



# Inference of Boolean networks from gene interaction graphs using a SAT solver

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Universidad Nacional Autónoma de México



# Outline

Boolean  
network  
inference

Griffin  
(prototype):  
Input and  
Output

Boolean  
network  
Interaction  
graph  
(topology)  
Block  
diagram

Boolean-  
network  
inference  
from  
interaction  
graph

Step 1: Entry  
variables  
Step 2:  
Incorporation  
of regulators  
Incremental  
SAT solver  
Step 3:  
Formulas

Practical

- 1 Griffin (prototype): Input and Output
  - Boolean network
  - Interaction graph (topology)
  - Block diagram
- 2 Boolean-network inference from interaction graph
  - Step 1: Entry variables
  - Step 2: Incorporation of regulators
  - Incremental SAT solver
  - Step 3: Formulas
- 3 Practical example
- 4 Concluding remarks



# 1. Griffin (prototype): Input and Output Boolean network (synchronous Boolean network)

Boolean  
network  
inference

synchronous *Boolean network* (next state)

$$\bar{x} = (x_1, x_2, x_3) \quad f(\bar{x}) = (f_1(\bar{x}), f_2(\bar{x}), f_3(\bar{x}))$$

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# 1. Griffin (prototype): Input and Output

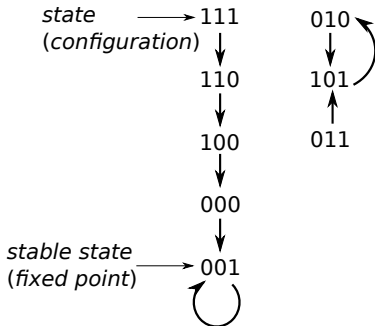
Boolean network (synchronous Boolean network)

Boolean  
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## 1 state graph (dynamics)



Griffin (prototype): Input and Output

Boolean network

Interaction graph (topology)

Block diagram

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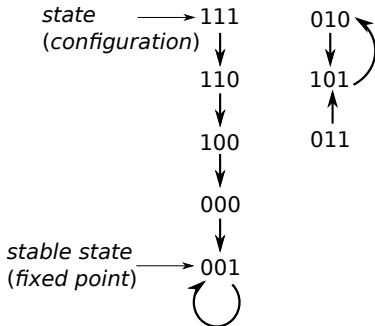
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① *state graph (dynamics)*

② *truth table* for next value of each bit



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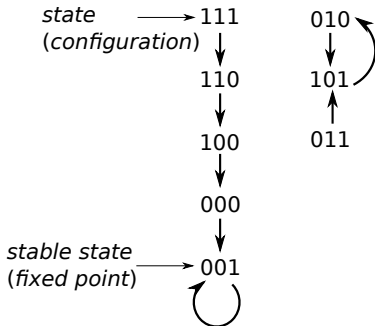
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- 1 *state graph (dynamics)*
- 2 *truth table* for next value of each bit



- 3 *Boolean formula* for next value of each bit

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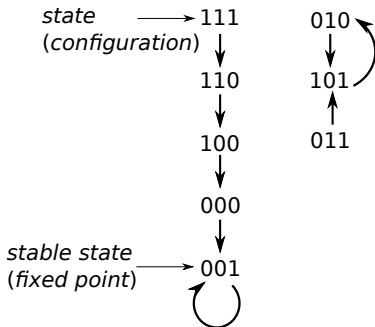
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- 1 *state graph (dynamics)*
- 2 *truth table* for next value of each bit



- 3 *Boolean formula* for next value of each bit

- gene regulatory networks
- pathway signaling networks
- cell cycles

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Interaction graph (topology)

Boolean  
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Input and  
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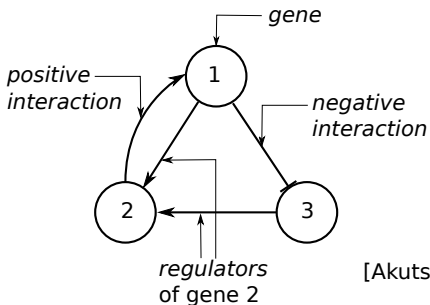
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## Interaction graph



[Akutsu et al. 1999]

meaning of interaction?

interaction graph, underspecification





# 1. Griffin: Input and Output

Meaning of positive and negative interactions

Boolean  
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*positive*

$i \rightarrow j$

same pattern  
of 0s and 1s

	$i$	$f_j$
	0	0
	1	1

truth table for gene  $j$

*negative*

$i \neg j$

	$i$	$f_j$
	0	1
	1	0

truth table for gene  $j$

possible to have both positive and  
negative regulation (*ambiguous*)



