

Modelling, Simulation, and Model Checking of Large Biological Regulatory Networks

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Stochastic simulation



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Biological Regulatory Networks

Overview

Computer science for systems biology

- Models for dynamical concurrent systems.
- Validation of the model / control of the system.
- We focus on Biological Regulatory Networks (BRNs).
- We introduce a new modelling framework: the Process Hitting.

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The Process Hitting [Paulevé, Magnin, Roux in TCSB 2011]

- Elementary framework for dynamical complex systems;
- Applied to BRNs; not limited to.
- Stochastic and Time dimensions (simulation + standard model checking).
- Software available (PINT - <http://process.hitting.free.fr>).

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The Process Hitting [Paulevé, Magnin, Roux in TCSB 2011]

- *Elementary* framework for **dynamical complex systems**;
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- **Stochastic and Time dimensions** (simulation + standard model checking).
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Large-scale model checking (dynamical properties)

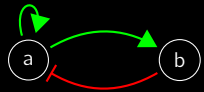
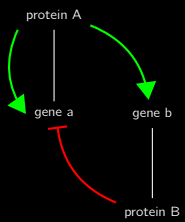
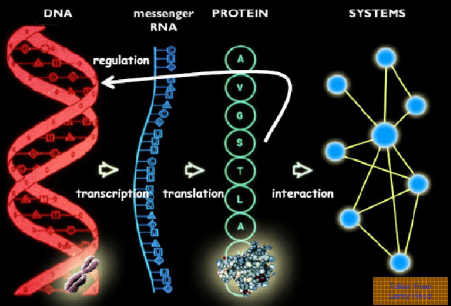
- Cope with state space explosion.
- Our approach: **Static Analysis** of the model.
- Static analysis by **Abstract Interpretation**.

Outline

- ① Introduction to BRNs
- ② The Process Hitting
- ③ Stochastic and Time Parameters
- ④ Static Analysis of Process Hitting
 - Fix Points
 - *Abstract Interpretation of Scenarios*
- ⑤ Applications
- ⑥ Outlook

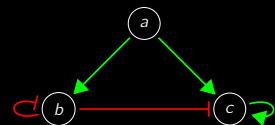
Biological Regulatory Networks (BRNs)

The interaction graph



Discrete Networks (BRNs)

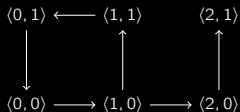
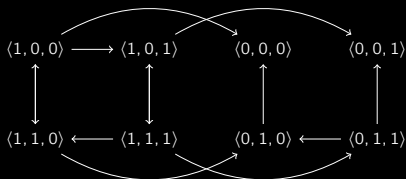
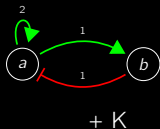
- Each component has a finite set of **qualitative levels**; e.g. $\{0, 1, 2\}$;
- may be seen as a quantization of the concentration of the component.



$$f^a(x) = 0$$

$$f^b(x) = x[a] \wedge \neg x[b]$$

$$f^c(x) = \neg x[b] \wedge (x[a] \vee x[c])$$



[René Thomas in Journal of Theoretical Biology, 1973] [A. Richard, J.-P. Comet, G. Bernot in Modern Formal Methods and Applications, 2006]

Hybrid Modelling

Continuous features governing discrete transitions

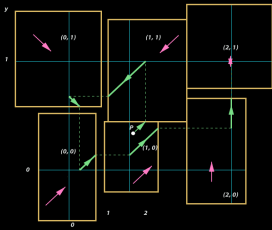
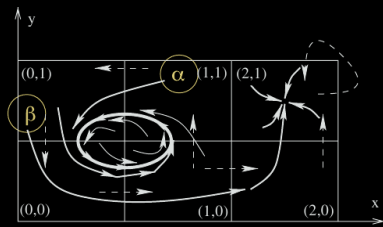
Introduce delays to actions

Stochastic Models

- Delays are **random variables** (generally exponential, i.e Markovian);
- \Rightarrow compute probabilities for observing behaviours.

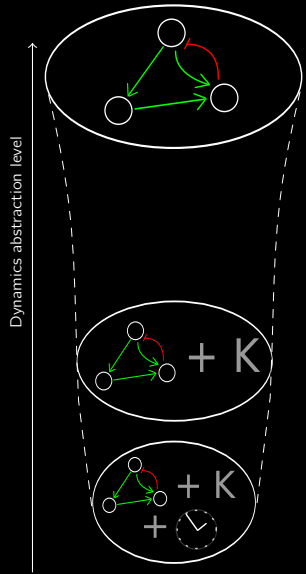
Stochastic Petri Nets / π -calculus, etc.

