Constraint-Based Methods for Bioinformatics Workshop

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

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Joint work with:
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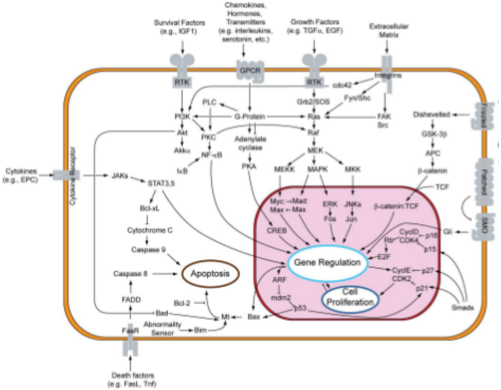
Context

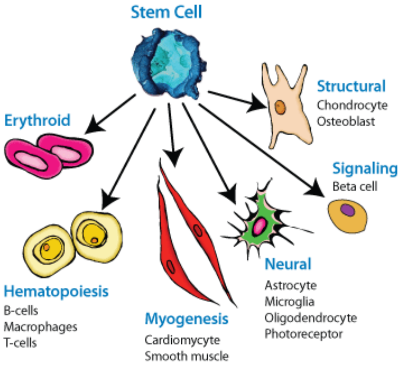


[Liu et al, in *Journal of Bioinformatics and Computational Biology*, 2012]

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

Motivation

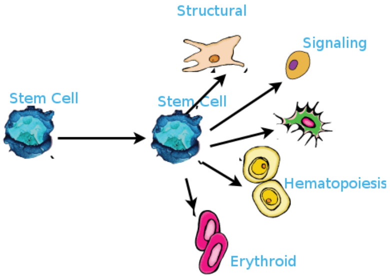
stem cell differentiation



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

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

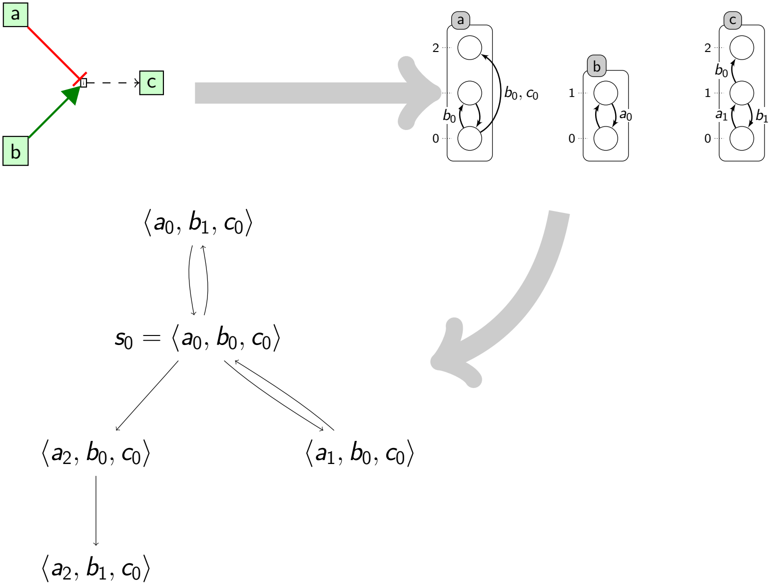
Contribution



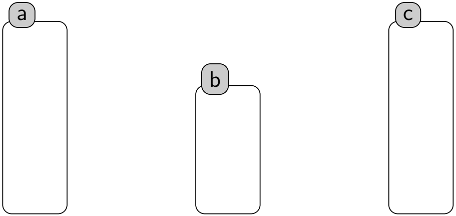
- Bifurcation transitions can be expressed in temporal logic formula
- **"Relaxation"** :
  verification of CTL formula (PSPACE (complete))
  $\Rightarrow$ NP problem which can be easily expressed in SAT/ASP
- **Contribution** :

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

Biological networks modelling

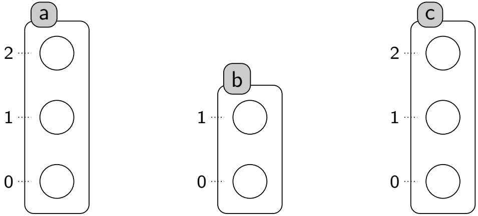


Automata Networks



Automata: components a, b, c

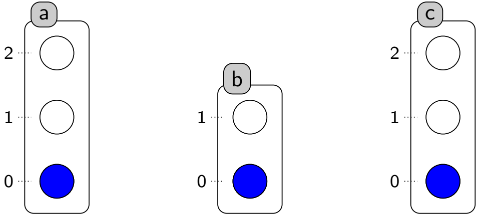
Automata Networks



Automata: components a, b, c

local states: levels of expression c_0, c_1, c_2

Automata Networks

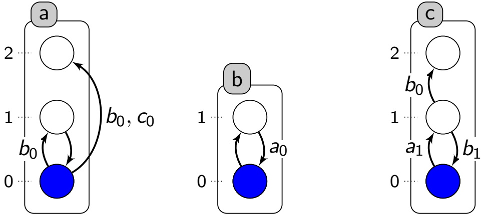


Automata: components a, b, c

local states: levels of expression c_0, c_1, c_2

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

Automata Networks



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

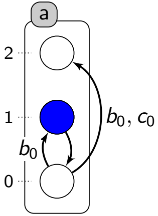
Automata: components $a,\ b,\ c$

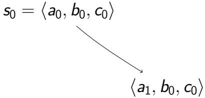
local states: levels of expression $c_0,\ c_1,\ c_2$

