

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



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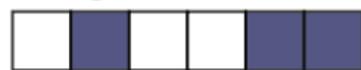
$f_1$

$f_n$

## Introduction

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

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↓ **synchronous** update



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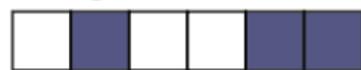
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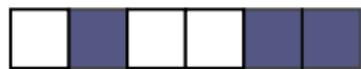


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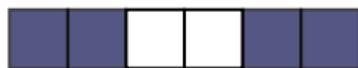


$f_1$

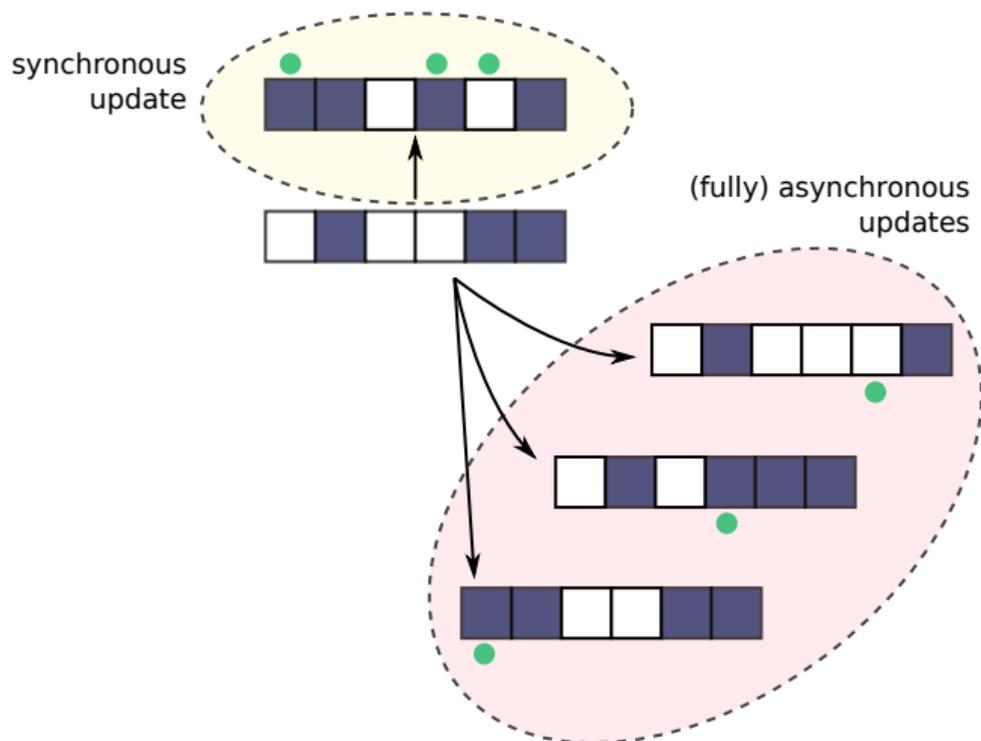
$f_n$



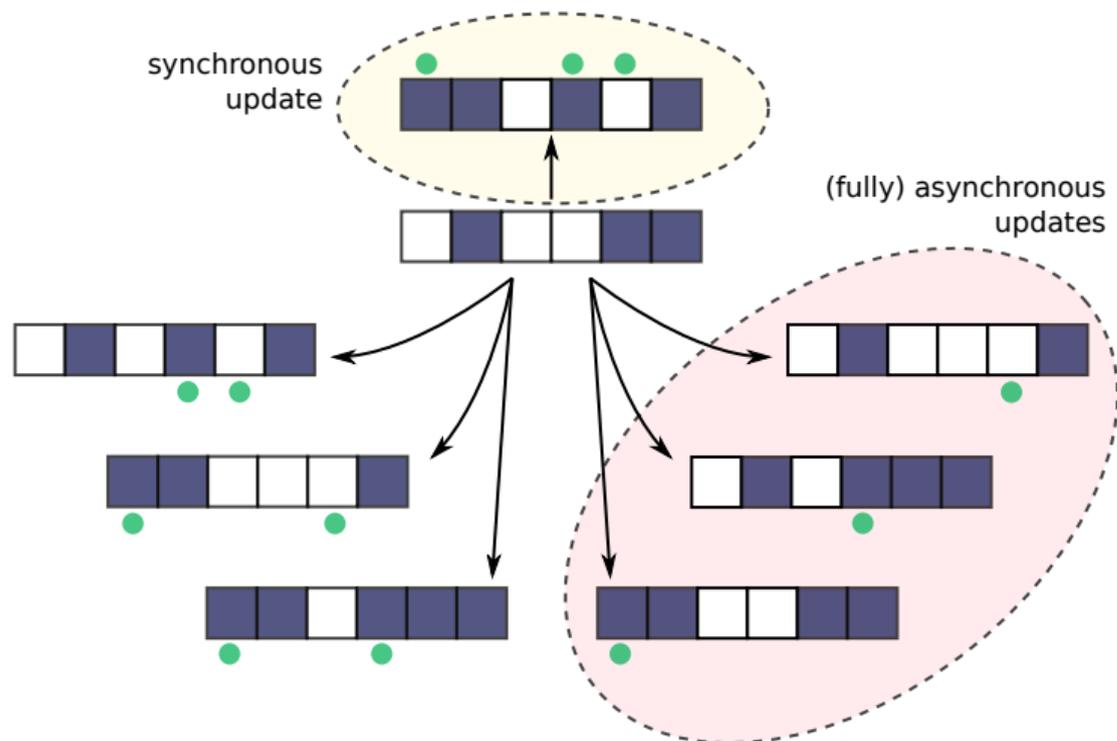
fully asynchronous updates



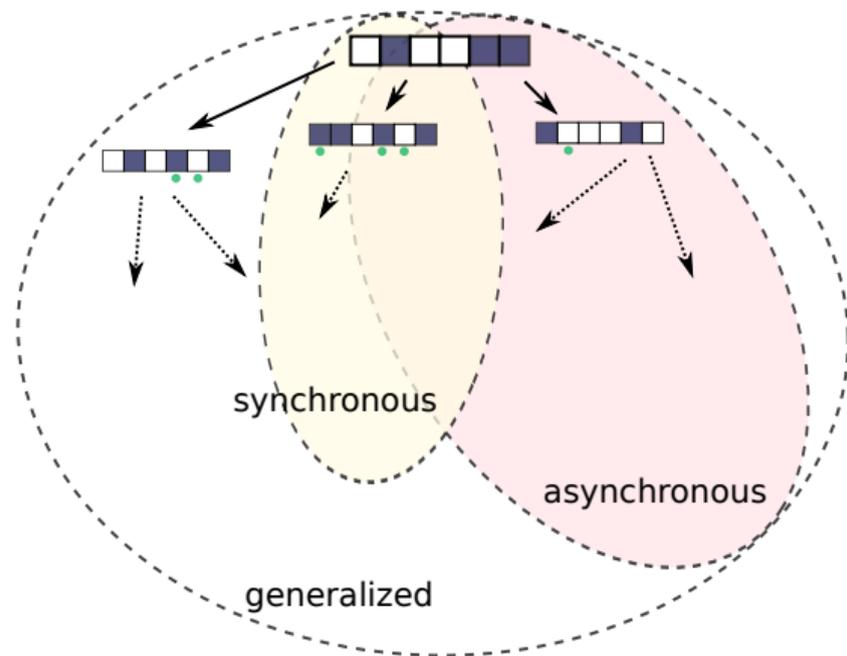
# Generalized Asynchronicity



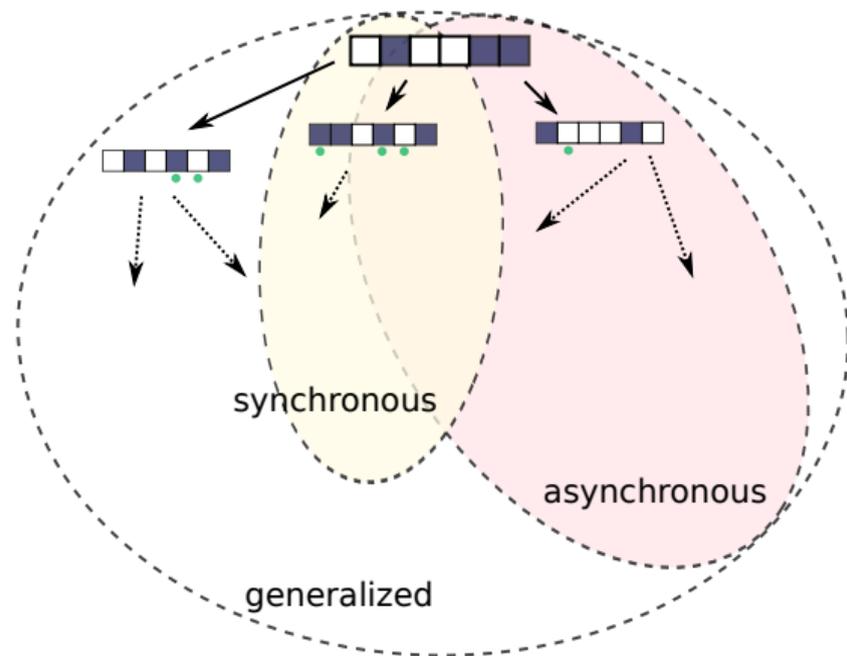
# Generalized Asynchronicity



# Reachable configurations



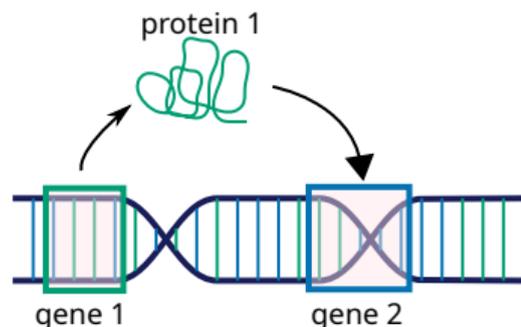
# Reachable configurations



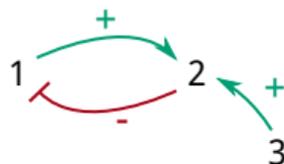