Timed Models

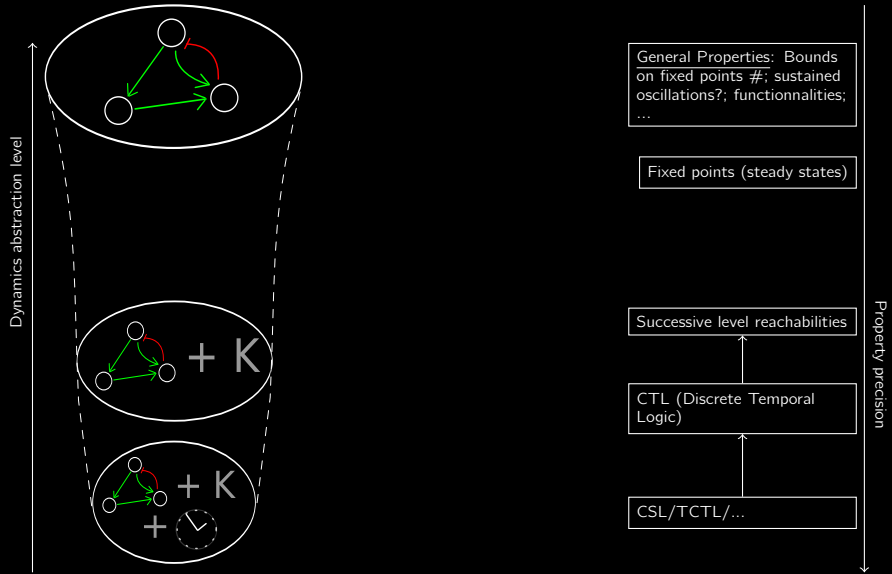


Timed / Hybrid Automata

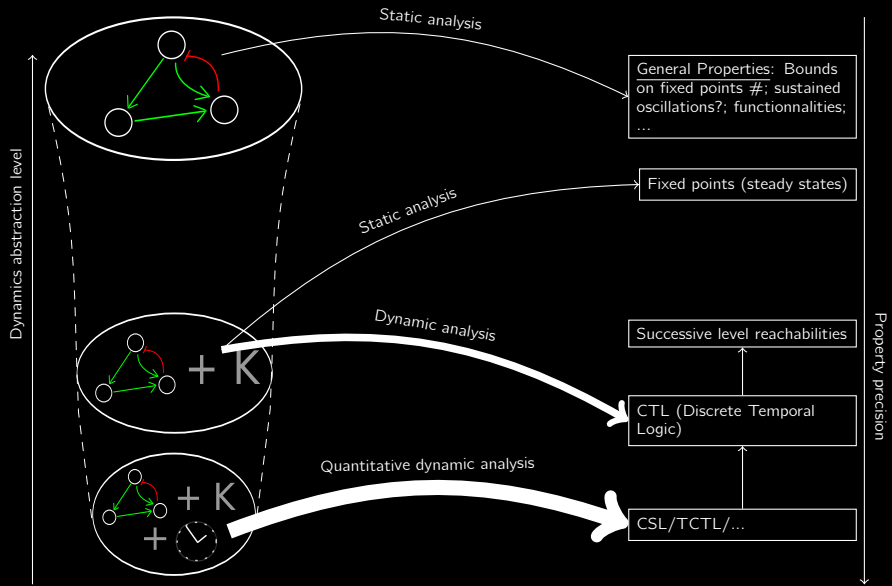
Summary and Contribution



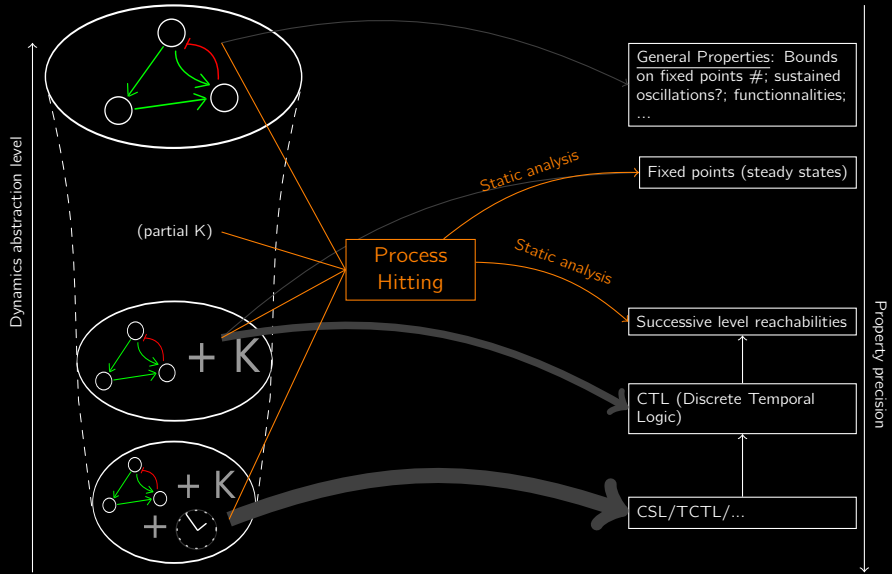
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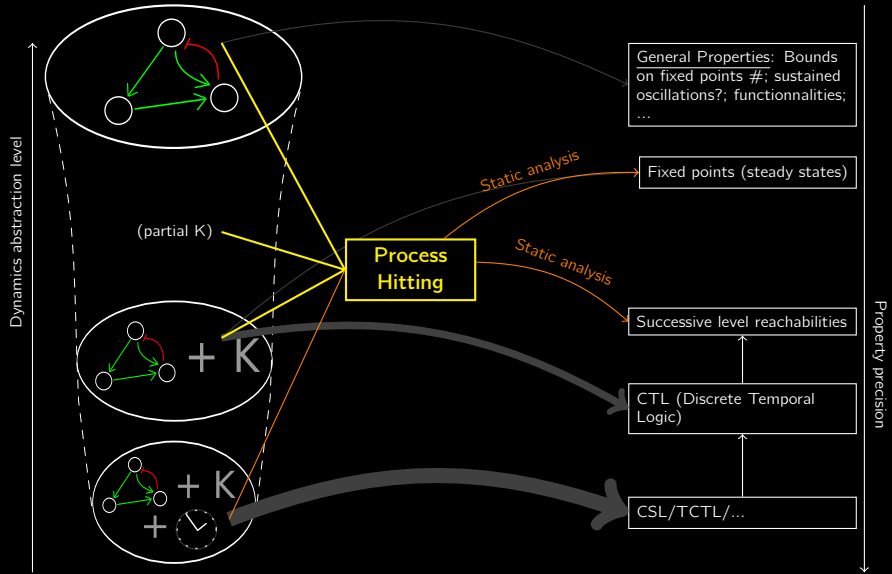
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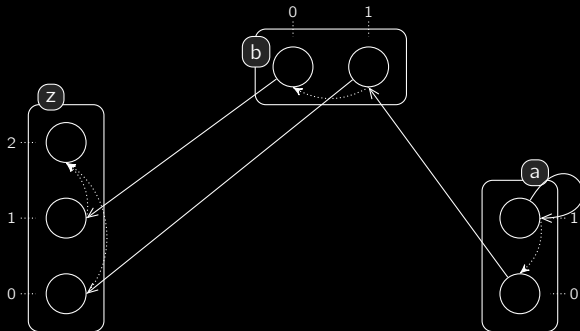


Outline



The Process Hitting Framework

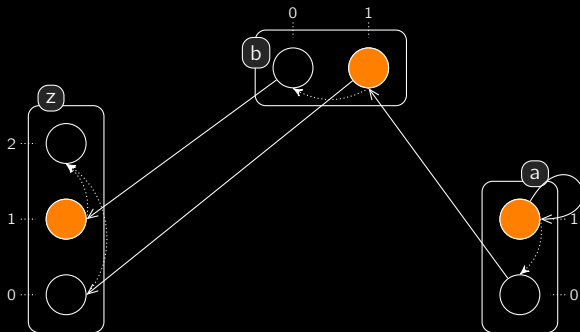
[Paulevé, Magnin, Roux in TCSB 2011]



- **Sorts:** a, b, z ; **Processes:** $a_0, a_1, b_0, b_1, z_0, z_1, z_2$;
- **Actions:** a_0 hits b_1 to make it bounce to b_0, \dots ;
- **States:** $\langle a_1, b_1, z_1 \rangle, \langle a_0, b_1, z_1 \rangle, \langle a_0, b_0, z_1 \rangle, \dots$;
- Restriction of Communicating Finite-State Machines (CFSM).

