

# Mixed-formalism hierarchical modeling and simulation with BioRica

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► **To cite this version:**

Alice Garcia, David James Sherman. Mixed-formalism hierarchical modeling and simulation with BioRica. 11th International Conference on Systems Biology, Oct 2010, Edimbourg, United Kingdom. pp.P02.446. <inria-00529669>

**HAL Id: inria-00529669**

**<https://hal.inria.fr/inria-00529669>**

Submitted on 27 Oct 2010

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# Mixed-formalism hierarchical modeling and simulation with BioRica

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## Challenge of reuse biological models

- Many models are produced for biological phenomena of different kinds. Models are complementary or competing.
- Each model is experimentally validated, and this is expensive.
- Numerical representations of biological systems are constructed using a number of different formalisms and simulation techniques.
- Need to reuse and compose existing models without revalidation.

## Hierarchical modelling

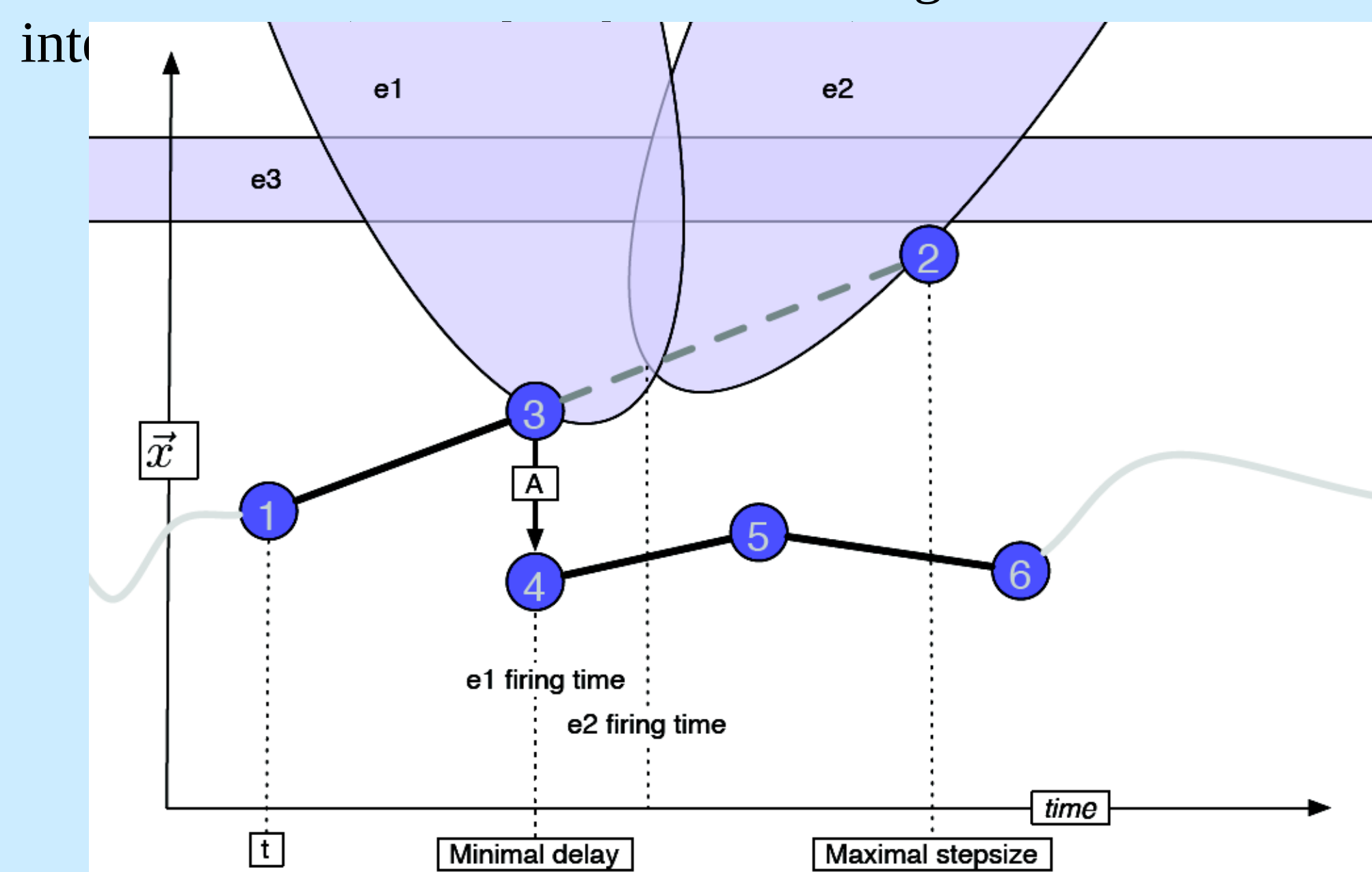
- Models are decomposed into hierarchical collection of nodes, each with its own local behaviour (De Alfaro et al. 2001).
- This gives a methodical way to combine existing models into a more complete one.
- Exists a rigorous semantics for hierarchical combination.
- BioRica (Soueidan et al. 2007) is a language for describing hierarchical modelling, derived from AltaRica (Arnold et al. 2000).
- Existing models in different formalisms (automata, ODE), and different notations like SBML (Hucka et al. 2010) can be combined into a BioRica model.

