### Constraint-Based Methods for Bioinformatics Workshop

## Identification of Bifurcations in Biological Regulatory Networks using Answer-Set Programming

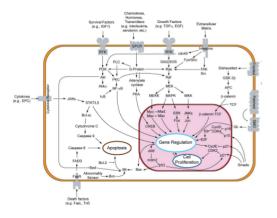
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#### Joint work with:

Olivier Roux<sup>1</sup>, Carito Guziolowski<sup>1</sup> and Loïc Paulevé<sup>2</sup>

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## Context

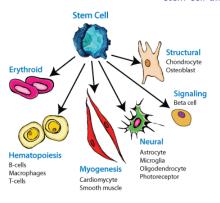


[Liu et al, in Journal of Bioinformatics and Computational Biology, 2012]

- Cellular processes are driven by networks of biological interactions.
- Formal modelling and analysis of Biological Regulatory Network.
- Static analysis of properties.

## Motivation

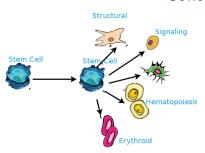
#### stem cell differentiation



[https://www.systembio.com/stem-cell-research/differentiation-reporters/overview]

- Loss of capability differentiation
- Which transitions (operations) are responsible of Bifurcations ?
- From which state?

## Contribution

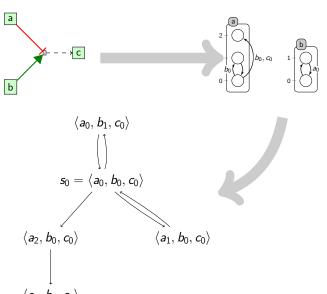


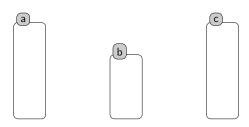
- Bifurcation transitions can be expressed in temporal logic formula
- "Relaxation":

  verification of CTL formula (PSPACE (complete))

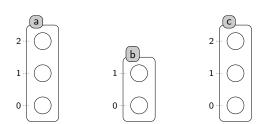
  ND problem which can be easily avgressed in SAT/A:
  - $\Rightarrow$  NP problem which can be easily expressed in SAT/ASP
- Contribution :

## Biological networks modelling



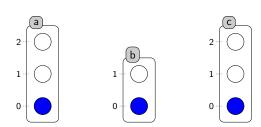


Automata: components a, b, c



Automata: components a, b, c

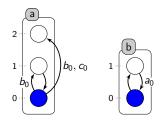
local states: levels of expression  $c_0$ ,  $c_1$ ,  $c_2$ 

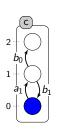


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States: sets of active local states  $\langle a_0, b_0, c_0 \rangle$ 





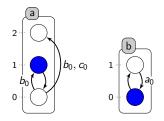
$$s_0=\langle a_0,b_0,c_0\rangle$$

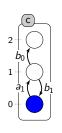
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local states: levels of expression  $c_0$ ,  $c_1$ ,  $c_2$ 

States: sets of active local states  $\langle a_0, b_0, c_0 \rangle$ 

Transitions: dynamics  $t_1=a_0\xrightarrow{b_0}a_1$ ,  $t_2=a_1\to a_0$ ,  $t_3=a_0\xrightarrow{b_0,c_0}a_2$ ,  $t_4=b_0\to b_1$ 





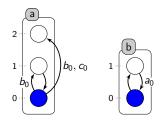


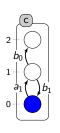
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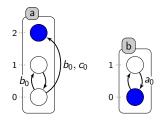
$$s_0 = \langle a_0, b_0, c_0 \rangle$$
  $\langle a_1, b_0, c_0 \rangle$ 

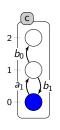
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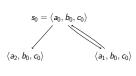
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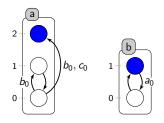


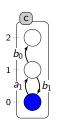
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Automata: components a, b, c
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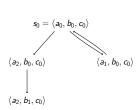
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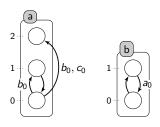


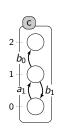
Automata: components a, b, c

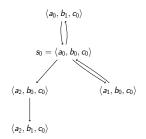
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Automata: components a, b, c