Should we reach configurations  
beyond generalized asynchronicity?

# Boolean networks for biological processes

Example with gene regulation

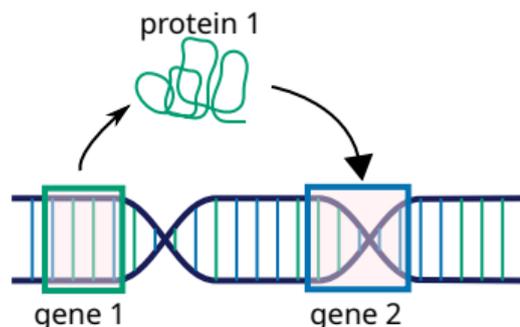


Influence graph

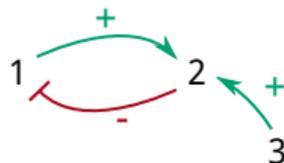


# Boolean networks for biological processes

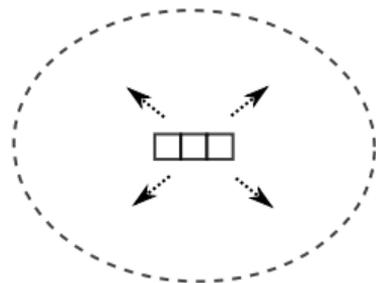
Example with gene regulation



Influence graph



Reachable configurations



Boolean network

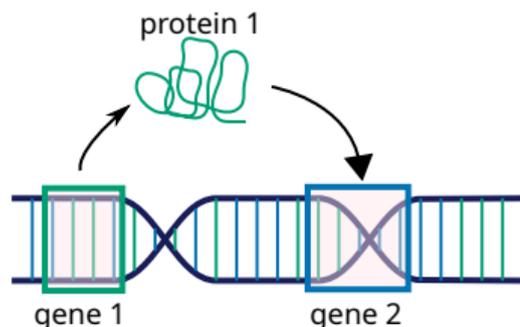
$$f_1(x) \triangleq \neg x_2$$

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

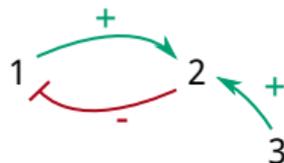
$$f_3(x) \triangleq \dots$$

# Boolean networks for biological processes

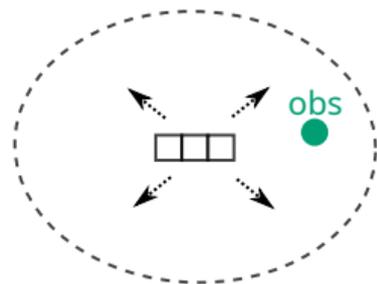
Example with gene regulation



Influence graph



Reachable configurations



Boolean network

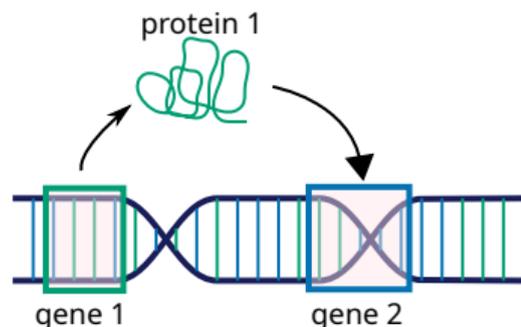
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Validation w.r.t. observations (e.g. time series data)

