

# Reconciling qualitative and abstract (and scalable) reasoning with Boolean networks

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# Overview

Discrete Dynamical Systems  
(Boolean Networks)



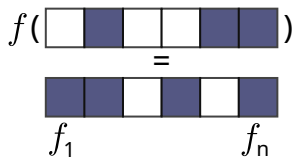
Systems Biology  
(Signalling/regulation networks)

Concurrency Theory  
(Semantics)

# Introduction

**Boolean Network (BN)**  $f : \mathbb{B}^n \rightarrow \mathbb{B}^n$

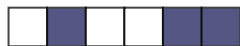
**Configuration:**  $x \in \mathbb{B}^n$



# Introduction

Boolean Network (BN)  $f : \mathbb{B}^n \rightarrow \mathbb{B}^n$

Configuration:  $x \in \mathbb{B}^n$



**synchronous** update



$$f\left(\begin{array}{|c|c|c|c|c|c|}\hline \square & \blacksquare & \square & \square & \blacksquare & \blacksquare \\ \hline\end{array}\right) =$$



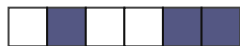
$f_1$

$f_n$

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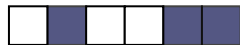


$$f\left(\begin{bmatrix} \text{white} & \text{dark blue} & \text{white} & \text{white} & \text{dark blue} & \text{dark blue} \end{bmatrix}\right) =$$