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Meaning of positive and negative interactions

Boolean network inference

Griffin (prototype): Input and Output

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possible to have both positive and negative regulation (*ambiguous*)

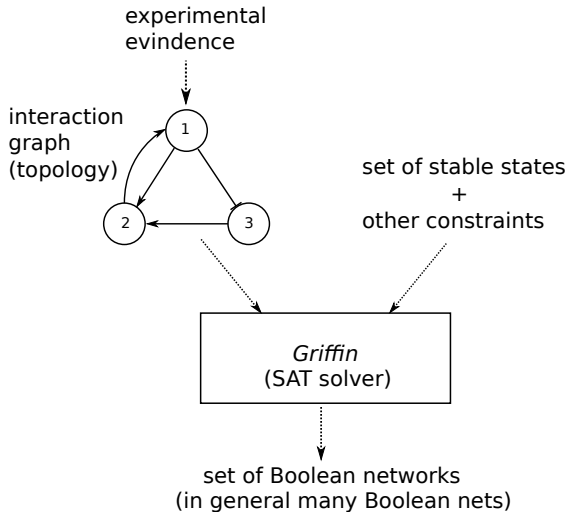
$$i \rightarrow j \text{ iff } \exists \bar{x}. f_j(\bar{x}[1/x_i]) > f_j(\bar{x}[0/x_i])$$

$$i \neg j \text{ iff } \exists \bar{x}. f_j(\bar{x}[1/x_i]) < f_j(\bar{x}[0/x_i])$$



# 1. Griffin: Input and Output

Block diagram



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## 2. Boolean-network inference

### Step 1: Entry variables

truth table of Boolean networks  $f$  represented with “entry” variables

$x$	$f(x)$		
0 0 0	$f_1000$	$f_2000$	$f_3000$
0 0 1	$f_1001$	$f_2001$	$f_3001$
0 1 0	$f_1010$	$f_2010$	$f_3010$
0 1 1	$f_1011$	$f_2011$	$f_3011$
1 0 0	$f_1100$	$f_2100$	$f_3100$
1 0 1	$f_1101$	$f_2101$	$f_3101$
1 1 0	$f_1110$	$f_2110$	$f_3110$
1 1 1	$f_1111$	$f_2111$	$f_3111$

Boolean  
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**Step 1: Entry  
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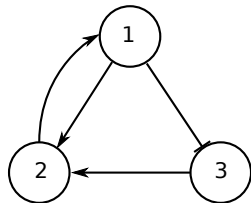


## 2. Boolean-network inference

### Step 2: Incorporation of regulators

$x$	$f(x)$		
0 0 0	$f_1000$	$f_2000$	$f_3000$
0 0 1	$f_1001$	$f_2001$	$f_3001$
0 1 0	$f_1010$	$f_2010$	$f_3010$
0 1 1	$f_1011$	$f_2011$	$f_3011$
1 0 0	$f_1100$	$f_2100$	$f_3100$
1 0 1	$f_1101$	$f_2101$	$f_3101$
1 1 0	$f_1110$	$f_2110$	$f_3110$
1 1 1	$f_1111$	$f_2111$	$f_3111$

interaction graph



**representative:**

for each column: same color = same variable

$\text{repr}(f_3000) = \text{repr}(f_3001) = \text{repr}(f_3010) = \text{repr}(f_3011)$

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## 2. Boolean-network inference incremental SAT solver

Boolean  
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Boolean formula in *conjunctive normal form*:

$$(l_0 \text{ or } \dots \text{ or } l_1) \text{ and } \dots \text{ and } (l_{r-1} \text{ or } \dots \text{ or } l_r)$$

where:

$$l_i = \begin{cases} x_i \text{ or} \\ \text{not } x_i \end{cases}$$

Find values (0, 1) for each  $x_i$  such that value of formula is 1.

Phenomenal progress in last 15 years: millions of variables for first solution.