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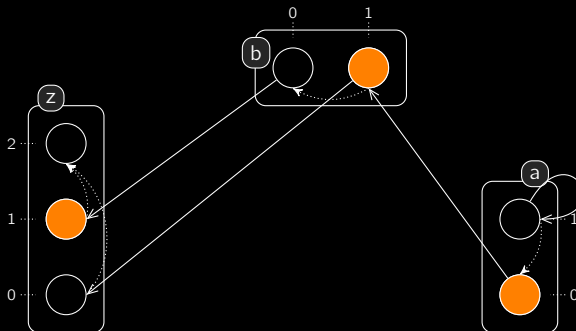
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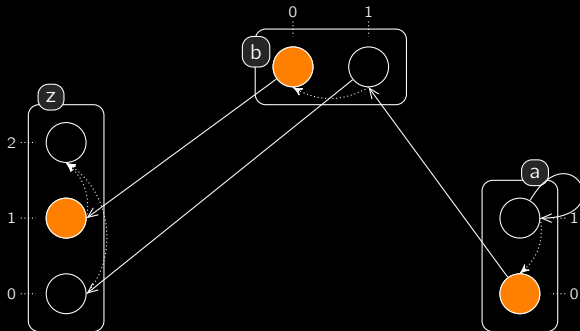
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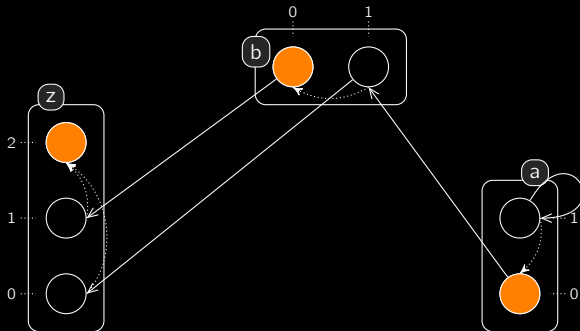
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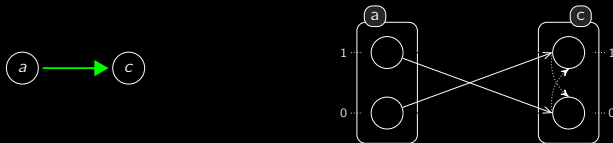
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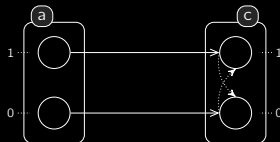
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From BRNs to Process Hittings



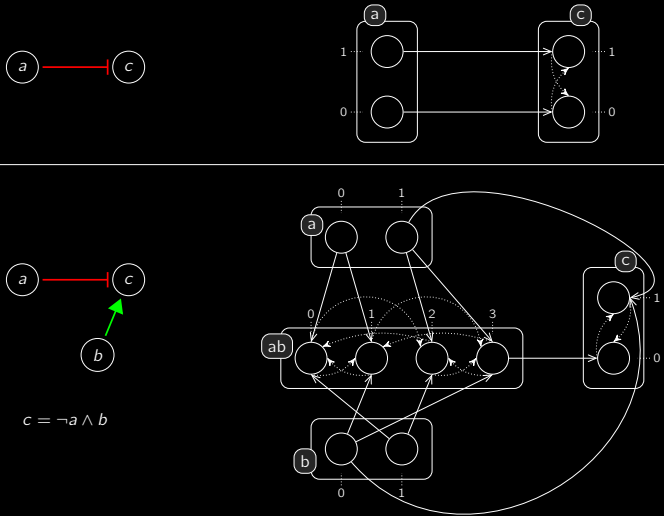
[Paulevé, Magnin, Roux in Trans. in Computational Systems Biology, 2011]

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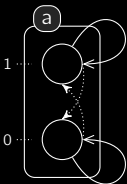
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Stochasticity Absorption Factor

[Paulevé, Magnin, Roux in IEEE TSE, 2010]

- Stochastic dimension is prominent... but no precise time features:
- (Markov) Exponential distribution: mean r^{-1} ; variance r^{-2} .
- At our level of abstraction, we need to tune time features.