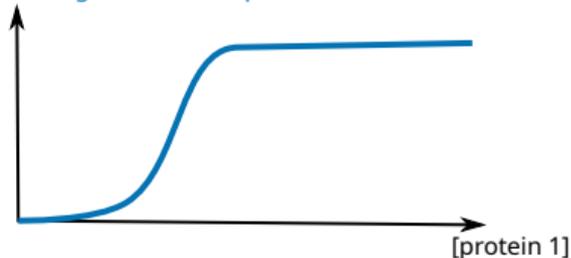
⇒ we expect **measurements match with reachable configurations**

# Gene expression is not Boolean

Qualitative modelling: Boolean vs multivalued networks

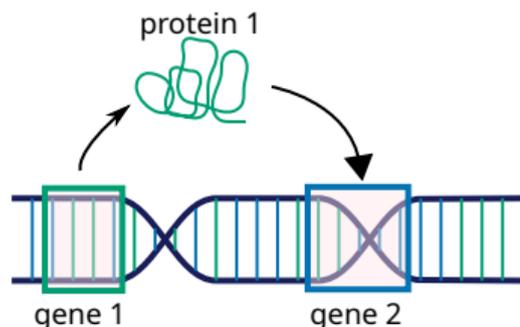


effect on gene 2 transcription



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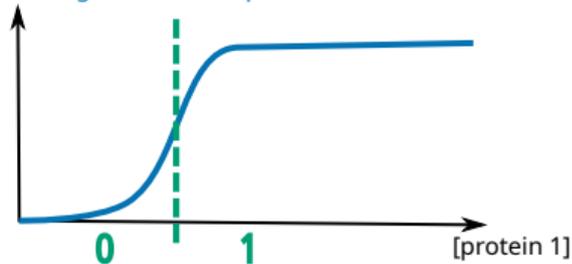
Qualitative modelling: Boolean vs multivalued networks



Boolean network

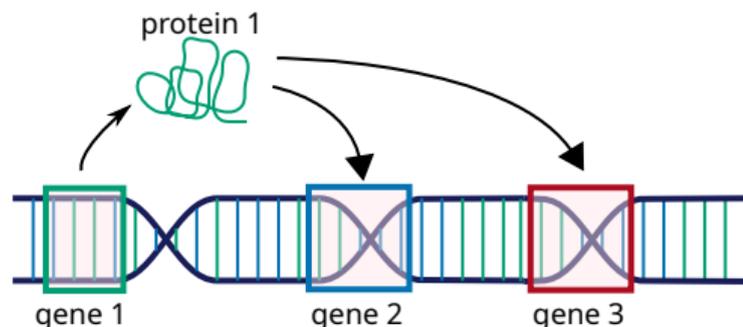
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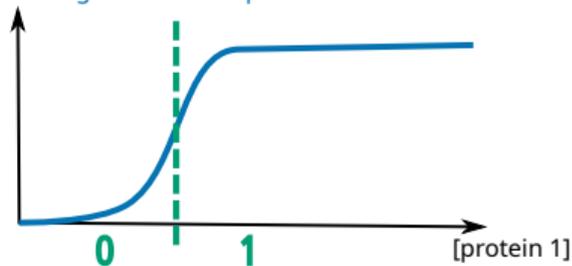


Boolean network

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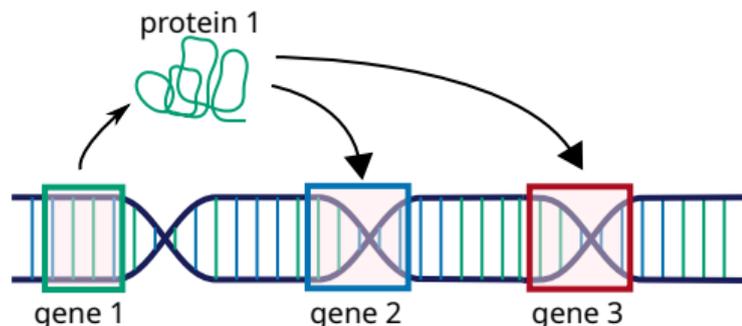
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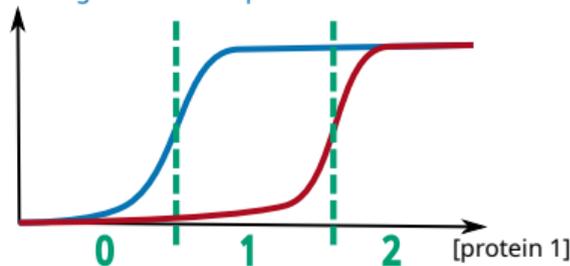


### Boolean network

$$f_2(x) \triangleq x_1$$

$$f_3(x) \triangleq x_1$$

effect on gene 3 transcription  
effect on gene 2 transcription



### Multivalued network

$$f_2(x) \triangleq (x_1 \geq 1)$$

$$f_3(x) \triangleq (x_1 \geq 2)$$

Remark: Multivalued models can require different thresholds for each target

# 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

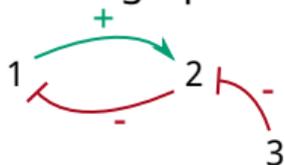
**Attractors** (smallest sets of conf. closed by transitions)

⇒ PSPACE-complete with update modes

Steady states/phenotypes

# Qualitative vs abstract modelling

Influence graph



**Boolean network**

- logic of activity w.r.t. regulators
- *update mode* (sync, async, etc.)

**Multilevel network**

- + define activation thresholds

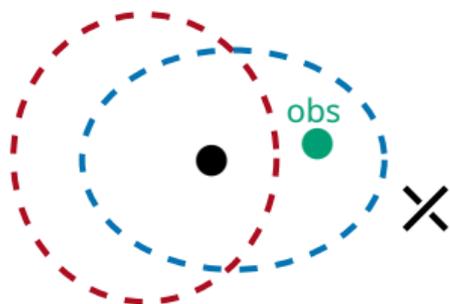
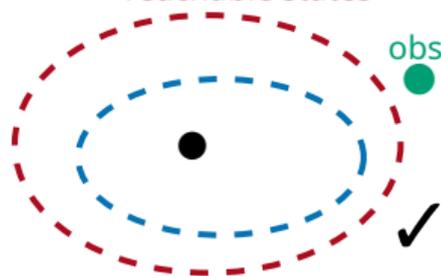
**Quantitative model**

information  
↓

## Consistency

analysis at Boolean level  
transposable to multilevel?

reachable states



**Update modes**  
of Boolean networks:  
**a bug...**

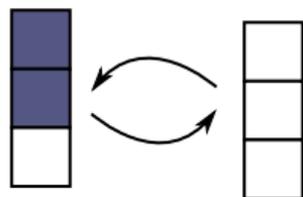
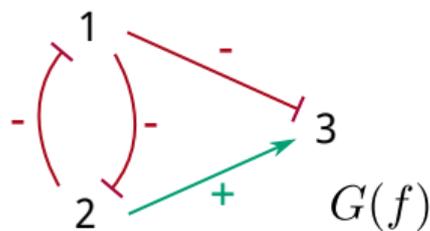
# Motivating example

