

École thématique BioRegul : Modélisation Formelle de Réseaux de Régulation
Biologique — 2023-06-06

Modélisation, analyse et inférence de paramètres pour les réseaux de régulation hybrides

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`http://maxime.folschette.name/`



Analysis of the Dynamics

- Centrale Nantes
PhD thesis
- 2011 → Efficient reachability analysis on large networks
 - 2014 → Dynamical patterns enumeration with answer set programming
- Univ Kassel
postdoc
- 2014 → Complex patterns enumeration with polyadic μ -calculus
 - 2015

Learning Models from Data

- Univ Nice
ATER
- 2015 → **Inference of constraints on hybrid parameters**
 - 2016
- Univ Nantes
ATER
- 2016 → Learning models from time series data
 - 2017

Learning New Knowledge from Models

- Univ Rennes
postdoc
- 2017 → Static computational model to study hepatocellular carcinoma progression
 - 2018
- CNRS/LS2N
postdoc
- 2018 → Integrate heterogeneous data with semantic web
 - 2019

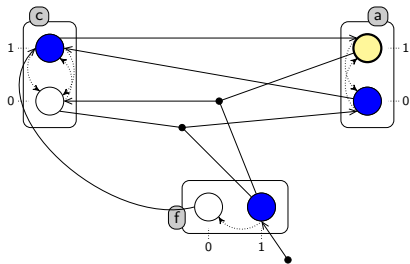
Today

- Centrale Lille
maître de conférences
- 2019 → Understand the role of glucose absorption in diabetes
 - ⋮ → Learn plankton food chains from measurements
 - **Formal verification of hybrid models**

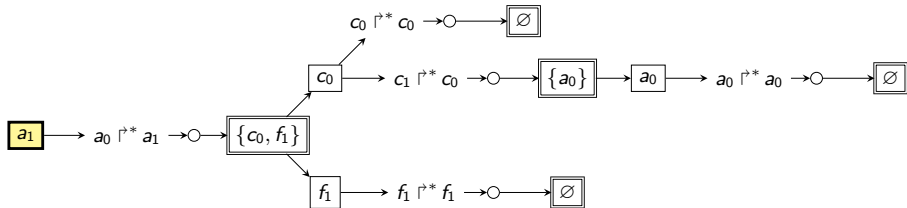
Other Works

Abstract Interpretation

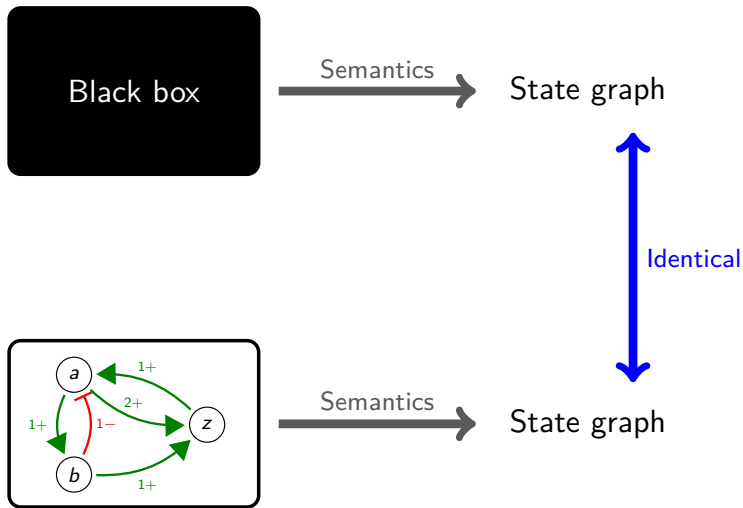
[Folschette *et al.*, *Theoretical Computer Science*, 2015b]



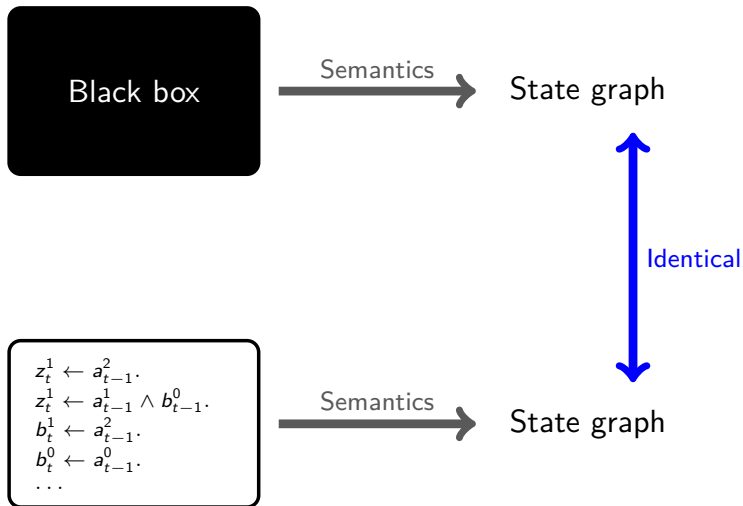
- No cycle
- No conflict
- All leaves are \emptyset



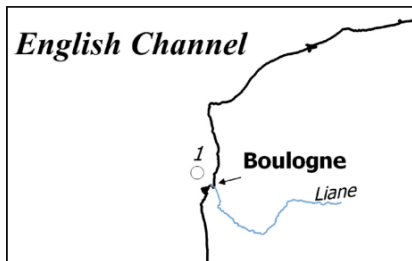
Learning Logic Programs for Explainability



Learning Logic Programs for Explainability



Long-term Datasets of Plankton Populations



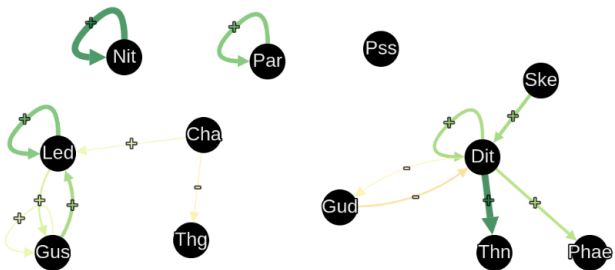
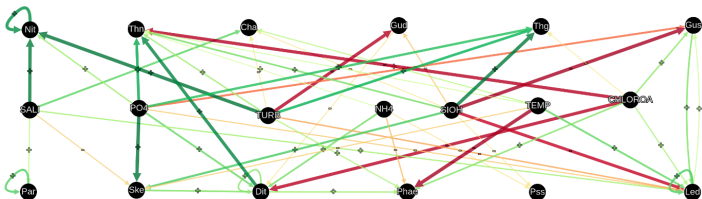
<https://www.seaonoe.org/data/00397/50832/>

Sampling location	Sampling date	Taxon	Value	Sampling depth
001-P-015	1992-05-18	CHLOROA	6.0	Surface (0-1m)
006-P-001	2019-12-02	Chaetoceros	1000.0	Surface (0-1m)
002-P-007	1994-05-25	Pleurosigma	100.0	Surface (0-1m)
002-P-030	2005-10-19	SALI	34.83	Surface (0-1m)
006-P-007	2015-09-28	Guinardia delicatula	11400.0	Surface (0-1m)

Environmental variables (7)

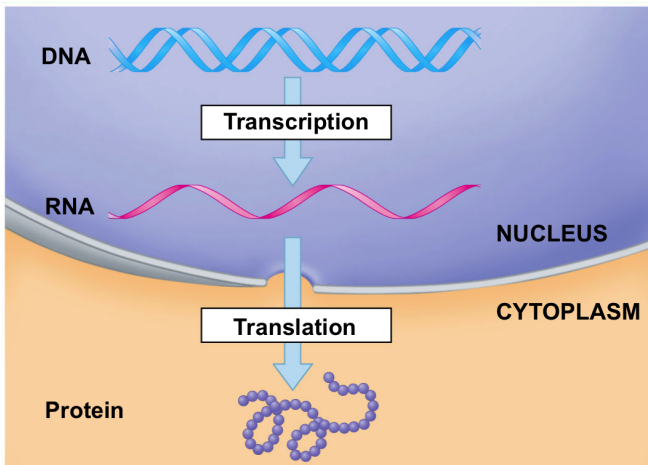
Phytoplankton species (12)

Learning Biotic Interactions



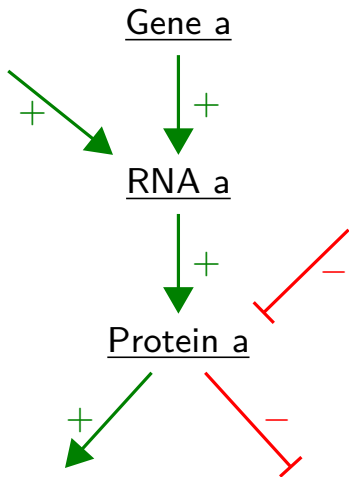
Discrete Models

Preliminary Abstraction

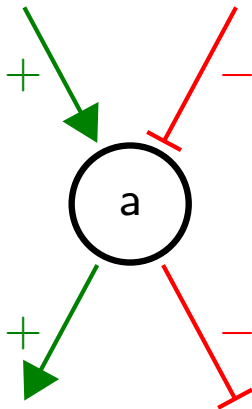


© 2012 Pearson Education, Inc.

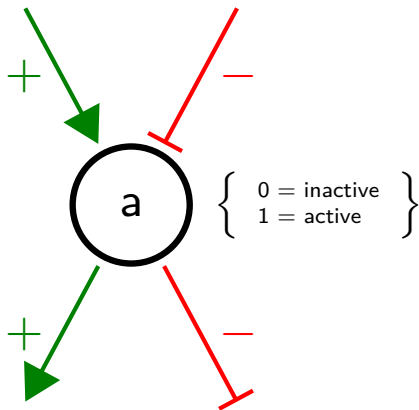
Preliminary Abstraction



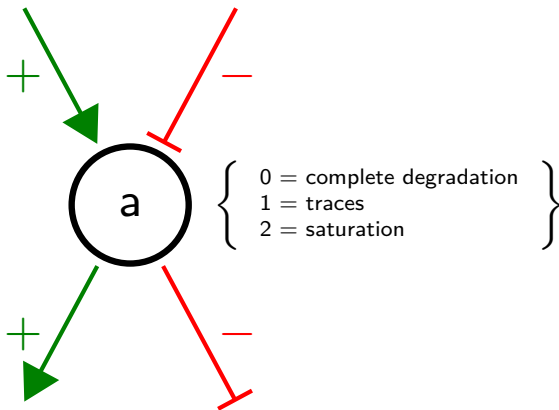
Preliminary Abstraction



Preliminary Abstraction



Preliminary Abstraction



State Graph

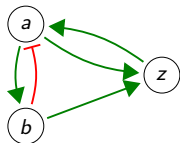
The state-graph depicts explicitly the whole dynamics

abz

000 010 001 011

100 110 101 111

200 210 201 211



State Graph

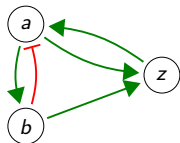
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100 \longrightarrow 110 101 111

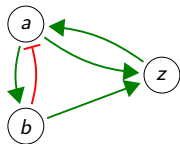
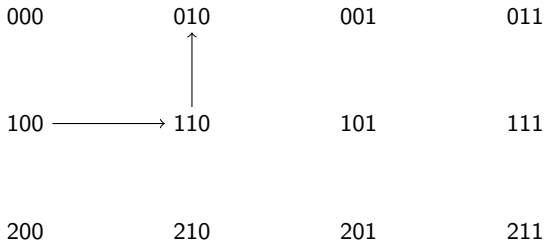
200 210 201 211



State Graph

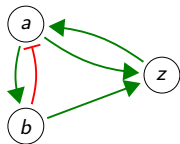
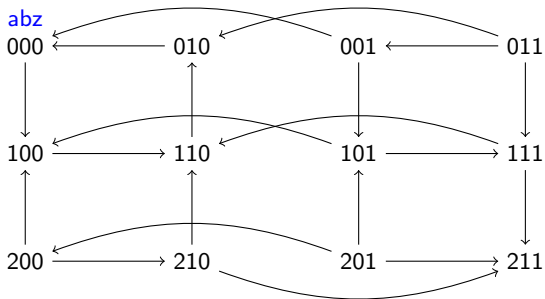
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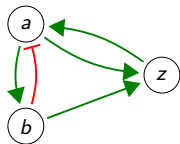
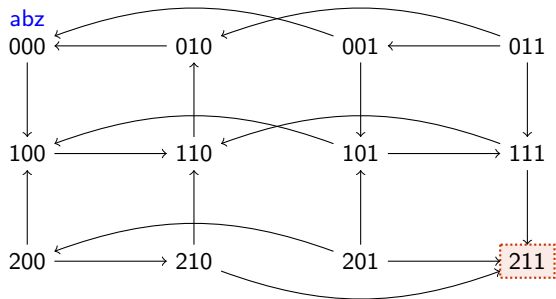
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State Graph

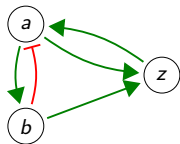
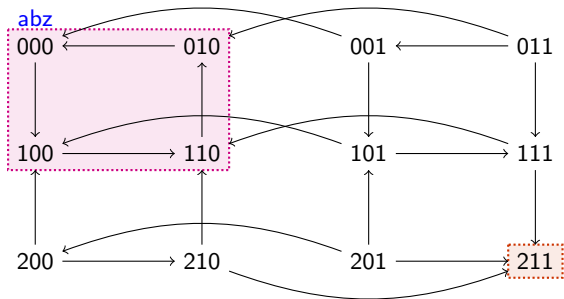
The state-graph depicts explicitly the whole dynamics



- **Stable state** = state with no successors

State Graph

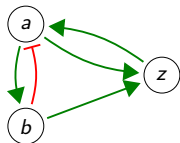
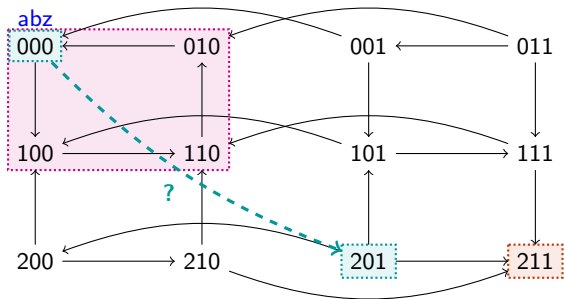
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- **Complex attractor** = minimal loop or composition of loops from which the dynamics cannot escape

State Graph

The state-graph depicts explicitly the whole dynamics



- **Stable state** = state with no successors
- **Complex attractor** = minimal loop or composition of loops from which the dynamics cannot escape
- **Reachability** = from **000**, can I reach **201**?

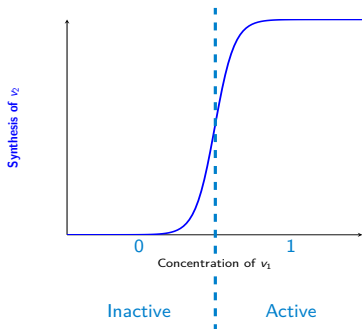
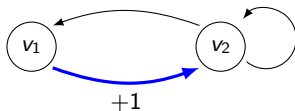
Origin of René Thomas Modeling

[Thomas, *Journal of Theoretical Biology*, 1973]



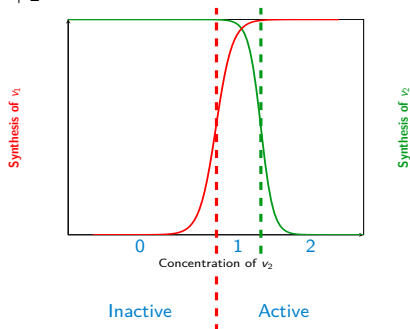
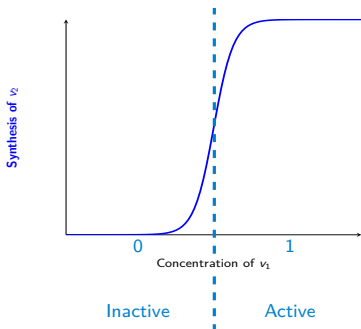
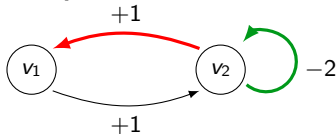
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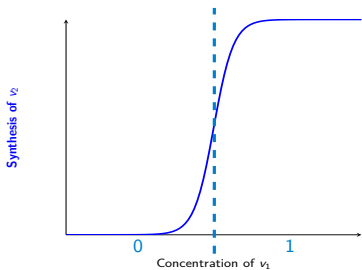
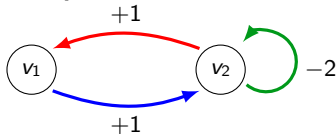
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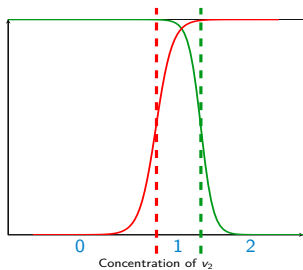


Inactive

Active

v_1 has no effect on v_2

v_1 triggers synthesis of v_2



Inactive

Active

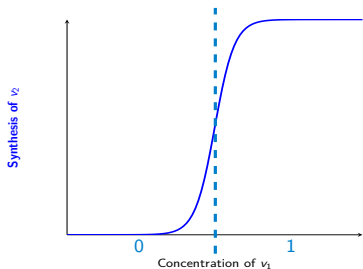
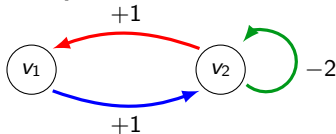
v_2 has no effect on v_1 or itself

v_2 triggers synthesis of v_1

v_2 triggers self-degradation

Origin of René Thomas Modeling

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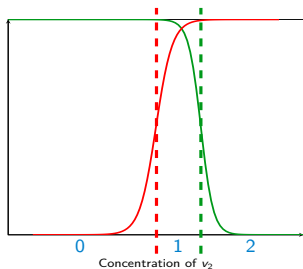


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Relative strength of $v_1 \xrightarrow{+1} v_2$ and $v_2 \xrightarrow{-2} v_2$? \implies Parameters!

Discrete Networks / Thomas Modeling

[Kauffman, *Journal of Theoretical Biology*, 1969]

[Thomas, *Journal of Theoretical Biology*, 1973]

- A set of components $N = \{a, b, z\}$

a

z

b

Discrete Networks / Thomas Modeling

[Kauffman, *Journal of Theoretical Biology*, 1969][Thomas, *Journal of Theoretical Biology*, 1973]

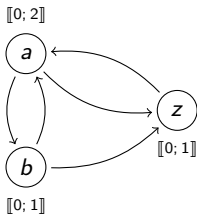
- A set of components $N = \{a, b, z\}$
- A discrete domain for each component $\text{dom}(a) = \llbracket 0; 2 \rrbracket$

 $\llbracket 0; 2 \rrbracket$ a z $\llbracket 0; 1 \rrbracket$ b $\llbracket 0; 1 \rrbracket$

Discrete Networks / Thomas Modeling

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- A set of components $N = \{a, b, z\}$
- A discrete domain for each component $\text{dom}(a) = \llbracket 0; 2 \rrbracket$
- Discrete parameters / evolution functions $K_{a,\omega}$



a	$K_{b,\omega}$
0	0
1	1
2	1

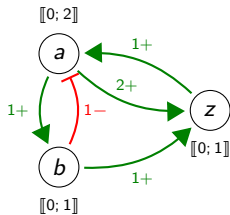
z	b	$K_{a,\omega}$
0	0	1
0	1	0
1	0	1
1	1	2

a	b	$K_{z,\omega}$
0	0	0
0	1	0
1	0	0
1	1	0
2	0	0
2	1	1

Discrete Networks / Thomas Modeling

[Kauffman, *Journal of Theoretical Biology*, 1969][Thomas, *Journal of Theoretical Biology*, 1973]

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- Signs & thresholds on the edges (redundant) $a \xrightarrow{2+} z$

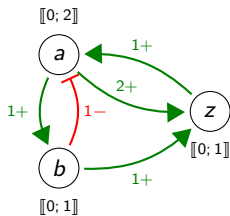


a	$K_{b,\omega}$	z	b	$K_{a,\omega}$	a	b	$K_{z,\omega}$
0	0	0	0	1	0	0	0
1	1	0	1	0	0	1	0
2	1	1	0	1	1	0	0
		1	1	2	1	1	0
					2	0	0
					2	1	1

Discrete Networks / Thomas Modeling

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a	$K_{b,\omega}$	z	b	$K_{a,\omega}$	a	b	$K_{z,\omega}$
0	0	0	0	1	0	0	0
1	1	0	1	0	0	1	0
2	1	1	0	1	1	0	0
		1	1	2	1	1	0
					2	0	0
					2	1	1

$$K_{v,\omega} \in \mathbb{N}$$

Parameters Identification

$$\text{Possible parametrizations} = \prod_{v \in N} |\text{dom}(v)| \left(\prod_{u \in \text{pred}(v)} |\text{dom}(u)| \right)$$



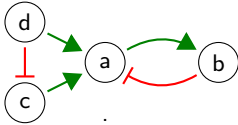
With Boolean variables, $|\text{dom}(v)| = 2$: $\left(2^{(2^{|\text{pred}(v)|})} \right)^{|N|}$

- Exponential in the number of nodes
- Double-exponential in the number of predecessors

To exhaustively and naively try out all parametrizations:

1. Assign one parametrization to the model
2. Compute the whole state-space (exponential !)
3. Test relevant properties (with model-checking)
4. Keep or discard this parametrization

Parameters Identification

Model	Possible parametrizations	Hypotheses
	16	$ \text{dom}(v) = 2$
	128	
	8192	
⋮	⋮	$ \text{pred}(v) = 1$
(10)	1048576	
(20)	1.1×10^{12}	
(100)	1.6×10^{60}	

Discrete Hoare Logic on Discrete Thomas Modeling

[Hoare, *Communications of the ACM*, 1969][Dijkstra, *Communications of the ACM*, 1975]**Hoare triple:** $\{ Pre \} p \{ Post \}$

- p is an imperative program
- Pre and $Post$ are properties (pre- and postcondition)

Meaning:“If Pre holds, then p can execute and $Post$ will hold after execution”**Example:** $\{ a = 0 \wedge b = 0 \} a+ ; b+ \{ a = 1 \wedge b > 0 \}$

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Meaning:“If Pre holds, then p can execute and $Post$ will hold after execution”**Example:** $\{ a = 0 \wedge b = 0 \} a+ ; b+ \{ a = 1 \wedge b > 0 \}$ **Weakest precondition calculus:**

Given p and $Post$, one can compute the weakest (most general) precondition $WPre$ so that $\{ WPre \} p \{ Post \}$ holds
 $WPre$ constrains the initial state of the system

Example: $\{ WPre \} a+ ; b+ \{ a = 1 \wedge b = 1 \}$
 $WPre \equiv a = 0 \wedge b = 0$

Discrete Hoare Logic on Discrete Thomas Modeling

[Hoare, *Communications of the ACM*, 1969][Dijkstra, *Communications of the ACM*, 1975][Bernot *et al.*, *Theoretical Computer Science*, 2015]**Hoare triple:** $\{ Pre \} p \{ Post \}$

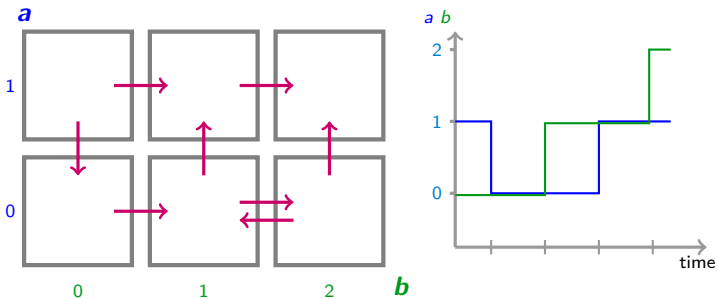
- p is an imperative program (known dynamical path from literature)
- Pre and $Post$ are properties (pre- and postcondition)

Meaning:"If Pre holds, then p can execute and $Post$ will hold after execution"**Example:** $\{ a = 0 \wedge b = 0 \} a+ ; b+ \{ a = 1 \wedge b > 0 \}$ **Weakest precondition calculus:**Given p and $Post$, one can compute the weakest (most general) precondition $WPre$ so that $\{ WPre \} p \{ Post \}$ holds $WPre$ constrains the initial state of the system and the parameters**Example:** $\{ WPre \} a+ ; b+ \{ a = 1 \wedge b = 1 \}$

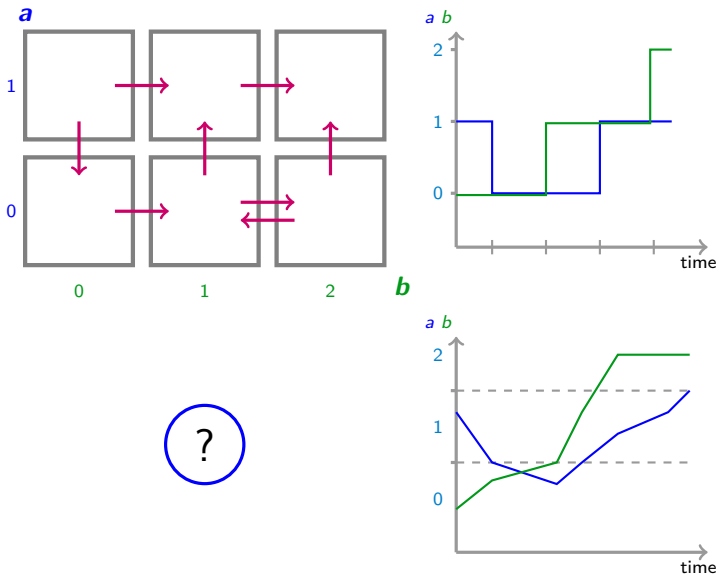
$$WPre \equiv a = 0 \wedge b = 0 \wedge K_{a,\{a=0,b=0\}} = 1 \wedge K_{b,\{a=1,b=0\}} = 1$$