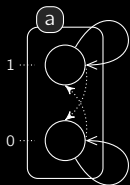


sa=1

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- **Stochastic dimension** is prominent... but **no precise time features**:
- (Markov) Exponential distribution: **mean** r^{-1} ; **variance** r^{-2} .
- At our level of abstraction, we need to **tune time features**.
- Provide a **stochasticity absorption factor** sa :
- duration follows the sum of sa exponentials of rate $r.sa$;
- **mean** r^{-1} ; **variance** $r^{-2}sa^{-1}$ (Erlang distribution).



$sa=1$



$sa=5$

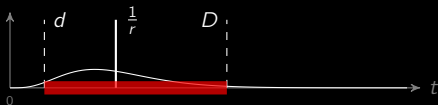


$sa=50$

Stochastic and Time Parameters

[Paulevé, Magnin, Roux in IEEE TSE, 2010]

- Specify either (r, sa) , or its **firing interval** $[d; D]$,
- which is the confidence interval at confidence coefficient $1 - \alpha$.
- We have **estimators** to translate $[d; D]$ to (r, sa) .

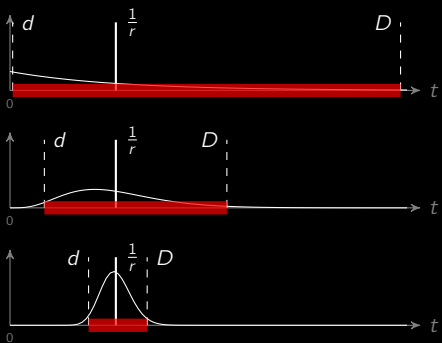



— action duration

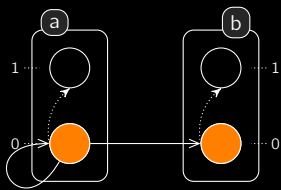
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 action duration



$\Rightarrow b_1$ is reached at a **very low probability**.

Simulation and Model Checking

Stochastic Model Checking

- Translation from the Erlang stochastic π -calculus to PRISM [Paulevé, Magnin, Roux in IEEE TSE, 2010].
- Applies to the Process Hitting as well.
- Not tractable with large stochasticity absorption factors;
- but there is hope in symmetry reductions, or abstractions of sequences of transitions, or ...

Simulation

- Non-Markovian simulation using the
- Generic abstract machine for stochastic process calculi [Paulevé, Youssef, Lakin, Phillips at CMSB 2010].
- Process Hitting simulator implemented in PINT.

Challenge: scalable inference of stochastic and time parameters...

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Challenge: scalable inference of stochastic and time parameters...
... still open; prior need for scalable qualitative analyses.

Static Analysis of Process Hitting

Intuition

- Simplicity of Process Hitting \Rightarrow simple structures.
- Efficient static derivation of dynamical features.

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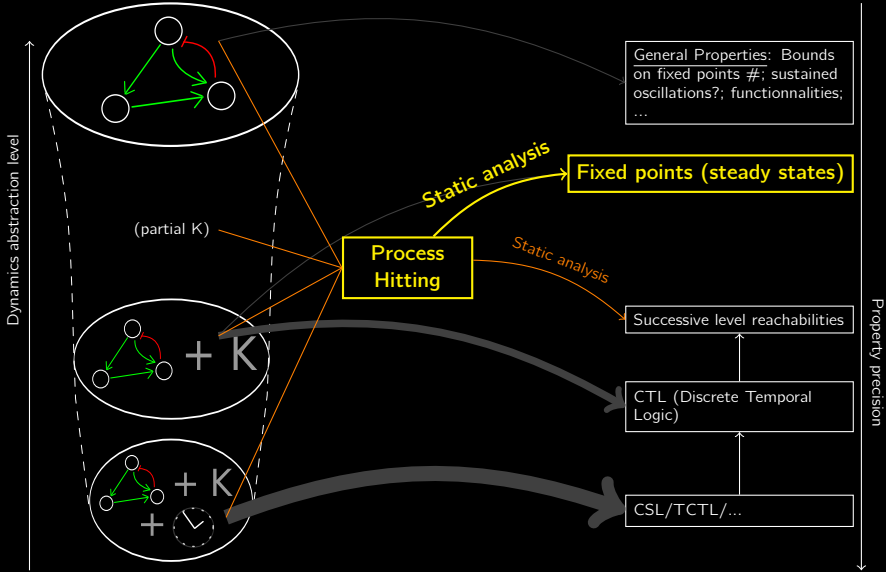
- Reduction to the n -cliques of a n -partite graph.