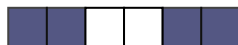


$f_1$

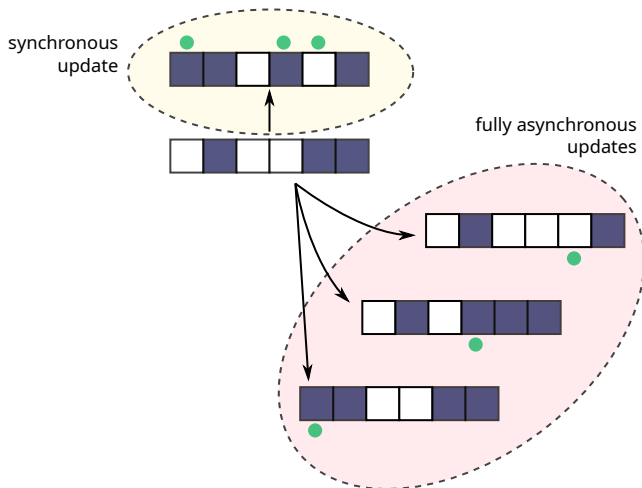
$f_n$



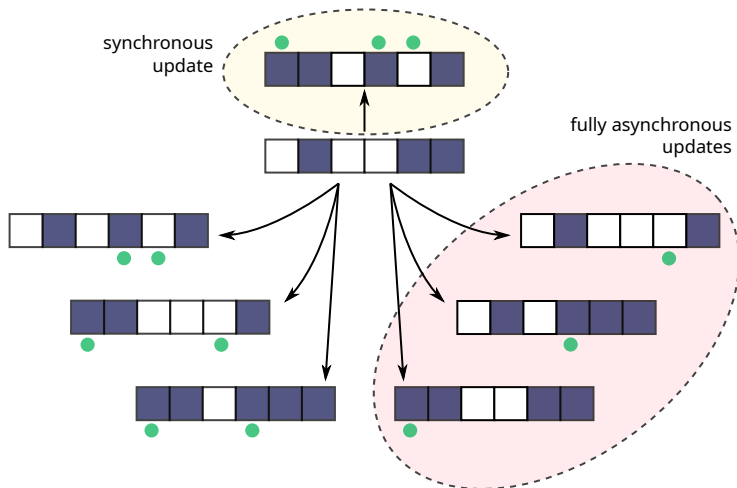
**fully**  
**asynchronous**  
updates



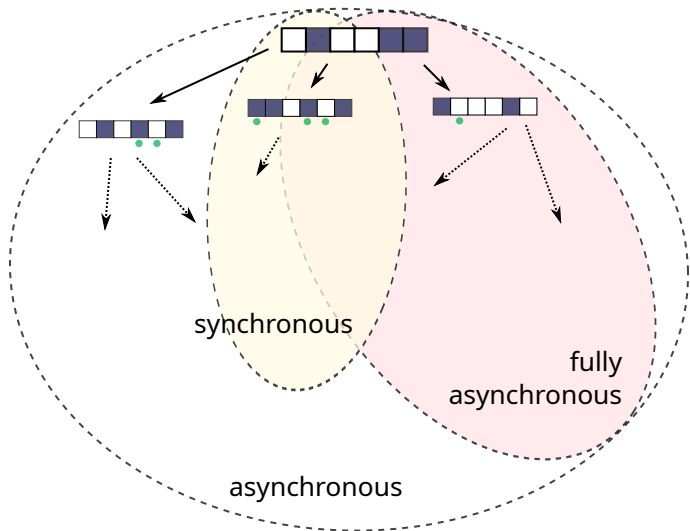
# Asynchronous updates



# Asynchronous updates



# Reachable configurations

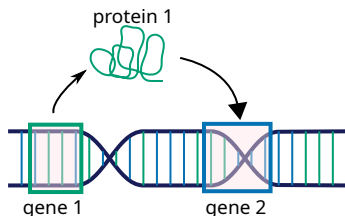




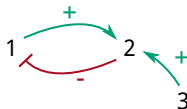
# Boolean networks for biological processes

Example with gene regulation

[Since 70's - Kauffman, Thomas]



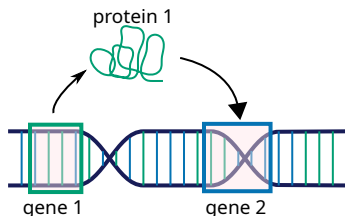
Influence graph



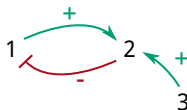
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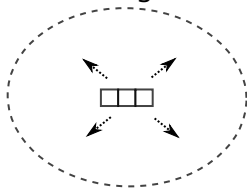
[Since 70's - Kauffman, Thomas]



Influence graph



Reachable configurations



Boolean  
network

$$f_1(x) \triangleq \neg x_2$$

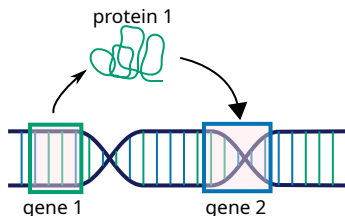
$$f_2(x) \triangleq x_1 \wedge x_3 \quad + \text{ update mode} =$$

$$f_3(x) \triangleq \dots$$

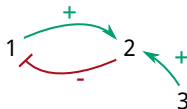
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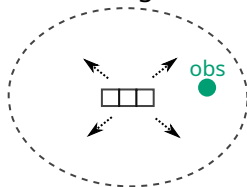
[Since 70's - Kauffman, Thomas]



Influence graph



Reachable configurations



Boolean network

$$\begin{aligned} f_1(x) &\triangleq \neg x_2 \\ f_2(x) &\triangleq x_1 \wedge x_3 \\ f_3(x) &\triangleq \dots \end{aligned} \quad + \text{ update mode} =$$

Validation w.r.t. observations (e.g. time series data)

⇒ we expect **measurements match with reachable configurations**

# Properties of Boolean networks for biology

Given a Boolean network  $f$  of dimension  $n$

**Reachability** (seq. of transitions from conf.  $x$  to  $y$ )

⇒ **PSPACE-complete** with update modes

Potential behaviours/capabilities of the cell

**Fixpoints** ( $f(x) = x$ )

⇒ **NP-complete** for sync/async/gasync

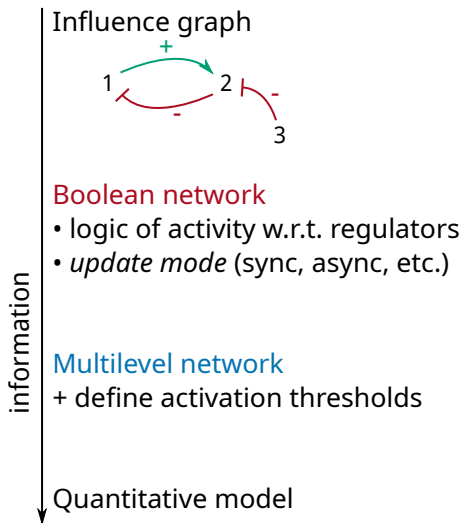
Steady states/phenotypes

**Attractors** (smallest sets of conf. closed by transitions)

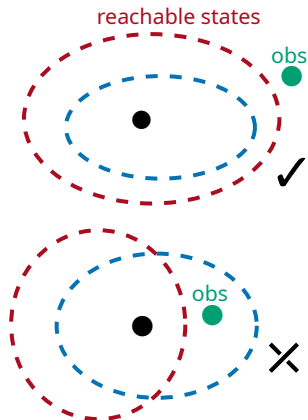
⇒ **PSPACE-complete** with update modes

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# Qualitative vs abstract modelling