Hybrid Models

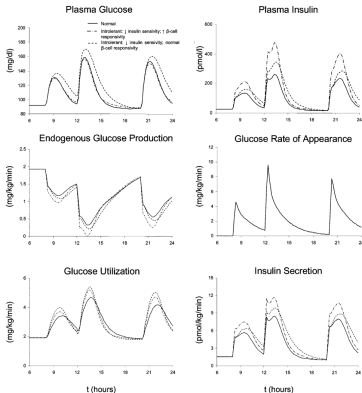
Objective of Hybrid Thomas Modeling



Objective of Hybrid Thomas Modeling



Differential Equations

[Dalla Man *et al.*, *IEEE Trans. Biomed. Eng.*, 2007]

$$\begin{cases} \dot{G}_p(t) = EGP(t) + Ra(t) - U_{ii}(t) - E(t) - k_1 \cdot G_p(t) + k_2 \cdot G_t(t) & G_p(0) = G_{pb} \\ \dot{G}_t(t) = -U_{id}(t) + k_1 \cdot G_p(t) - k_2 \cdot G_t(t) & G_t(0) = G_{tb} \\ G(t) = \frac{G_p}{V_G} & G(0) = G_b \end{cases}$$

$$\begin{cases} Q_{sto}(t) = Q_{sto1}(t) + Q_{sto2}(t) & Q_{sto}(0) = 0 \\ Q_{sto1}(t) = -k_{gri} \cdot Q_{sto1}(t) + D \cdot d(t) & Q_{sto1}(0) = 0 \\ Q_{sto2}(t) = -k_{empt}(Q_{sto}) \cdot Q_{sto2}(t) + k_{gri} \cdot Q_{sto1}(t) & Q_{sto2}(0) = 0 \\ \dot{Q}_{gut} = -k_{abs} \cdot Q_{gut}(t) + k_{empt}(Q_{sto}) \cdot Q_{sto2}(t) & Q_{gut}(0) = 0 \\ Ra(t) = \frac{f \cdot k_{abs} \cdot Q_{gut}(t)}{BW} & Ra(0) = 0 \end{cases}$$

$$\begin{cases} \dot{I}_l(t) = -(m_1 + m_3(t)) \cdot I_l(t) + m_2 I_p(t) + S(t) & I_l(0) = I_{lb} \\ \dot{I}_p(t) = -(m_2 + m_4) \cdot I_p(t) + m_1 \cdot I_l(t) & I_p(0) = I_{pb} \\ I(t) = \frac{I_p}{V_I} & I(0) = I_b \end{cases}$$

Comparison of Frameworks

Differential equations

- + Very precise simulations
- Difficult to tune (equations/parameters)
- Sometimes too complex to check simple properties

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- Abstraction: No continuous time/levels

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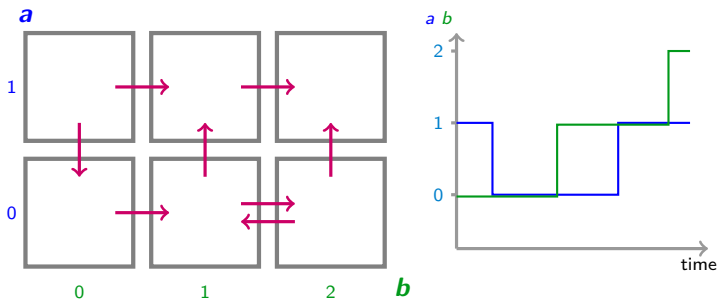
Hybrid models

- + Middle-ground: continuous time/levels
- + Not too complex in terms of parameters/simulation
- New formalism: requires new tools
 - To find parameters
 - To check dynamical properties

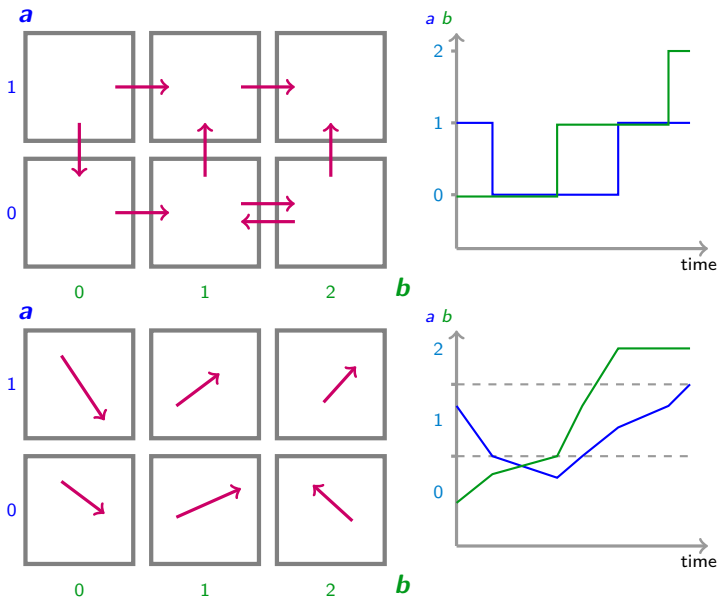
Discrete models

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Idea of Hybrid Thomas Modeling



Idea of Hybrid Thomas Modeling



Definitions

[Cornillon et al., *Mod. Complex Biol. Syst. in the Context of Genomics*, 2016]

Discrete state:

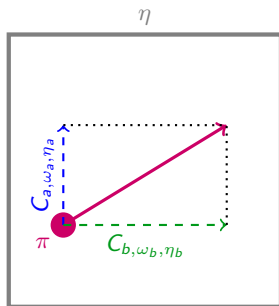
$$\eta = (\eta_a, \eta_b)$$

$$\eta \in \times_{u \in N} \text{dom}(u)$$

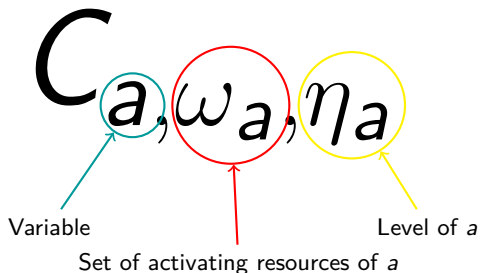
Fractional part in discrete state:

$$\pi = (\pi_a, \pi_b)$$

$$\pi \in [0, 1]^{|M|}$$



Celerity:



Hybrid Gene Regulatory Network (HGRN)

[Cornillon *et al.*, *Mod. Complex Biol. Syst. in the Context of Genomics*, 2016]

- A set of components $N = \{a, b, z\}$

a

z

b

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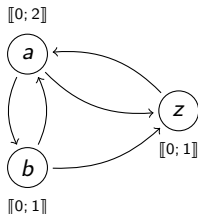
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 $\llbracket 0; 2 \rrbracket$ a z $\llbracket 0; 1 \rrbracket$ b $\llbracket 0; 1 \rrbracket$

Hybrid Gene Regulatory Network (HGRN)

[Cornillon et al., *Mod. Complex Biol. Syst. in the Context of Genomics*, 2016]

- A set of components $N = \{a, b, z\}$
- A discrete domain for each component $\text{dom}(a) = \llbracket 0; 2 \rrbracket$
- Real parameters C_{a, ω_a, η_a}

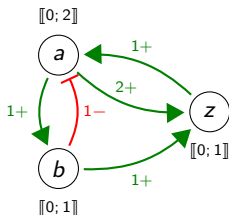


a	b	C_{b, ω_b, η_b}	a	b	z	C_{a, ω_a, η_a}	...
0	0	$C_{b, \emptyset, 0}$	0	0	0	$C_{a, \{b\}, 0}$	
0	1	$C_{b, \emptyset, 1}$	1	0	0	$C_{a, \{b\}, 1}$	
1, 2	0	$C_{b, \{a\}, 0}$	2	0	0	$C_{a, \{b\}, 2}$	
1, 2	1	$C_{b, \{a\}, 1}$	0	0	1	$C_{a, \{b, z\}, 0}$	
			1	0	1	$C_{a, \{b, z\}, 1}$	
			2	0	1	$C_{a, \{b, z\}, 2}$	
	0		0	1	0	$C_{a, \emptyset, 0}$	
	1		1	1	0	$C_{a, \emptyset, 1}$	
	2		2	1	0	$C_{a, \emptyset, 2}$	
	0		0	1	1	$C_{a, \{z\}, 0}$	
	1		1	1	1	$C_{a, \{z\}, 1}$	
	2		2	1	1	$C_{a, \{z\}, 2}$	

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- A set of components $N = \{a, b, z\}$
- A discrete domain for each component $\text{dom}(a) = \llbracket 0; 2 \rrbracket$
- Real parameters C_{a,ω_a,η_a}
- Signs & thresholds on the edges (redundant) $a \xrightarrow{2+} z$

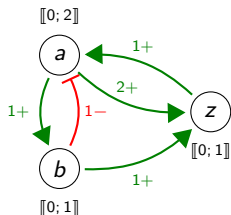


a	b	C_{b,ω_b,η_b}	a	b	z	C_{a,ω_a,η_a}	...
0	0	$C_{b,\emptyset,0}$	0	0	0	$C_{a,\{b\},0}$	
0	1	$C_{b,\emptyset,1}$	1	0	0	$C_{a,\{b\},1}$	
1, 2	0	$C_{b,\{a\},0}$	2	0	0	$C_{a,\{b\},2}$	
1, 2	1	$C_{b,\{a\},1}$	0	0	1	$C_{a,\{b,z\},0}$	
			1	0	1	$C_{a,\{b,z\},1}$	
			2	0	1	$C_{a,\{b,z\},2}$	
	0		0	1	0	$C_{a,\emptyset,0}$	
	1		1	1	0	$C_{a,\emptyset,1}$	
	2		2	1	0	$C_{a,\emptyset,2}$	
	0		0	1	1	$C_{a,\{z\},0}$	
	1		1	1	1	$C_{a,\{z\},1}$	
	2		2	1	1	$C_{a,\{z\},2}$	

Hybrid Gene Regulatory Network (HGRN)

[Cornillon et al., Mod. Complex Biol. Syst. in the Context of Genomics, 2016]

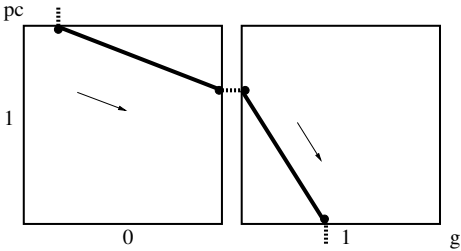
- A set of components $N = \{a, b, z\}$
- A discrete domain for each component $\text{dom}(a) = \llbracket 0; 2 \rrbracket$
- Real parameters C_{a,ω_a,η_a}
- Signs & thresholds on the edges (redundant) $a \xrightarrow{2+} z$



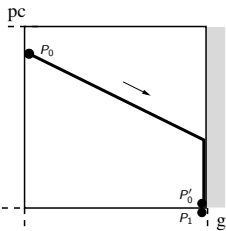
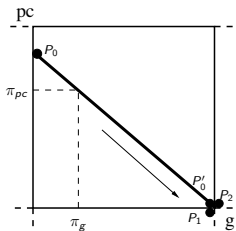
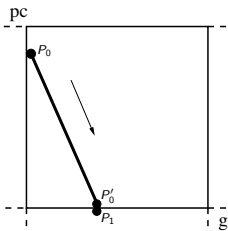
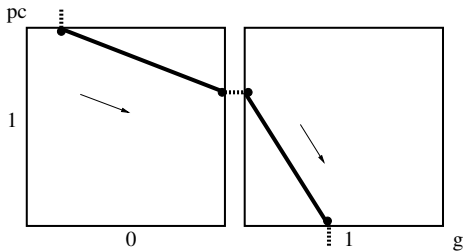
a	b	C_{b,ω_b,η_b}	a	b	z	C_{a,ω_a,η_a}	...
0	0	$C_{b,\emptyset,0}$	0	0	0	$C_{a,\{b\},0}$	
0	1	$C_{b,\emptyset,1}$	1	0	0	$C_{a,\{b\},1}$	
1, 2	0	$C_{b,\{a\},0}$	2	0	0	$C_{a,\{b\},2}$	
1, 2	1	$C_{b,\{a\},1}$	0	0	1	$C_{a,\{b,z\},0}$	
			1	0	1	$C_{a,\{b,z\},1}$	
			2	0	1	$C_{a,\{b,z\},2}$	
	0		0	1	0	$C_{a,\emptyset,0}$	
	1		1	1	0	$C_{a,\emptyset,1}$	
	2		2	1	0	$C_{a,\emptyset,2}$	
	0		0	1	1	$C_{a,\{z\},0}$	
	1		1	1	1	$C_{a,\{z\},1}$	
	2		2	1	1	$C_{a,\{z\},2}$	

$$C_{a,\omega_a,\eta_a} \in \mathbb{R}$$

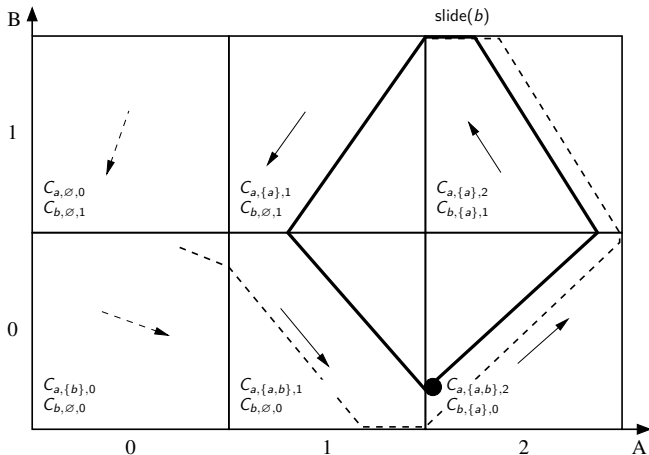
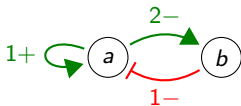
Semantics



Semantics

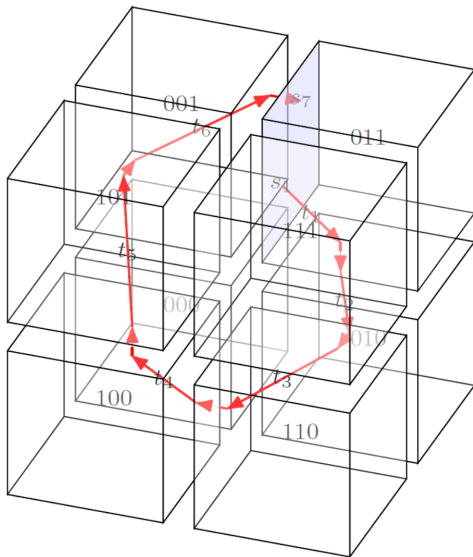
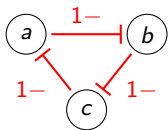


Example (2 dimensions)



Example (3 dimensions)

[Honglu Sun *et al.*, *BIOINFORMATICS*, 2023]



Possible Parametrizations

[Cornillon *et al.*, *Mod. Complex Biol. Syst. in the Context of Genomics*, 2016]

$$C_{a, \omega_a, \eta_a} \in \mathbb{R}$$

Possible Parametrizations

[Cornillon et al., *Mod. Complex Biol. Syst. in the Context of Genomics*, 2016]

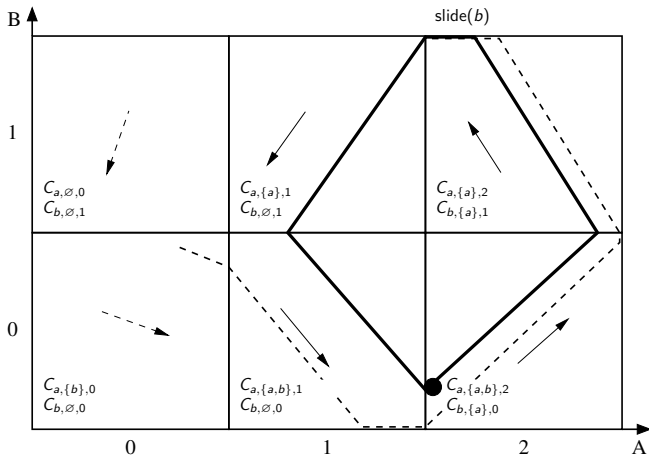
$$C_{a, \omega_a, \eta_a} \in \mathbb{R}$$

Possible parametrizations = ∞

Hybrid Hoare Logic

Hybrid Hoare Logic to Infer Parameters

$$\left\{ \begin{matrix} ??? \\ ??? \end{matrix} \right\} \left(\begin{matrix} T_4 \\ \top \\ b+ \end{matrix} \right); \left(\begin{matrix} T_3 \\ \text{slide}^+(b) \\ a- \end{matrix} \right); \left(\begin{matrix} T_2 \\ \top \\ b- \end{matrix} \right); \left(\begin{matrix} T_1 \\ \top \\ a+ \end{matrix} \right) \left\{ \begin{matrix} \eta_a = 2 \wedge \eta_b = 0 \\ \pi_{\text{initial}} = \pi_{\text{final}} \end{matrix} \right\}$$



Hybrid Hoare Logic

[Behaegel *et al.*, *TIME'17*, 2017]

Hoare triple: $\left\{ \begin{array}{c} D' \\ H' \end{array} \right\} \left(\begin{array}{c} \Delta t \\ \text{assert} \\ v_{\pm} \end{array} \right) \left\{ \begin{array}{c} D \\ H \end{array} \right\}$

Hoare triple: $\left\{ \begin{array}{l} D' \\ H' \end{array} \right\} \left(\begin{array}{c} \Delta t \\ \text{assert} \\ v_{\pm} \end{array} \right) \left\{ \begin{array}{l} D \\ H \end{array} \right\}$

Instruction:

$$\left(\begin{array}{c} \Delta t \\ \text{assert} \\ v_{\pm} \end{array} \right) \begin{array}{l} \leftarrow \text{Time spent in the qualitative state} \\ \leftarrow \text{Biological knowledge (saturation, celerity, ...)} \\ \leftarrow \text{Qualitative instruction (} v_{+} \text{ ou } v_{-} \text{)} \end{array}$$

- $\Delta t \in \mathbb{R}^{+}$
- *assert* is a property using the following predicates:
 - $\text{slide}^{\pm/+/-}(u)$ \leftarrow Variable u “slides” on a border (e.g., saturation)
 - $\text{noslide}^{\pm/+/-}(u)$ \leftarrow Variable u does not “slide”
 - $C_u > 0$ \leftarrow Constraints on the celerities of the current qualitative state

Hybrid Hoare Logic

[Behaegel *et al.*, *TIME'17*, 2017]

Hoare triple: $\left\{ \begin{array}{l} D' \\ H' \end{array} \right\} \left(\begin{array}{c} \Delta t \\ \text{assert} \\ v_{\pm} \end{array} \right) \left\{ \begin{array}{l} D \\ H \end{array} \right\}$

Properties (pre- and post-conditions):

$$\left\{ \begin{array}{l} D \\ H \end{array} \right\} \leftarrow \begin{array}{l} \text{Qualitative/discrete part} \\ \text{Hybrid/real part} \end{array}$$

D and H are properties on:

- $\eta_u \in \mathbb{N}$ \leftarrow Qualitative states of the variables
- $\pi_u \in [0..1]$ \leftarrow Fractional parts (position in the hybrid state)
- $C_{u,\omega,n}$ \leftarrow Celerities
- Δt \leftarrow Time

Weakest Precondition in Hybrid Hoare Logic

[Behaegel *et al.*, *TIME'17*, 2017]

Hoare triple:
$$\left\{ \begin{array}{l} \text{WP}(D) \\ \text{WP}(H) \end{array} \right\} \left(\begin{array}{c} \Delta t \\ \text{assert} \\ \nu \pm \end{array} \right) \left\{ \begin{array}{l} D \\ H \end{array} \right\}$$

$$\text{WP}(D) \equiv D[\eta_\nu \setminus \eta_\nu \pm 1]$$

$$\begin{aligned} \text{WP}(H) \equiv & H[\eta_\nu \setminus \eta_\nu \pm 1] \wedge \Phi_\nu^\pm(\Delta t) \\ & \wedge \mathcal{F}_\nu(\Delta t) \wedge \neg \mathcal{W}_\nu^\pm \wedge \mathcal{A}(\Delta t, \text{assert}) \wedge \mathcal{J}_\nu \end{aligned}$$

Weakest Precondition in Hybrid Hoare Logic

[Behaegel *et al.*, *TIME'17*, 2017]

$$\text{Hoare triple: } \left\{ \begin{array}{l} \text{WP}(D) \\ \text{WP}(H) \end{array} \right\} \left(\begin{array}{c} \Delta t \\ \text{assert} \\ v \pm \end{array} \right) \left\{ \begin{array}{l} D \\ H \end{array} \right\}$$

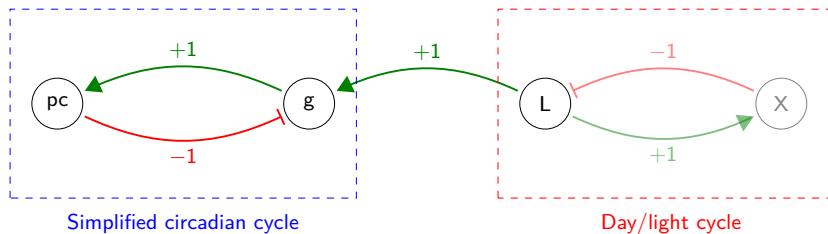
$$\text{WP}(D) \equiv D[\eta_v \setminus \eta_v \pm 1]$$

$$\begin{aligned} \text{WP}(H) \equiv & H[\eta_v \setminus \eta_v \pm 1] \wedge \Phi_v^\pm(\Delta t) \\ & \wedge \mathcal{F}_v(\Delta t) \wedge \neg \mathcal{W}_v^\pm \wedge \mathcal{A}(\Delta t, \text{assert}) \wedge \mathcal{J}_v \end{aligned}$$

- $H[\eta_v \setminus \eta_v \pm 1]$ ← Final hybrid state with substitutions
- $\Phi_v^\pm(\Delta t)$ ← Current celerity of v makes it change discrete state
- $\mathcal{F}_v(\Delta t)$ ← v is the first variable to change discrete state
- \mathcal{W}_v^\pm ← v does not face a black wall (opposing celerity)
- $\mathcal{A}(\Delta t, \text{assert})$ ← Δt and *assert* must be true in current state
- \mathcal{J}_v ← Connect successive steps (if several instructions)

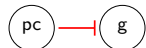
Applications

A Simplified Circadian Cycle Model


[Behaegel *et al.*, *TIME*'17, 2017]


pc = PER/CRY complex
 g = *per* and *cry* genes

L = light of the day
 X = Modeling artifact (clock)

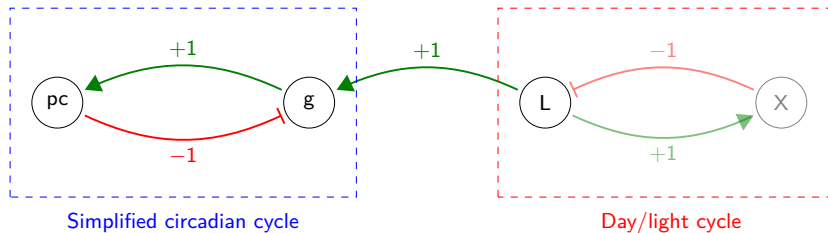
 = PER/CRY complex inhibits *per* and *cry*

 = transcription and complexation

 = light makes BMAL1/CLOCK complex activate *per* and *cry*

 = 12h day/night oscillation

A Simplified Circadian Cycle Model

[Behaegel *et al.*, *TIME'17*, 2017]

$$\left\{ \begin{array}{l} D_8 \\ H_8 \end{array} \right\} \left(\begin{array}{c} 0.9 \\ \top \\ pc- \end{array} \right); \left(\begin{array}{c} 4.5 \\ \top \\ g+ \end{array} \right); \left(\begin{array}{c} 0.6 \\ \top \\ X+ \end{array} \right); \left(\begin{array}{c} 5.53 \\ \text{slide}^+(g) \\ pc+ \end{array} \right); \left(\begin{array}{c} 0.47 \\ \top \\ L- \end{array} \right); \left(\begin{array}{c} 5.4 \\ \text{slide}^+(pc) \\ g- \end{array} \right); \left(\begin{array}{c} 0.6 \\ \text{slide}^-(L) \\ X- \end{array} \right); \left(\begin{array}{c} 6 \\ \top \\ L+ \end{array} \right) \left\{ \begin{array}{l} D_0 \\ H_0 \end{array} \right\}$$

where

$$\left\{ \begin{array}{l} D_0 \\ H_0 \end{array} \right\} \equiv \left(\eta_g = 0 \right) \wedge \left(\eta_{pc} = 1 \right) \wedge \left(\eta_L = 1 \right) \wedge \left(\eta_X = 0 \right) \\ \equiv \left(\pi_g = 0.12 \right) \wedge \left(\pi_{pc} = 0.12 \right) \wedge \left(\pi_L = 0 \right) \wedge \left(\pi_X = 0 \right)$$