Reachability properties $\text{EF } a_i \wedge (\text{EF } b_j \wedge \dots)$

- **Limited complexity**: \approx polynomial in $\# \text{sorts}$; exponential in $\# \text{processes per sort}$.
- May be inconclusive: **Yes/No/Jocker**.
- Abstract interpretation techniques.
- \Rightarrow address **very large systems**.
- + extraction of **key processes** (towards control).

[Paulevé, Magnin, Roux in SASB'10 + Technical report]

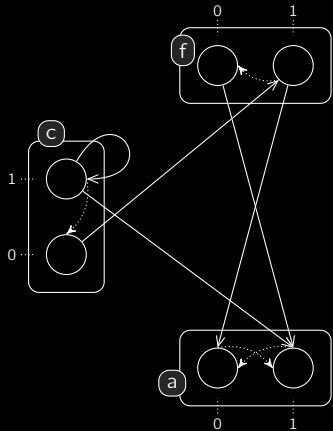
Outline



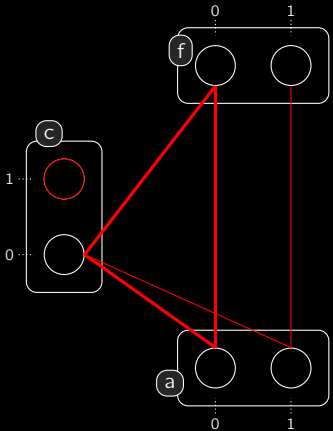
Fix Points

[Paulevé, Magnin, Roux in TCSB 2011]

Process Hitting



Hitless graph

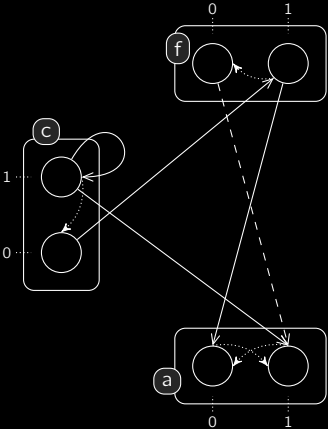


n -cliques are fix points

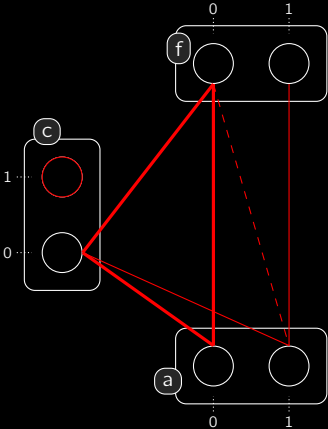
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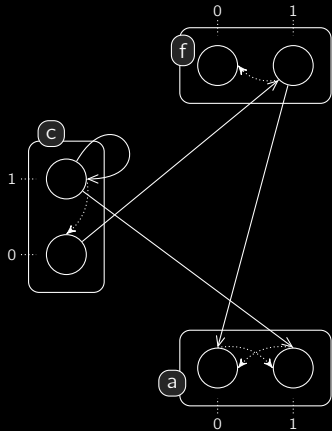


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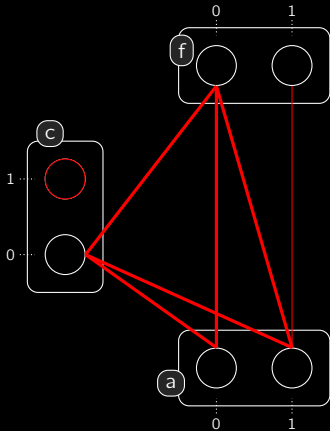
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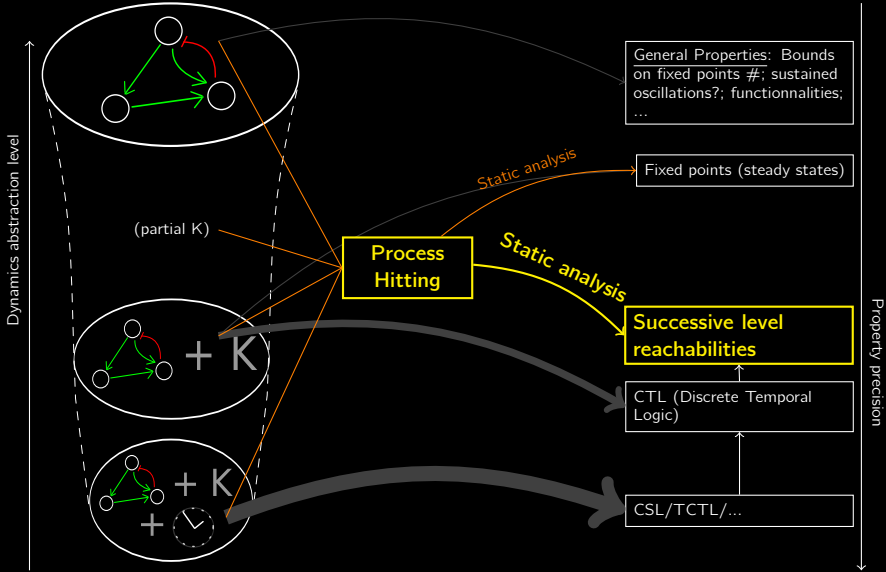