## 2. Boolean-network inference

### Interaction formulas

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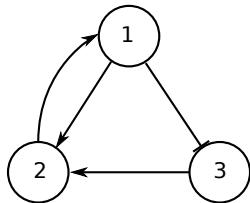
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$x$	$f(x)$		
0 0 0	$f_1000$	$f_2000$	$f_3000$
0 0 1	$f_1000$	$f_2001$	$f_3000$
0 1 0	$f_1010$	$f_2000$	$f_3000$
0 1 1	$f_1010$	$f_2001$	$f_3000$
1 0 0	$f_1000$	$f_2100$	$f_3100$
1 0 1	$f_1000$	$f_2101$	$f_3100$
1 1 0	$f_1010$	$f_2100$	$f_3100$
1 1 1	$f_1010$	$f_2101$	$f_3100$

interaction graph



$2 \rightarrow 1$ :

$$\varphi_{RG}^+(2, 1) = (\text{not } f_1000) \text{ and } f_1010$$



## 2. Boolean-network inference

### Interaction formulas

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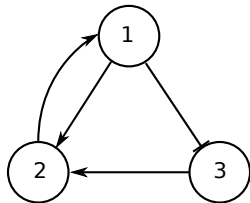
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0 1 1	$f_1010$	$f_2001$	$f_3000$
1 0 0	$f_1000$	$f_2100$	$f_3100$
1 0 1	$f_1000$	$f_2101$	$f_3100$
1 1 0	$f_1010$	$f_2100$	$f_3100$
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interaction graph



$2 \rightarrow 1:$

$$\varphi_{RG}^+(2, 1) = (\text{not } f_1000) \text{ and } f_1010$$





## 2. Boolean-network inference

### Interaction formulas

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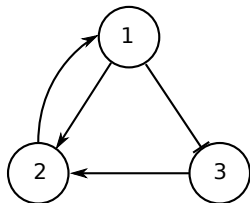
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interaction graph



$2 \rightarrow 1$ :

$$\varphi_{RG}^+(2, 1) = (\text{not } f_1000) \text{ and } f_1010$$



## 2. Boolean-network inference

### Interaction formulas

$$\varphi_{R_G}^+(2, 1) = (\text{not } f_1000) \text{ and } f_1010$$

$$\varphi_{R_G}^+(1, 2) = ((\text{not } f_2000) \text{ and } f_2100)$$

$$\text{or } ((\text{not } f_2001) \text{ and } f_2101)$$

$$\varphi_{R_G}^+(3, 2) = (\text{not } f_2000) \text{ and } f_2001$$

$$\text{or } ((\text{not } f_2100) \text{ and } f_2101)$$

$$\varphi_{R_G}^-(1, 3) = f_3000 \text{ and } (\text{not } f_3100)$$

$$\varphi_{R_G} = \varphi_{R_G}^+(1, 2) \text{ and } \varphi_{R_G}^+(2, 1) \text{ and } \varphi_{R_G}^+(3, 2) \text{ and } \varphi_{R_G}^-(1, 3)$$

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## 2. Boolean-network inference

### Interaction formulas

$$\varphi_{R_G}^+(2, 1) = (\text{not } f_1000) \text{ and } f_1010$$

$$\varphi_{R_G}^+(1, 2) = ((\text{not } f_2000) \text{ and } f_2100)$$

$$\text{or } ((\text{not } f_2001) \text{ and } f_2101)$$

$$\varphi_{R_G}^+(3, 2) = (\text{not } f_2000) \text{ and } f_2001$$

$$\text{or } ((\text{not } f_2100) \text{ and } f_2101)$$

$$\varphi_{R_G}^-(1, 3) = f_3000 \text{ and } (\text{not } f_3100)$$

$$\varphi_{R_G} = \varphi_{R_G}^+(1, 2) \text{ and } \varphi_{R_G}^+(2, 1) \text{ and } \varphi_{R_G}^+(3, 2) \text{ and } \varphi_{R_G}^-(1, 3)$$

*symbolic* method (represents the set of all desired networks as a Boolean formula)

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## 2. Boolean-network inference

### Stationary-states formula

Boolean  
network  
inference

001 is a stationary state:

$$\varphi_{\text{FP}}(G, x) = (\text{not } f_1000) \text{ and } (\text{not } f_2001) \text{ and } f_3000$$

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## 2. Boolean-network inference

### Stationary-states formula

Boolean  
network  
inference

001 is a stationary state:

$$\varphi_{\text{FP}}(G, x) = (\text{not } f_1000) \text{ and } (\text{not } f_2001) \text{ and } f_3000$$

001 is the only stationary state:

$$\varphi_{\text{NFP}} = \varphi_{\text{FP}} \text{ and}$$

**not** ((**not**  $f_1000$ ) **and** (**not**  $f_2000$ ) **and** (**not**  $f_3000$ )) **and**

**not** ((**not**  $f_1010$ ) **and**  $f_2000$  **and** (**not**  $f_3000$ )) **and**

**not** ((**not**  $f_1010$ ) **and**  $f_2001$  **and**  $f_3000$ ) **and**

**not** ( $f_1000$  **and** (**not**  $f_2100$ ) **and** (**not**  $f_3100$ )) **and**

...