Constraints

$$\begin{aligned}
& ((((((((((\pi_g^0 = 0.12) \wedge ((\pi_{pc}^0 = 0.12) \wedge (\pi_L^0 = 0)) \wedge (((\pi_L^1 = 1) \wedge ((C_{L,\{m5\},0} > 0) \wedge (\pi_L^1 = \\
& (\pi_L^1 - (C_{L,\{m5\},0} \times 6.6)))) \wedge ((\neg((C_{g,\emptyset,0} > 0) \wedge (\pi_g^1 > (\pi_g^1 - (C_{g,\emptyset,0} \times 6.6)))) \wedge (\neg((C_{pc,\emptyset,1} < 0) \wedge (\pi_{pc}^1 < \\
& (\pi_{pc}^1 - (C_{pc,\emptyset,1} \times 6.6)))) \wedge (\neg((C_{X,\emptyset,0} > 0) \wedge (\pi_X^1 > (\pi_X^1 - (C_{X,\emptyset,0} \times 6.6)))))) \wedge ((\pi_L^1 = (1 - \pi_L^0)) \wedge ((\pi_g^1 = \\
& \pi_g^0) \wedge ((\pi_{pc}^1 = \pi_{pc}^0) \wedge (\pi_X^1 = \pi_X^0)))))) \wedge (((\pi_X^2 = 0) \wedge ((C_{X,\emptyset,1} < 0) \wedge (\pi_X^2 = (\pi_X^2 - (C_{X,\emptyset,1} \times 0.6)))) \wedge ((\neg((C_{g,\emptyset,0} > \\
& 0) \wedge (\pi_g^2 > (\pi_g^2 - (C_{g,\emptyset,0} \times 0.6)))) \wedge (\neg((C_{pc,\emptyset,1} < 0) \wedge (\pi_{pc}^2 < (\pi_{pc}^2 - (C_{pc,\emptyset,1} \times 0.6)))) \wedge (\neg((C_{L,\emptyset,0} > 0) \wedge (\pi_L^2 > \\
& (\pi_L^2 - (C_{L,\emptyset,0} \times 0.6)))))) \wedge (((\pi_L^2 = 0) \wedge ((C_{L,\emptyset,0} < 0) \Rightarrow (\pi_L^2 < (\pi_L^2 - (C_{L,\emptyset,0} \times 0.6)))) \wedge ((\pi_X^2 = (1 - \pi_X^1)) \wedge ((\pi_g^2 = \\
& \pi_g^1) \wedge ((\pi_{pc}^2 = \pi_{pc}^1) \wedge (\pi_L^2 = \pi_L^1)))))) \wedge (((\pi_g^3 = 0) \wedge ((C_{g,\emptyset,1} < 0) \wedge (\pi_g^3 = (\pi_g^3 - (C_{g,\emptyset,1} \times 5.4)))) \wedge (\neg((C_{pc,\{m2\},1} < \\
& 0) \wedge (\pi_{pc}^3 < (\pi_{pc}^3 - (C_{pc,\{m2\},1} \times 5.4)))) \wedge (\neg((C_{L,\emptyset,0} > 0) \wedge (\pi_L^3 > (\pi_L^3 - (C_{L,\emptyset,0} \times 5.4)))) \wedge (\neg((C_{X,\emptyset,1} < 0) \wedge (\pi_X^3 < \\
& (\pi_X^3 - (C_{X,\emptyset,1} \times 5.4)))))) \wedge (((\pi_{pc}^3 = 1) \wedge ((C_{pc,\{m2\},1} > 0) \Rightarrow (\pi_{pc}^3 > (\pi_{pc}^3 - (C_{pc,\{m2\},1} \times 5.4)))) \wedge ((\pi_g^3 = \\
& (1 - \pi_g^2) \wedge ((\pi_{pc}^3 = \pi_{pc}^2) \wedge ((\pi_L^3 = \pi_L^2) \wedge (\pi_X^3 = \pi_X^2)))))) \wedge (((\pi_L^4 = 0) \wedge ((C_{L,\emptyset,1} < 0) \wedge (\pi_L^4 = \\
& (\pi_L^4 - (C_{L,\emptyset,1} \times 0.47)))) \wedge (\neg((C_{g,\{m3\},1} < 0) \wedge (\pi_g^4 < (\pi_g^4 - (C_{g,\{m3\},1} \times 0.47)))) \wedge (\neg((C_{pc,\{m2\},1} < \\
& 0) \wedge (\pi_{pc}^4 < (\pi_{pc}^4 - (C_{pc,\{m2\},1} \times 0.47)))) \wedge (\neg((C_{X,\{m4\},1} < 0) \wedge (\pi_X^4 < (\pi_X^4 - (C_{X,\{m4\},1} \times 0.47)))))) \wedge ((\pi_L^4 = \\
& (1 - \pi_L^3) \wedge ((\pi_g^4 = \pi_g^3) \wedge ((\pi_{pc}^4 = \pi_{pc}^3) \wedge (\pi_X^4 = \pi_X^3)))))) \wedge (((\pi_{pc}^5 = 1) \wedge ((C_{pc,\{m2\},0} > 0) \wedge (\pi_{pc}^5 = \\
& (\pi_{pc}^5 - (C_{pc,\{m2\},0} \times 5.53)))) \wedge (\neg((C_{g,\{m1,m3\},1} < 0) \wedge (\pi_g^5 < (\pi_g^5 - (C_{g,\{m1,m3\},1} \times 5.53)))) \wedge (\neg((C_{L,\emptyset,1} < \\
& 0) \wedge (\pi_L^5 < (\pi_L^5 - (C_{L,\emptyset,1} \times 5.53)))) \wedge (\neg((C_{X,\{m4\},1} < 0) \wedge (\pi_X^5 < (\pi_X^5 - (C_{X,\{m4\},1} \times 5.53)))))) \wedge (((\pi_g^5 = \\
& 1) \wedge ((C_{g,\{m1,m3\},1} > 0) \Rightarrow (\pi_g^5 > (\pi_g^5 - (C_{g,\{m1,m3\},1} \times 5.53)))) \wedge ((\pi_{pc}^5 = (1 - \pi_{pc}^4)) \wedge ((\pi_g^5 = \pi_g^4) \wedge ((\pi_L^5 = \\
& \pi_L^4) \wedge (\pi_X^5 = \pi_X^4)))))) \wedge (((\pi_X^6 = 1) \wedge ((C_{X,\{m4\},0} > 0) \wedge (\pi_X^6 = (\pi_X^6 - (C_{X,\{m4\},0} \times 0.6)))) \wedge (\neg((C_{g,\{m1,m3\},1} < \\
& 0) \wedge (\pi_g^6 < (\pi_g^6 - (C_{g,\{m1,m3\},1} \times 0.6)))) \wedge (\neg((C_{pc,\{m2\},0} > 0) \wedge (\pi_{pc}^6 > (\pi_{pc}^6 - (C_{pc,\{m2\},0} \times 0.6)))) \wedge (\neg((C_{L,\{m5\},1} < \\
& 0) \wedge (\pi_L^6 < (\pi_L^6 - (C_{L,\{m5\},1} \times 0.6)))))) \wedge ((\pi_X^6 = (1 - \pi_X^5)) \wedge ((\pi_g^6 = \pi_g^5) \wedge ((\pi_{pc}^6 = \pi_{pc}^5) \wedge (\pi_L^6 = \\
& \pi_L^5)))))) \wedge (((\pi_g^7 = 1) \wedge ((C_{g,\{m1,m3\},0} > 0) \wedge (\pi_g^7 = (\pi_g^7 - (C_{g,\{m1,m3\},0} \times 4.5)))) \wedge (\neg((C_{pc,\emptyset,0} > 0) \wedge (\pi_{pc}^7 > \\
& (\pi_{pc}^7 - (C_{pc,\emptyset,0} \times 4.5)))) \wedge (\neg((C_{L,\{m5\},1} < 0) \wedge (\pi_L^7 < (\pi_L^7 - (C_{L,\{m5\},1} \times 4.5)))) \wedge (\neg((C_{X,\{m4\},0} > 0) \wedge (\pi_X^7 > \\
& (\pi_X^7 - (C_{X,\{m4\},0} \times 4.5)))))) \wedge ((\pi_g^7 = (1 - \pi_g^6)) \wedge ((\pi_{pc}^7 = \pi_{pc}^6) \wedge ((\pi_L^7 = \pi_L^6) \wedge (\pi_X^7 = \pi_X^6)))))) \wedge (((\pi_{pc}^8 = \\
& 0) \wedge ((C_{pc,\emptyset,1} < 0) \wedge (\pi_{pc}^8 = (\pi_{pc}^8 - (C_{pc,\emptyset,1} \times 0.9)))) \wedge (\neg((C_{g,\{m3\},0} > 0) \wedge (\pi_g^8 > \\
& (\pi_g^8 - (C_{g,\{m3\},0} \times 0.9)))) \wedge (\neg((C_{L,\{m5\},1} < 0) \wedge (\pi_L^8 < (\pi_L^8 - (C_{L,\{m5\},1} \times 0.9)))) \wedge (\neg((C_{X,\{m4\},0} > 0) \wedge (\pi_X^8 > \\
& (\pi_X^8 - (C_{X,\{m4\},0} \times 0.9)))))) \wedge ((\pi_{pc}^8 = (1 - \pi_{pc}^7)) \wedge ((\pi_g^8 = \pi_g^7) \wedge ((\pi_L^8 = \pi_L^7) \wedge (\pi_X^8 = \pi_X^7))))))
\end{aligned}$$

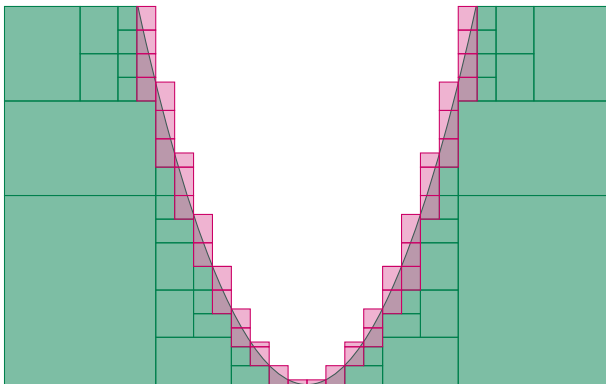
Constraints After Simplification

Constraints After Simplification

$$\begin{aligned}
& (((((((((\pi_g^0 = 0.12) \wedge ((\pi_{pc}^0 = 0.12) \wedge (\pi_L^0 = 0)) \wedge (((\pi_L^1 = 1) \wedge ((C_{L,\{m5\},0} > 0) \wedge (\pi_L^1 = \\
& (\pi_L^1 - (C_{L,\{m5\},0} \times 6.6)))) \wedge ((\neg((C_{g,\emptyset,0} > 0) \wedge (\pi_g^1 > (\pi_g^1 - (C_{g,\emptyset,0} \times 6.6)))) \wedge (\neg((C_{pc,\emptyset,1} < 0) \wedge (\pi_{pc}^1 < \\
& (\pi_{pc}^1 - (C_{pc,\emptyset,1} \times 6.6)))) \wedge \neg((C_{X,\emptyset,0} > 0) \wedge (\pi_X^1 > (\pi_X^1 - (C_{X,\emptyset,0} \times 6.6)))))) \wedge ((\pi_L^1 = (1 - \pi_L^0)) \wedge ((\pi_g^1 = \\
& \pi_g^0) \wedge ((\pi_{pc}^1 = \pi_{pc}^0) \wedge (\pi_X^1 = \pi_X^0)))))) \wedge (((\pi_X^2 = 0) \wedge ((C_{X,\emptyset,1} < 0) \wedge (\pi_X^2 = (\pi_X^2 - (C_{X,\emptyset,1} \times 0.6)))) \wedge ((\neg((C_{g,\emptyset,0} > \\
& 0) \wedge (\pi_g^2 > (\pi_g^2 - (C_{g,\emptyset,0} \times 0.6)))) \wedge (\neg((C_{pc,\emptyset,1} < 0) \wedge (\pi_{pc}^2 < (\pi_{pc}^2 - (C_{pc,\emptyset,1} \times 0.6)))) \wedge (\neg((C_{L,\emptyset,0} > 0) \wedge (\pi_L^2 > \\
& (\pi_L^2 - (C_{L,\emptyset,0} \times 0.6)))))) \wedge (((\pi_L^2 = 0) \wedge ((C_{L,\emptyset,0} < 0) \Rightarrow (\pi_L^2 < (\pi_L^2 - (C_{L,\emptyset,0} \times 0.6)))) \wedge ((\pi_X^2 = (1 - \pi_X^1)) \wedge ((\pi_g^2 = \\
& \pi_g^1) \wedge ((\pi_{pc}^2 = \pi_{pc}^1) \wedge (\pi_L^2 = \pi_L^1)))))) \wedge (((\pi_g^3 = 0) \wedge ((C_{g,\emptyset,1} < 0) \wedge (\pi_g^3 = (\pi_g^3 - (C_{g,\emptyset,1} \times 5.4)))) \wedge (\neg((C_{pc,\{m2\},1} < \\
& 0) \wedge (\pi_{pc}^3 < (\pi_{pc}^3 - (C_{pc,\{m2\},1} \times 5.4)))) \wedge (\neg((C_{L,\emptyset,0} > 0) \wedge (\pi_L^3 > (\pi_L^3 - (C_{L,\emptyset,0} \times 5.4)))) \wedge (\neg((C_{X,\emptyset,1} < 0) \wedge (\pi_X^3 < \\
& (\pi_X^3 - (C_{X,\emptyset,1} \times 5.4)))))) \wedge (((\pi_{pc}^3 = 1) \wedge ((C_{pc,\{m2\},1} > 0) \Rightarrow (\pi_{pc}^3 > (\pi_{pc}^3 - (C_{pc,\{m2\},1} \times 5.4)))) \wedge ((\pi_g^3 = \\
& (1 - \pi_g^2) \wedge ((\pi_{pc}^3 = \pi_{pc}^2) \wedge ((\pi_L^3 = \pi_L^2) \wedge (\pi_X^3 = \pi_X^2)))))) \wedge (((\pi_L^4 = 0) \wedge ((C_{L,\emptyset,1} < 0) \wedge (\pi_L^4 = \\
& (\pi_L^4 - (C_{L,\emptyset,1} \times 0.47)))) \wedge (\neg((C_{g,\{m3\},1} < 0) \wedge (\pi_g^4 < (\pi_g^4 - (C_{g,\{m3\},1} \times 0.47)))) \wedge (\neg((C_{pc,\{m2\},1} < \\
& 0) \wedge (\pi_{pc}^4 < (\pi_{pc}^4 - (C_{pc,\{m2\},1} \times 0.47)))) \wedge (\neg((C_{X,\{m4\},1} < 0) \wedge (\pi_X^4 < (\pi_X^4 - (C_{X,\{m4\},1} \times 0.47)))))) \wedge ((\pi_L^4 = \\
& (1 - \pi_L^3) \wedge ((\pi_g^4 = \pi_g^3) \wedge ((\pi_{pc}^4 = \pi_{pc}^3) \wedge (\pi_X^4 = \pi_X^3)))))) \wedge (((\pi_{pc}^5 = 1) \wedge ((C_{pc,\{m2\},0} > 0) \wedge (\pi_{pc}^5 = \\
& (\pi_{pc}^5 - (C_{pc,\{m2\},0} \times 5.53)))) \wedge (\neg((C_{g,\{m1,m3\},1} < 0) \wedge (\pi_g^5 < (\pi_g^5 - (C_{g,\{m1,m3\},1} \times 5.53)))) \wedge (\neg((C_{L,\emptyset,1} < \\
& 0) \wedge (\pi_L^5 < (\pi_L^5 - (C_{L,\emptyset,1} \times 5.53)))) \wedge (\neg((C_{X,\{m4\},1} < 0) \wedge (\pi_X^5 < (\pi_X^5 - (C_{X,\{m4\},1} \times 5.53)))))) \wedge (((\pi_g^5 = \\
& 1) \wedge ((C_{g,\{m1,m3\},1} > 0) \Rightarrow (\pi_g^5 > (\pi_g^5 - (C_{g,\{m1,m3\},1} \times 5.53)))) \wedge ((\pi_{pc}^5 = (1 - \pi_{pc}^4)) \wedge ((\pi_g^5 = \pi_g^4) \wedge ((\pi_L^5 = \\
& \pi_L^4) \wedge (\pi_X^5 = \pi_X^4)))))) \wedge (((\pi_X^6 = 1) \wedge ((C_{X,\{m4\},0} > 0) \wedge (\pi_X^6 = (\pi_X^6 - (C_{X,\{m4\},0} \times 0.6)))) \wedge (\neg((C_{g,\{m1,m3\},1} < \\
& 0) \wedge (\pi_g^6 < (\pi_g^6 - (C_{g,\{m1,m3\},1} \times 0.6)))) \wedge (\neg((C_{pc,\{m2\},0} > 0) \wedge (\pi_{pc}^6 > (\pi_{pc}^6 - (C_{pc,\{m2\},0} \times 0.6)))) \wedge (\neg((C_{L,\{m5\},1} < \\
& 0) \wedge (\pi_L^6 < (\pi_L^6 - (C_{L,\{m5\},1} \times 0.6)))))) \wedge ((\pi_X^6 = (1 - \pi_X^5)) \wedge ((\pi_g^6 = \pi_g^5) \wedge ((\pi_{pc}^6 = \pi_{pc}^5) \wedge (\pi_L^6 = \\
& \pi_L^5)))))) \wedge (((\pi_g^7 = 1) \wedge ((C_{g,\{m1,m3\},0} > 0) \wedge (\pi_g^7 = (\pi_g^7 - (C_{g,\{m1,m3\},0} \times 4.5)))) \wedge (\neg((C_{pc,\emptyset,0} > 0) \wedge (\pi_{pc}^7 > \\
& (\pi_{pc}^7 - (C_{pc,\emptyset,0} \times 4.5)))) \wedge (\neg((C_{L,\{m5\},1} < 0) \wedge (\pi_L^7 < (\pi_L^7 - (C_{L,\{m5\},1} \times 4.5)))) \wedge (\neg((C_{X,\{m4\},0} > 0) \wedge (\pi_X^7 > \\
& (\pi_X^7 - (C_{X,\{m4\},0} \times 4.5)))))) \wedge ((\pi_g^7 = (1 - \pi_g^6)) \wedge ((\pi_{pc}^7 = \pi_{pc}^6) \wedge ((\pi_L^7 = \pi_L^6) \wedge (\pi_X^7 = \pi_X^6)))))) \wedge (((\pi_{pc}^8 = \\
& 0) \wedge ((C_{pc,\emptyset,1} < 0) \wedge (\pi_{pc}^8 = (\pi_{pc}^8 - (C_{pc,\emptyset,1} \times 0.9)))) \wedge (\neg((C_{g,\{m3\},0} > 0) \wedge (\pi_g^8 > \\
& (\pi_g^8 - (C_{g,\{m3\},0} \times 0.9)))) \wedge (\neg((C_{L,\{m5\},1} < 0) \wedge (\pi_L^8 < (\pi_L^8 - (C_{L,\{m5\},1} \times 0.9)))) \wedge (\neg((C_{X,\{m4\},0} > 0) \wedge (\pi_X^8 > \\
& (\pi_X^8 - (C_{X,\{m4\},0} \times 0.9)))))) \wedge ((\pi_{pc}^8 = (1 - \pi_{pc}^7)) \wedge ((\pi_g^8 = \pi_g^7) \wedge ((\pi_L^8 = \pi_L^7) \wedge (\pi_X^8 = \pi_X^7))))))
\end{aligned}$$

Nonlinear Solver: AbSolute

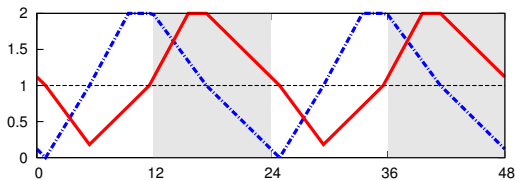
[Pelleau *et al.*, VMCAI, 2013]



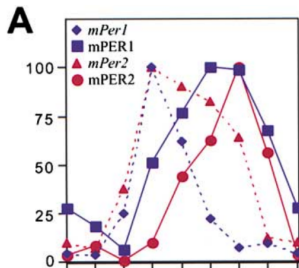
Solving the constraint: $y \leq x^2$

Results

- **Simplifications** of the constraints → Not very effective
- Using a non-linear solver: **AbSolute** → We obtain solutions
- Results checked with a simulation:



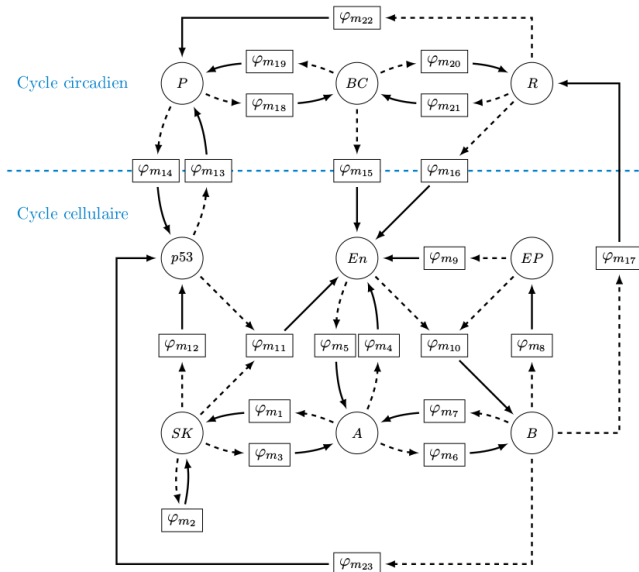
Simulation with 1 set of compatible values



Experiments

Coupling of Cell and Circadian Cycles

[Behaegel, PhD thesis, 2018]



Coupling of Cell and Circadian Cycles

[Behaegel, PhD thesis, 2018]

$$\begin{pmatrix} 0.59 \\ \top \\ BC+ \end{pmatrix}; \begin{pmatrix} 2.44 \\ \text{slide}^+(BC) \wedge \text{slide}^-(P) \\ R+ \end{pmatrix}; \begin{pmatrix} 0.52 \\ \text{slide}^-(EP) \\ SK+ \end{pmatrix}; \begin{pmatrix} 1.77 \\ \top \\ p53+ \end{pmatrix}; \begin{pmatrix} 1.78 \\ \top \\ SK+ \end{pmatrix}; \begin{pmatrix} 2.05 \\ \text{slide}^+(R) \\ P+ \end{pmatrix};$$

$$\begin{pmatrix} 1.51 \\ \text{slide}^+(SK) \\ En- \end{pmatrix}; \begin{pmatrix} 1.93 \\ \top \\ BC- \end{pmatrix}; \begin{pmatrix} 0.20 \\ \text{slide}^-(En) \\ A+ \end{pmatrix}; \begin{pmatrix} 2.13 \\ \top \\ SK- \end{pmatrix}; \left(\begin{pmatrix} 0.11 \\ \text{slide}^-(BC) \wedge \\ \text{slide}^+(P) \\ R- \end{pmatrix} \right); \begin{pmatrix} 2.02 \\ \text{slide}^+(A) \\ SK- \end{pmatrix};$$

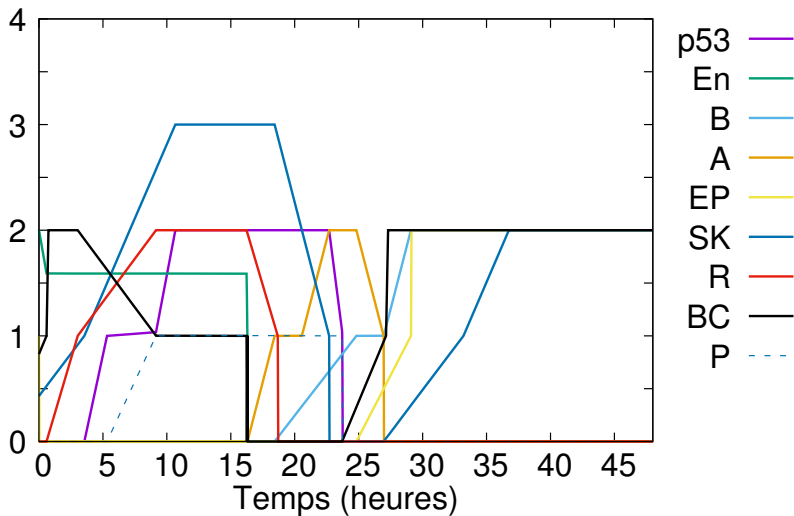
$$\begin{pmatrix} 1.06 \\ \top \\ p53- \end{pmatrix}; \begin{pmatrix} 1.07 \\ \text{slide}^-(SK) \\ B+ \end{pmatrix}; \begin{pmatrix} 1.97 \\ \top \\ P- \end{pmatrix}; \begin{pmatrix} 0.16 \\ \top \\ A- \end{pmatrix}; \begin{pmatrix} 2.14 \\ \text{slide}^+(B) \\ EP+ \end{pmatrix}; \begin{pmatrix} 0.18 \\ \text{slide}^+(EP) \\ En+ \end{pmatrix};$$

$$\begin{pmatrix} 0.18 \\ \text{slide}^-(A) \wedge \text{slide}^+(En) \\ B- \end{pmatrix}; \begin{pmatrix} 0.19 \\ \text{slide}^-(B) \\ EP- \end{pmatrix} \left\{ \left(\begin{pmatrix} (\eta_P = 0) \wedge (\eta_{BC} = 0) \wedge \\ (\eta_R = 0) \wedge (\eta_{SK} = 0) \wedge \\ (\eta_{EP} = 0) \wedge (\eta_A = 0) \wedge \\ (\eta_B = 0) \wedge (\eta_{En} = 1) \wedge \\ (\eta_{PP} = 0) \\ \top \end{pmatrix} \right) \right\}$$

1536 constraints on 152 celerities and 378 fractional parts →
10 minutes → unsure solutions only

Coupling of Cell and Circadian Cycles

[Behaegel, PhD thesis, 2018]

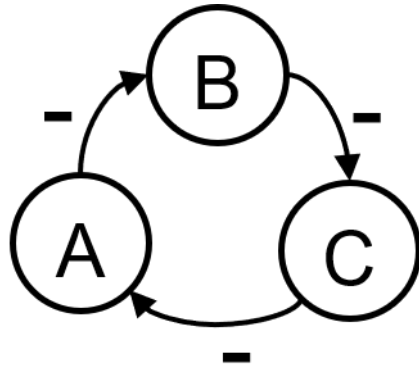


Inference with Optimization Methods



Problem definition

Parameters identification of a hybrid model of **repressilator**



The influence graph of **repressilator**

All parameters:

V_{ac0a0}	V_{ba0b0}	V_{cb0c0}	
V_{ac0a1}	V_{ba0b1}	V_{cb0c1}	
V_{ac1a0}	V_{ba1b0}	V_{cb1c0}	$\theta_{ab} \theta_{bc} \theta_{ca}$
V_{ac1a1}	V_{ba1b1}	V_{cb1c1}	

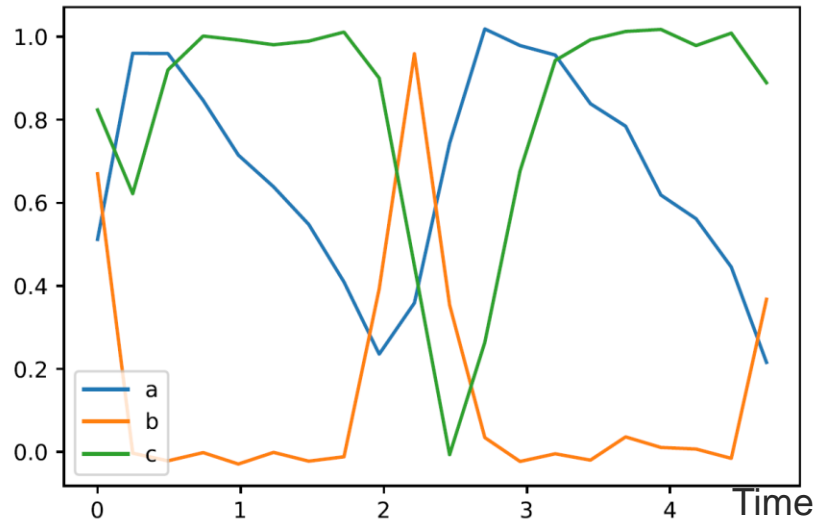
$\underbrace{\hspace{15em}}$ Temporal derivatives (V_i) $\underbrace{\hspace{3em}}$ Thresholds (θ_i)



Problem definition

Parameters identification of a hybrid model of **repressilator**

Expressions of genes



Time series data (by simulation)

Parameters identification



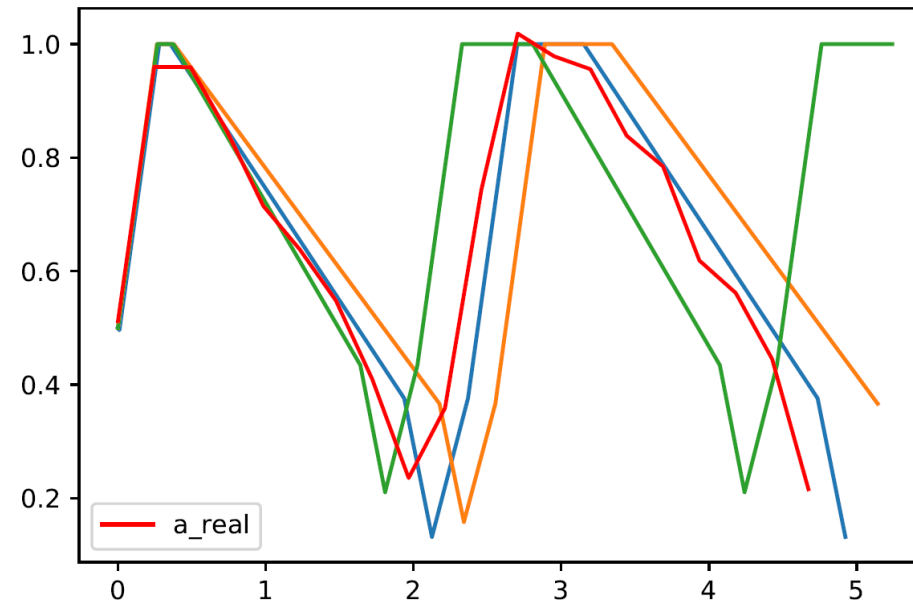
Constraint I (Parameter set)

$$V_i \in I_i^v = [\underline{I}_i^v, \overline{I}_i^v]$$

$$\theta_i \in I_i^\theta = [\underline{I}_i^\theta, \overline{I}_i^\theta]$$

Idea of the parameters identification method

Parameters identification by optimization:
Find parameter set which minimizes the difference
between simulations and real data



Simulations of several models (**model1**, **model2**, **model3**) in a parameter set



Objective function

Optimize the function:

$$\frac{1}{n_1} \sum_{model_i \in M} \frac{1}{n_0} \sum_{j=1}^{n_0} weight^j * (real\ data^j - simulation_i^j)^2$$

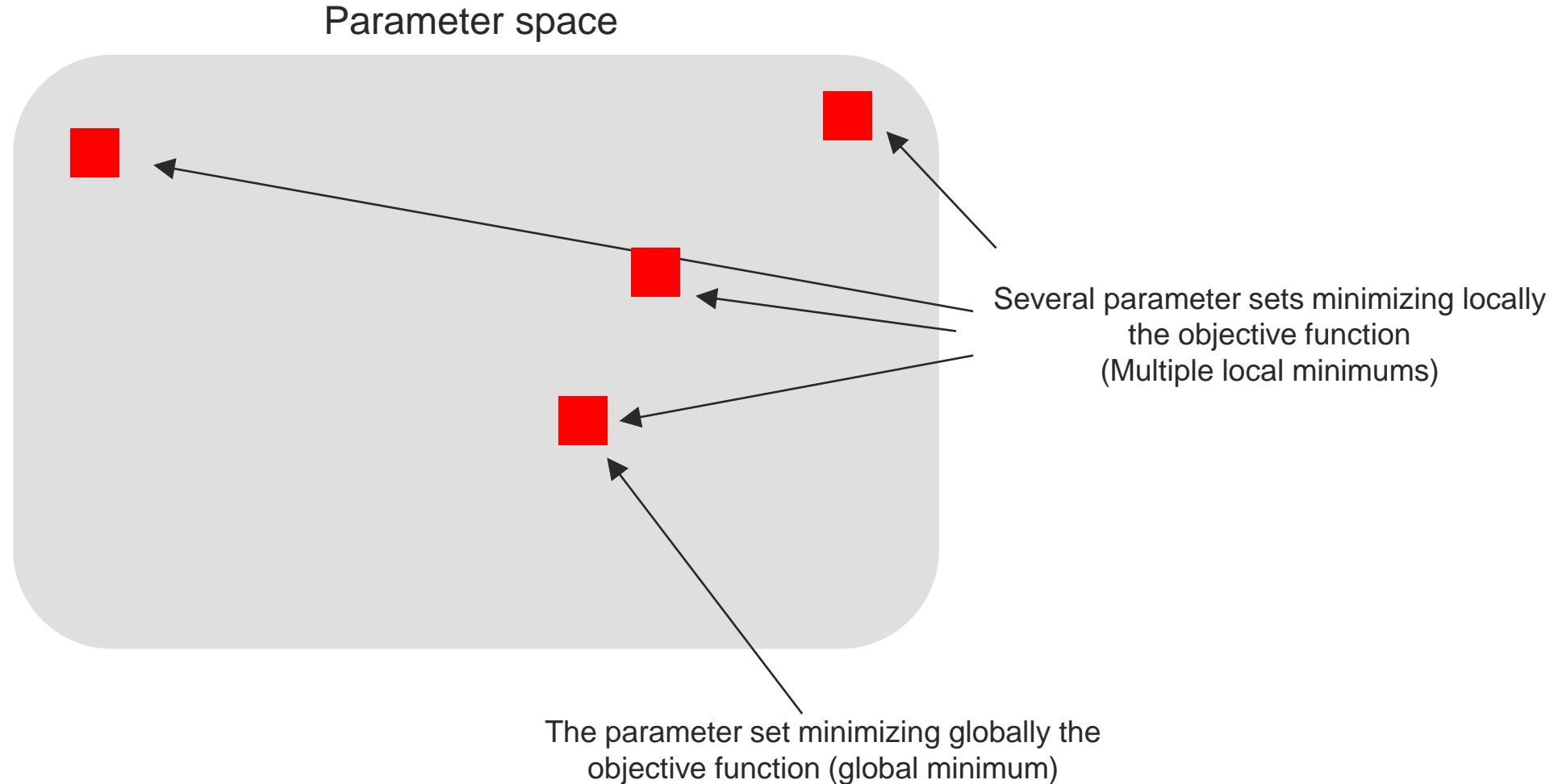
M is a set of n_1 models sampled from a parameter set.

$simulation_i$ is the simulation of $model_i$.

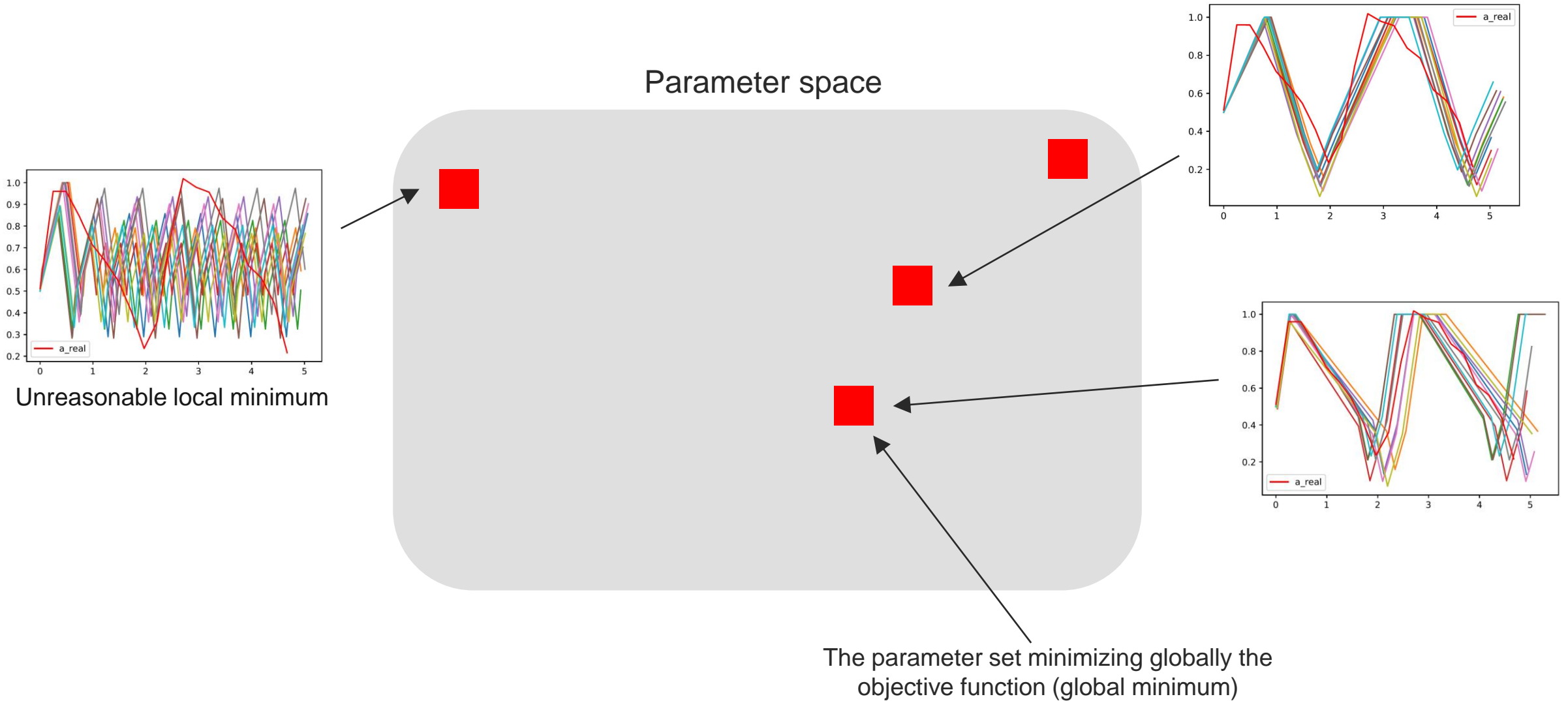
n_0 is the number of data.

$weight^j$ is proportional to the estimated temporal derivative.

Optimization based on genetic algorithm

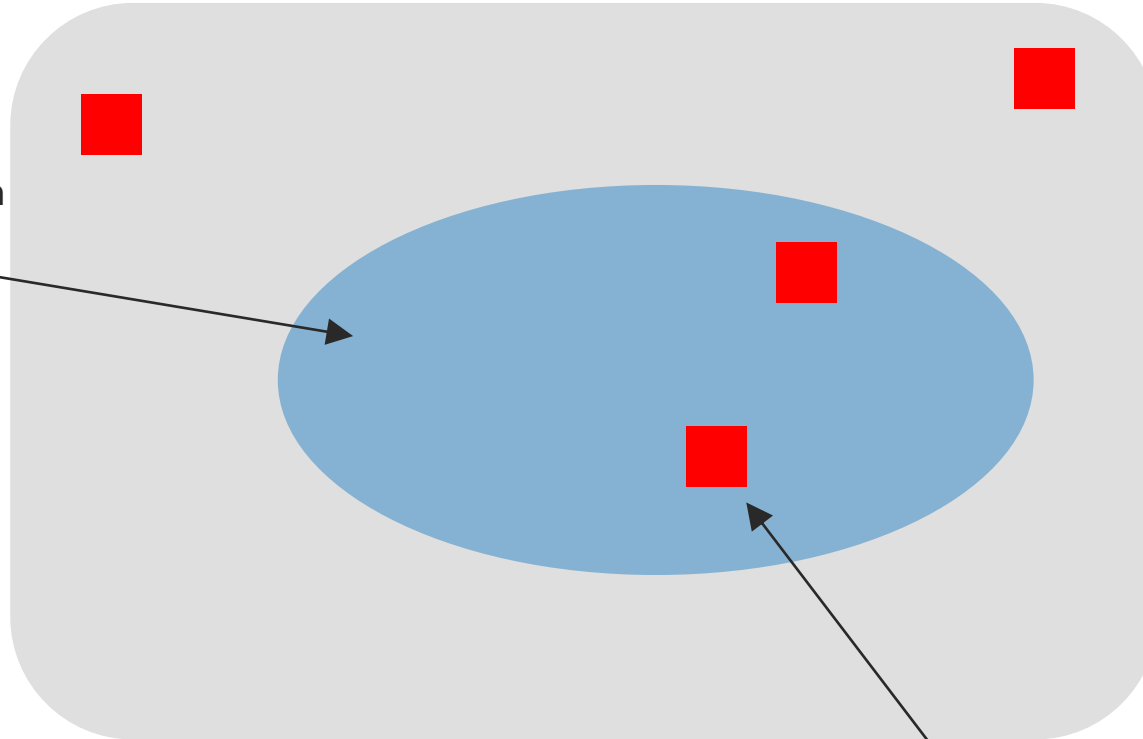


Optimization based on genetic algorithm



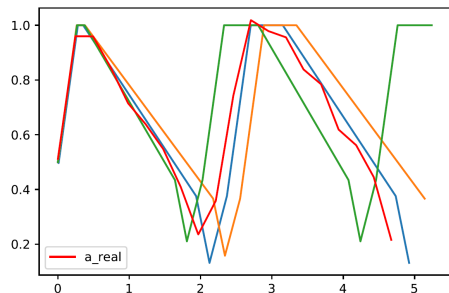
Optimization based on genetic algorithm

Parameter space



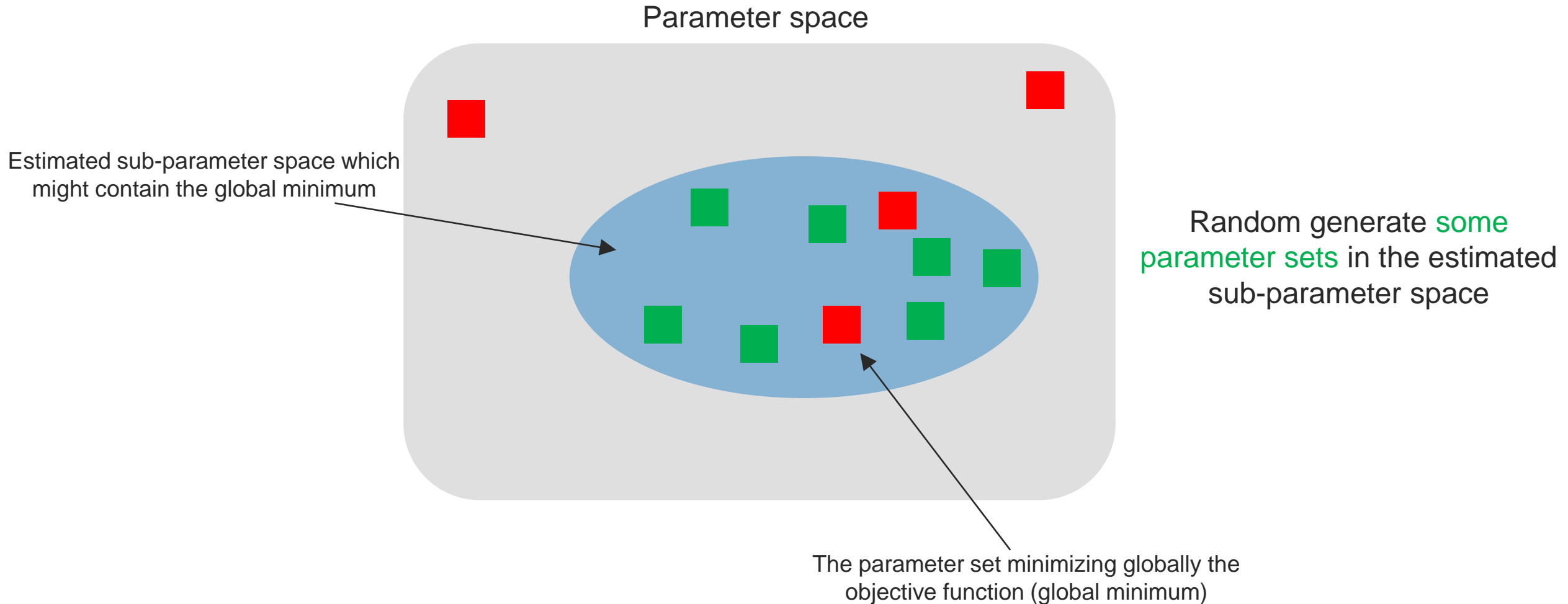
Estimated sub-parameter space which might contain the global minimum

- Based on the estimated temporal derivatives
- Based on biological knowledge (Oscillation? How many periods?)
- Based on symbolic analysis of this model

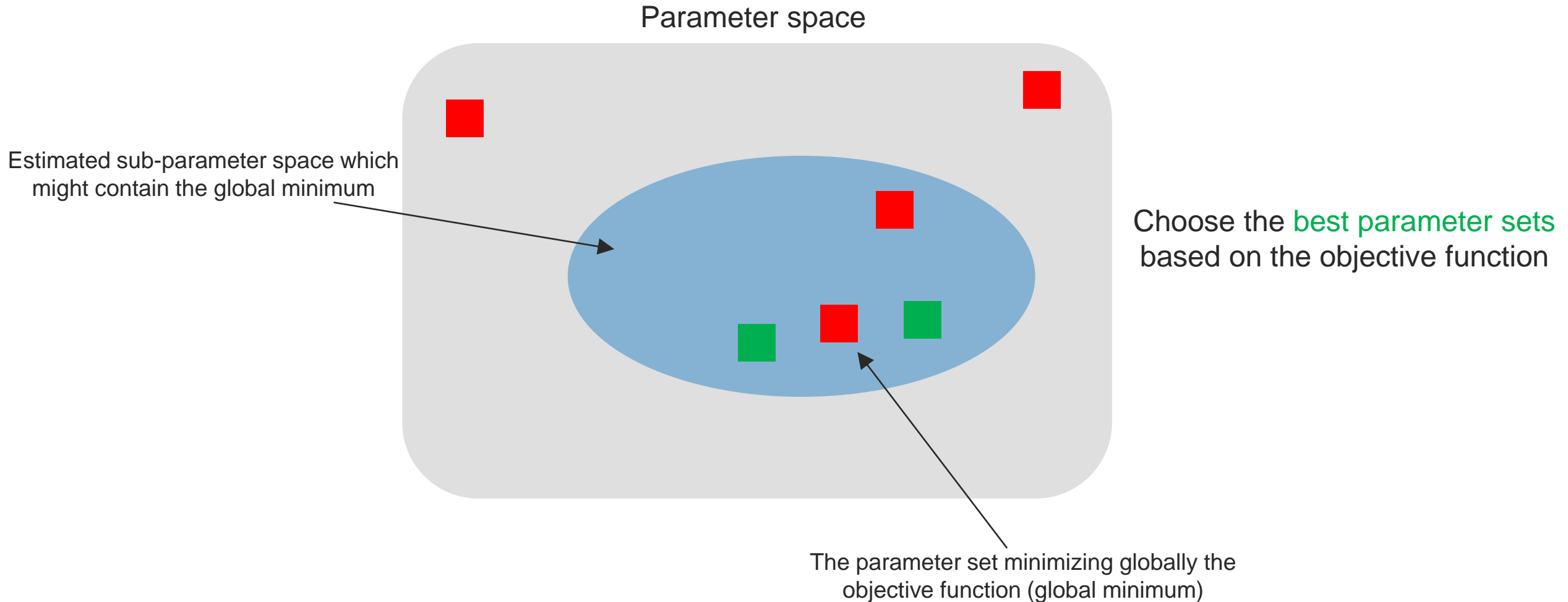


The parameter set minimizing globally the objective function (global minimum)