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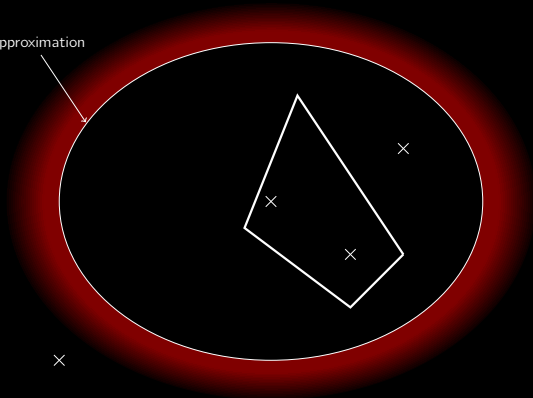
Approximation of Reachability Properties

- Successive reachability of processes (reach a_0 then b_1 then ...);
- Approach using abstract interpretation techniques;
- Results in both over- and under-approximations;
- Limited complexity at the cost of potentially being inconclusive.

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Over-approximation



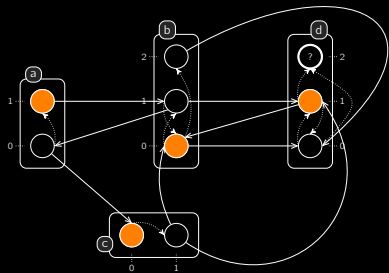
[Paulevé, Magnin, Roux at SASB 2010 + MSCS submitted]

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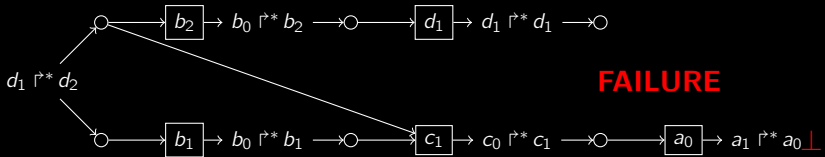
- Loïc Paulevé, Morgan Magnin, Olivier Roux



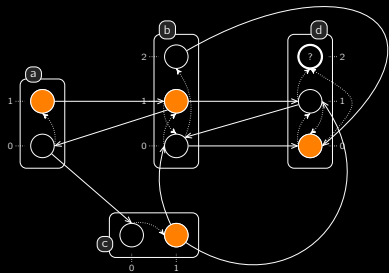
Un-ordered Over-approximation



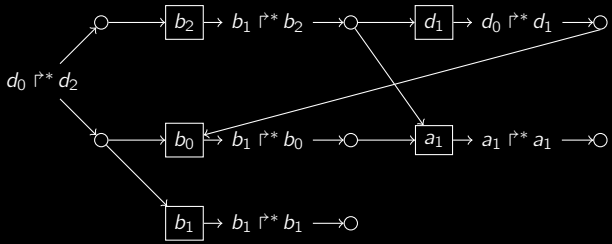
- **Necessary condition:** there always exists a solution ending with a trivial objective.