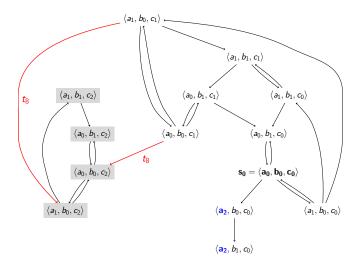
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States: sets of active local states

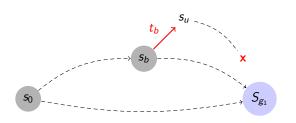
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$$t_4 = b_0 \rightarrow b_1$$

## Illustration of bifurcations



## Definition of bifurcation



## Definition

 $t_b$  is a bifurcation transition from  $s_0$  to  $g_1$ 

 $\iff$  there exists  $s_b$  such that:

(C1) 
$$s_u \not\rightarrow^* g_1$$
 (C2)  $s_b \rightarrow^* g_1$  (C3)  $s_0 \rightarrow^* s_b$ 

## Formal approximation of reachability

Static analysis by abstract interpretation

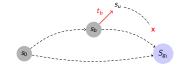
The reachability approximations for ANs introduced in [Paulevé, Magnin, Roux in *Mathematical Structures in Computer Science*, 2012]

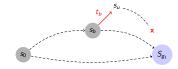
$$\mathsf{UA}(s \to^* s') \Rightarrow \mathbf{s} \to^* \mathbf{s}' \Rightarrow \mathsf{OA}(s \to^* s')$$

- OA (over-approximations): necessary conditions for  $s \to^* s'$
- UA (under-approximation): sufficient conditions for  $s \to^* s'$  (but the converse does not hold in general)

#### Interest:

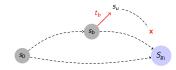
- Avoid state space explosion
- Decide in an efficient way reachability properties





(C1) 
$$s_b \cdot t_b \not\rightarrow^* g_1$$

(C1) 
$$s_b \cdot t_b \not\rightarrow^* g_1$$
 (I1<sup>#</sup>)  $\neg OA(s_u \rightarrow^* g_1)$  (I1<sup>#</sup>)  $\Rightarrow$  (C1)



(C1) 
$$s_b \cdot t_b \not\rightarrow^* g_1$$

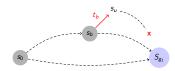
$$(\mathsf{I}\mathsf{1}^\#) \neg \mathsf{OA}(s_u \to^* g_1) \qquad \qquad (\mathsf{I}\mathsf{1}^\#) \Rightarrow (\mathsf{C}\mathsf{1})$$

$$(\mathsf{I}1^\#)\Rightarrow (\mathsf{C}1)$$

(C2) 
$$s_b \rightarrow^* g_1$$

$$(\mathsf{I2}^\#) \ \mathsf{UA}(s_b \to^* g_1)$$

$$(12^{\#}) \Rightarrow (C2)$$



(C1) 
$$s_b \cdot t_b \not\rightarrow^* g_1$$

$$(\mathsf{I}\mathsf{1}^\#) \neg \mathsf{OA}(s_u \to^* g_1)$$

$$(11^{\#}) \Rightarrow (C1)$$

(C2) 
$$s_b \rightarrow^* g_1$$

$$(\mathsf{I2}^\#) \ \mathsf{UA}(\mathsf{s}_b \to^* \mathsf{g}_1)$$

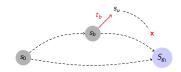
$$(12^{\#}) \Rightarrow (C2)$$

(C3) 
$$s_0 \rightarrow^* s_b$$

(13) 
$$s_b \in \operatorname{reach}(s_0)$$

$$\text{(I3$^\#$)} \ \ \mathsf{UA}(s_0 \to^* s_b)$$

$$(13^{\#}) \Rightarrow (C3)$$



(C1) 
$$s_b \cdot t_b \not\rightarrow^* g_1$$

$$(\mathsf{I}\mathsf{1}^\#) \neg \mathsf{OA}(s_u \to^* g_1)$$

$$(I1^{\#}) \Rightarrow (C1)$$

(C2) 
$$s_b \rightarrow^* g_1$$

$$(12^\#) \quad \mathsf{UA}(s_b \to^* g_1)$$

$$(12^{\#}) \Rightarrow (C2)$$

(C3) 
$$s_0 \rightarrow^* s_b$$

(13) 
$$s_b \in \operatorname{reach}(s_0)$$

(
$$13^{\#}$$
) UA( $s_0 \rightarrow^* s_b$ )

$$(13^{\#}) \Rightarrow (C3)$$

$$(11^{\#})$$
 and  $(12^{\#})$  and  $((13)or(13^{\#})) \Rightarrow t_b$  is a bifurcation.

