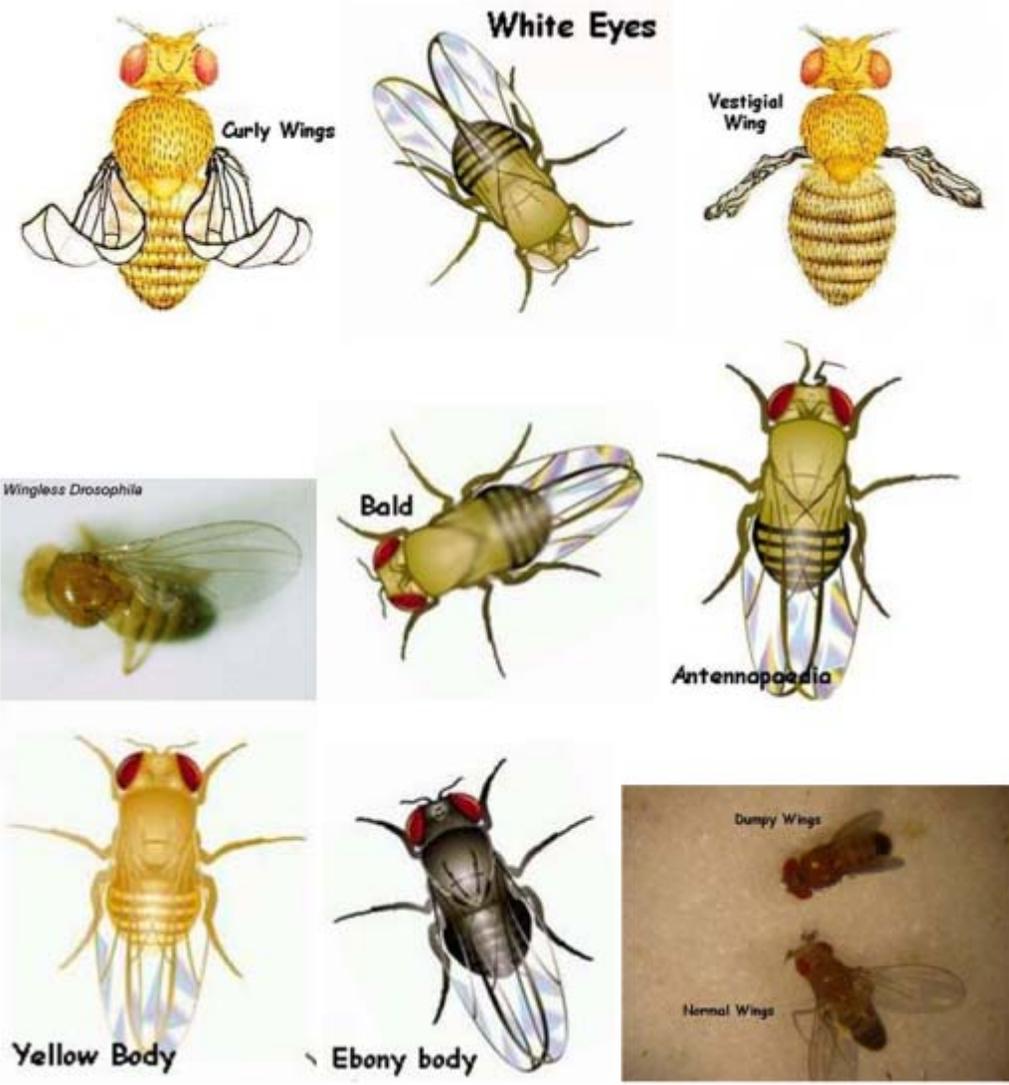
$$(x = 2) \Rightarrow AX AF(x = 2)$$

$$\neg E(\neg(x = 2) U (x = 1))$$

Can also be used for **network inference**.



IV. Application: Bio-Logic



Understand the regulatory logic underlying developmental and other biological processes

Source: www.chaossience.org.uk



- Molecular systems biology
- Logical modeling of regulatory structures
 - ❖ Boolean logic
 - ❖ Multi-valued logic
- Logical analysis of the dynamics
 - ❖ Non-determinism
 - ❖ Temporal logic
 - ❖ Model checking
- Bio-Logic