Un-ordered Over-approximation

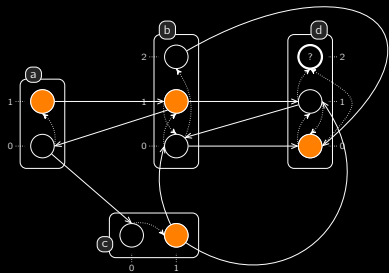


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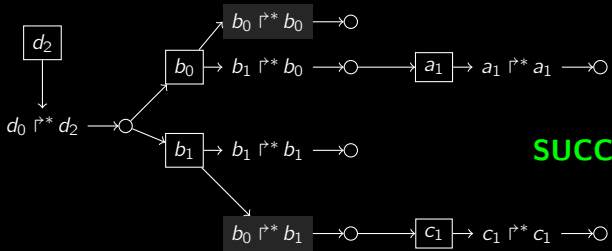
OK

Under-approximation



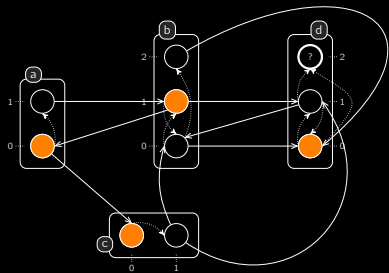
Sufficient condition:

- $\lceil \mathcal{B}_\xi^\omega \rceil$ has no cycle; and
- all referenced objectives have at least one solution.



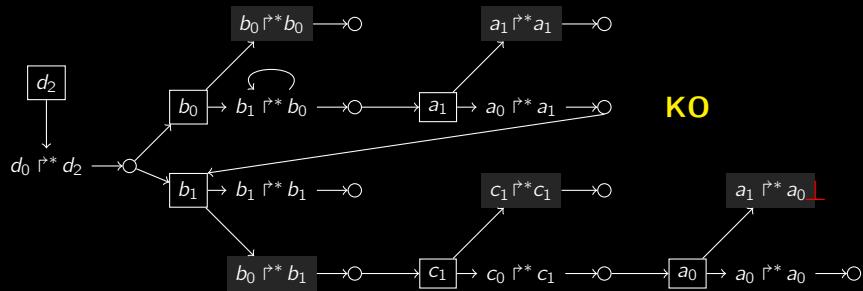
SUCCESS

Under-approximation



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Abstract Interpretation of Process Hitting

Summary

- **Reachability properties** of the form $\text{EF } a_i \wedge (\text{EF } b_j \wedge \dots)$;
- **Static computation** of abstract structures from Process Hitting models.
- **Recursive and iterative** reasonments.
- Over- and under-approximations (**Yes/No/Jocker**).
- Extraction of **key processes** (that are necessary): **towards control**.

Complexities

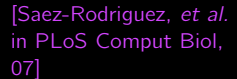
- \approx **polynomial in #sorts**; exponential in **#processes per sort**.

\Rightarrow efficient with a **small number of processes per sort**; while a **very large number of sorts** can be handled.

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(94 components)

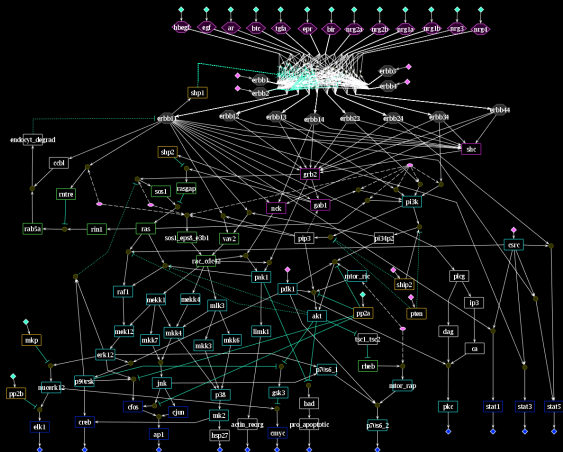


133 sorts,
448 processes,
1124 actions:
 $\approx 2 \cdot 10^{58}$ states.

Loïc Paulevé, Morgan Magnin, Olivier Roux

EGFR/ErbB Signalling

(104 components)



[Samaga, *et al.* in
PLoS Comput Biol,
2009]

Process Hitting

193 sorts,
748 processes,
2356 actions:
 $\approx 2 \cdot 10^{96}$ states.

Reachability analysis **always conclusive**; around **0.05s** (compared to *libddd*: out of memory). [<http://ddd.lip6.fr>]

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Static Analysis of Process Hitting

- Fix points by topological analysis;
- Very efficient over- and under-approximations of process reachability;
- Extract necessary processes for achieving reachabilities: towards control.
- Brings new insight to derive precise dynamical properties from BRNs.

Outlook

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- etc.

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Application to BRNs

- Address **bigger BRNs** (E. Coli, etc.);
- Focus on **properties of interest** for BRNs analysis;
- Suggestions are very welcome.