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

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

Automata Networks





Automata: components $a,\ b,\ c$

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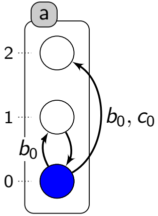

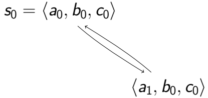

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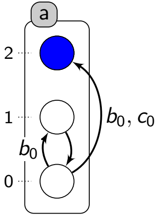

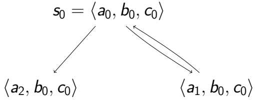

Automata: components $a,\ b,\ c$

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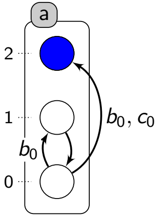

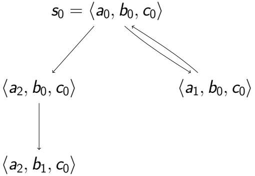

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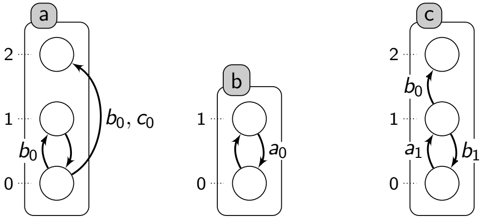

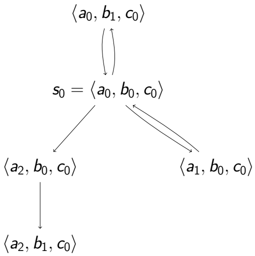

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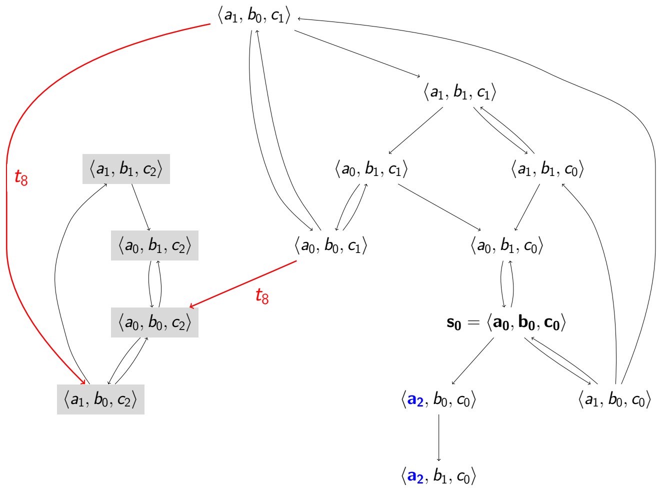
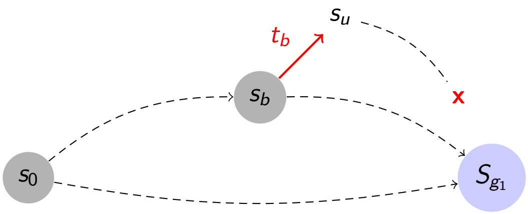
Illustration of bifurcations



Definition of bifurcation



Definition

t_b is a bifurcation transition from s_0 to g_1 if and only if 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]

$$\mathrm{UA}(s \rightarrow^* s') \Rightarrow \mathbf{s} \rightarrow^* \mathbf{s}' \Rightarrow \mathrm{OA}(s \rightarrow^* s')$$

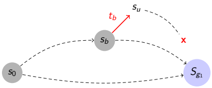
- OA (over-approximations): necessary conditions for $s \rightarrow^* s'$
- UA (under-approximation): sufficient conditions for $s \rightarrow^* s'$ (but the converse does not hold in general)

Interest:

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

General scheme for identification of bifurcation

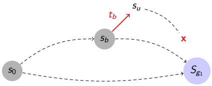
concepts and tools



#: sufficient conditions.

General scheme for identification of bifurcation

concepts and tools

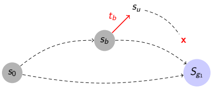


#: sufficient conditions.

(C1) $s_b \cdot t_b \not\rightarrow^* g_1$ (I1#) $\neg\,\mathrm{OA}(s_u \rightarrow^* g_1)$ (I1#) $\Rightarrow$ (C1)

General scheme for identification of bifurcation

concepts and tools



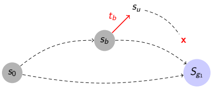
#: sufficient conditions.