(embedded in many actual biological networks)

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

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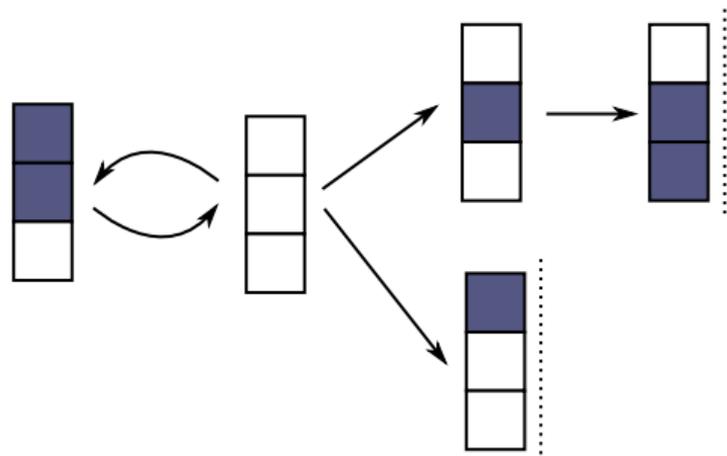
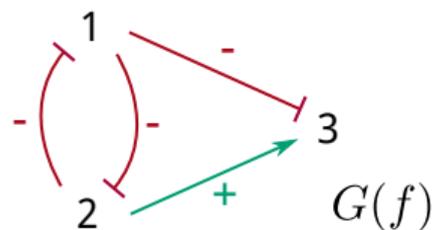


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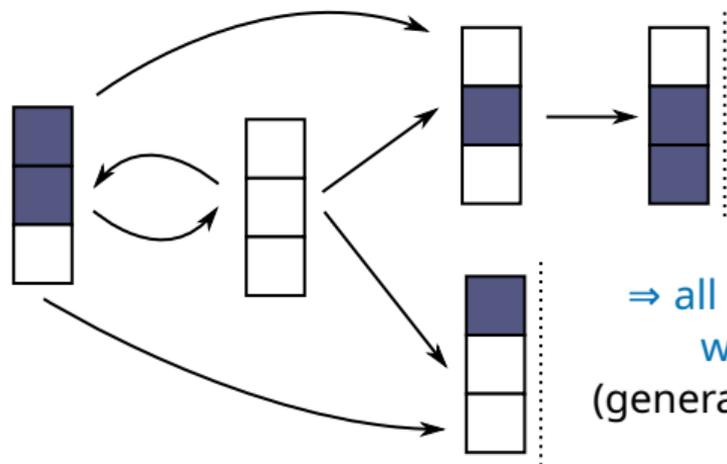
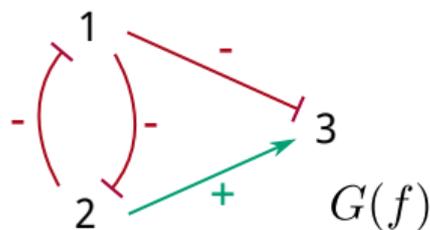


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$\Rightarrow$  all configurations reachable  
with any update mode  
(generalized) asynchronous mode

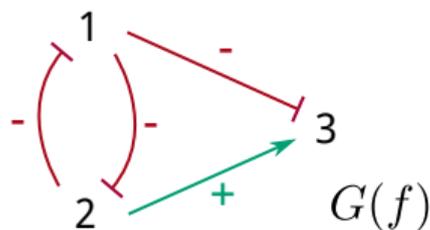
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Compatible **continuous/multilevel** dynamics:

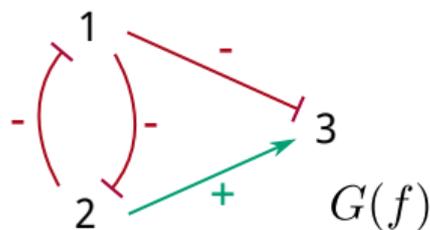


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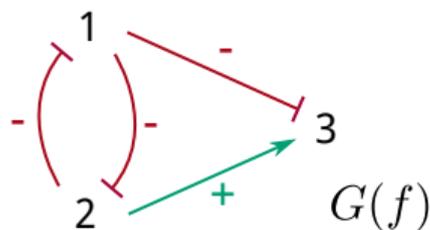


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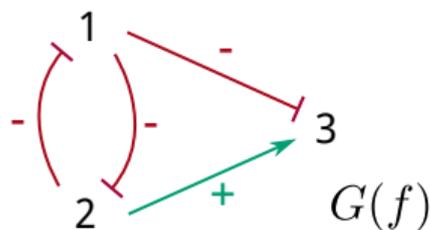
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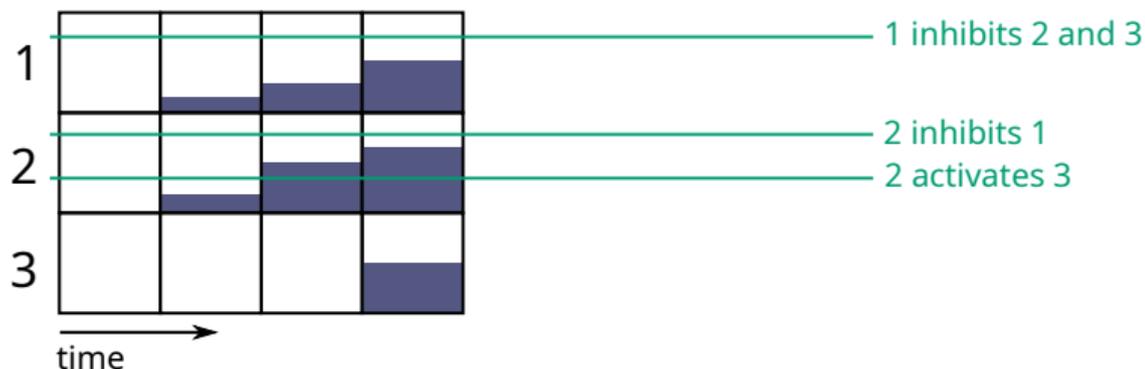
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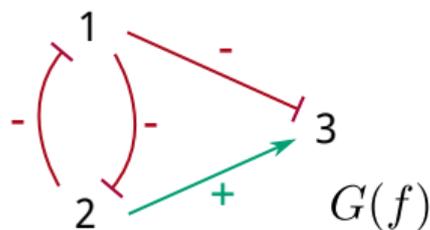