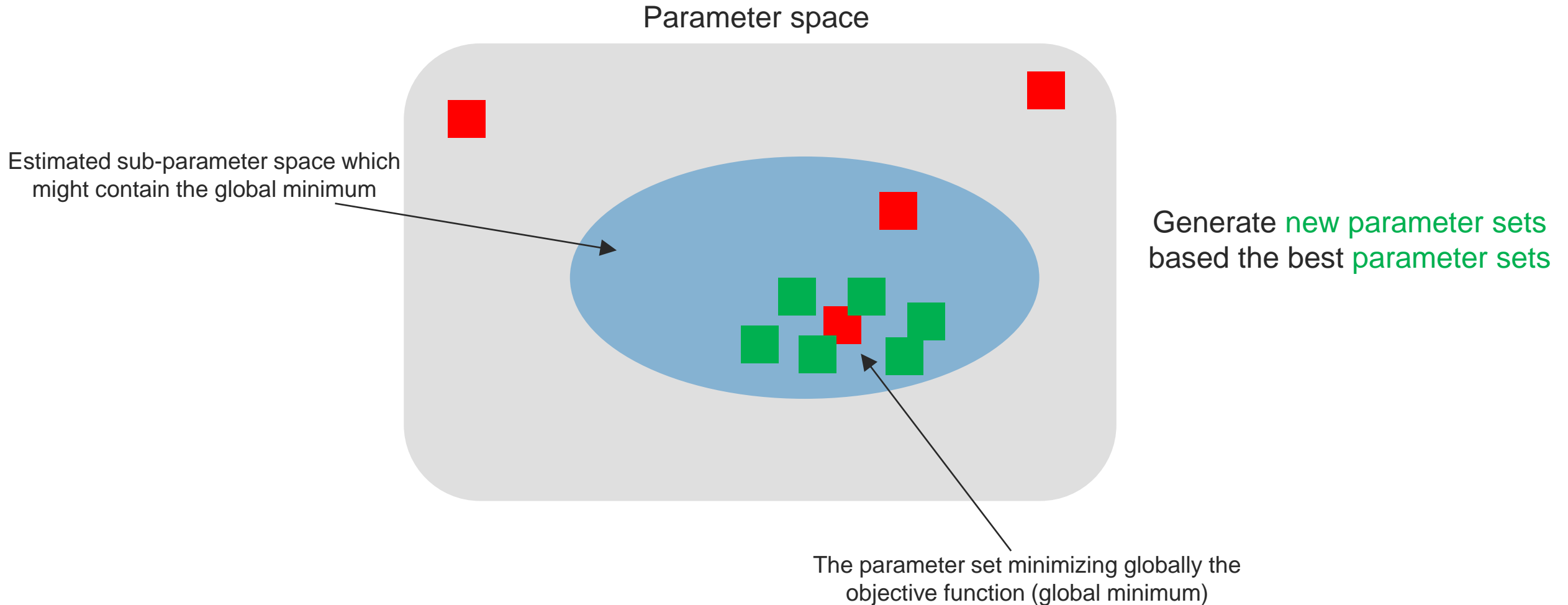
Optimization based on genetic algorithm



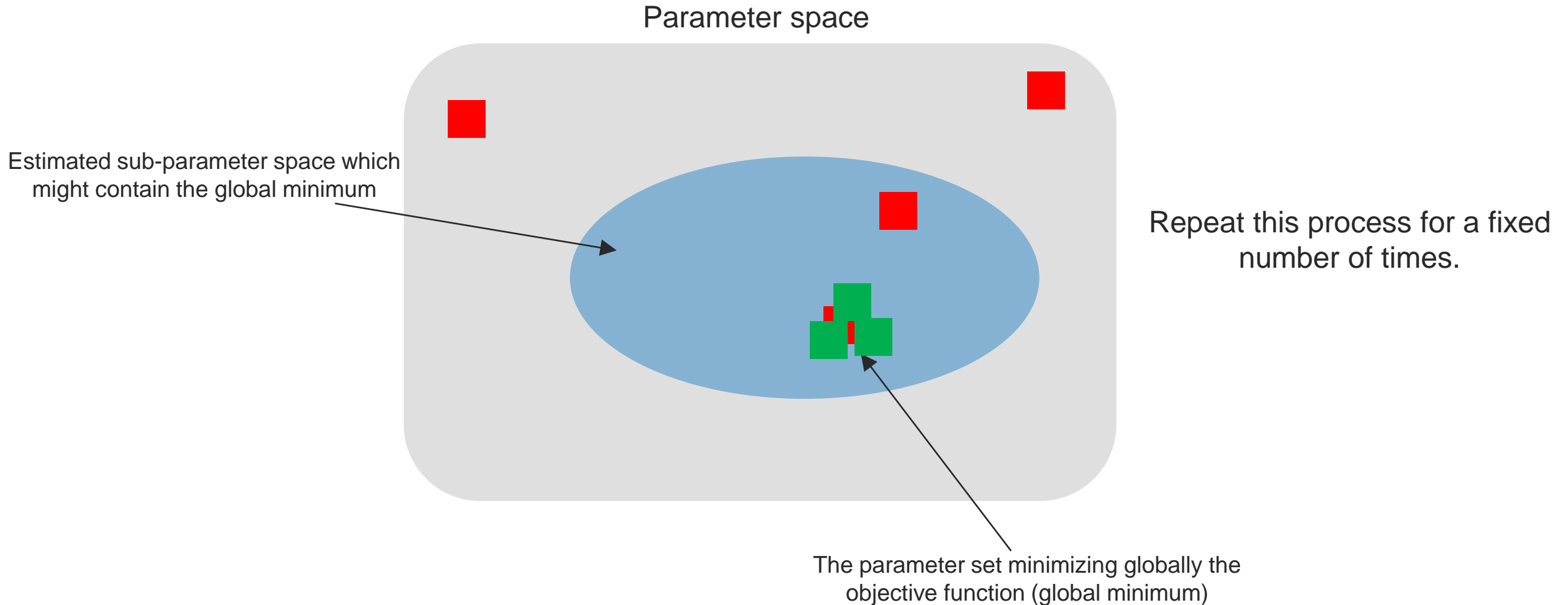
Optimization based on genetic algorithm



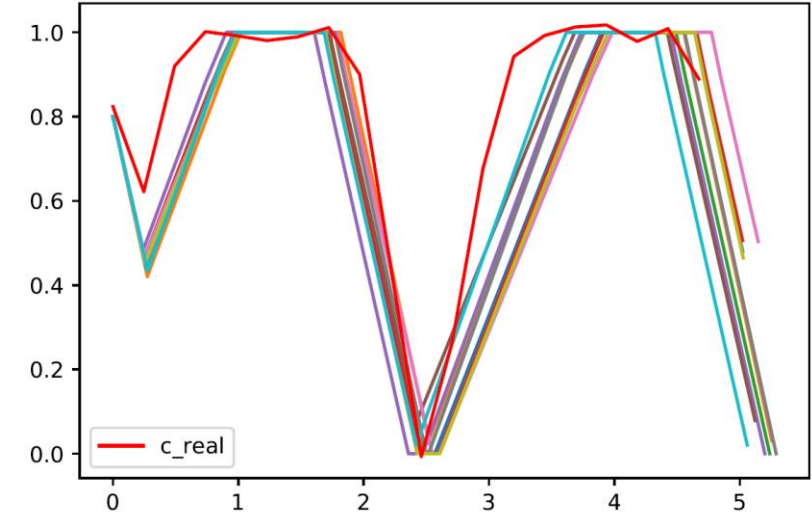
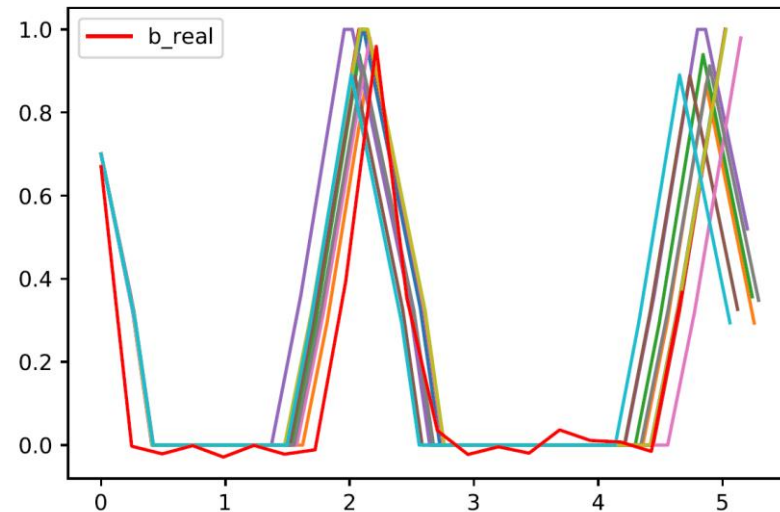
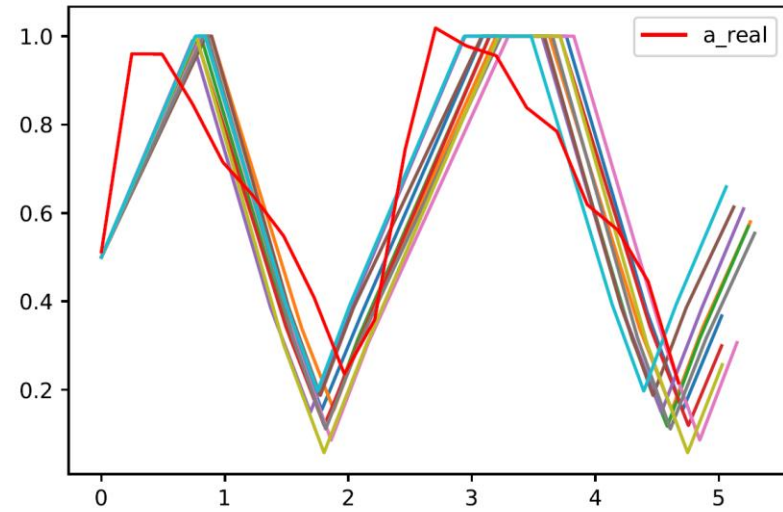
Optimization based on genetic algorithm



Optimization based on genetic algorithm



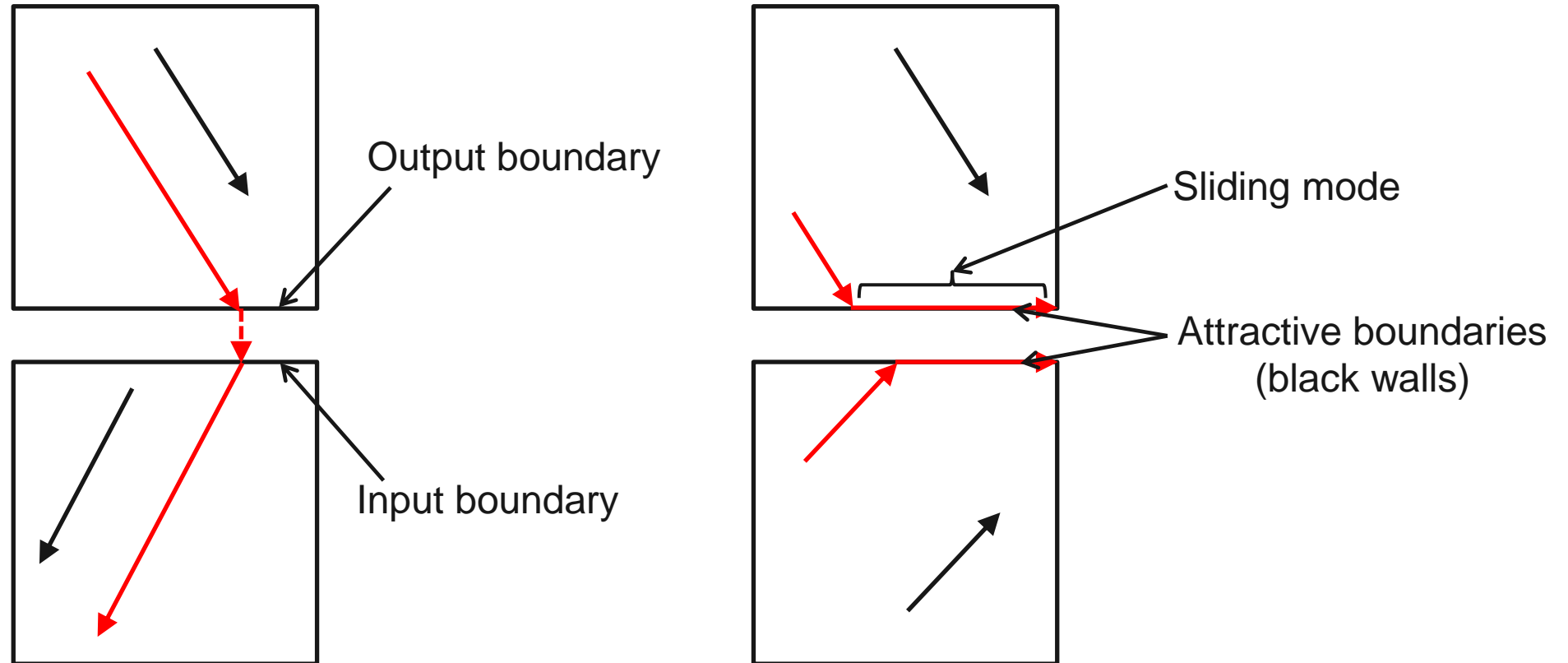
Simulation of the best parameter set found by the algorithm



Dynamical Analysis

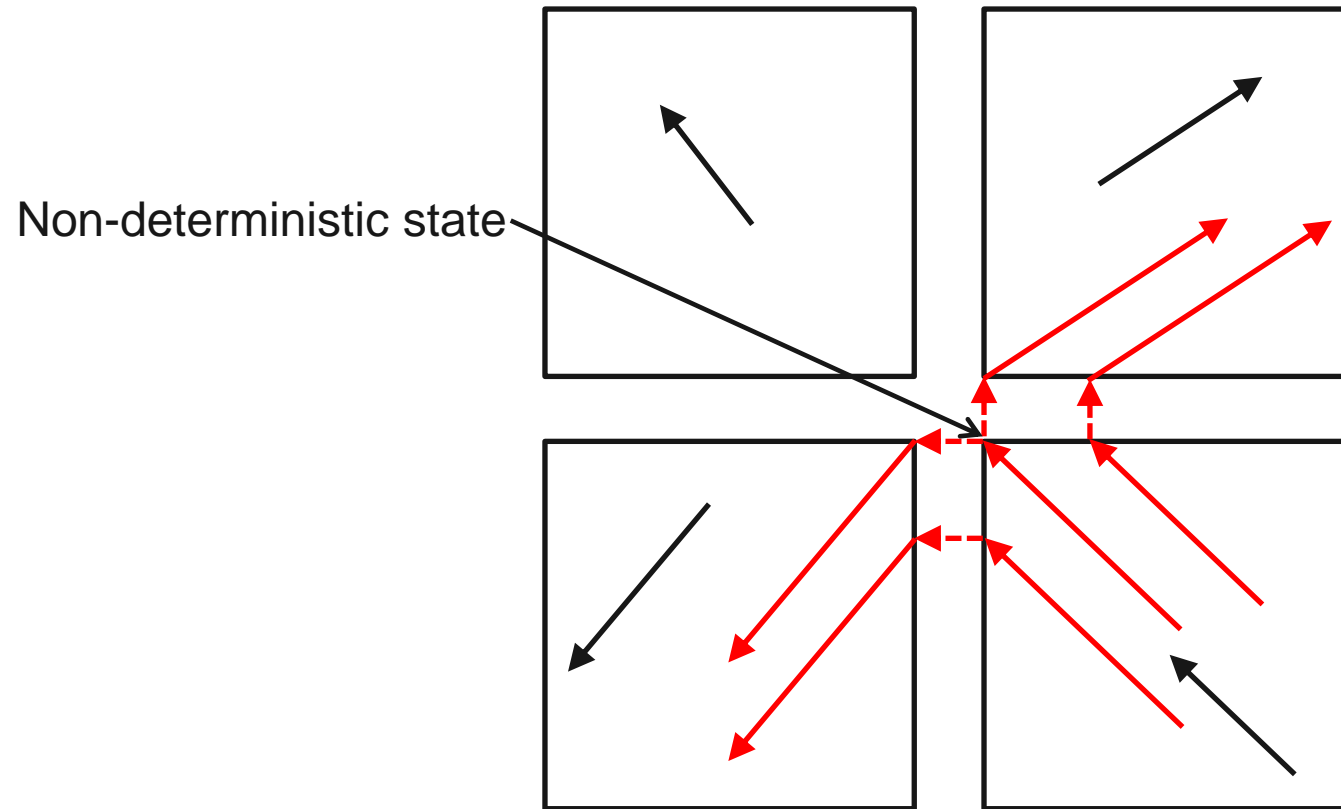


Boundaries and simulation of HGRN



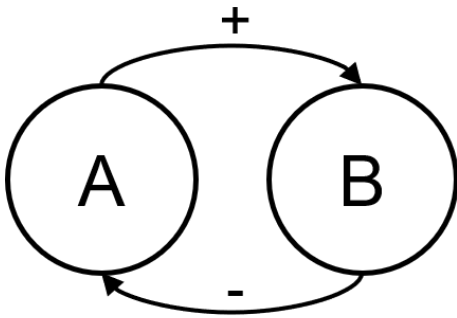


Boundaries and simulation of HGRN

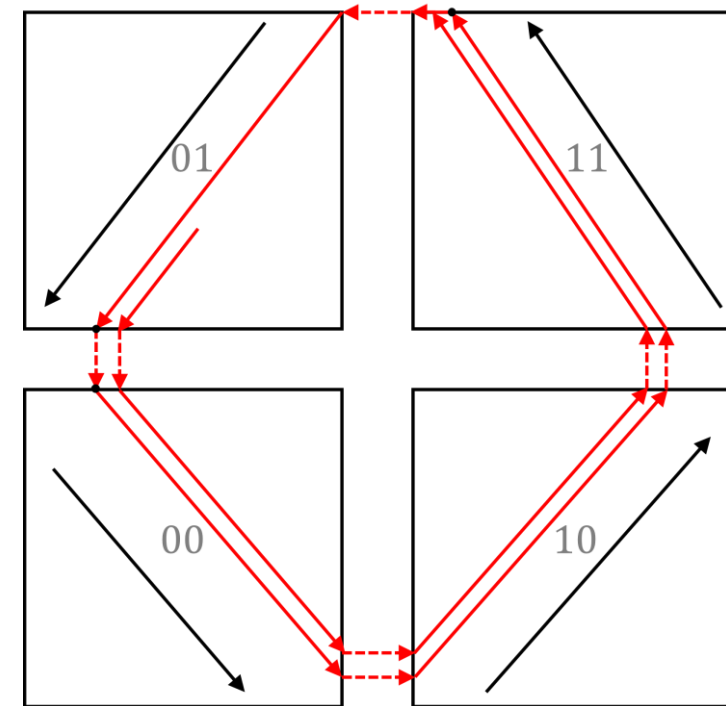


Objective

- Find limit cycles
- Analyze the stability of limit cycles



A	B	C_A	C_B
0	0	0.6	-0.7
0	1	-0.7	-0.9
1	0	0.7	0.8
1	1	-0.6	0.9



Related works:

[Belgacem et al., IEEE Conference on Decision and Control, 2020], [Firippi et al., Chaos: Interdisc. J. Nonlinear Sci., 2020], [Edwards, Physica D: Nonlinear Phenomena, 2000], [Edwards et al., Advances in chemical physics, 2006], etc.



Search for limit cycles

1. Abstraction of the model with discrete domain. →
2. Find cycles of discrete domains which contain continuous trajectories. →
3. Search for limit cycle(s) inside cycles found in step 2.

Search for limit cycles

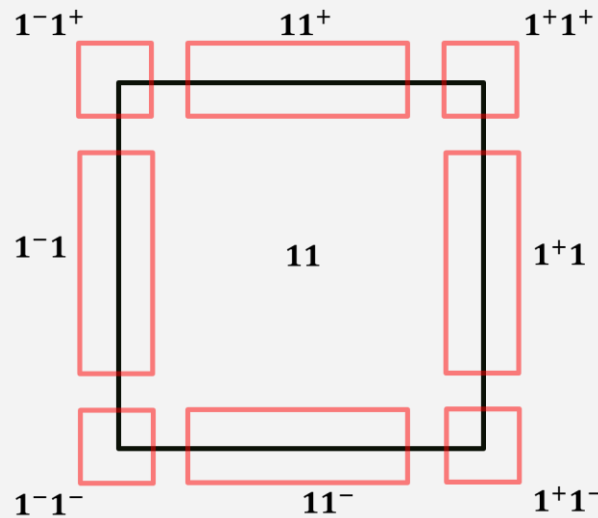
1. Abstraction of the model with discrete domain.



2. Find cycles of discrete domains which contain continuous trajectories.



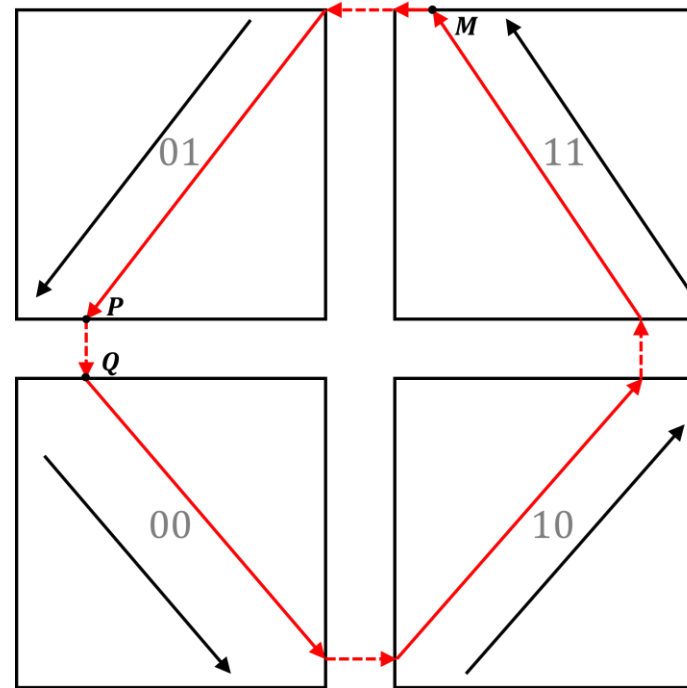
3. Search for limit cycle(s) inside cycles found in step 2.



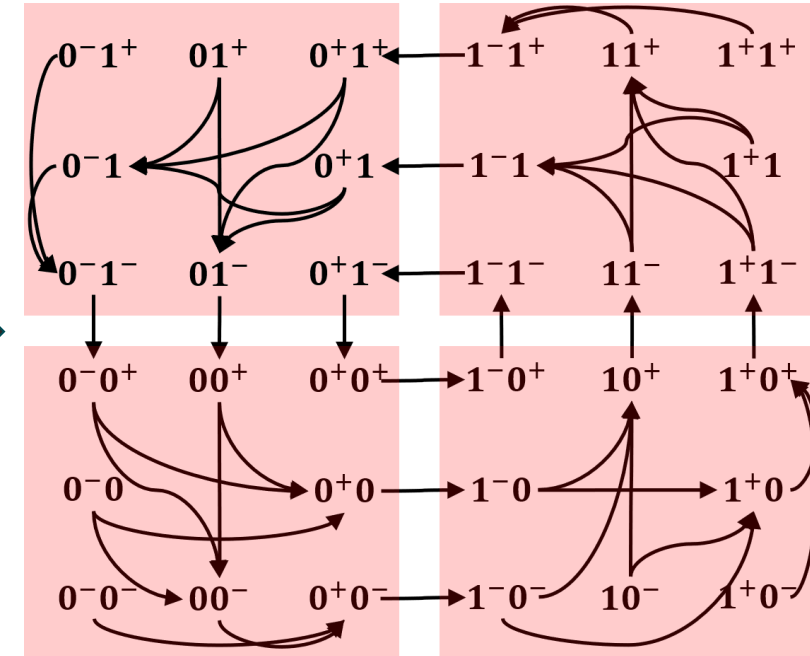
$$(1^+, 1^+) = \{(\pi, 11) \mid \pi^1 = 1, \pi^2 = 1\}$$

$$(1, 1^+) = \{(\pi, 11) \mid \pi^1 \in]0, 1[, \pi^2 = 1\}$$

Examples of **discrete domain**



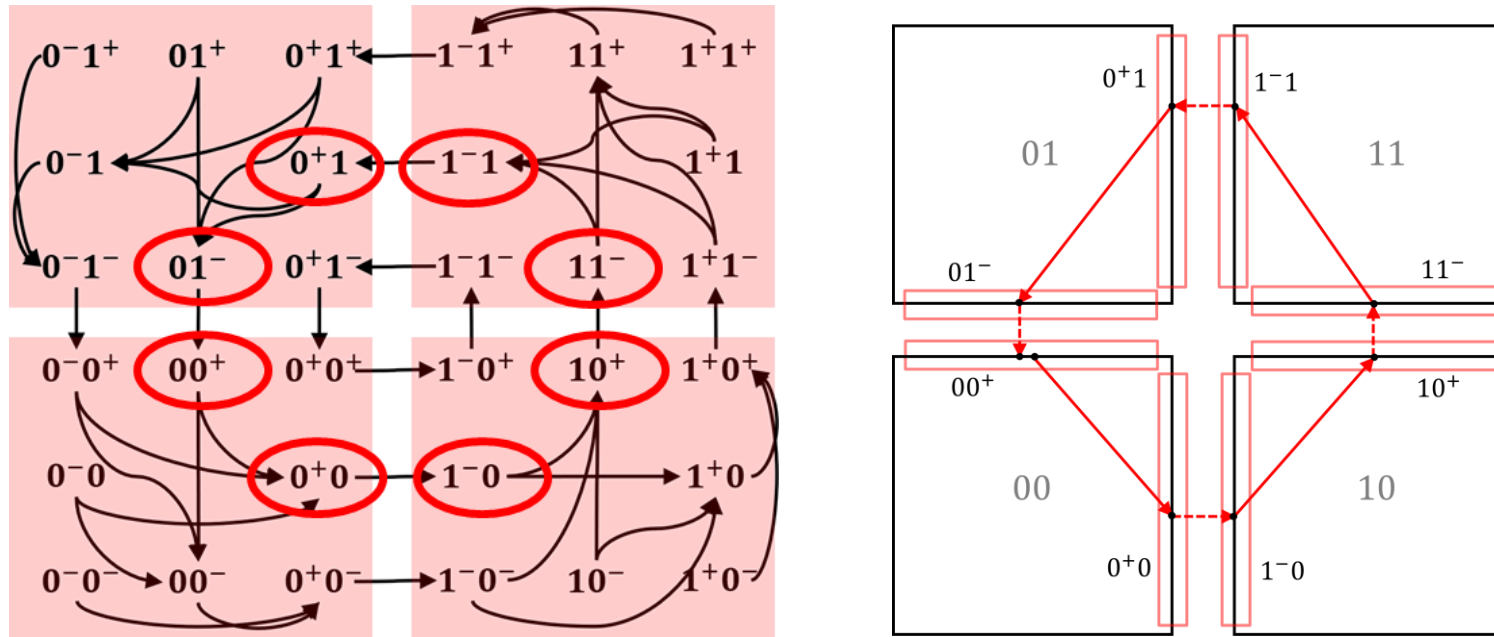
Original system



Graph of discrete domains

Search for limit cycles

1. Abstraction of the model with discrete domain.
2. Find cycles of discrete domains which contain continuous trajectories.
3. Search for limit cycle(s) inside cycles found in step 2.