(C1) $s_b \cdot t_b \not\rightarrow^* g_1$ (I1$^\#$) $\neg\,\mathrm{OA}(s_u \rightarrow^* g_1)$ (I1$^\#$) $\Rightarrow$ (C1)

(C2) $s_b \rightarrow^* g_1$ (I2$^\#$) $\mathrm{UA}(s_b \rightarrow^* g_1)$ (I2$^\#$) $\Rightarrow$ (C2)

General scheme for identification of bifurcation

concepts and tools



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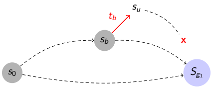
(C1) $s_b \cdot t_b \not\rightarrow^* g_1$	(I1#) $\neg\,\mathrm{OA}(s_u \rightarrow^* g_1)$	(I1#) $\Rightarrow$ (C1)
(C2) $s_b \rightarrow^* g_1$	(I2#) $\mathrm{UA}(s_b \rightarrow^* g_1)$	(I2#) $\Rightarrow$ (C2)
(C3) $s_0 \rightarrow^* s_b$	(I3) $s_b \in \mathrm{reach}(s_0)$	(I3) $\Leftrightarrow$ (C3)
	(I3#) $\mathrm{UA}(s_0 \rightarrow^* s_b)$	(I3#) $\Rightarrow$ (C3)

General scheme for identification of bifurcation

concepts and tools



$\#$: sufficient conditions.

(C1) $s_b \cdot t_b \not\to^* g_1$ (I1$^\#$) $\neg\,\mathrm{OA}(s_u \to^* g_1)$ (I1$^\#$) $\Rightarrow$ (C1)

(C2) $s_b \to^* g_1$ (I2$^\#$) $\mathrm{UA}(s_b \to^* g_1)$ (I2$^\#$) $\Rightarrow$ (C2)

(C3) $s_0 \to^* s_b$ (I3) $s_b \in \mathrm{reach}(s_0)$ (I3) $\Leftrightarrow$ (C3)

(I3$^\#$) $\mathrm{UA}(s_0 \to^* s_b)$ (I3$^\#$) $\Rightarrow$ (C3)

(I1$^\#$) and (I2$^\#$) and ((I3)or(I3$^\#$)) $\Rightarrow$ t_b is a bifurcation.

Automata Network modelling of Biological Networks

Transition-centered specification

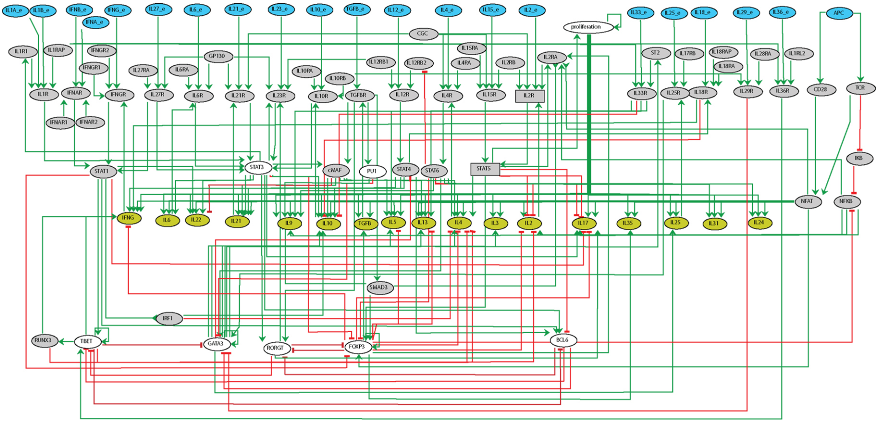
- ...in opposition to function-centered of Boolean/Thomas networks
- explicit context/ causality of state changes
- closely related to (safe) Petri nets
- step semantics (purely async, purely sync, mixed)

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)		(I3)		(I3$^{\#}$)	
		$\lvert t_b \rvert$	Time	$\lvert t_b \rvert$	Time	$\lvert t_b \rvert$	Time
Lambda phage	CI_2	10	$0.1s$	6	$0.1s$	0	$0.2s$
$\lvert\Sigma\rvert = 4 \quad \lvert T\rvert = 11$	Cro_2	3	$0.1s$	3	$0.1s$	2	$0.3s$
EGF/TNF	NFkB_0	5	$0.2s$	4	$0.1s$	2	$0.1s$
$\lvert\Sigma\rvert = 28 \quad \lvert T\rvert = 55$	IKB_1	5	$0.2s$	3	$0.1s$	2	$0.1s$
Th_th17	RORGT_1	18	$48s$	9	$23s$	8	$26s$
$\lvert\Sigma\rvert = 101 \quad \lvert T\rvert = 381$	$\mathrm{BCL6}_1$	7	$26s$	5	$23s$	4	$24s$
Th_HTG	$\mathrm{BCL6}_1$	out-of-time		out-of-time		6	$61s$
$\lvert\Sigma\rvert = 101 \quad \lvert T\rvert = 381$	$\mathrm{GATA3}_1$					7	$34s$

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

Conclusions & Perspectives

Summary

- $$\left\{\begin{array}{l}\text{AI (Artificial inteligence)}\\ +\\ \text{AI (Abstract interpretation)}\end{array}\right. \Longrightarrow \text{formal approximation of bifurcations}$$
- 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!