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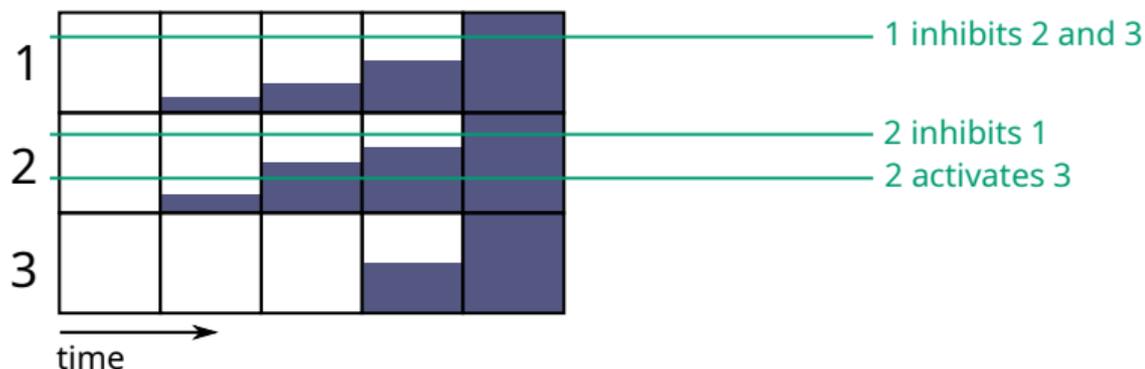
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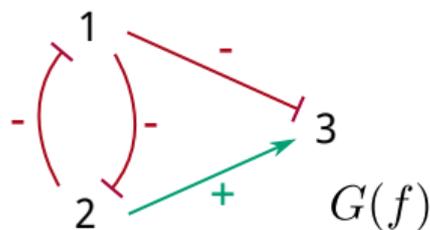


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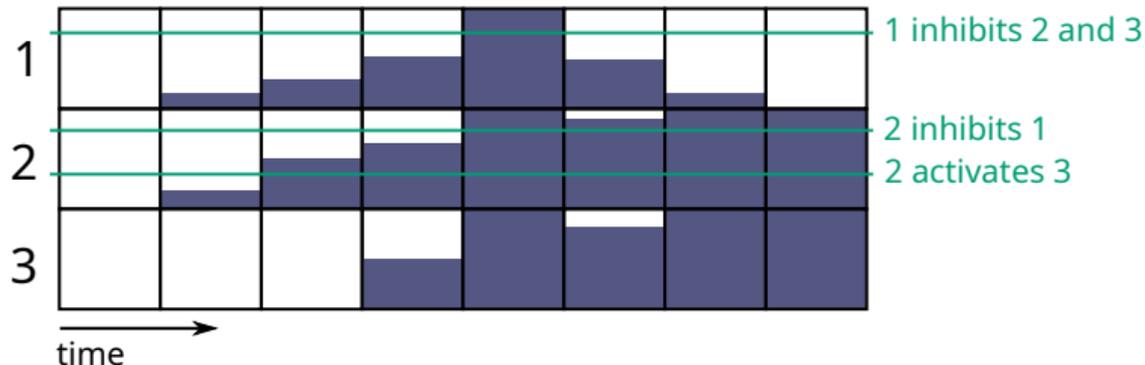
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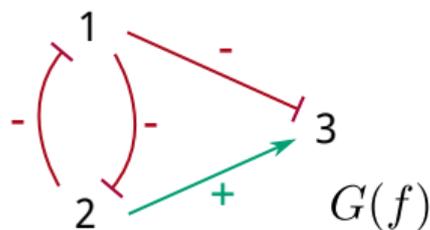


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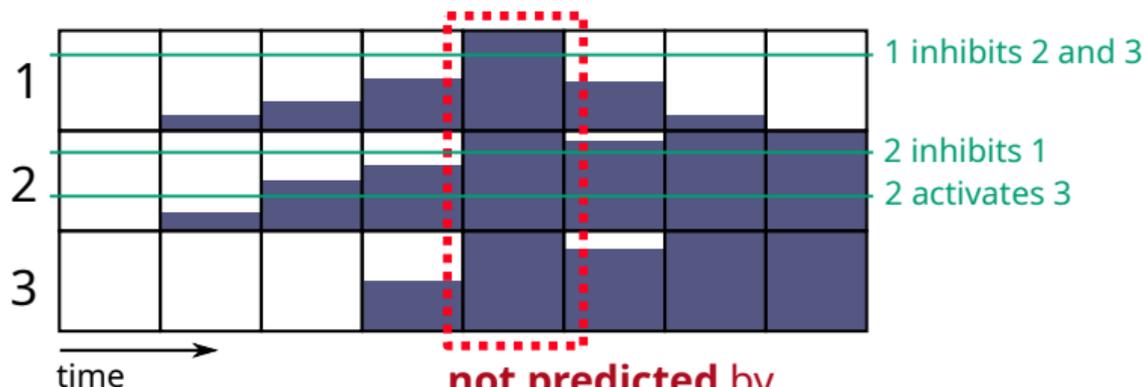
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Compatible **continuous/multilevel** dynamics:



## Practical implications

Update modes can miss admissible transitions

### Model synthesis from observations

⇒ Reject valid solutions (false negatives)

(wrongly concludes on reachability)

### Prediction for reprogramming (control)

Find mutations such that

1.  $y$  (goal phenotype) is reachable from  $x$

⇒ False negatives

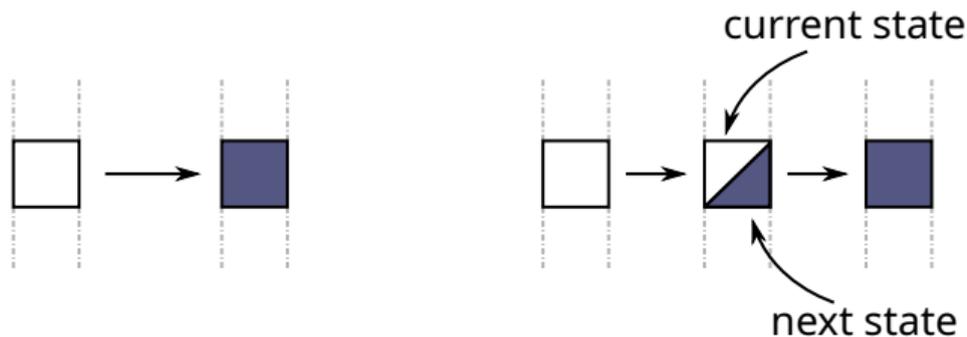
2.  $z$  (bad phenotype) is not reachable from  $x$