## Stochastic transition systems (De Alfaro et al. 2001)

- Biological systems are not deterministic.
- Stochastic transition systems allows timed events and non deterministic transition relations.
- Extend transition system with random timers : random variable with a distribution specified by the stochastic labelling.
- Add stochastic behaviour to describe the likelihood that an event fires when it is activated concurrently with another event.

## Combining continuous and discrete behaviours

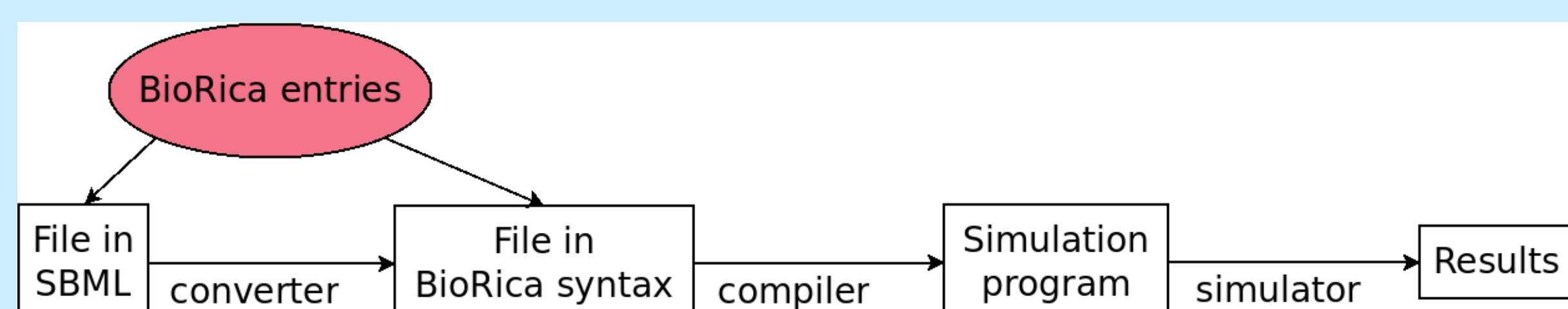
- Challenge is efficiently combining the computation of the behaviours.
- Simulation deals with continuous time and allows for discrete events that roll back the time according to these discrete intervals.



- The numerical integrator advances between  $t$  (point 1) and the maximal stepsize defined by an adaptive integration algorithm (2).
- The guards of events  $e1$ ,  $e2$  are satisfied.
- The regions where guards are satisfied are shaded.
- The firing time of  $e1$  (3) is used to reset the simulator after the discrete transition A (4)

## BioRica

- A platform for hierarchical mixed-formalism modeling of complex systems in biology.
- Contained a converter, a compiler and a simulator.