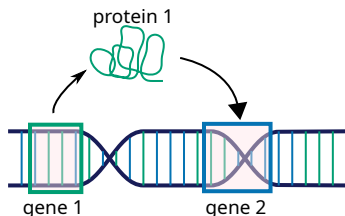


**Consistency**  
analysis at Boolean level  
transposable to multilevel?

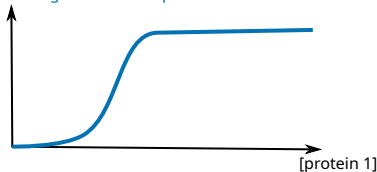


# Gene expression is not Boolean

Qualitative modelling: Boolean vs multivalued networks

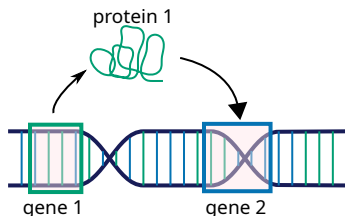


effect on gene 2 transcription



# Gene expression is not Boolean

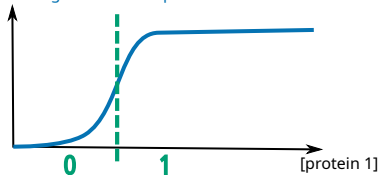
Qualitative modelling: Boolean vs multivalued networks



Boolean network

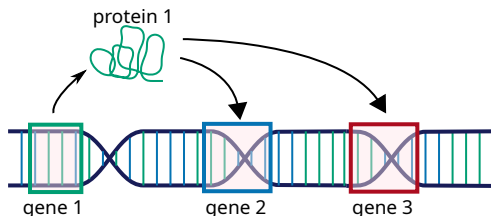
$$f_2(x) \triangleq x_1$$

effect on gene 2 transcription



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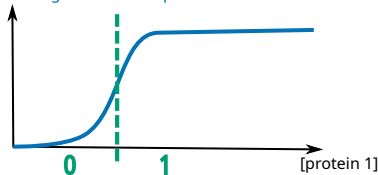


Boolean network

$$f_2(x) \triangleq x_1$$

$$f_3(x) \triangleq x_1$$

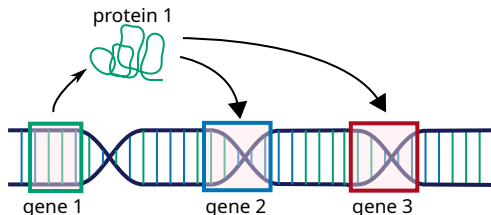
effect on gene 2 transcription





# Gene expression is not Boolean

Qualitative modelling: Boolean vs multivalued networks

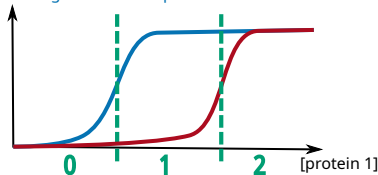


Boolean network

$$f_2(x) \triangleq x_1$$

$$f_3(x) \triangleq x_1$$

effect on gene 3 transcription  
effect on gene 2 transcription



Multivalued network

$$F_2(x) \triangleq (x_1 \geq 1)$$

$$F_3(x) \triangleq (x_1 \geq 2)$$

# Multivalued Networks

$$F : \mathbb{M}^n \rightarrow \{-1, 0, 1\}^n \quad \mathbb{M} \triangleq \{0, 1, \dots, m\}$$

Asynchronous semantics

$$x \in \mathbb{M}^n \xrightarrow[\text{a}]{F} y \in \mathbb{M}^n$$

$$\Delta$$

$$\forall i \in \Delta(x, y), y_i = x_i + F_i(x)$$

To ease notations,

$$F_i(x) \triangleq 1 \text{ if } P(x) \text{ else } -1 \quad \equiv \quad F_i(x) \triangleq P(x)$$

Remark: very **similar to ODEs** ( $\mathbb{R}_{\geq 0}^n \rightarrow \mathbb{R}^n$ )

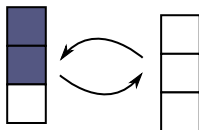
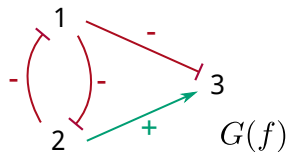
**Asynchronous  
Boolean networks:  
a bug...**