⇒ False positives

**Most permissive semantics  
of Boolean networks**  
enabling new behaviours

# Most permissive semantics

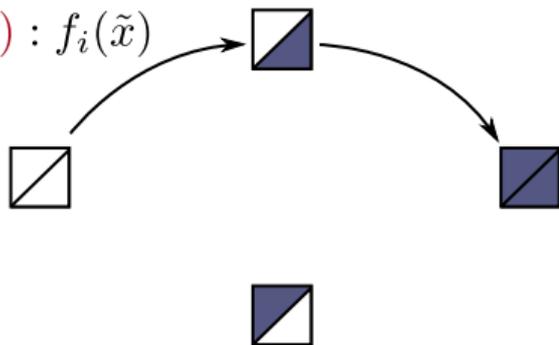
- **delay between firing and application** of state change  
⇒ allow interleaving other state changes
- in "intermediate" states 
- **other components choose what they see**



# Most permissive semantics

Rules for state of component  $i$ :

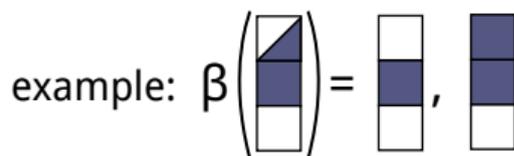
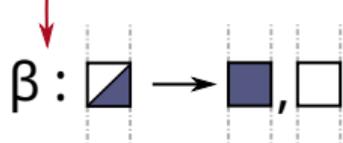
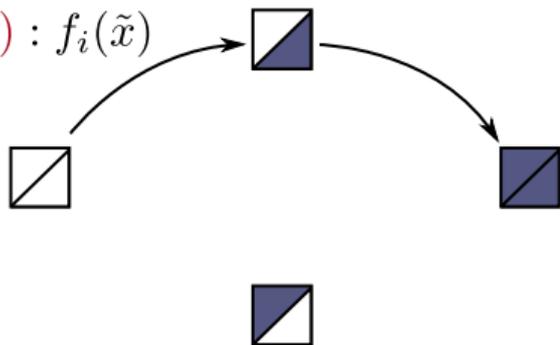
$$\exists \tilde{x} \in \beta(x) : f_i(\tilde{x})$$



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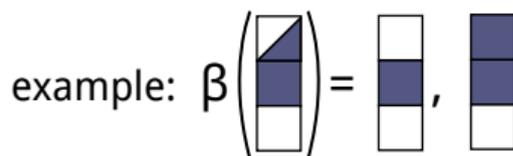
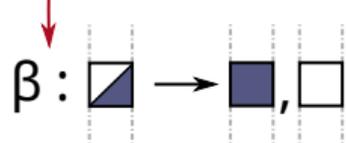
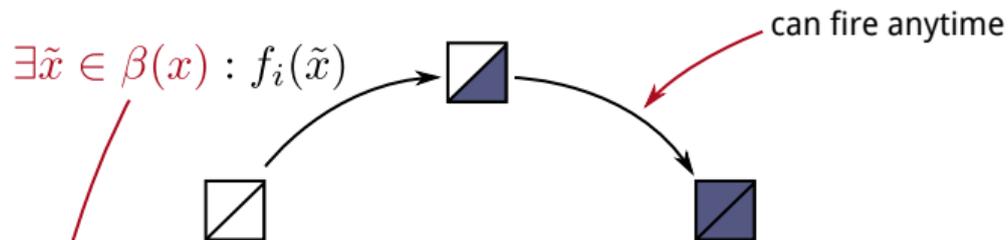
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Choose value of "changing" components  
(act as choosing an activation threshold)

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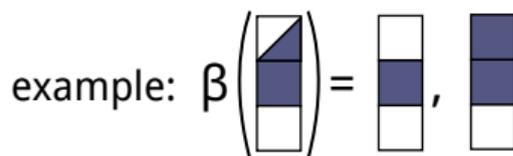
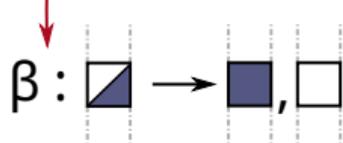
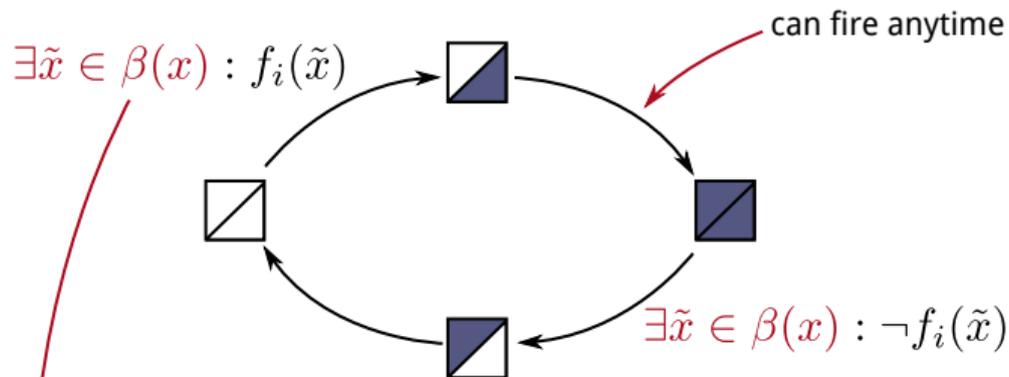
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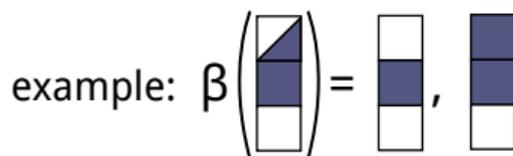
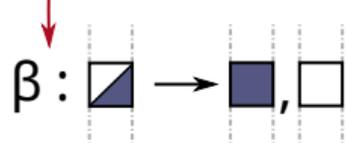
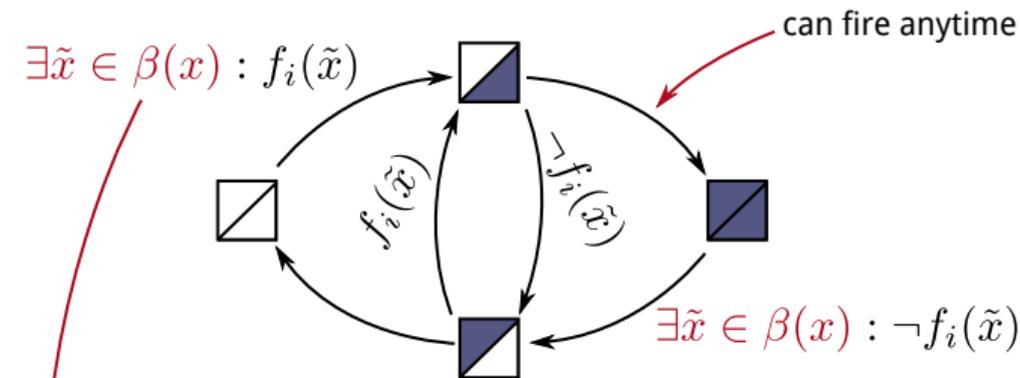
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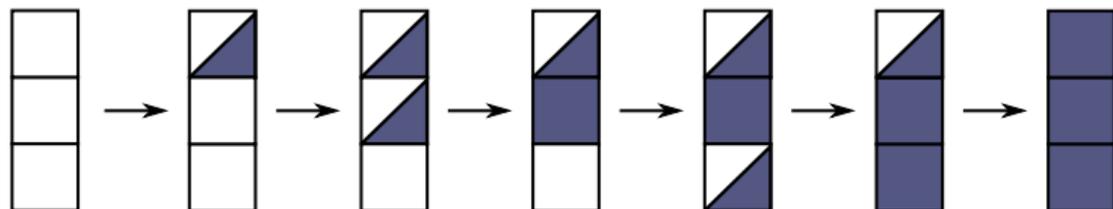
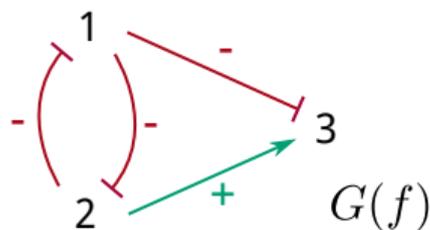
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## Application to motivating example