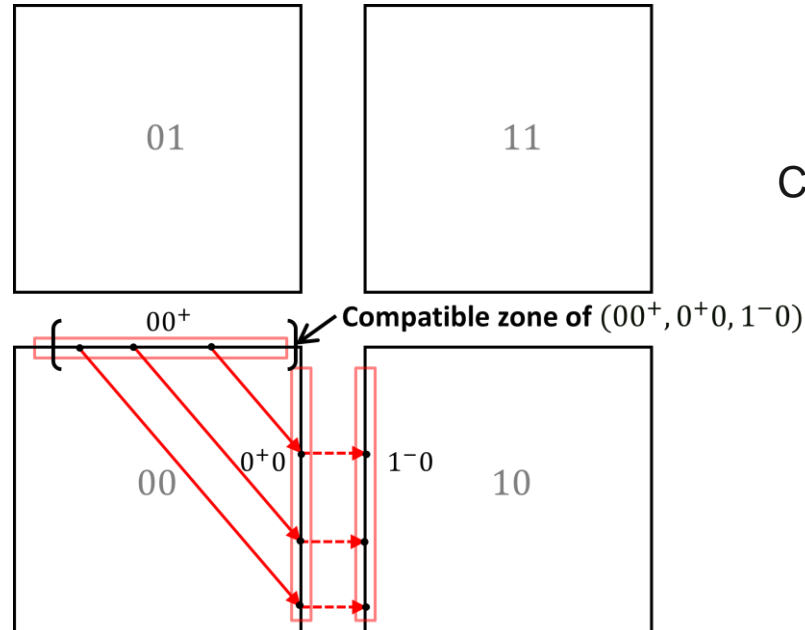
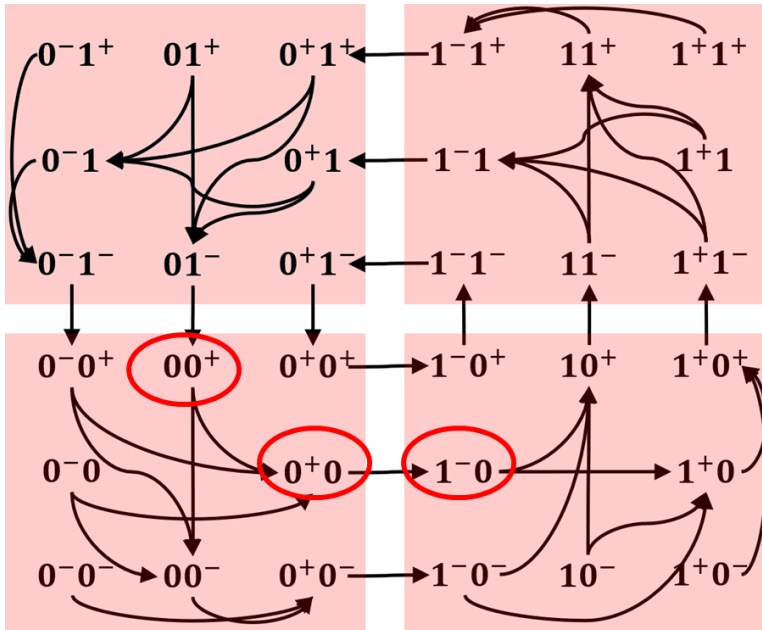
Example of a cycle of discrete domains which contains continuous trajectories

Search for limit cycles

1. Abstraction of the model with discrete domain.

2. Find cycles of discrete domains which contain continuous trajectories.

3. Search for limit cycle(s) inside cycles found in step 2.



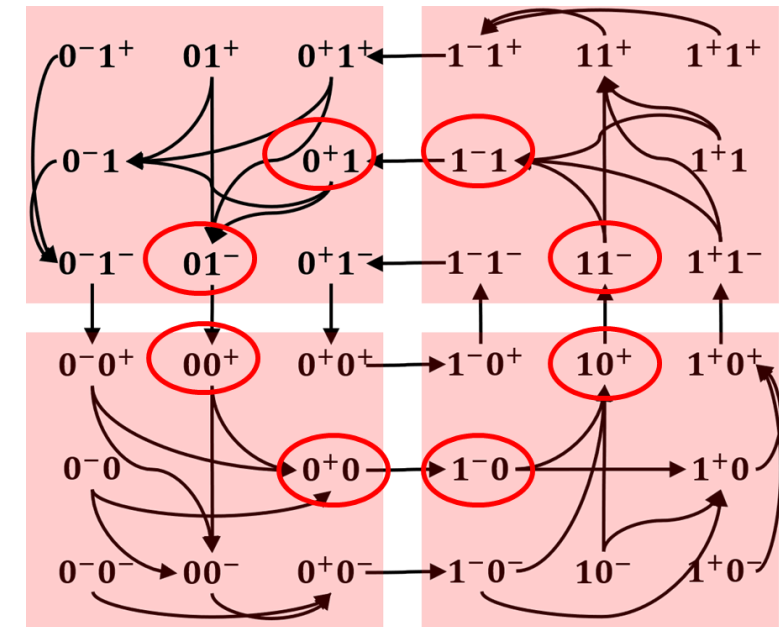
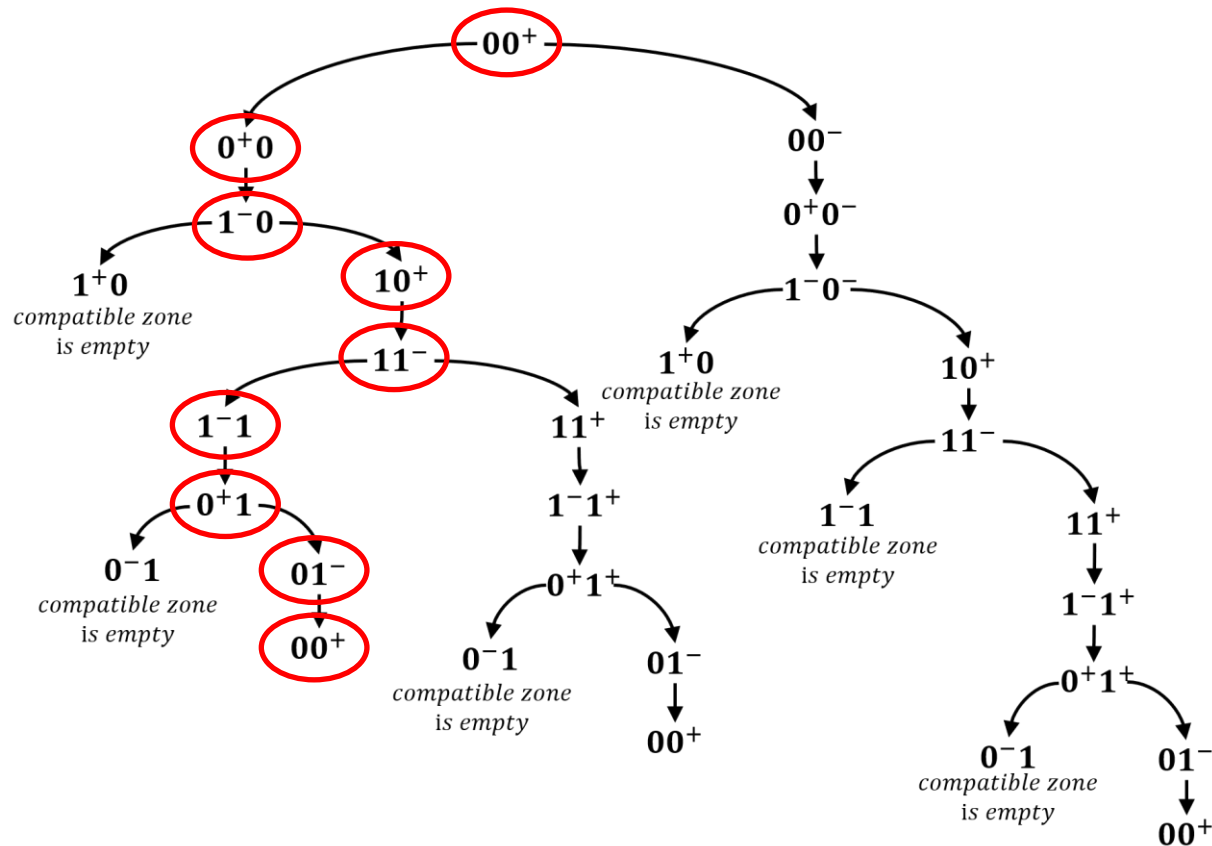
Compatible zone S of $(00^+, 0^+0, 1^-0)$:
 $S \subseteq 00^+$ s.t. $\forall h \in S$, any trajectory from h stays inside $(00^+, 0^+0, 1^-0)$.

Example of **compatible zone**

Search for limit cycles

1. Abstraction of the model with discrete domain.
2. Find cycles of discrete domains which contain continuous trajectories.
3. Search for limit cycle(s) inside cycles found in step 2.

Find cycles of discrete domains of which the **compatible zones** are not empty.



Search for limit cycles

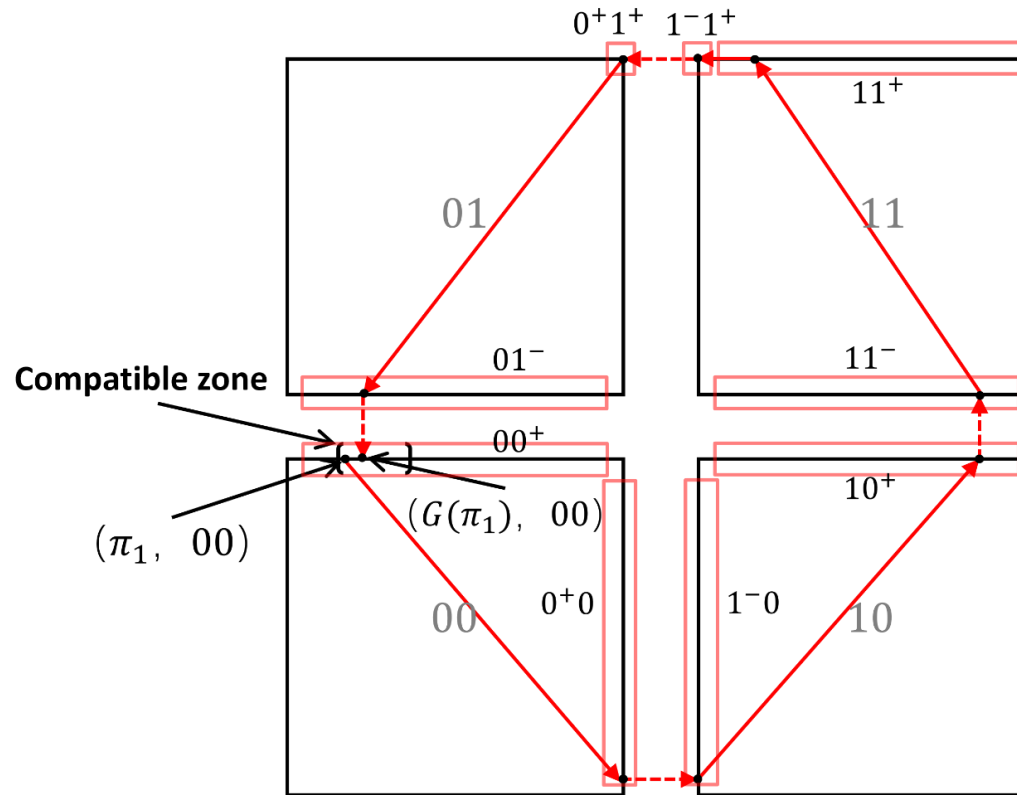
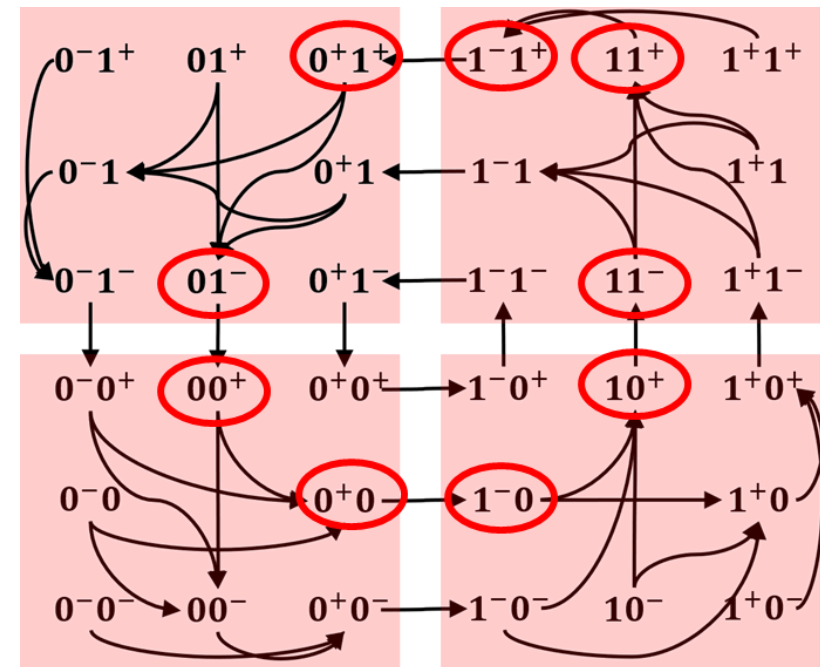
1. Abstraction of the model with discrete domain.



2. Find cycles of discrete domains which contain continuous trajectories.



3. Search for limit cycle(s) inside cycles found in step 2.



$$G(\pi_1) = M \times \pi_1 + b$$

G : Poincaré map

Method: Search $(\pi_0, 00)$ in the compatible zone such that $G(\pi_0) = \pi_0$.

Search for limit cycles

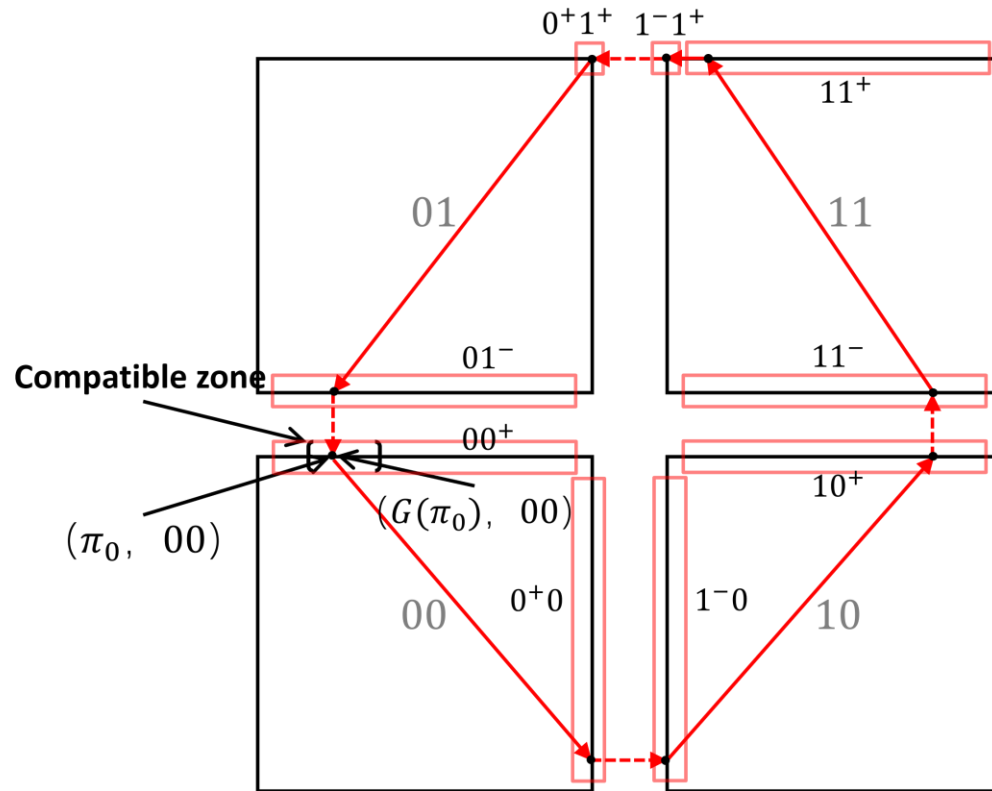
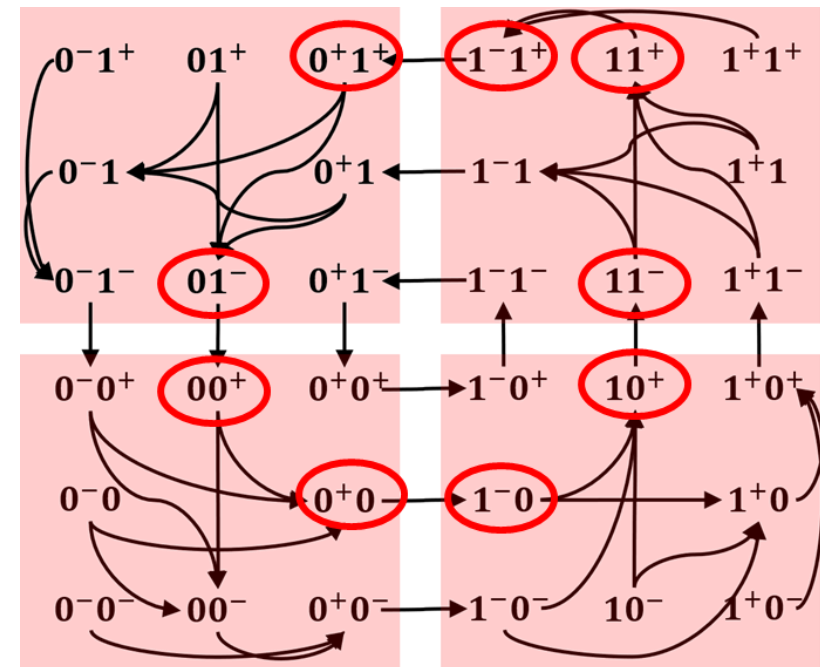
1. Abstraction of the model with discrete domain.



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$$G(\pi_0) = M \times \pi_0 + b$$

G : Poincaré map

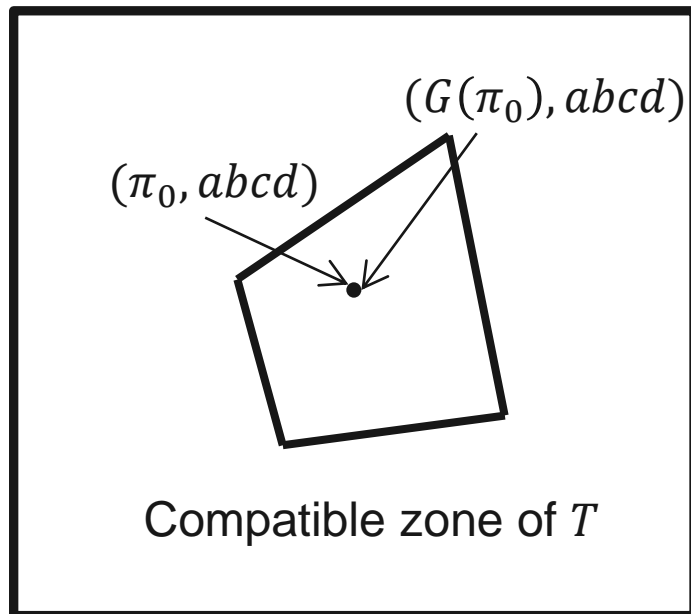
Method: Search $(\pi_0, 00)$ in the compatible zone such that $G(\pi_0) = \pi_0$.



Stability analysis of limit cycle

A cycle of discrete domains $T: a^+b^+cd \rightarrow \dots \rightarrow a^+b^+cd$

Discrete domain a^+b^+cd



$$\pi_0 = \begin{pmatrix} 1 \\ 1 \\ x_0 \\ y_0 \end{pmatrix}$$

$$G(\pi_0) = \pi_0 \Leftrightarrow \begin{pmatrix} x_0 \\ y_0 \end{pmatrix} = A \begin{pmatrix} x_0 \\ y_0 \end{pmatrix} + b$$

Two eigenvalues of A : λ_1, λ_2

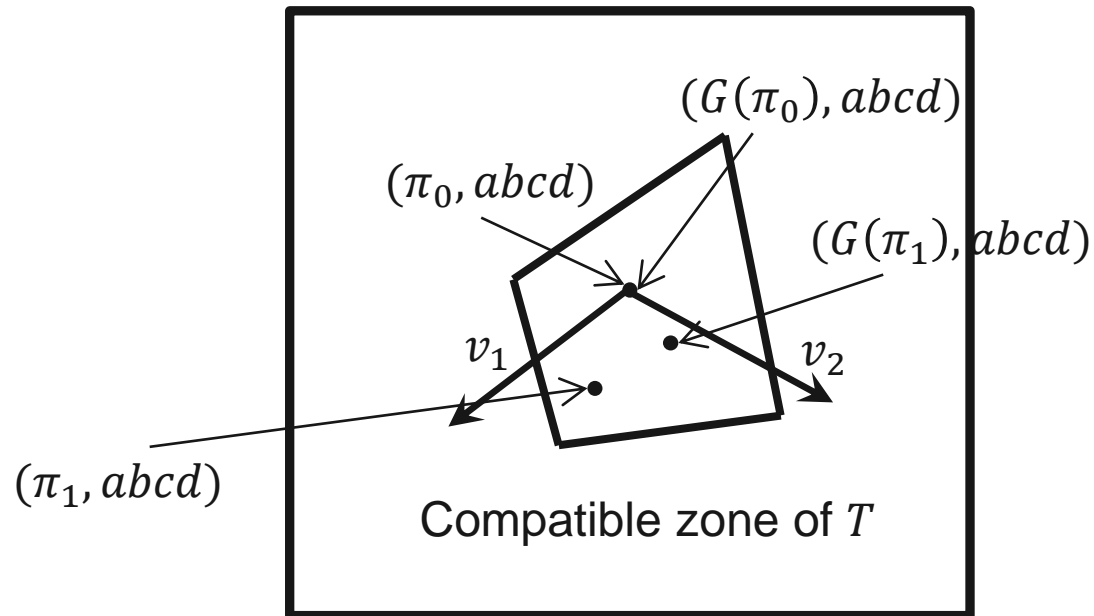
It is a stable limit cycle $\Leftrightarrow |\lambda_i| < 1, i \in \{1,2\}$

Example of stability analysis

Stability analysis of limit cycle

A cycle of discrete domains $T: a^+b^+cd \rightarrow \dots \rightarrow a^+b^+cd$

Discrete domain a^+b^+cd



$$\pi_0 = \begin{pmatrix} 1 \\ 1 \\ x_0 \\ y_0 \end{pmatrix}$$

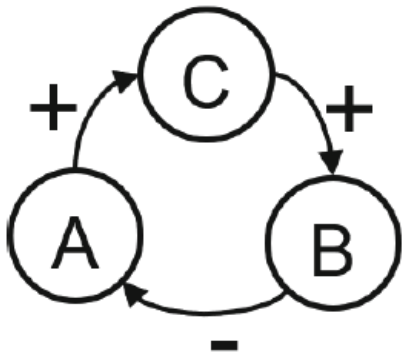
$$G(\pi_0) = \pi_0 \Leftrightarrow \begin{pmatrix} x_0 \\ y_0 \end{pmatrix} = A \begin{pmatrix} x_0 \\ y_0 \end{pmatrix} + b$$

Two eigenvalues of A : $\lambda_1, \lambda_2 \in]0, 1[$

Example of stability analysis



Application



A	B	C	C_A	C_B	C_C
0	0	0	1	-0.6	-0.7
0	0	1	1	0.7	-0.9
0	1	0	-0.8	-0.8	-0.7
0	1	1	-0.8	0.6	-0.9
1	0	0	0.7	-0.6	0.6
1	0	1	0.7	0.7	0.5
1	1	0	-0.9	-0.8	0.6
1	1	1	-0.9	0.6	0.5

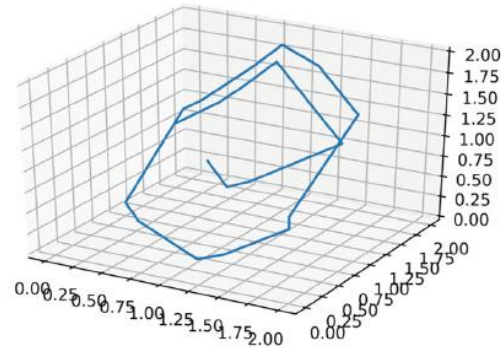
A	B	C	C_A	C_B	C_C
0	0	0	3	-0.6	-0.7
0	0	1	3	0.7	-2.9
0	1	0	-2.8	-0.8	-0.7
0	1	1	-2.8	0.6	-2.9
1	0	0	2.7	-0.6	2.6
1	0	1	2.7	0.7	0.5
1	1	0	-2.9	-0.8	2.6
1	1	1	-2.9	0.6	0.5

A	B	C	C_A	C_B	C_C
0	0	0	3	-0.6	-0.7
0	0	1	3	0.7	-2.9
0	1	0	-0.8	-0.8	-0.7
0	1	1	-0.8	0.6	-2.9
1	0	0	0.7	-0.6	2.6
1	0	1	0.7	0.7	0.5
1	1	0	-2.9	-0.8	2.6
1	1	1	-2.9	0.6	0.5

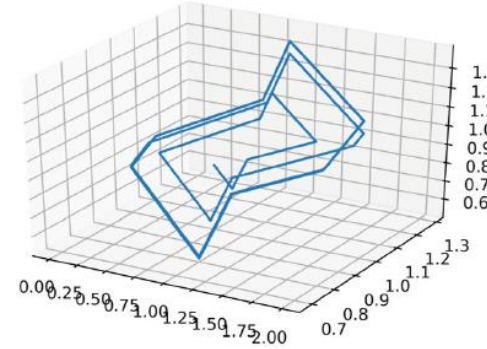
Three HGRNs of negative feedback loop in 3 dimensions