# Motivating example (embedded in many actual biological networks)

$$f_1(x) \triangleq \neg x_2$$

$$f_2(x) \triangleq \neg x_1$$

$$f_3(x) \triangleq \neg x_1 \wedge x_2$$

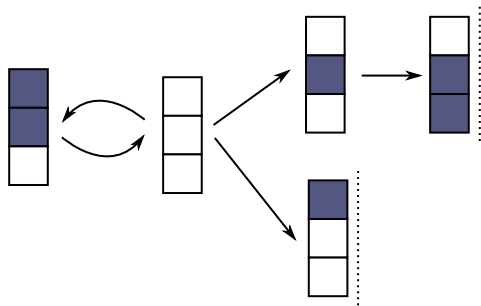
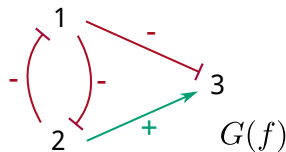


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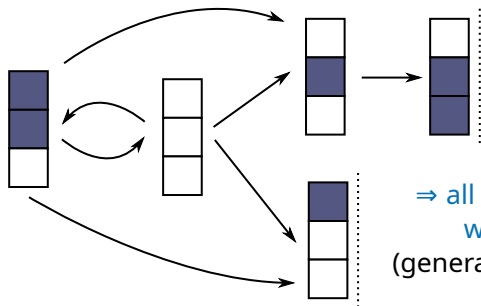
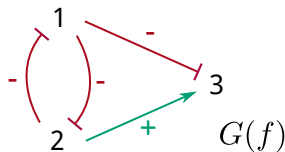


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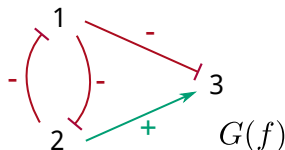
$\Rightarrow$  all configurations reachable  
with any update mode  
(generalized) asynchronous mode

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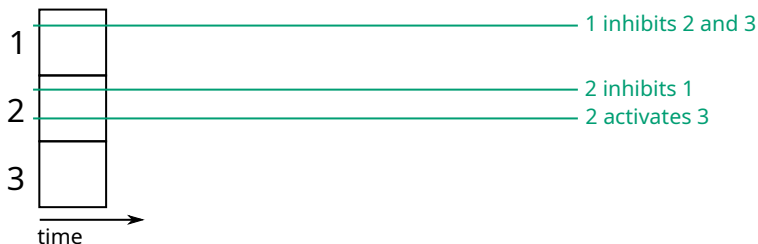
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$$F_2(x) \triangleq \neg x_1$$

$$F_3(x) \triangleq \neg x_1 \wedge (x_2 \geq 1)$$



Compatible continuous/multilevel model:

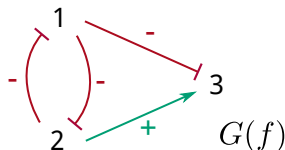


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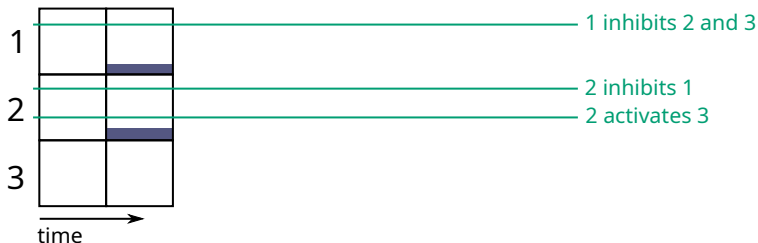
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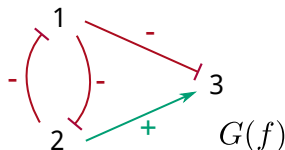


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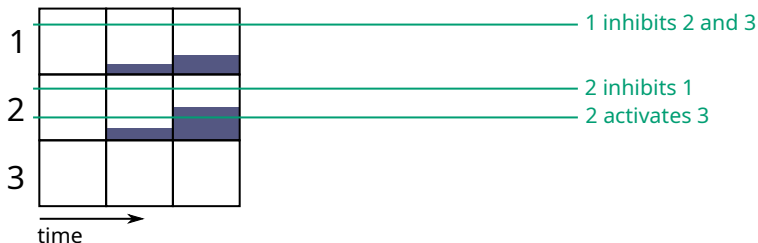
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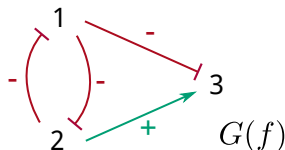