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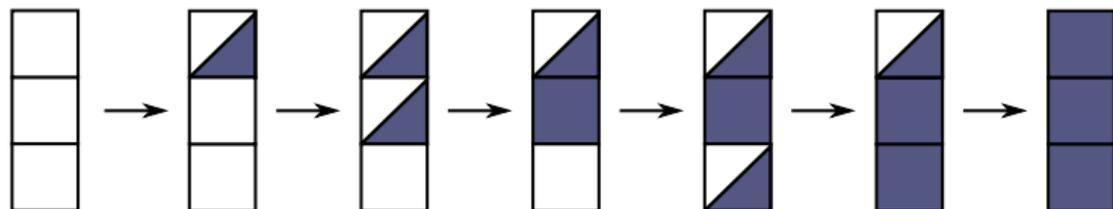
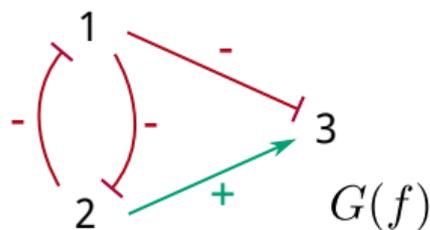


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$\Rightarrow$  valid with respect to multivalued refinements

# 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

$$F : \mathbb{M}^n \rightarrow \{\uparrow, -, \downarrow\}$$

is a refinement of a Boolean network  $f$  iff

$$F_i(\begin{array}{|c|c|c|} \hline \square & \blacksquare & \square \\ \hline \blacksquare & \blacksquare & \square \\ \hline \end{array}) = \uparrow \implies \exists \begin{array}{|c|c|c|} \hline \text{---} & \text{---} & \text{---} \\ \hline \blacksquare & \blacksquare & \text{---} \\ \hline \end{array} : f_i(\begin{array}{|c|c|c|} \hline \square & \blacksquare & \square \\ \hline \end{array}) = 1$$

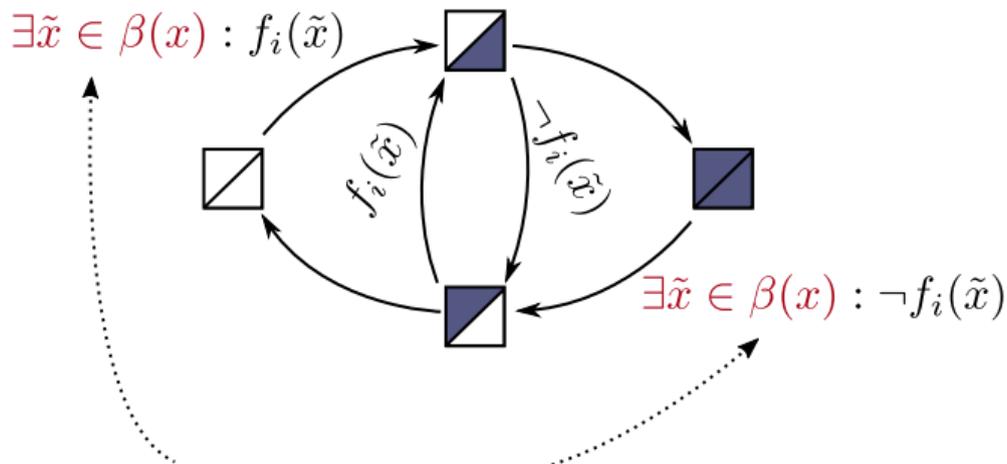
$$F_i(\begin{array}{|c|c|c|} \hline \blacksquare & \square & \square \\ \hline \blacksquare & \square & \square \\ \hline \end{array}) = \downarrow \implies \exists \begin{array}{|c|c|c|} \hline \text{---} & \text{---} & \text{---} \\ \hline \text{---} & \text{---} & \blacksquare \\ \hline \end{array} : f_i(\begin{array}{|c|c|c|} \hline \blacksquare & \square & \square \\ \hline \end{array}) = 0$$

**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

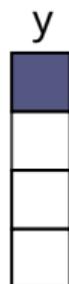
- locally-monotonic networks ( $f_i$  are monotone)
- when  $f$  is encoded as BDDs/Petri nets/...

# Reachability with the most permissive semantics

## Deciding reachability requires quadratic nb of transitions

Main property:

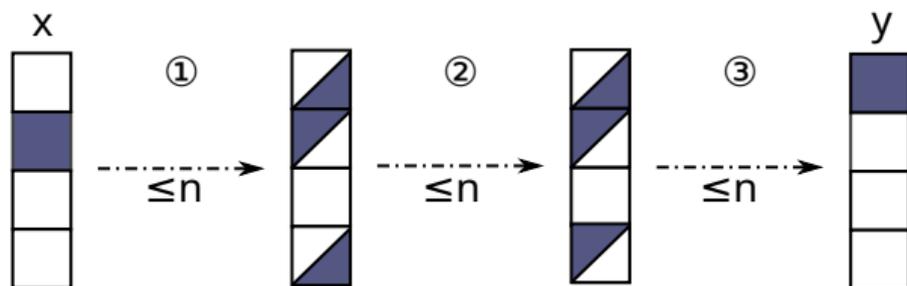
$y$  reachable from  $x \Leftrightarrow$  there exists a path of length  $\leq 3n$  transitions



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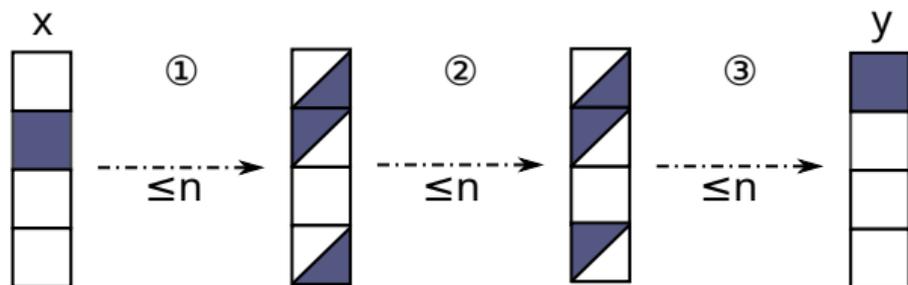
- ① - only transitions to "in-between" states\*
- ② - orient towards final states
- ③ - converge to final states

\* : some components must not be updated!