## **Implementation**

#### Formal approximation of reachability

- Analysis of local causality of transitions
- Computation of a so called Local Causality Graph
- OA/UA: particular patterns in LCG
- LCG size: poly(#automata),exp(|single automaton|)

#### **ASP** implementation

• Encode NP problem: find  $s_b$ ,  $t_b$  such that

$$\neg \mathsf{OA}(s_b \cdot t_b \to^* g_1)$$
 and  $\mathsf{UA}(s_b \to^* g_1)$  and  $\mathsf{UA}(s_0 \to^* s_b)$ 

• enumeration with clingo

## Automata Network modelling of Biological Networks

#### Transition-centered specification

- ...in opposition to function-centered of Boolean/Thomas networks
- explicit context/ causality of state changes
- closely related to (safe) Petri nets
- step semantics (purely async, purely sync, mixed)

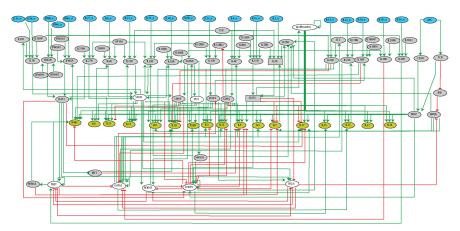
#### Modelling

- any Boolean/Thomas networks can be encoded;
- in case of logical rules uncertainty: model the union of Boolean/Thomas networks (over-approximation of behaviours)

## Case study

Lambda phage: (4 components and 11 interactions); EGF/TNF: (28 components and 55 interactions);

t\_helper differentiation: (101 components and 381 interactions).



[Abou-Jaoudé et al, in Frontiers in Bioengineering and Biotechnology, 2015]

## Results of identification of bifurcations

Automata Network	Goal	M-C (NuSMV)		with (I3)		with (I3 <sup>#</sup> )	
		$ t_b $	Time	$ t_b $	Time	$ t_b $	Time
Lambda phage	${ m CI_2}$	10	0.1 <i>s</i>	6	0.1 <i>s</i>	0	0.2s
$ \Sigma  = 4   T  = 11$	$Cro_2$	3	0.1 <i>s</i>	3	0.1 <i>s</i>	2	0.3 <i>s</i>
EGF/TNF	$NFkB_0$	5	0.2s	4	0.1 <i>s</i>	2	0.1 <i>s</i>
$ \Sigma  = 28   T  = 55$	$IKB_1$	5	0.2s	3	0.1 <i>s</i>	2	0.1 <i>s</i>
Th_th17	$RORGT_1$	18	48 <i>s</i>	9	23 <i>s</i>	8	26 <i>s</i>
$ \Sigma  = 101   T  = 381$	BCL6 <sub>1</sub>	7	26 <i>s</i>	5	23 <i>s</i>	4	24 <i>s</i>
Th_HTG	BCL6 <sub>1</sub>	out-of-time		out-of-time		6	61 <i>s</i>
$ \Sigma  = 101   T  = 381$	GATA3 <sub>1</sub>					7	34 <i>s</i>

Implemented in ASP (Answer Set Programming) and solve with clingo 4.5.4.

## Conclusions & Perspectives

#### Summary

•

```
 \left\{ \begin{array}{l} \mathsf{AI} \ (\mathsf{Artificial} \ \mathsf{inteligence}) \\ + \\ \mathsf{AI} \ (\mathsf{Abstract} \ \mathsf{interpretation}) \end{array} \right. \Longrightarrow \mathsf{formal} \ \mathsf{approximation} \ \mathsf{of} \ \mathsf{bifurcations}
```

- Tractable on large networks (compared to model-checking)
- Under-approximation: some bifurcations are not returned.

#### **Perspectives**

- Over-approximation of bifurcations
- Use bifurcations for the analysis of probability
- Applications for predicting targets for cellular reprogramming.

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Thank you for your attention!