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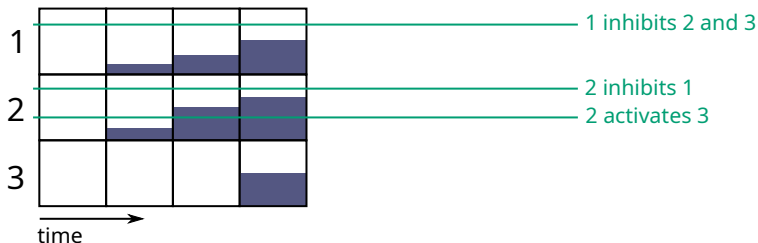
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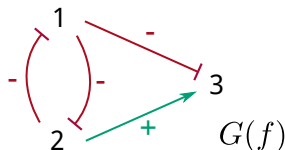


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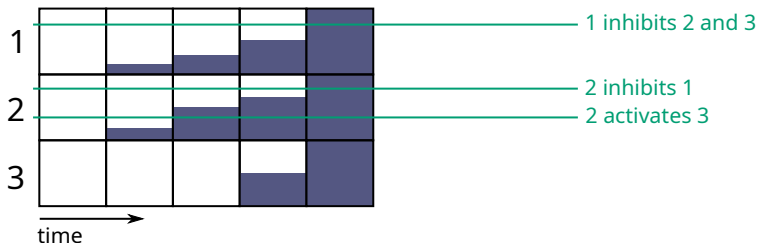
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Compatible **continuous/multilevel** model:

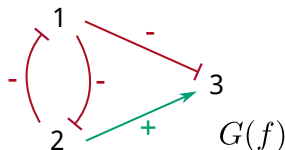


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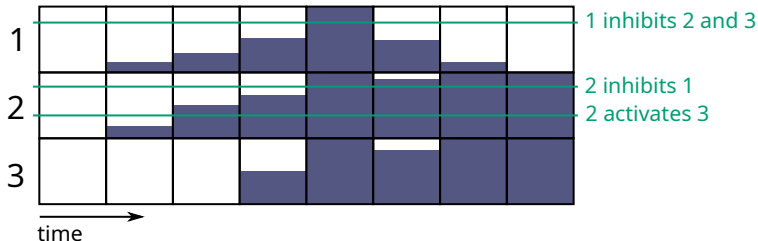
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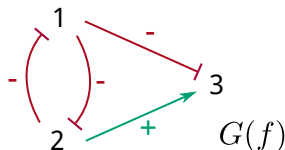


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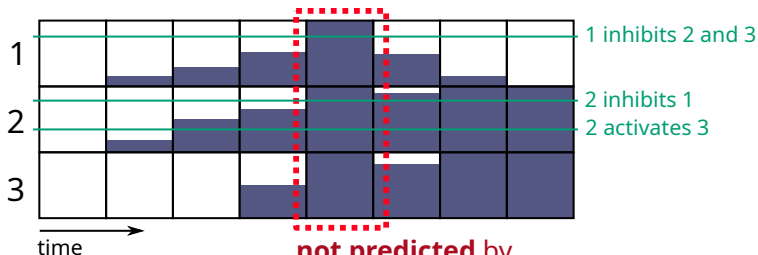
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Compatible continuous/multilevel model:





**not predicted by  
update modes in Boolean**

**Most Permissive**  
**Boolean networks**  
enabling new behaviours

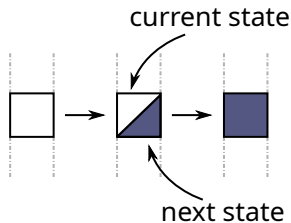
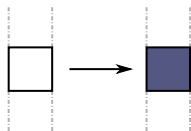
# Most permissive semantics

- **delay between firing and application** of state change

⇒ allow interleaving other state changes

- in "intermediate" states  

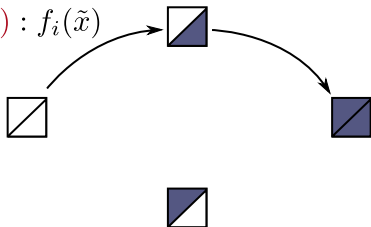
**other components choose what they see**



# Most permissive semantics

Rules for state of component i:

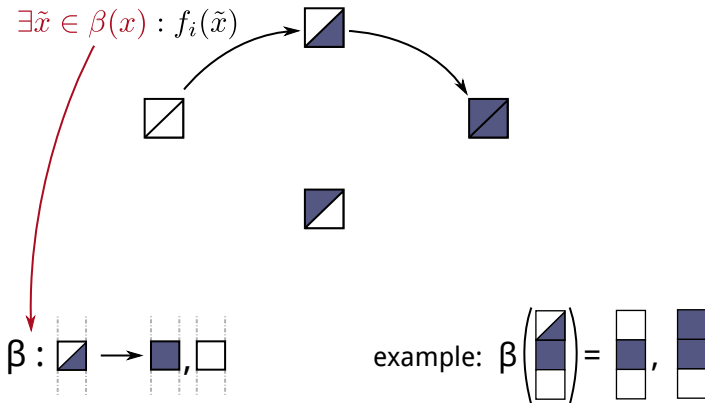
$$\exists \tilde{x} \in \beta(x) : f_i(\tilde{x})$$





# Most permissive semantics

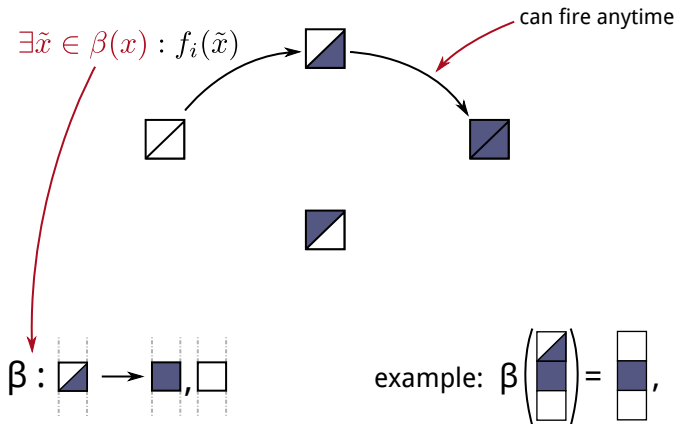
Rules for state of component  $i$ :



Choose value of "changing" components  
(act as choosing an activation threshold)

# Most permissive semantics

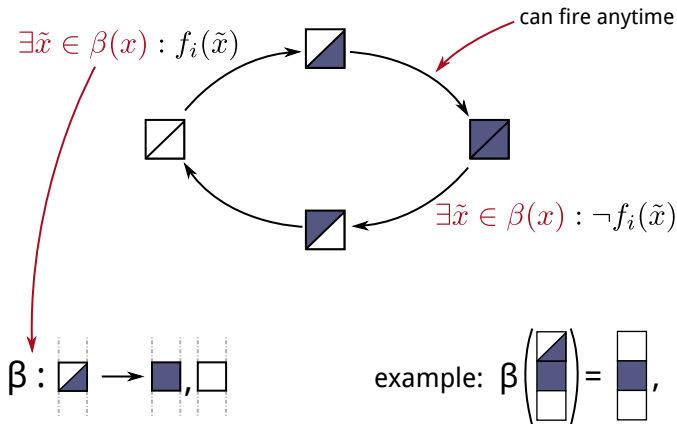
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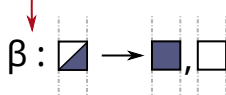
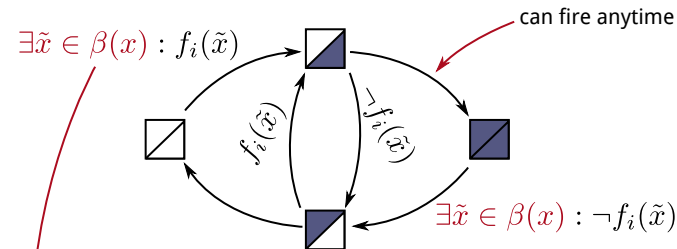
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# Most permissive semantics

Rules for state of component  $i$ :



example:  $\beta \left( \begin{array}{|c|} \hline \text{white} \\ \hline \text{dark blue} \\ \hline \text{white} \\ \hline \end{array} \right) = \begin{array}{|c|} \hline \text{dark blue} \\ \hline \end{array}, \begin{array}{|c|} \hline \text{dark blue} \\ \hline \end{array}$

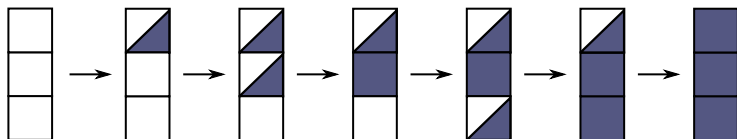
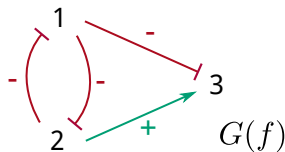
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## Application to motivating example

$$f_1(x) \triangleq \neg x_2$$

$$f_2(x) \triangleq \neg x_1$$

$$f_3(x) \triangleq \neg x_1 \wedge x_2$$

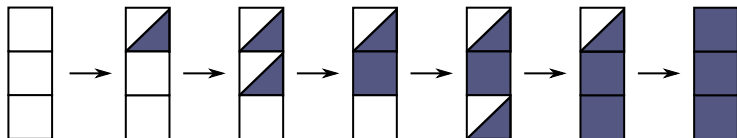
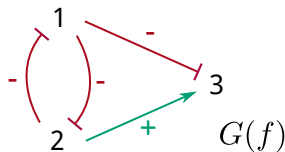