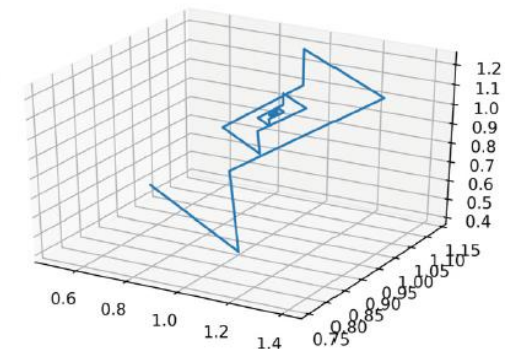
Our analysis method



A stable limit cycle

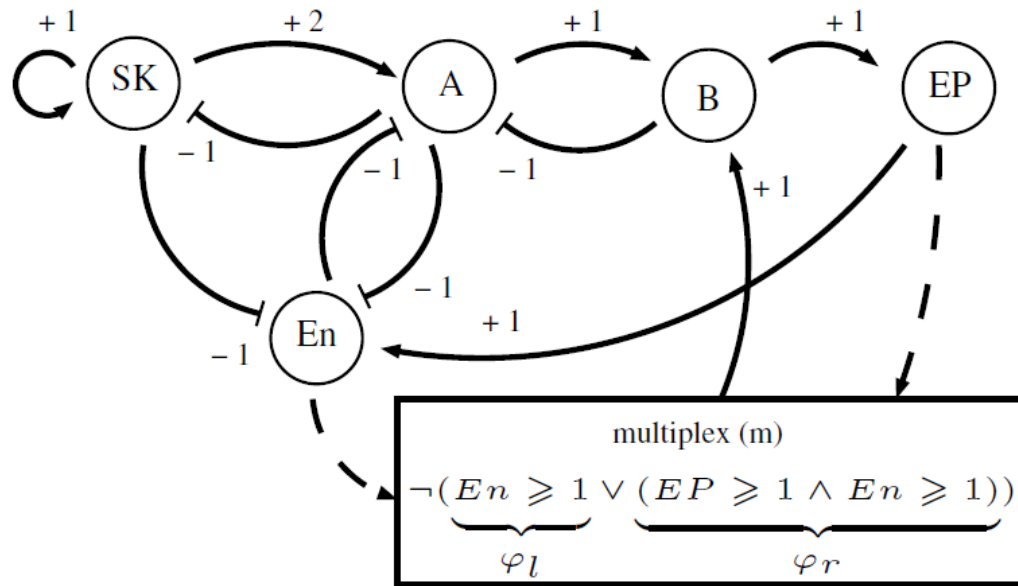


A stable limit cycle



No limit cycle

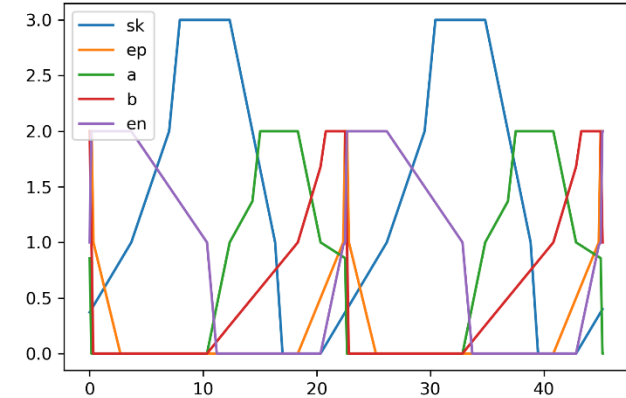
Application



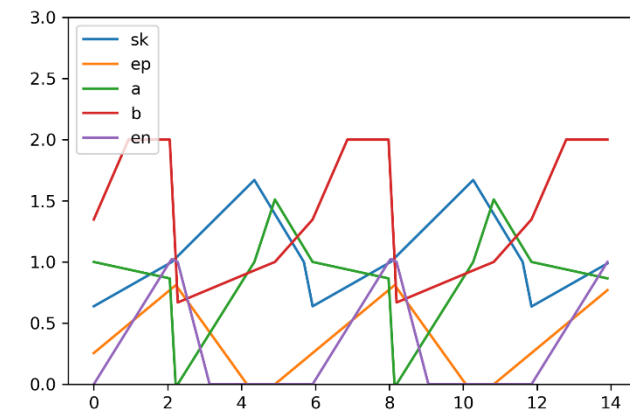
A HGRN of cell cycle

[Behaegel et al., Journal of bioinformatics and computational biology, 2016]

Our analysis method



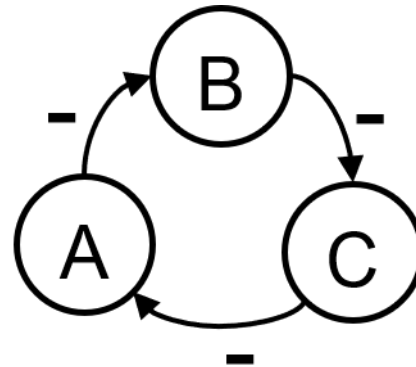
A stable limit cycle



An unstable limit cycle

Repressilator

The specific gene regulatory network studied in this work:



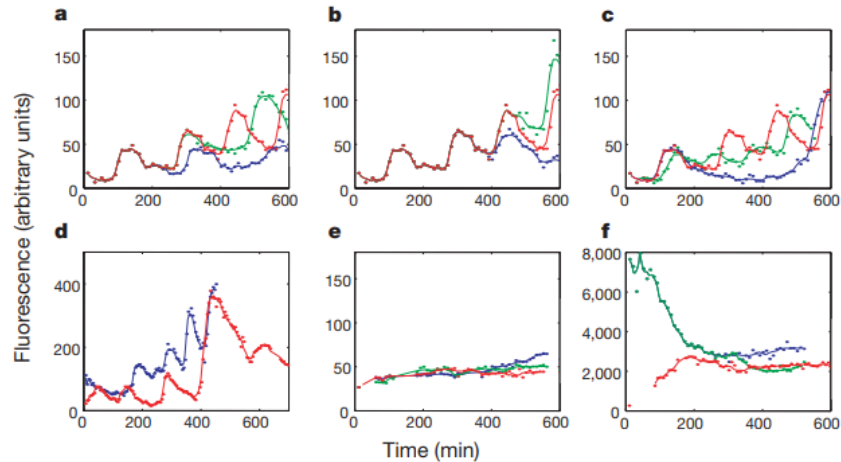
The influence graph of the canonical repressilator



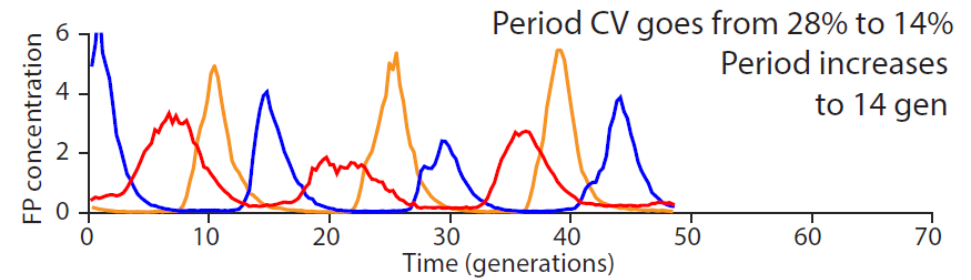
Repressilator

Why study oscillations in repressilator?

- Human-made genetic oscillations have potential medical applications
- Existence of synthetic repressilator with sustained oscillation (but not stable)

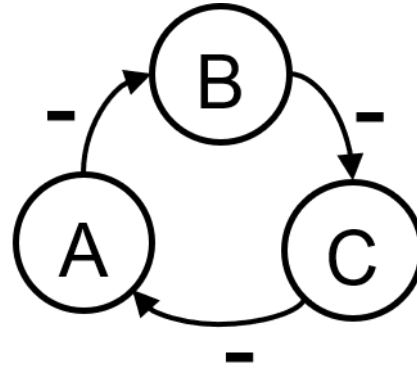


[Elowitz, M. B. et al., Nature, 2000]



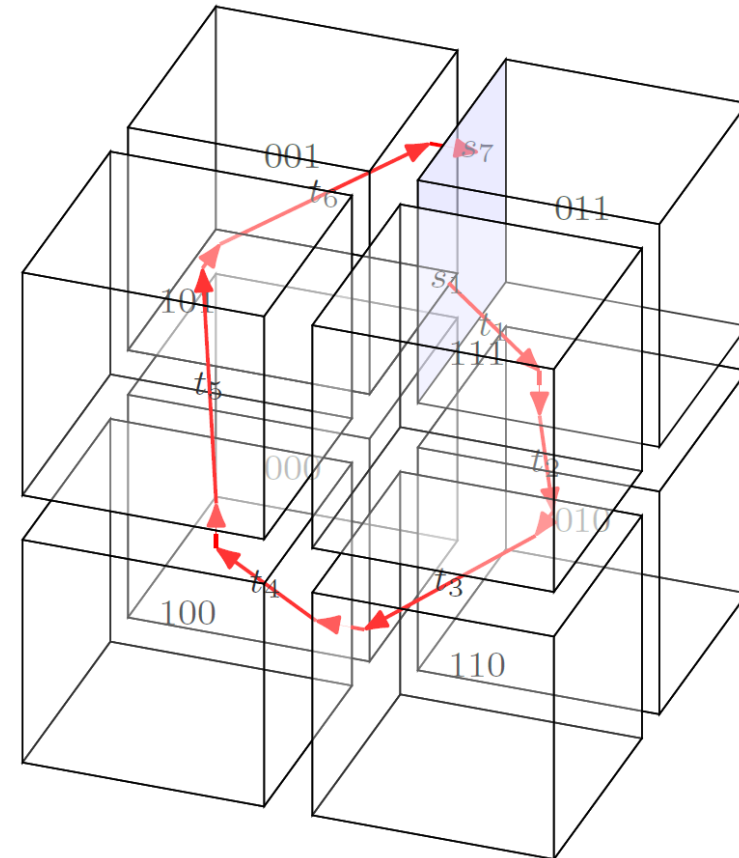
[Potvin-Trottier, L. et al, Nature, 2016]

HGRN of repressilator



A	B	C	C_A	C_B	C_C
0	0	0	C_{ac0a0}	C_{ba0b0}	C_{cb0c0}
0	0	1	$-C_{ac1a0}$	C_{ba0b0}	C_{cb0c1}
0	1	0	C_{ac0a0}	C_{ba0b1}	$-C_{cb1c0}$
0	1	1	$-C_{ac1a0}$	C_{ba0b1}	$-C_{cb1c1}$
1	0	0	C_{ac0a1}	$-C_{ba1b0}$	C_{cb0c0}
1	0	1	$-C_{ac1a1}$	$-C_{ba1b0}$	C_{cb0c1}
1	1	0	C_{ac0a1}	$-C_{ba1b1}$	$-C_{cb1c0}$
1	1	1	$-C_{ac1a1}$	$-C_{ba1b1}$	$-C_{cb1c1}$

All parameters of this HGRN of repressilator



An example of this HGRN of repressilator

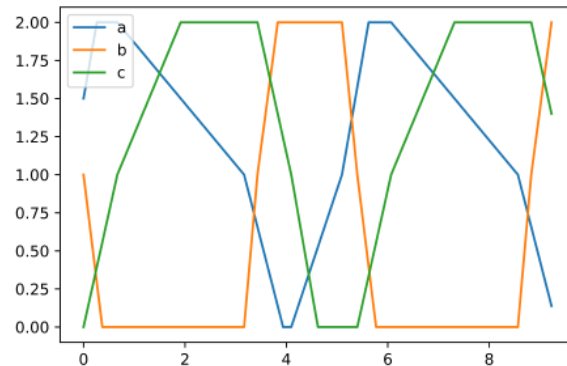


Objective of this work

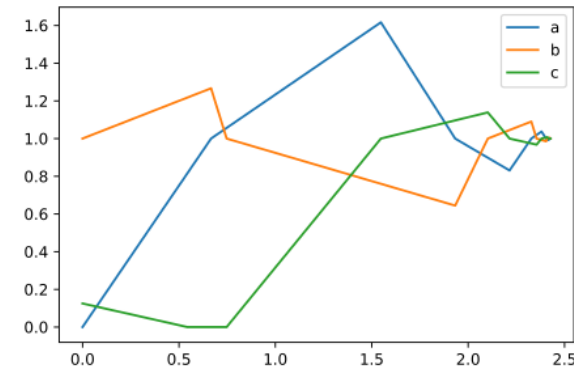
Find constraints on parameters for the existence of sustained oscillations

A	B	C	C_A	C_B	C_C
0	0	0	C_{ac0a0}	C_{ba0b0}	C_{cb0c0}
0	0	1	$-C_{ac1a0}$	C_{ba0b0}	C_{cb0c1}
0	1	0	C_{ac0a0}	C_{ba0b1}	$-C_{cb1c0}$
0	1	1	$-C_{ac1a0}$	C_{ba0b1}	$-C_{cb1c1}$
1	0	0	C_{ac0a1}	$-C_{ba1b0}$	C_{cb0c0}
1	0	1	$-C_{ac1a1}$	$-C_{ba1b0}$	C_{cb0c1}
1	1	0	C_{ac0a1}	$-C_{ba1b1}$	$-C_{cb1c0}$
1	1	1	$-C_{ac1a1}$	$-C_{ba1b1}$	$-C_{cb1c1}$

All parameters of this HGRN of repressilator



A model with sustained oscillations



A model without sustained oscillation

Related work

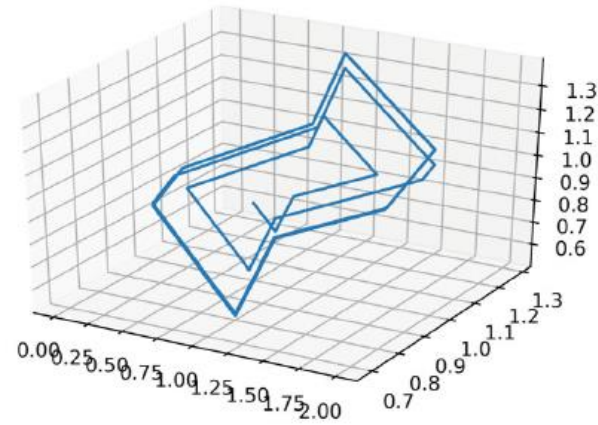
This work relies on our previous work [Sun, H. et al., CMSB, 2022]

Contributions of [Sun, H. et al., CMSB, 2022]:

- Automatic search of limit cycle
- Stability analysis of limit cycle

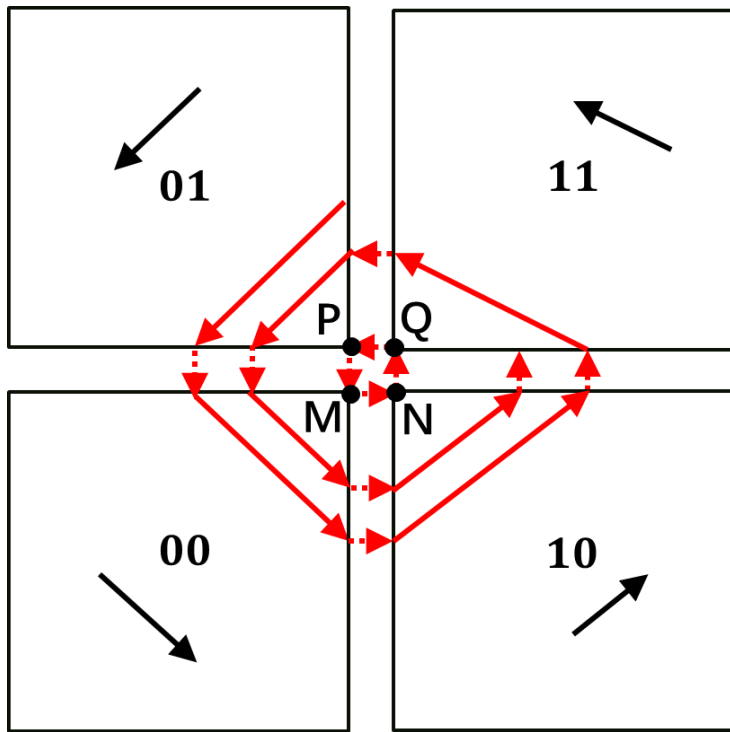
A	B	C	C_A	C_B	C_C
0	0	0	3	-0.6	-0.7
0	0	1	3	0.7	-2.9
0	1	0	-2.8	-0.8	-0.7
0	1	1	-2.8	0.6	-2.9
1	0	0	2.7	-0.6	2.6
1	0	1	2.7	0.7	0.5
1	1	0	-2.9	-0.8	2.6
1	1	1	-2.9	0.6	0.5

Method of
[Sun, H. et al., CMSB, 2022]

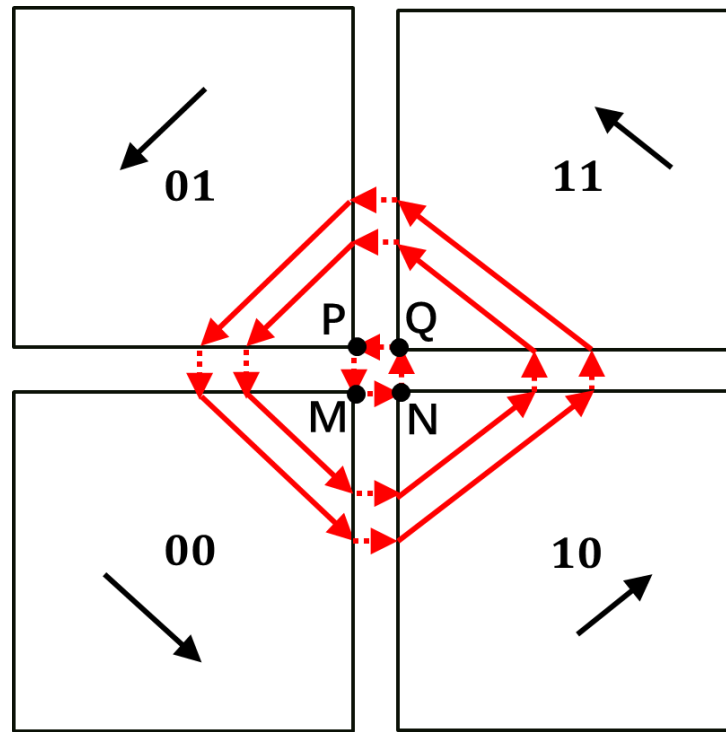


Different qualitative behaviors of this HGRN

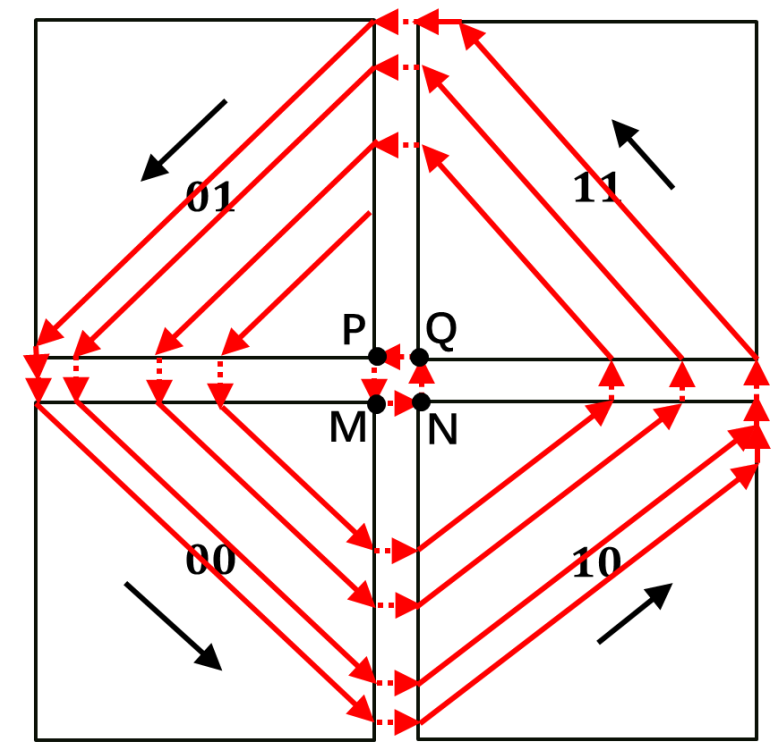
Illustration of different qualitative behaviors by a 2-Dim model



Damped oscillations



Parallel cycles
(special case of sustained oscillations)

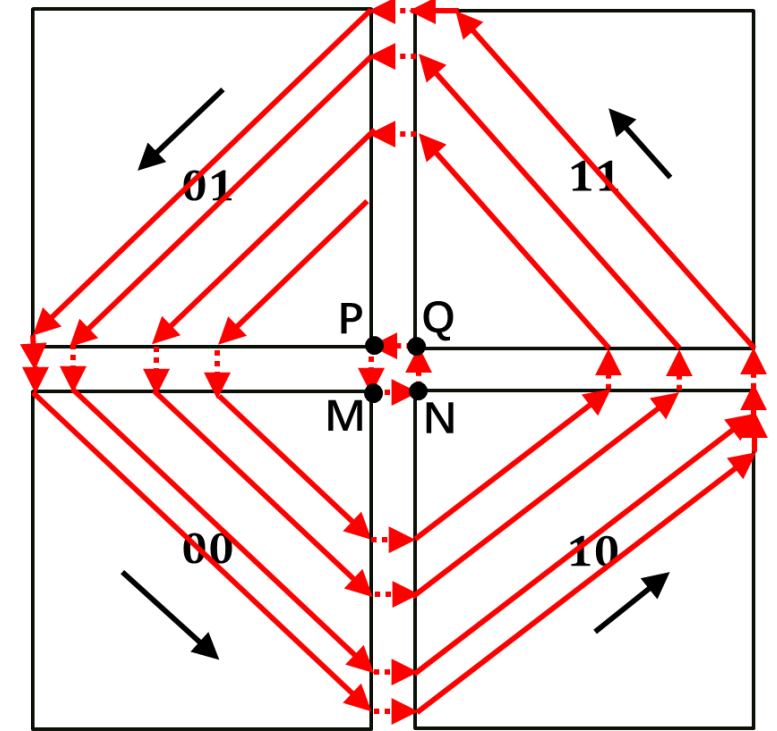
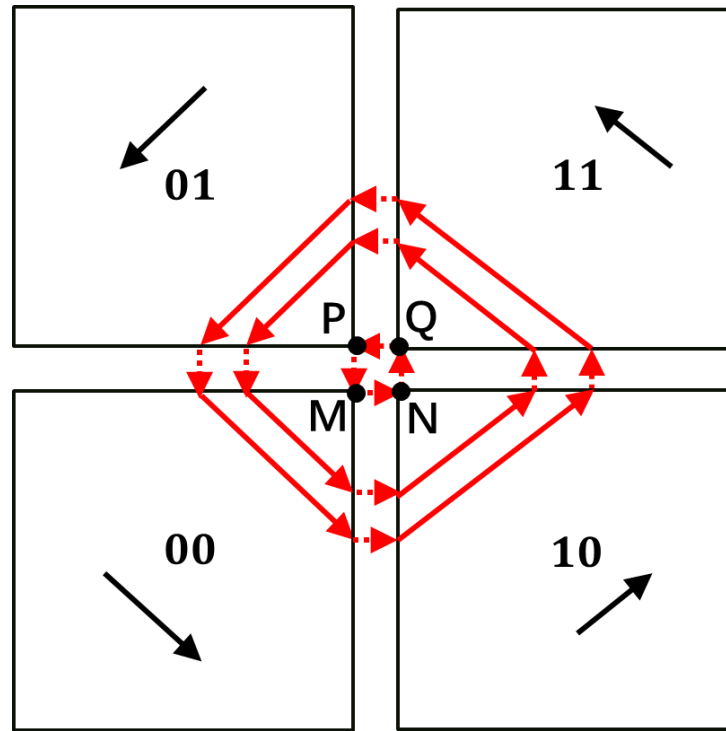
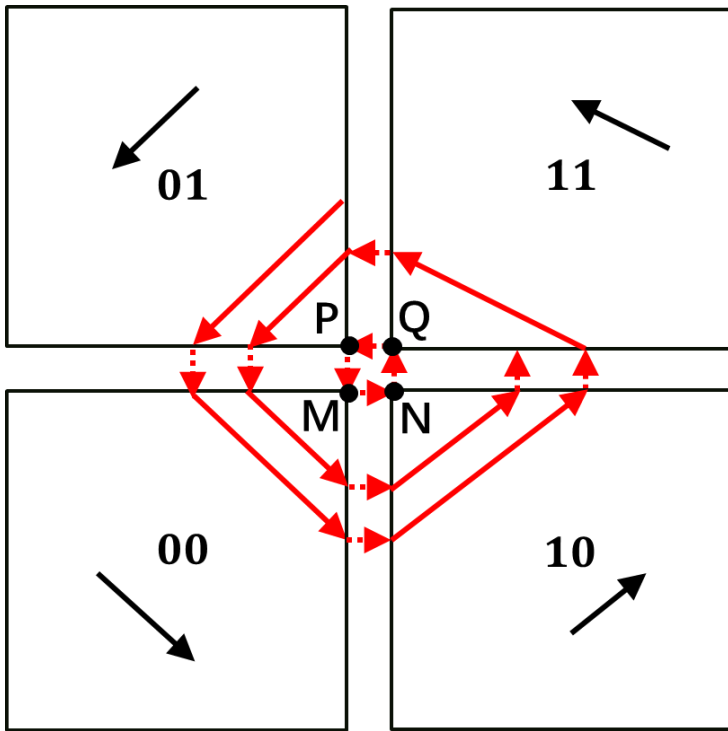


Sustained oscillations

Different qualitative behaviors of this HGRN

How to determine different qualitative behaviors?

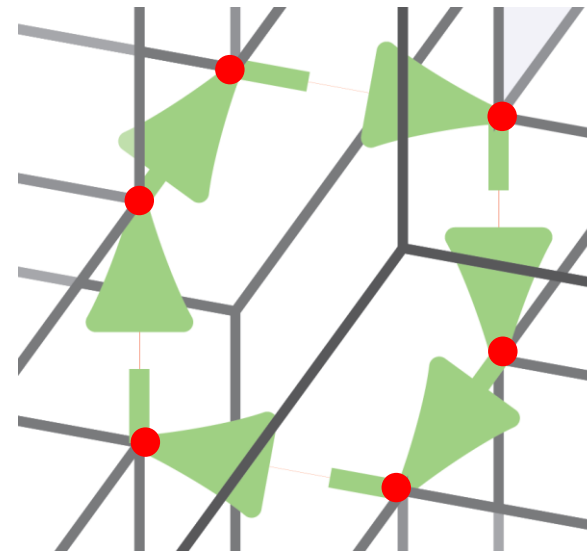
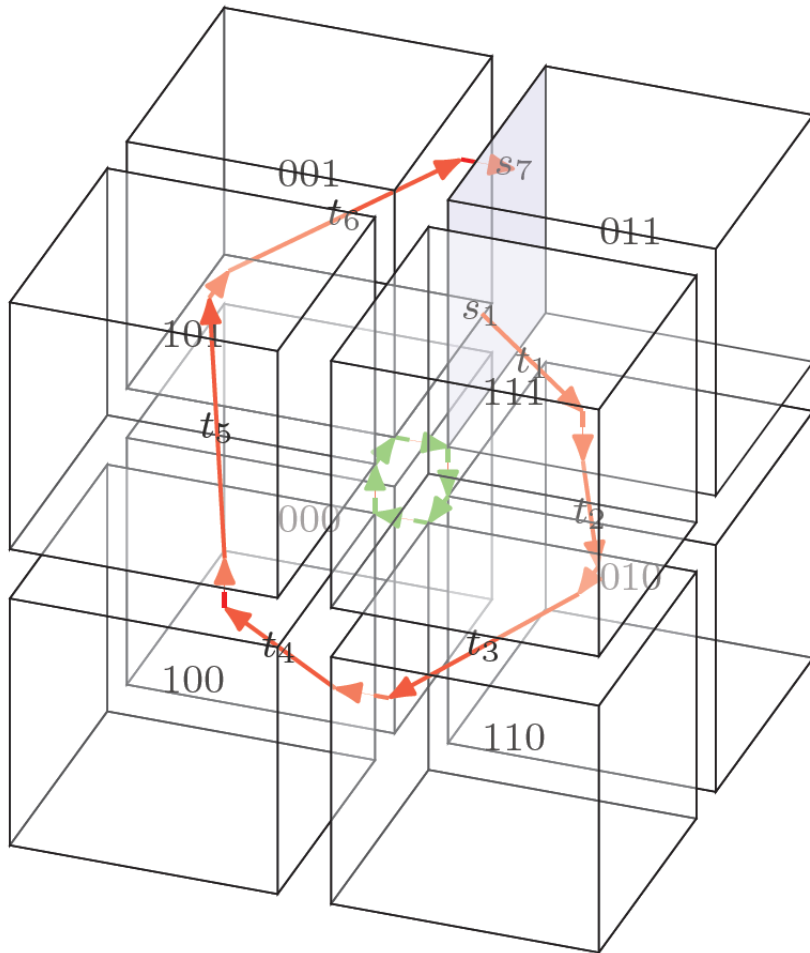
Existence of sustained oscillation \Leftrightarrow Stability of the characteristic state



Different qualitative behaviors of this HGRN

How to determine different qualitative behaviors?

