

# Reaction networks with delays applied to toxicity analysis

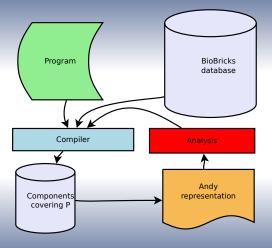
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BioPPN 2014



# The SYNBIOTIC project





### Motivation

- Main goal: design of artificial bio-systems
- How: development of computer-aided tools
- What: specification and analysis of cellular regulation networks (i.e., genetic and signalization networks and metabolic pathways)



# Requirements

#### We want to build a model where:

- different regulatory networks can be expressed
- safety properties can be guaranteed

#### Safety

- ullet in general  $\Rightarrow$  nothing bad can happen
- ullet in a bio-framework  $\Rightarrow$  the system do not exhibit toxic behaviors



# Toxicology

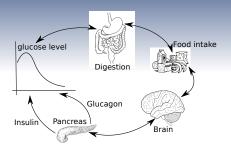
- The toxicity process is a sequence of physiological events that causes the abnormal behavior of a living organism with respect to its healthy state.
- Healthy physiological states generally correspond to homeostasis.
- Toxicity highly depends on the exposure time and the thresholds dosage delimiting the ranges of safe and hazardous effects.

#### Definition (Toxicity)

Toxicity is the deregulation of the homeostasis processes



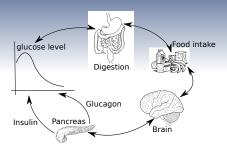
# Blood glucose regulation



- Glucose regulation is a homeostatic process.
- Glycemia is regulated by insulin and glucagon.
- Assimilation of sugars vs aspartame.



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#### Toxic!

Assimilation of food (even if it contains aspartame) should calm hunger and induce satiety, not the opposite!



### **Features**

#### Our model features

- An explicit notion of discrete time
- Species with expression levels and decay
- Reactions with duration



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#### ANDy

An ANDy network is a set of species  ${\mathcal S}$  governed by a set of reactions  ${\mathcal R}$ 



# Species

- Species have a finite number  $\mathcal{L}_s$  of expression levels.
- Each species  ${\it s}$  is initialized at level  $\eta_{\it s}$  and it decays gradually as time passes by.
- Duration of decay vary among levels:

$$\delta_s: [0..\mathcal{L}_s - 1] \to \mathbb{N}^+ \cup \{\omega\}.$$

$$\delta_s(0) = \omega.$$



#### Reactions

Reactions govern evolution of species

$$\rho ::= A_{\rho} ; I_{\rho} \xrightarrow{\Delta} R_{\rho}$$

- $A_{
  ho}$ ,  $I_{
  ho}$  are sets of pairs  $(m{s},\eta_{m{s}})$
- $R_{
  ho}$  is a set of pairs  $(s, \pm n)$
- Each reaction has a response time

$$\Delta:\mathcal{R}\to\mathbb{N}^+$$

Time required for yielding increase (+) and/or decrease (-) of levels of results.



# **Dynamics**

#### A reaction of response time $\Delta$ can take place if

- each activator/reactant stays at least at a given level
- each involved inhibitor is at most at a given level during the whole reaction time.
  - Outcome: the level of results of the reaction can be increased or decreased.



#### **Formalization**

The dynamics of ANDy is formalized using high-level Petri nets.

- Time is explicitly represented.
- Places: Species + 1 place for time
- Transition: Reaction + 1 transition for time



## **Places**

- We assume a unique discrete global clock that starts at zero and always shows the current date (timestamp).
- Each species is represented by a place
- The state of a species s is a tuple  $\langle I_s, u_s, \lambda_s \rangle$ 
  - Is stores the current level;
  - $u_s$  is a timestamp recording the last date when the level has been updated;
  - $\lambda_s$  is a tuple of timestamps with  $\mathcal{L}_s$  fields;



#### Transitions - 1

ANDy networks can evolve in two ways:

- lacktriangle as effect of an enabled reaction  $\rho$
- as an effect of the clock:



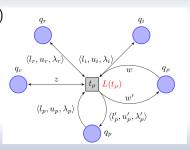
#### Transition: reaction

#### Transition guard:

$$w < z \wedge w' = z \wedge \wedge \\ \bigwedge_{(a,\eta_a) \in A_{\rho}} (l_a \ge \eta_a \wedge z - \lambda[\eta_a] \ge \Delta(\rho) \\ \bigwedge_{(i,\eta_i) \in I_{\rho}} (l_i < \eta_i \wedge z - \lambda[\eta_i] \ge \Delta(\rho))$$

Result: a result r at level  $I_r$  and the clock at time t

$$(r,+1)$$
  
 $\langle I_r, u_r, \lambda_r \rangle \rightarrow \langle I_r + 1, t, \lambda_r \{ {}^t\!/I_r + 1 \} \rangle$   
 $(r,-1)$   
 $\langle I_r, u_r, \lambda_r \rangle \rightarrow \langle I_r - 1, t, \lambda_r \{ {}^t\!/I_r \} \rangle$ 

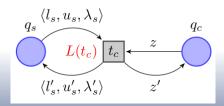




#### Transitions – Clock

- as an effect of the clock:
  - The timestamp t stored in the clock is incremented by one (t+1).
  - A species may stay at level I for  $\delta(I)$  time units. Decay happens as soon as the interval  $\delta(I)$  is elapsed ,

$$\langle I, u, \lambda \rangle \rightarrow \langle I - 1, t + 1, \lambda \{t + 1/I\} \rangle$$





