# Towards the use of Process Hitting to tackle biological observations inconsistent with background knowledge

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# Motivations

- Given an existing network (background knowledge) and a (new) observation that is inconsistent with this network, how can we automatically design the minimal set of missing actions that can mimic the observation?
- Process Hitting is an efficient framework to cope with large networks (~ 100 components)

# Motivations

- Our proposition: design a method taking advantage of the Process Hitting methods to address the completion of networks with inconsistent observations
- Restrictions w.r.t. current work:
  - Consider only addition of actions, not removal of actions
  - Modeling of the evolution of a gene expression in case of ko w.r.t. wild type, under steady state assumption

# Overview

- Motivating example
- Reminder about the Process Hitting framework
- 4-level based logics and associated truth tables
- Translating 4-level based models into Process Hitting
- Further discussions

# Motivating example

Background theory B: Boolean network consisting of the three Boolean functions

- Mig1p = not GRR1
- Rgt1p = not (Mig1p & RGT1)
- YGL157w = not Rgt1p
- Observation O:



When GRR1 is ko, then the gene expression of YGL157w decreases, i.e.:

When the gene expression of GRR1 decreases, the gene expression of YGL157w also decreases. ( we write it by *promoted(ygl157w, grr1)* )

# Inconsistency between B and O

 □ Given the following initial state, we meet the fact that the gene expression of YGL157w decreases
 < GRR1 = -1, Mig1p = 0, Rgt1p = 0, RGT1 = 0, YGL157w = 0 >
 ⇒ This is *inconsistent* with the observation...



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[Paulevé et al., Transactions on Computational Systems Biology, 2011]



**Sorts**: components *a*, *b*, *z* 

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[Paulevé et al., Mathematical Structures in Computer Science, 2012]



 $\begin{array}{l} \rightarrow \text{ Concretization of the objective} = \text{scenario} \\ \underline{a_0 \rightarrow c_0 \ l^{\circ} \ c_1} :: b_0 \rightarrow d_0 \ l^{\circ} \ d_1 :: c_1 \rightarrow b_0 \ l^{\circ} \ b_1 :: b_1 \rightarrow d_1 \ l^{\circ} \ d_2 \end{array}$ 

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Computing *P* or *Q* is much simpler (roughly **polynomial**)  $\rightarrow$  Efficient for big models  $\rightarrow$  **Hundredths of seconds** 

#### Enrichment of the Process Hitting

[Folschette et al., CS2Bio'13, 2013]

Several additions to improve the expressiveness of the Process Hitting:

- Priorities
  - Groups of actions with similar temporal/probabilistic parameters
- Neutralizing edges
  - Atomistic delay relations between actions
- Synchronous actions
  - Multiple reactants and products  $\rightarrow$  Biochemical reactions

All these formalisms can be translated to a canonical form

A new static analysis has been developed to check reachability properties  $\rightarrow$  Efficient dynamic analysis on large models

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- Each action is linked to a class of priority
- An action is playable only if no action with a higher priority is playable



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• Allows to model classes of actions with similar temporal/stochastic parameters



#### Neutralizing edges



- Allows to integrate temporal data about relative reaction delays
- Atomistic preemptions

$$c_0 
ightarrow d_0 
vert^{
ightarrow} d_1$$
 cannot be played while

$$a_0 \rightarrow b_0 
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#### Synchronous actions

- Synchronization between actions:
  - Presence of catalysts
  - Consumption of reactants
  - Creation of products
- Convenient for biochemical equations:  $X \xrightarrow{Y} Z$  in the following form:  $\{X_1, Y_1, Z_0\} \rightarrowtail \{X_0, Z_1\}$

 $\begin{aligned} h_1 &= \{c_1\} \rightarrowtail \{c_0\} \\ h_2 &= \{a_0, b_1, c_0, d_0\} \rightarrowtail \{c_1, d_1\} \end{aligned}$ 

All processes of Amust be present to play  $A \rightarrow B$ 

After playing  $A \rightarrow B$ , all processes of B are active



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# Modeling ideas

- 4 cases to consider:
  - The concentration of a component c in ko of a given gene g is **higher** than its concentration in Wild Type (which will be denoted  $\uparrow$ )
  - The concentration of a component c in case of ko of a given gene g is **lower** than its concentration in Wild Type (which then will be denoted  $\downarrow$ )
  - The concentration of a component c in case of ko of a given gene g is stable compared to Wild Type (which then will be denoted -)
  - When the evolution of the concentration of a component c between ko and wild type is **unknown**: add a fourth level « unknown" in the logical framework, but not necessary in the Process Hitting final representation.

# Our stoichiometric modeling

- A and B: denoting the effect by the complex of A and B
  - $\Rightarrow$  Strength: depending on the amount of the complex



❑ A or B: denoting the (individual) effects by A and B ⇒ Strength: depending on the amount of both A and B



# Truth table in 4 valued logic (1/2)

1: increase. ↓: decrease. -: unchanged.

Α	В	A and B	A or B
Ť	Ť	Ť	Ť
Ť	Ļ	Ļ	unknown
Ť	-	-	1
1	unknown	unknown	unknown
Ļ	Ļ	Ļ	Ļ
Ļ	-	Ļ	Ļ
Ļ	unknown	Ļ	unknown
-	-	-	-
-	unknown	unknown	unknown
unknown	unknown	unknown	unknown

# Truth table in 4 values logic (2/2)

↑: increase. ↓: decrease. -: unchanged.

Α	٦A	
1	Ļ	
Ļ	1	
-	-	
unknown	unknown	

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# Principle of the translation of « 4 valued logics » into PH

- When A has more than one regulator, use a cooperative sort to update A according to the state of regulators -> need to use priorities in PH
- « unknown » is modeled by modeling every potential underlying behavior

### Translating 4 valued logics into Process Hitting: A=B

A = B
 A = not B
 A = B and C
 A = B or C



# Translating 4 valued logics into Process Hitting

A = B
 A = not B
 A = B and C
 A = B or C



 Maybe add a slide with the translation of A = B and C, but the resulting PH is quite complex ?

### Back to the example



### Back to the example: one execution



### Back to the example: one execution



### And with synchronous semantics?



# **Our question**

In case that the dynamics of the model does not encompass the observation into any playable scenario of actions... how to **detect missing actions as few as possible** that can lead the goal state?

# **Related discussions**

- Asynchronous versus synchronous semantics, w.r.t. the addition of priorities
- Compare 4-valued logics with existing approaches with ODEs
- Interest for a cut-sets based approach

# **Cut-sets in Process Hitting**

- Sets of necessary processes that, should they be disabled, would prevent the considered reachability
- Useful to **refute a model**: if a cut set computed from the model does not prevent the reachability in the concrete (modeled) system, then it is a proof that there exists concrete behaviors that are not reproducible by the model.
- See (Paulevé et al., 2014) and Loïc's talk last year

Finding actions for explaining the observation with the background theory (Boolean network)



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### Research plan and future work

- Formalize an algorithmic approach to tackle this completion problem
- Study models with **feedback loops** and extend the principle of 4-valued logics
- Tackle models with time series data as input