Existence of sustained oscillation \Leftrightarrow Stability of the characteristic state

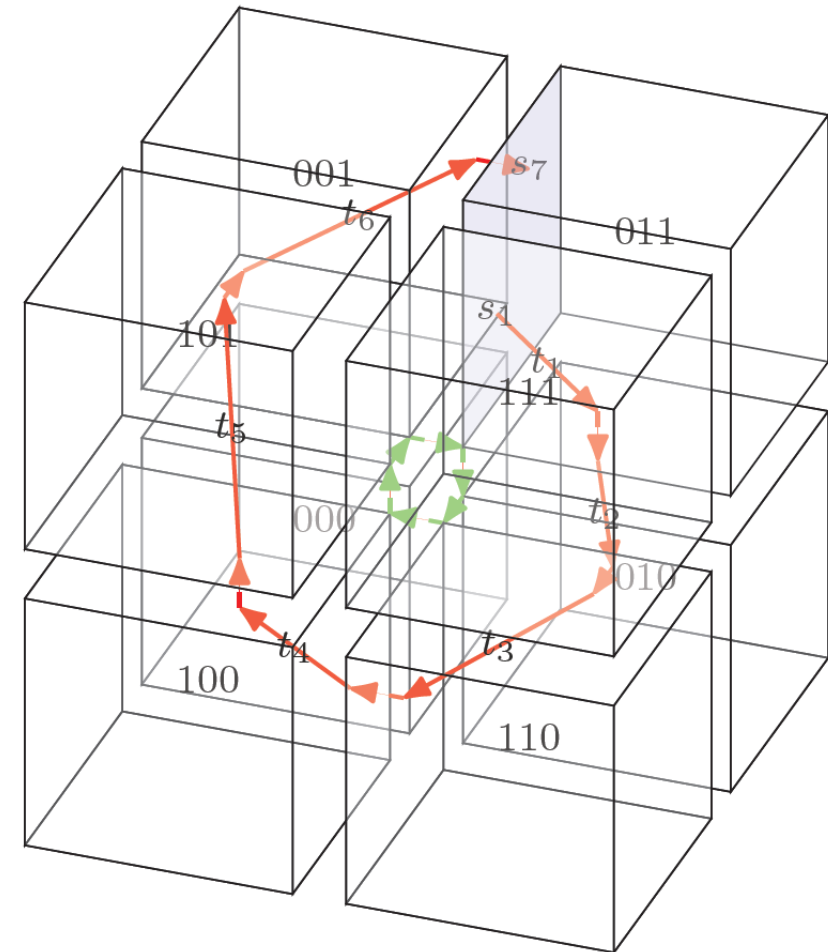


The characteristic state of this HGRN of repressilator

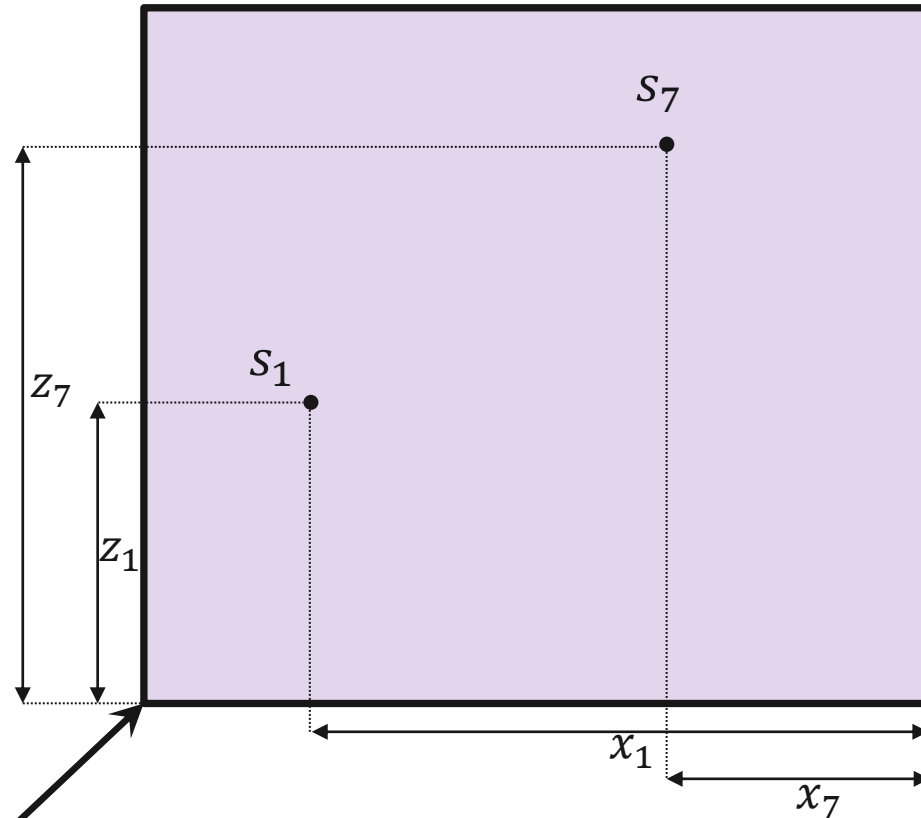


Determine the stability of the characteristic state

Use the Poincaré map (first return map) to determine the stability of the characteristic state



characteristic state



$$\begin{pmatrix} x_7 \\ z_7 \end{pmatrix} = A \begin{pmatrix} x_1 \\ z_1 \end{pmatrix} + b$$



Determine the stability of the characteristics state

Use the Poincaré map (first return map) to determine the stability of the characteristic state

The two eigenvalues of A : λ_1, λ_2 ($\lambda_1 > \lambda_2$)

The characteristic state is unstable $\Leftrightarrow \lambda_1 \geq 1$

Illustration of the case $\lambda_1 \geq 1$

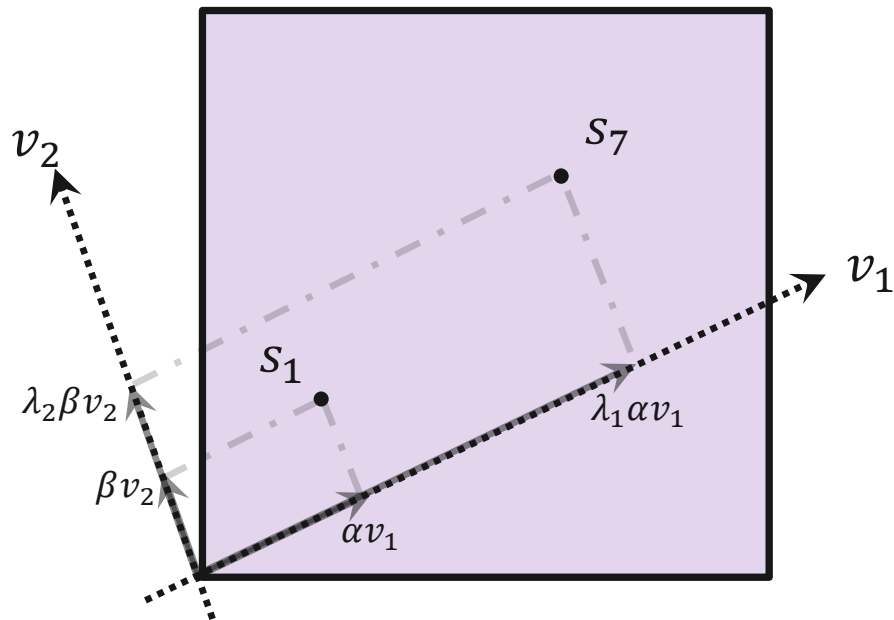
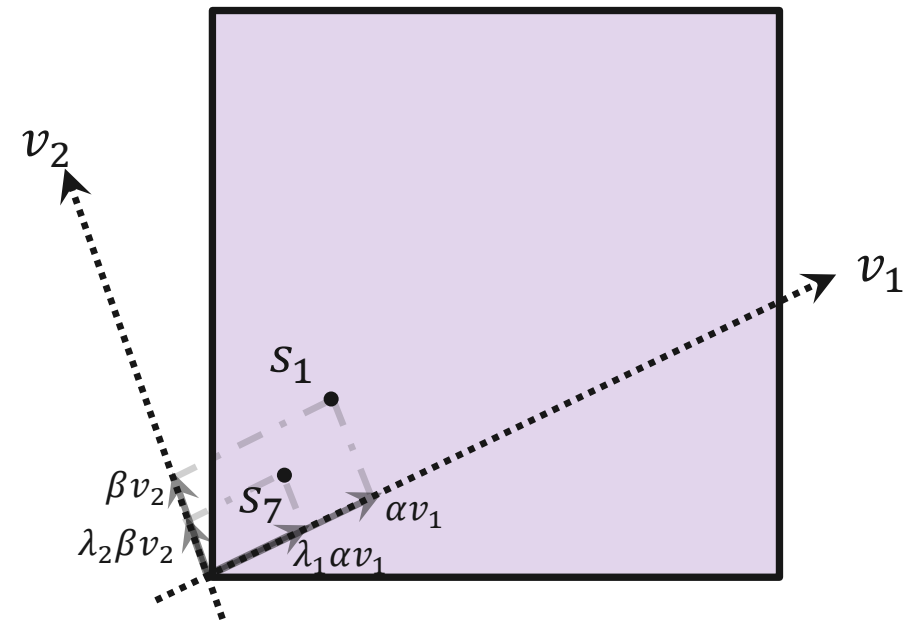


Illustration of the case $\lambda_1 < 1$





Condition for sustained oscillations

Sustained oscillations exist



The characteristic state is unstable



$$\lambda_1 \geq 1$$



By computing the expression of λ_1

$$(P_1 \geq 0) \vee (P_2 \geq 0)$$

P_1 and P_2 are polynomials on parameters

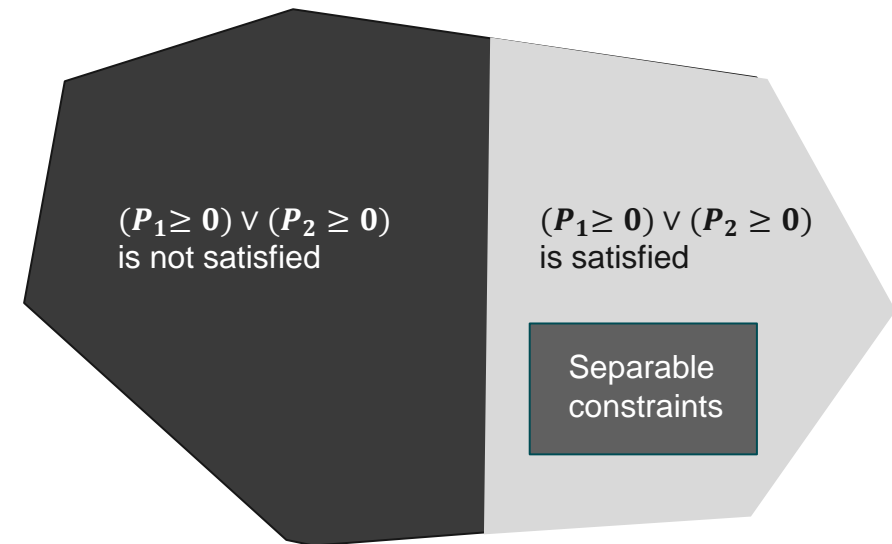


Separable constraints on parameters

Separable constraints: $C_{xyixj} \in [\underline{C_{xyixj}}, \overline{C_{xyixj}}]$

The objective is to find separable constraints such that under these constraints $(P_1 \geq 0) \vee (P_2 \geq 0)$ is always satisfied.

A	B	C	C_A	C_B	C_C
0	0	0	C_{ac0a0}	C_{ba0b0}	C_{cb0c0}
0	0	1	$-C_{ac1a0}$	C_{ba0b0}	C_{cb0c1}
0	1	0	C_{ac0a0}	C_{ba0b1}	$-C_{cb1c0}$
0	1	1	$-C_{ac1a0}$	C_{ba0b1}	$-C_{cb1c1}$
1	0	0	C_{ac0a1}	$-C_{ba1b0}$	C_{cb0c0}
1	0	1	$-C_{ac1a1}$	$-C_{ba1b0}$	C_{cb0c1}
1	1	0	C_{ac0a1}	$-C_{ba1b1}$	$-C_{cb1c0}$
1	1	1	$-C_{ac1a1}$	$-C_{ba1b1}$	$-C_{cb1c1}$



All parameters of this HGRN of repressilator

Illustration of separable constraints in the parameter space



Bernstein coefficients

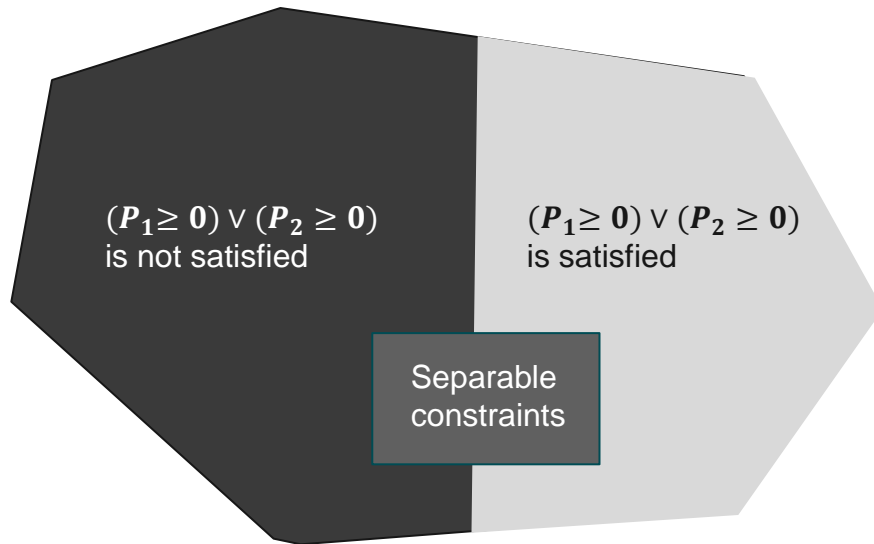
One property of Bernstein coefficients:

For any polynomial $f: \mathbb{R}^n \rightarrow \mathbb{R}$, we can compute the Bernstein coefficients $\{b_i\}$, $b_i \in \mathbb{R}$, such that:

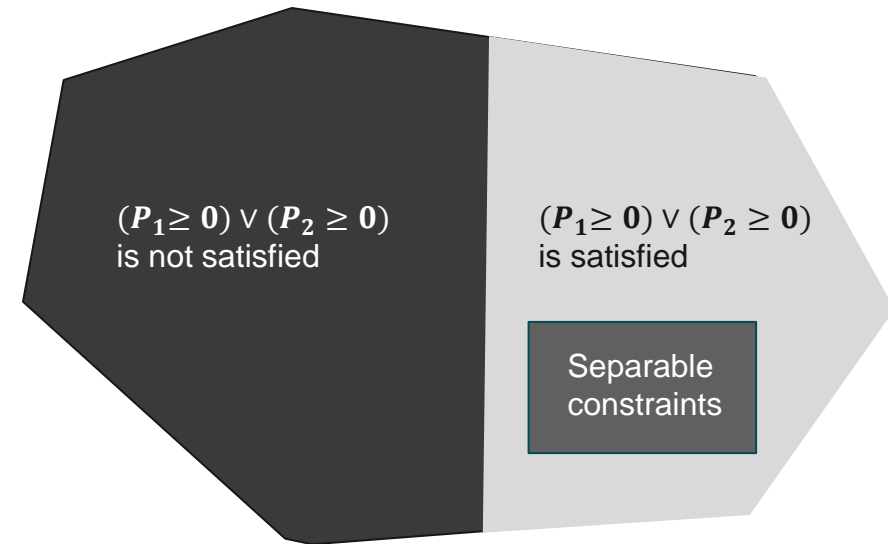
$$\text{Min}(\{b_i\}) \leq f(x) \leq \text{Max}(\{b_i\}), \forall x \in [0,1]^n$$

Satisfiability under separable constraints

Given separable constraints: $C_{xyixj} \in [\underline{C_{xyixj}}, \overline{C_{xyixj}}]$, $(P_1 \geq 0) \vee (P_2 \geq 0)$ is always satisfied?



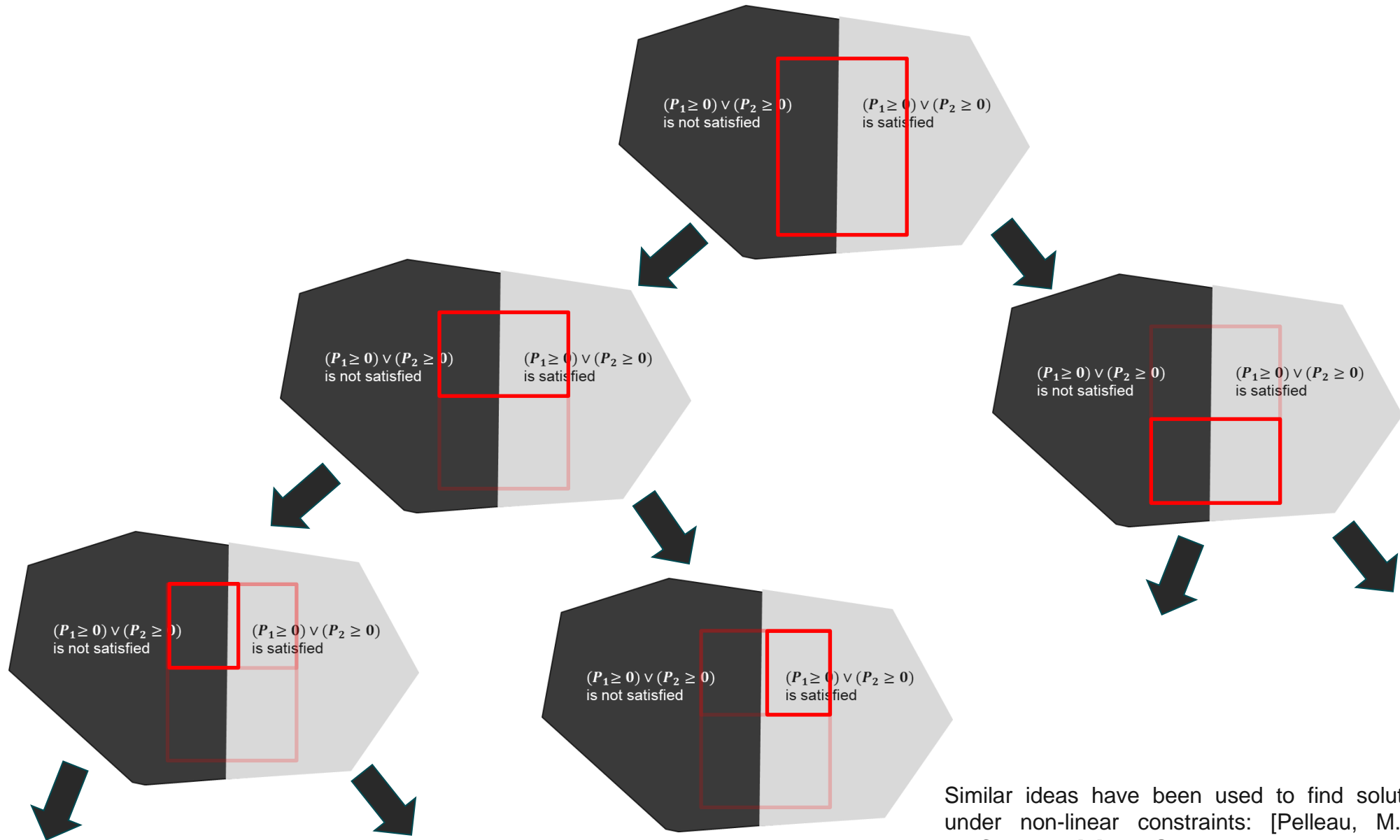
Example of separable constraints which do not satisfy $(P_1 \geq 0) \vee (P_2 \geq 0)$



Example of separable constraints which satisfy $(P_1 \geq 0) \vee (P_2 \geq 0)$



Search of separable constraints by a splitting strategy



Similar ideas have been used to find solution sets under non-linear constraints: [Pelleau, M. et al., VMCAI, 2013], [Ziat, G. et al., Formal Methods, 2019]



An example of separable constraints

A	B	C	C_A	C_B	C_C
0	0	0	C_{ac0a0}	C_{ba0b0}	C_{cb0c0}
0	0	1	$-C_{ac1a0}$	C_{ba0b0}	C_{cb0c1}
0	1	0	C_{ac0a0}	C_{ba0b1}	$-C_{cb1c0}$
0	1	1	$-C_{ac1a0}$	C_{ba0b1}	$-C_{cb1c1}$
1	0	0	C_{ac0a1}	$-C_{ba1b0}$	C_{cb0c0}
1	0	1	$-C_{ac1a1}$	$-C_{ba1b0}$	C_{cb0c1}
1	1	0	C_{ac0a1}	$-C_{ba1b1}$	$-C_{cb1c0}$
1	1	1	$-C_{ac1a1}$	$-C_{ba1b1}$	$-C_{cb1c1}$

$$\begin{array}{lll} C_{ac0a0} \in [0.0, 0.5] & C_{ba0b0} \in [0.0, 0.5] & C_{cb0c0} \in [0.0, 0.5] \\ C_{ac0a1} \in [0.0, 0.5] & C_{ba0b1} \in [0.0, 0.5] & C_{cb0c1} \in [0.0, 0.5] \\ C_{ac1a0} \in [0.5, 1.0] & C_{ba1b0} \in [0.5, 1.0] & C_{cb1c0} \in [0.5, 1.0] \\ C_{ac1a1} \in [0.0, 1.0] & C_{ba1b1} \in [0.0, 1.0] & C_{cb1c1} \in [0.0, 1.0] \end{array}$$

All parameters of this HGRN of repressilator

Conclusion

State of the art

- Hybrid formalism: between discrete networks and ODEs
- How to find the parameters?
- How to formally analyze the dynamics?

Parameters inference

- Hybrid Hoare logic with hybrid Dijkstra predicate calculus
- Optimization algorithm (parameters and thresholds)

Formal analysis of the dynamics

- Enumeration of simple cycles
- Formal verification of cycle stability with Poincaré map

Collaborations & Bibliography



**Jonathan
BEHAEGEL**



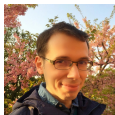
**Jean-Paul
COMET**

- Jonathan Behaegel, Jean-Paul Comet, Maxime Folschette. [Constraint Identification Using Modified Hoare Logic on Hybrid Models of Gene Networks](#). *International Symposium on Temporal Representation and Reasoning (TIME'17)*, 2017.

Collaborations & Bibliography

Differential equations
 System biology
 Stability analysis
 Circadian rhythm
 Formal verification
 Limit cycle
 Time series data
 Piecewise affine system
 Hybrid model
 Gene regulatory networks
 Cell cycle
 Boolean networks
 Poincaré map
 Repressilator
 Parameter identification
 Invariant sets
 Sum-of-squares
 Bi-inspired heuristics
 Model checking

**Honglu
SUN**



**Morgan
MAGNIN**



**Jean-Paul
COMET**

- Honglu Sun, Maxime Folschette, Morgan Magnin. [Limit cycle analysis of a class of hybrid gene regulatory networks](#). 20th International Conference on Computational Methods in Systems Biology (CMSB'22), *Lecture Notes in Computer Science*, vol. 13447, 2022.
- Honglu Sun, Jean-Paul Comet, Maxime Folschette, Morgan Magnin. [Condition for sustained oscillations in repressilator based on a hybrid modeling of gene regulatory networks](#). *International Conference on Bioinformatics Models, Methods and Algorithms (BIOINFORMATICS 2023)*, 2023.

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- Émilien Cornillon, Jean-Paul Comet, Gilles Bernot, Gilles Énée. [Hybrid gene network: a new formalism and a software environment](#). *Modelling Complex Biological Systems in the Context of Genomics*, 2016.
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- Marie Pelleau, Antoine Miné, Charlotte Truchet, Frédéric Benhamou. [A Constraint Solver based on Abstract Domains](#). 14th International Conference on Verification, Model Checking, and Abstract Interpretation (VMCAI'13), *Lecture Notes in Computer Science*, volume 7737, 2013.