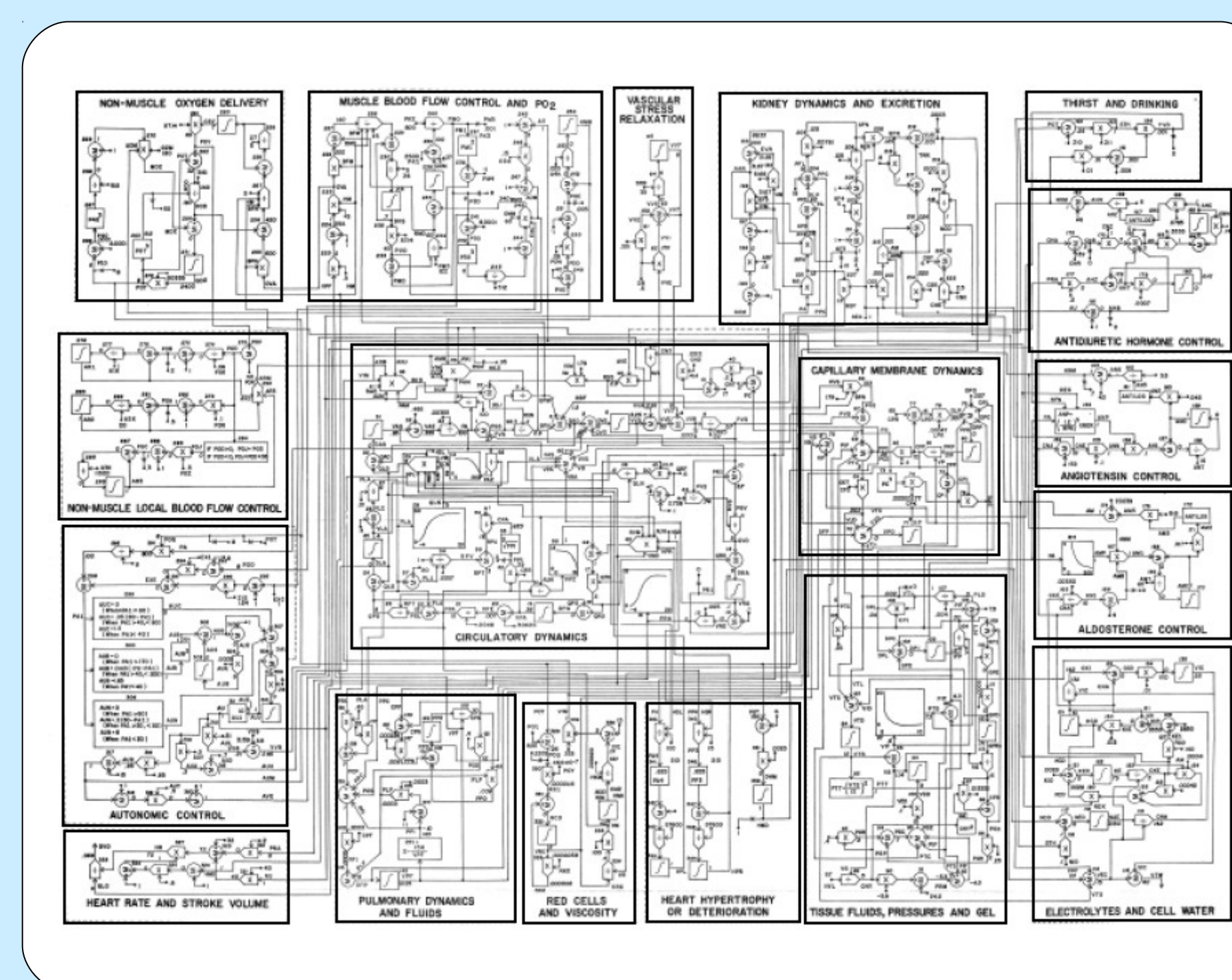