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$\Rightarrow$  valid with respect to multivalued refinements

# Properties of the most permissive semantics

**Correct abstraction** of multilevel/quantitative systems:

- includes all the **transitions of every update mode**
- multilevel **refinements only remove behaviours**
- **Reachability** can be decided in **quadratic nb of transitions**  
(PTIME with locally-monotonic networks, or encoded as BDDs/Petri nets/...;  
NP-complete otherwise; instead of PSPACE-complete with update modes)
- **Attractors are hypercubes** (minimal trap spaces)
  - ⇒ finding **attractors is NP-complete** (instead of PSPACE-complete)
  - ⇒ fixpoints are the same as with update modes

# Refinements of Boolean Networks

A **multivalued network**

$$F : \mathbb{M}^n \rightarrow \{-1, 0, 1\}^n$$

is a **refinement of a Boolean network**  $f$  iff

$$F_i(\begin{array}{|c|c|c|} \hline \text{white} & \text{blue} & \text{white} \\ \hline \end{array}) > 0 \implies \exists \begin{array}{|c|c|c|} \hline \text{white} & \text{blue} & \text{white} \\ \hline \end{array} : f_i(\begin{array}{|c|c|c|} \hline \text{white} & \text{blue} & \text{white} \\ \hline \end{array}) = 1$$

$$F_i(\begin{array}{|c|c|c|} \hline \text{blue} & \text{white} & \text{white} \\ \hline \end{array}) < 0 \implies \exists \begin{array}{|c|c|c|} \hline \text{blue} & \text{white} & \text{white} \\ \hline \end{array} : f_i(\begin{array}{|c|c|c|} \hline \text{blue} & \text{white} & \text{white} \\ \hline \end{array}) = 0$$

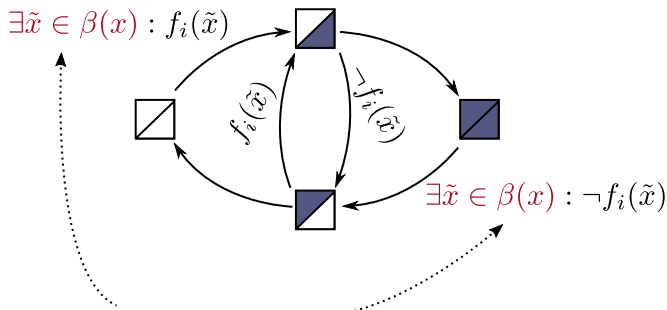
**Most permissive semantics weakly simulates  
any multivalued refinement with *any update mode***

(can be extended to ODEs)



# Reachability with the most permissive semantics

**Cost of one transition** in component  $i$



**NP** (SAT) in general;

**Linear** with

- locally-monotonic networks ( $f_i$  are monotone)
- when  $f$  is encoded as BDDs/Petri nets/...

# Reachability with the most permissive semantics

**Deciding reachability requires quadratic nb of transitions**

Main property:

y reachable from x  $\Leftrightarrow$  there exists a **path of length  $\leq 3n$  transitions**

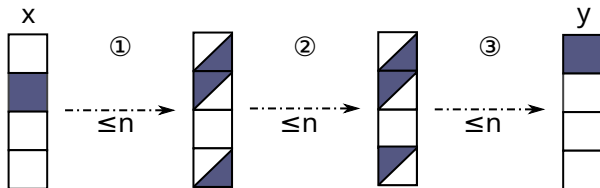


# Reachability with the most permissive semantics

## Deciding reachability requires quadratic nb of transitions

Main property:

y reachable from x  $\Leftrightarrow$  there exists a path of length  $\leq 3n$  transitions



- ① - only transitions to "in-between" states\*
- ② - orient towards final states
- ③ - converge to final states

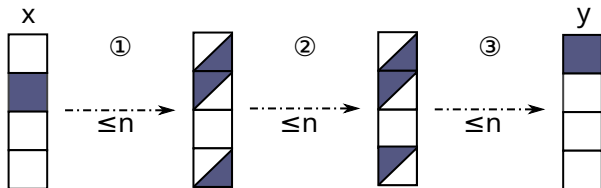
\*: some components must not be updated!