**not** ( $f_1010$  **and**  $f_2101$  **and**  $f_3100$ )

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## 2. Boolean-network inference

### Formulas

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Formulas:

- exact set of stationary states (exponential in number of genes)
- possibly other (hypothetical) stationary states
- exact set of interactions
- possibly other (hypothetical) interactions
- nonambiguity (no positive and negative interactions)
- stationary states of mutations



## 2. Boolean-network inference

### Formulas

#### Boolean network inference

Griffin (prototype):  
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Boolean-network inference from interaction graph

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Formulas:

- exact set of stationary states (exponential in number of genes)
- possibly other (hypothetical) stationary states
- exact set of interactions
- possibly other (hypothetical) interactions
- nonambiguity (no positive and negative interactions)
- stationary states of mutations

For each interaction:

- one of {mandatory, hypothetical},
- one of {positive, negative, unknown sign},
- one of {ambiguous, nonambiguous}.



### 3. Practical example

*Arabidopsis thaliana* root stem cell niche

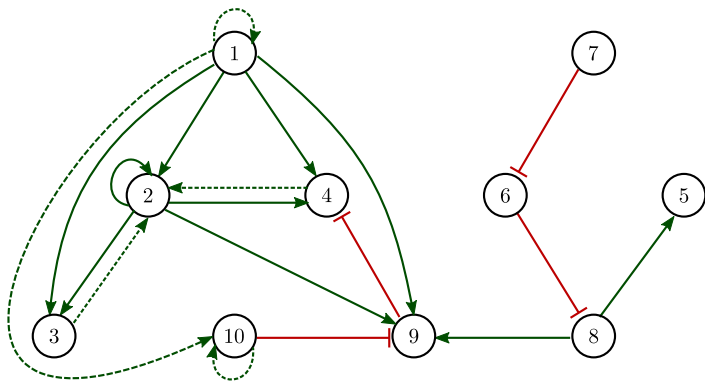
Boolean network inference

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no mutations: By hand, 2 networks. *Griffin*, 74 networks.

mutations: By hand, 1 network. *Griffin*, 3 networks.





## 4. Concluding remarks

### Related work

#### Boolean network inference

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Practical

- 1 [Akutsu, Miyano and Kuhara, 1999]
  - entry variables
  - input: set of “state/next-state” pairs ( $\sim$  time series)
  - output: Boolean networks
  - exhaustive search (nonsymbolic)
- 2 [Azpeitia, Weinstein, Benítez, Mendoza and Alvarez-Buylla, 2013]
  - input: interaction graph (topology)
  - output: Boolean networks
  - search with random component (nonsymbolic)



## 4. Concluding remarks

### Boolean-network inference and symbolic methods

- 1 interaction graph (topology)  $\rightarrow$  Boolean network is important
  - 1 root stem cell niche *A. thaliana*
  - 2 flower *A. thaliana*
  - 3 yeast cell cycle
  - 4 mammalian cell cycle
  - 5 T-cell receptor signaling
  - 6 T-cell differentiation
  - 7 vulva *C. elegans*
- 2 search space is vast
- 3 only possible with symbolic methods (SAT solvers, binary decision diagrams (BDDs))

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graph  
(topology)  
Block  
diagram

Boolean-  
network  
inference  
from  
interaction  
graph

Step 1: Entry  
variables  
Step 2:  
Incorporation  
of regulators  
Incremental  
SAT solver  
Step 3:  
Formulas

Practical



# 4. Concluding remarks

## Future work

Boolean  
network  
inference

Griffin  
(prototype):  
Input and  
Output

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## Future work

- 1 multi-valued genes
- 2 more efficient method for exact set of stable states
- 3 asynchronous networks



# 4. Concluding remarks

## Summary

Boolean  
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Griffin  
(prototype):  
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## Summary

- 1 Boolean-network inference from interaction graph is important
- 2 search space is vast
- 3 forthcoming networks will be large
- 4 symbolic methods (representing set of desired solutions as Boolean formula)
- 5 we presented a method for symbolically finding synchronous Boolean networks having a number of properties
- 6 we wish to extend our method to:
  - 1 multi-valued genes
  - 2 asynchronous networks