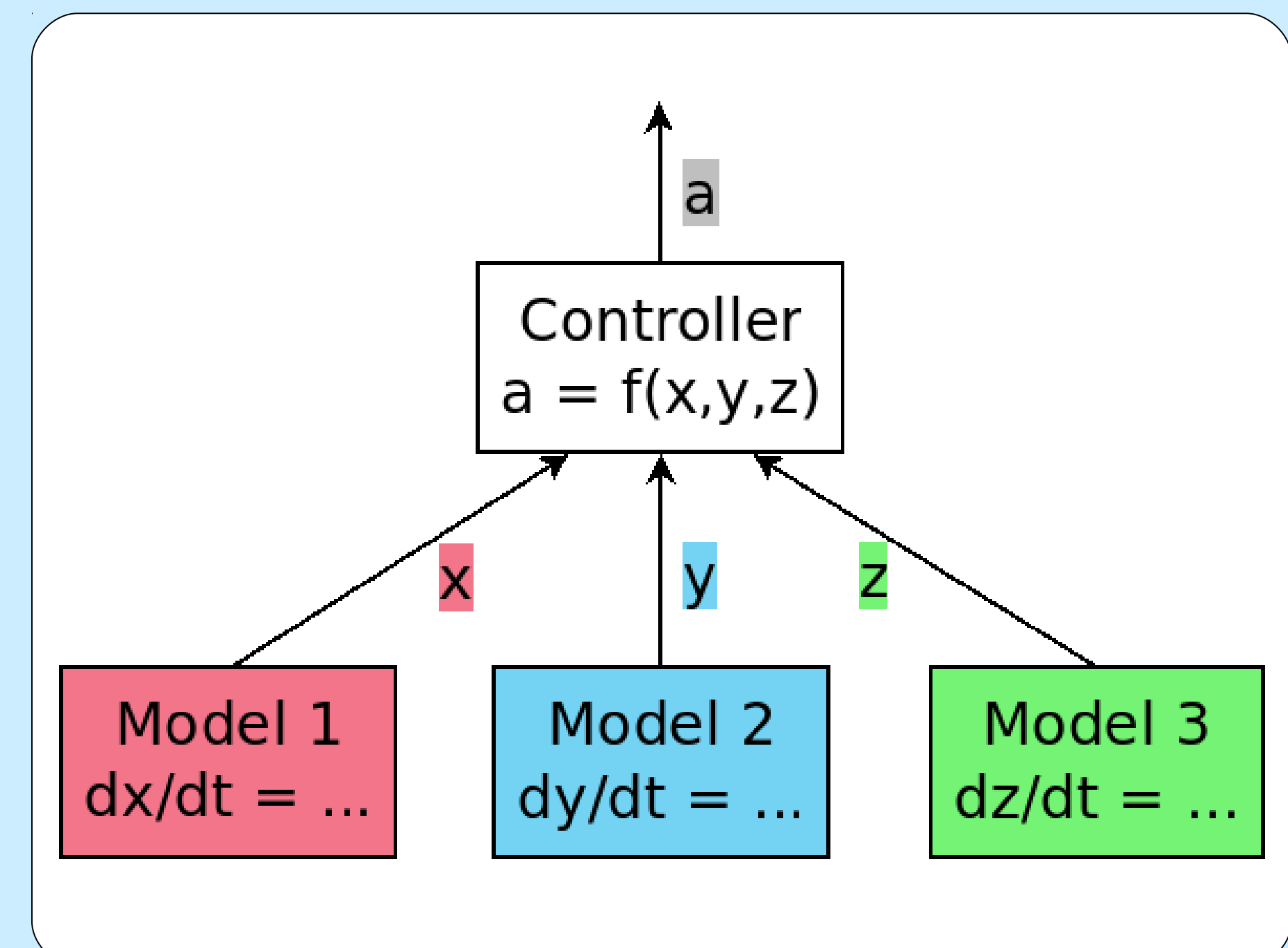