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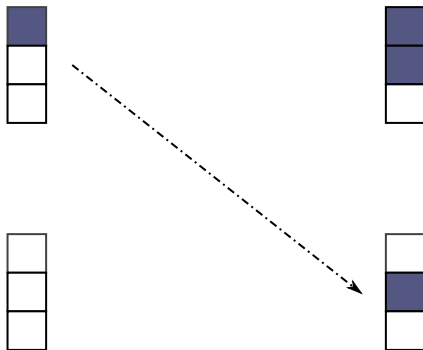
\*: some components must not be updated!

NP in general  
PTIME w/  
locally-monotonic;  
BDDs; Petri nets..

# Attractors with the most permissive semantics

Attractor: smallest set of configurations closed by transitions

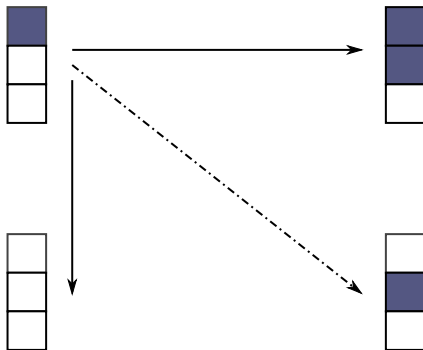
**Attractors are hypercubes**



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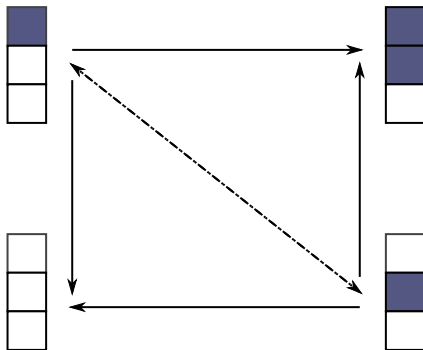
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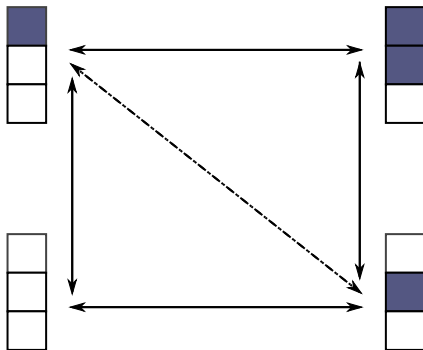
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# Attractors with the most permissive semantics

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## Attractors are hypercubes

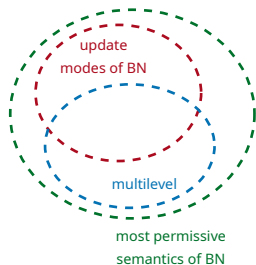


Attractors of most permissive semantics = minimal trap spaces

Deciding if trap space is minimal = 2 reachability checking



# Is most permissive semantics restrictive?



## Minimality of abstraction

to any "most permissive" transition,  
there is corresponding multilevel transition  
(work in progress w/ "most permissive" paths:  
non-minimal, but tricky counter-examples)

- **fixpoints** (stable states) are **preserved (identical)**
  - **trap spaces**: known to be relevant for reasoning with **attractors**  
[Klarner et al in Nat. Comp. 2015] [Naldi in Front. Phys. 2018]
- ⇒ most permissive semantics seems still adequate to model  
**differentiation processes** !

# Boolean network synthesis

WiP (PhD studies of [Stéphanie Chevalier](#) LRI-LaBRI-Curie)

## Constraints

- Putative influences (domain of Boolean functions)
- +/- reachability between partial configurations
- existence of (reachable) attractors

## Output

- Exhaustive enumeration of compatible Boolean networks

## SAT approach (Answer-Set Programming)

- Application to [cellular differentiation](#) data
- Scalable to >100 nodes (>1000 w/o neg. reachability)

## Conclusion

Update modes of Boolean networks (sync, async, etc.):

- difficult to justify for biology (strong implications on dynamics)
  - can miss important behaviours [CHP at AUTOMATA'18]
- ⇒ lead to reject valid models of biological systems...
- have limited tractability (model-checking, ...)

Most permissive semantics:

- correct abstraction: guarantees that multilevel/quantitative refinements only remove behaviours
  - simpler complexity: reachability PTIME, attractors NP
- ⇒ genome-scale tractability

WiP: assess non-minimality (Juraj Kolčák, LSV-LaBRI PhD student)