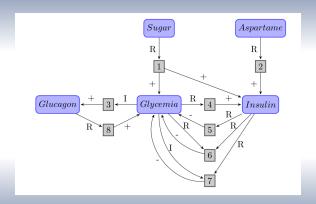
# Glucose regulation – 1

#### The set of species involved:

Sugar	$\mathcal{L}_{sugar} = \{0, 1\}$	$\delta_{sugar}(1)=2$
Aspartame	$\mathcal{L}_{\textit{aspartame}} = \{ 0, 1 \}$	$\delta_{aspartame}(1) = 2$
Glycemia	$\mathcal{L}_{\textit{glycemia}} = \{0, 1, 2, 3\}$	$\delta_{glycemia}(1)=8$
		$\delta_{glycemia}(2)=8$
		$\delta_{glycemia}(3)=8$
Glucagon	$\mathcal{L}_{ extit{glucagon}} = \{ extit{0},  exttt{1}\}$	$\delta_{glucagon}(1)=3$
Insulin	$\mathcal{L}_{\textit{insulin}} = \{0, 1, 2\}$	$\delta_{\textit{insulin}}(1) = 3$
		$\delta_{\textit{insulin}}(2) = 3$



# Glucose regulation – 2





# Glucose regulation – 3

#### The set of reactions:

$\rho_{\mathbf{k}}$	Activators A <sub>k</sub>	Inhibitors I <sub>k</sub>	Results R <sub>k</sub>	$\Delta_k$
$\rho_1$	{(Sugar, 1)}	Ø	$\{(\mathit{Insulin}, +),$	
			(Glycemia, +)	1
$ ho_{2}$	{(Aspartame, 1)}	Ø	$\{(\mathit{Insulin}, +)\}$	1
$ ho_{3}$	Ø	{(Glycemia, 1)}	$\{(Glucagon, +)\}$	1
$ ho_{4}$	{(Glycemia, 3)}	Ø	$\{(\mathit{Insulin}, +)\}$	1
$ ho_{5}$	{(Insulin, 2)}	Ø	$\{(Glycemia, -)\}$	2
$ ho_{6}$	$\{(Insulin, 1),$			
	(Glycemia, 3)}	Ø	$\{(Glycemia, -)\}$	2
$\rho_7$	{( <i>Insulin</i> , 1)}	{(Glycemia, 2)}	$\{(Glycemia, -)\}$	2
$ ho_8$	{(Glucagon, 1)}	Ø	$\{(Glycemia, +)\}$	2



#### **Observation**

- Decay and reactions are different types of behaviors
- Decay is synchronous it corresponds to an abstraction of the action of the environment
- Reactions are asynchronous their duration corresponds to the time required to observe an effect
- Execution time vs Simulation time
   More reactions are enabled less probable is the execution of time



# Toxicity analysis

- ANDy can be used to detect and predict toxic behaviors related to the dynamics of bio-molecular networks.
- We resort to temporal logics and model checking techniques.
- We use computation tree logic (CTL)
- We provide an abstraction of ANDy into Kripke structures



# **Examples of questions**

We are interested in checking whether the inner equilibrium of an organism is maintained when administrating drugs or applying stressors.

#### Toxicology properties can be classified into:

- properties checking for the appearance of symptoms,
- properties characterizing causal relations between events.



# Glucose regulation

#### Causality:

Does assimilation of sweeteners cause hypoglycemia?

 $\mathbf{EF}[((Sugar, 1) \lor (Aspartame, 1)) \land (Glycemia, 1)] \rightarrow$ AF(Glycemia, 2)



# Paths for glucose regulation

$$\textbf{EF}[((Sugar,1) \lor (Aspartame,1)) \land (Glycemia,1)] \rightarrow \\ \textbf{AF}(Glycemia,2)$$

Toxicity analysis

Path that satisfies

```
(Sugar, 1), (Aspartame, 0), (Glycemia, 1), (Insulin, 0), (Glucagon, 0) \xrightarrow{\rho_1}
(Sugar, 1), (Aspartame, 0), (Glycemia, 2), (Insulin, 1), (Glucagon, 0)
```

Path that contradicts

```
(Sugar, 0), (Aspartame, 1), (Glycemia, 1), (Insulin, 0), (Glucagon, 0) \xrightarrow{\rho_2}
(Sugar, 0), (Aspartame, 1), (Glycemia, 1), (Insulin, 1), (Glucagon, 0) \stackrel{\rho_7}{\longrightarrow}
(Sugar, 0), (Aspartame, 0), (Glycemia, 0), (Insulin, 1), (Glucagon, 0)
```



# Sound and completeness

#### Theorem

Given an ANDy network  $(\mathcal{S},\mathcal{R})$ , its encoding into

- Kripke structures
- Timed Automata

is sound and complete.





# Summing up

- ANDy, a high-level Petri net framework for cellular regulation networks.
- Species that can degrade as time passes by governed by a set of reactions.
- Toxicity properties can be expressed via a temporal logic.
- Properties can be verified thanks to a sound and complete abstraction.







Motivation

#### Final remarks

• Comparison with stochastic models à la Gillespie

Toxicity analysis

- Refinement of the abstraction
- Implementation: Snakes, Snoopy + Marcie





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