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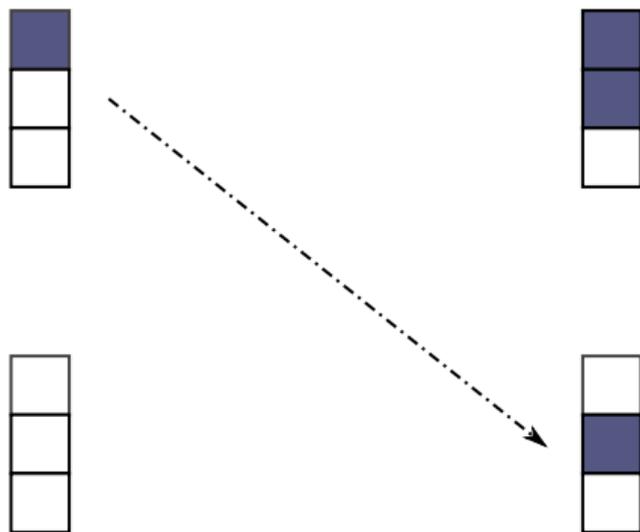
\*: some components must not be updated!

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

# Attractors with the most permissive semantics

Attractor: smallest set of configurations closed by transitions

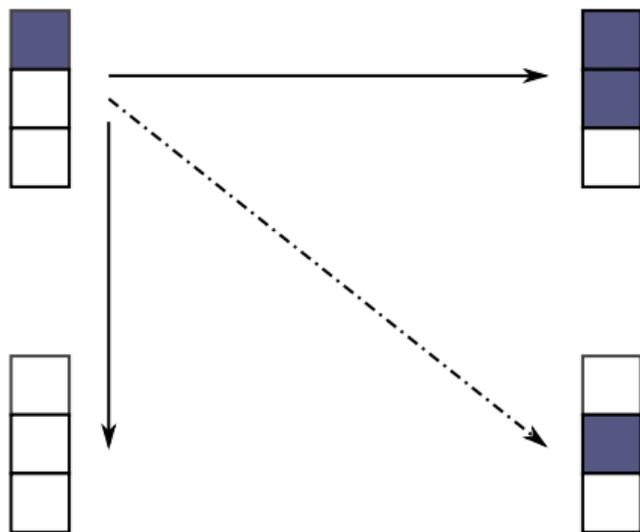
## Attractors are hypercubes



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Attractor: smallest set of configurations closed by transitions

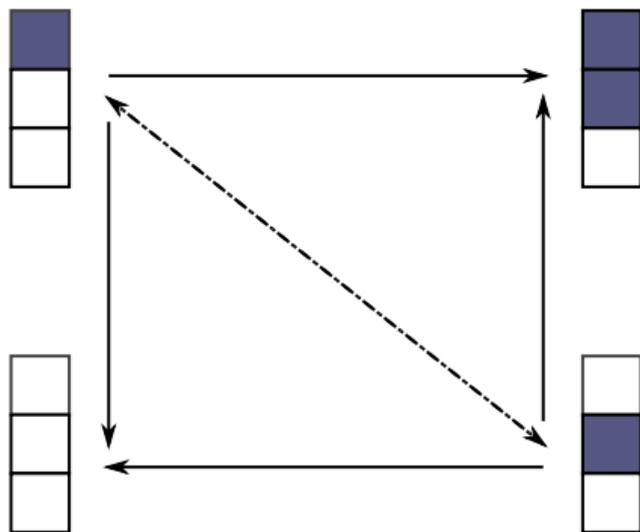
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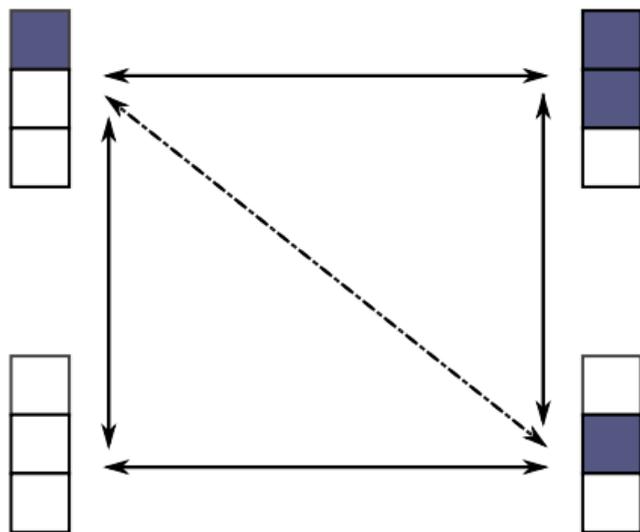
## Attractors are hypercubes



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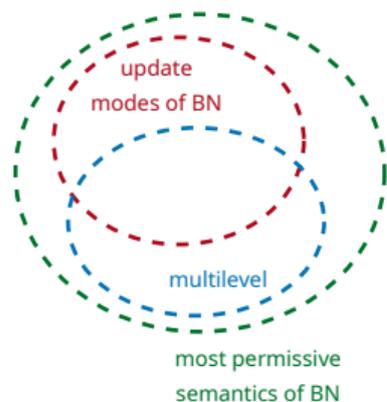
## Attractors are hypercubes



Attractors of most permissive semantics = minimal trap spaces

Existence of attractor within hypercube is NP-complete

# 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** !

## Applications

Prototype python library + ASP (SAT) implementation

<https://github.com/pauleve/mpbn>

### **Boolean network synthesis from reachability properties**

⇒ becomes NP

⇒ CaspoTS implements most permissive reachability (ASP)

<https://github.com/bioasp/caspots>

### **Computation of reachable attractors**

⇒ In the order of ms for networks tested so far (~100 nodes)

WiP with most permissive semantics:

- model synthesis from differentiation data [Stéphanie Chevalier]
- prediction for cellular reprogramming

## Conclusion

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

- difficult to justify (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 adding information (multilevel, thresholds) will only remove behaviours
  - **simpler complexity**: reachability PTIME, attractors NP
- ⇒ much higher tractability

Future work: most permissive for multilevel networks