## Practical examples

Needs :

**Intelligent combination of models without refinement.**

Solution : Numerical models and a discrete controller.



Needs :

**Integrate modules with different time scales.**

Solution : Hierarchical representation of the model.

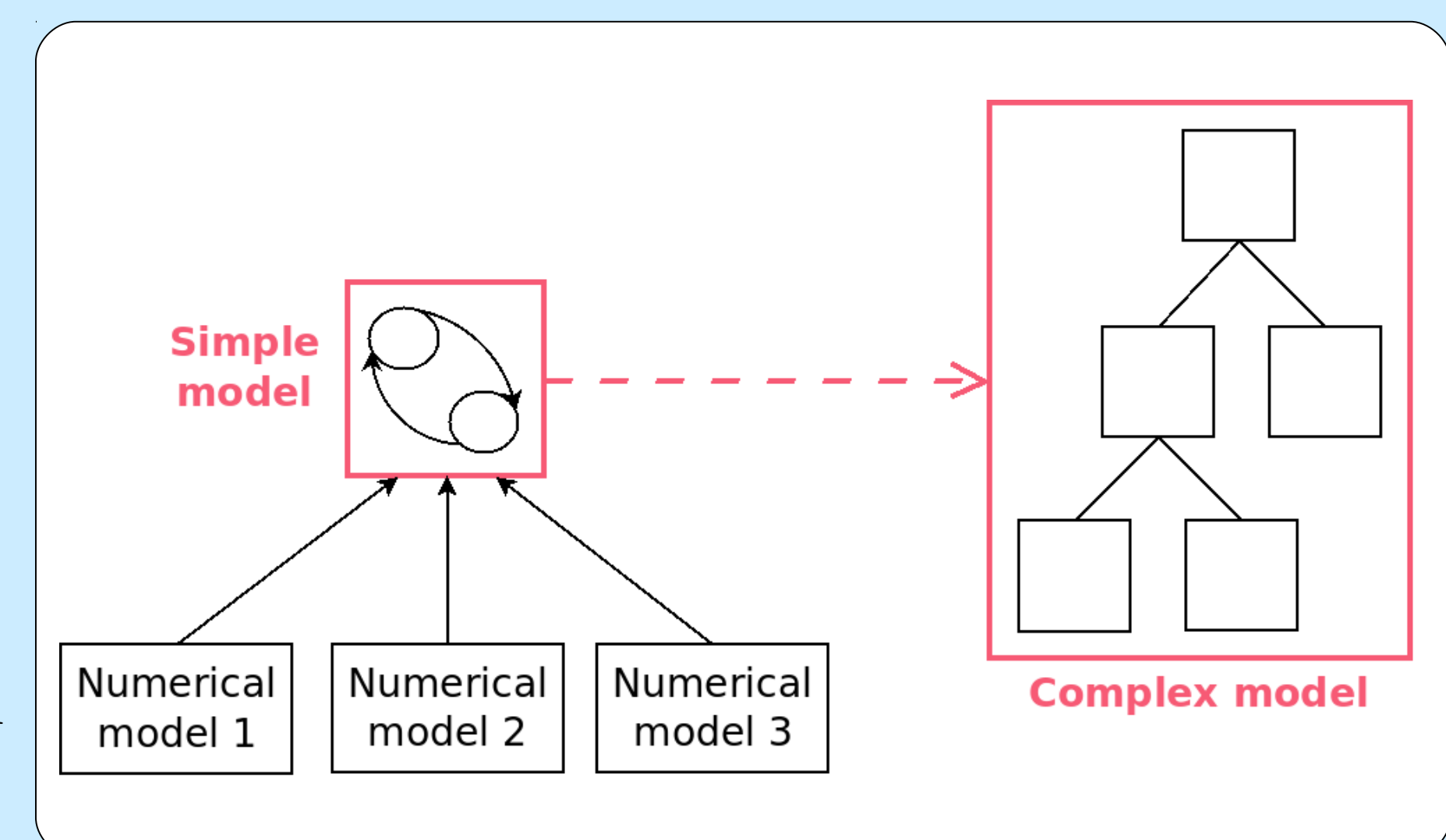
Canonical example : Guyton circulatory model (Guyton et al. 1972) : an extensive mathematical model of human circulatory physiology.

Needs :

**Successive refinement.**

Solution : Convert a simple automate to a complex model.

Example : Replace simple physical switch by model of a cell signalling pathway



## Availability

- SBML converter, compiler, simulator and examples.
- Available soon : <http://biorica.gforge.inria.fr/>

## References

- L. De Alfaro , T. A. Henzinger , R. Jhala. **Compositional Methods for Probabilistic Systems**. Proceedings of the 12th International Conference on Concurrency Theory, p.351-365, August 20-25, 2001
- H. Soueidan, M. Nikolski, D. Sherman. **BioRica: A multi model description and simulation system**. In: *Proceedings of Foundations of Systems Biology in Engineering (FOSBE)*, Stuttgart, Germany, F. Allgöwer, M. Reuss (editors), ISBN 978-3-8167-7436-5, September 2007.
- A. Arnold, A. Griffault, G. Point and A. Rauzy. **The AltaRica formalism for describing concurrent systems**. *Fundam. Inf.* 40, issue 2-3, pages 109-124, IOS Press, 2000.
- Hucka, Michael, Bergmann, Frank, Hoops, Stefan, Keating, Sarah, Sahle, Sven, and Wilkinson, Darren. **The Systems Biology Markup Language (SBML): Language Specification for Level 3 Version 1 Core (Release 1 Candidate)**. *Nature Precedings*; 2010.
- H. Soueidan. **Discrete event modeling and analysis for Systems Biology models**. Thesis, Bordeaux 1 university, 2009.
- A. C. Guyton, T. G. Coleman, H. J. Granger. **Circulation: Overall Regulation**. *Annual Review of Physiology*, March 1972, Vol. 34, Pages 13-44.

