

# **Modelling, Simulation, and Model Checking of Large-Scale Biological Regulatory Networks**

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LIX / AMIB team - 1st June 2011





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*Stochastic simulation*



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I3S & CNRS, Nice, France

*Biological Regulatory Networks*

## 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).
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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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## Large-scale model checking (dynamical properties)

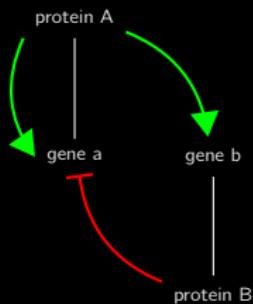
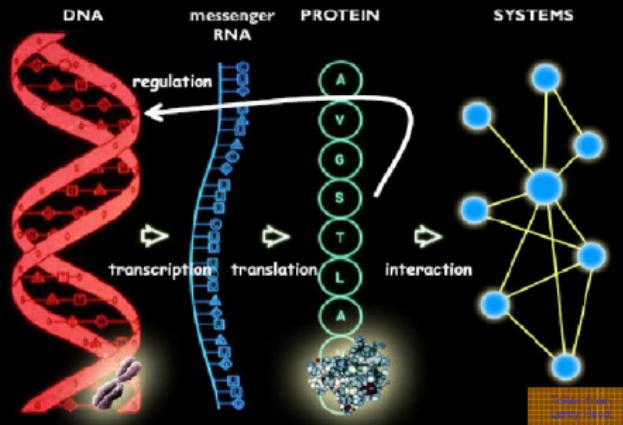
- Cope with state space explosion.
- Our approach: **Static Analysis** of the model.
- Static analysis by **Abstract Interpretation**.

## 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

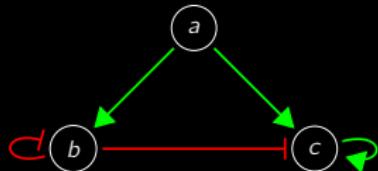
# Biological Regulatory Networks (BRNs)

The interaction graph



# Discrete Networks (BRNs)

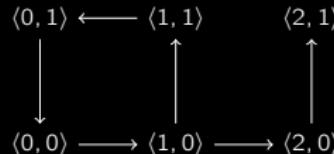
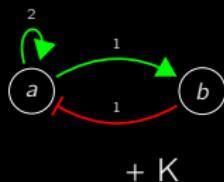
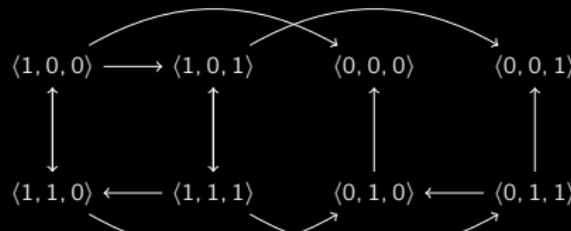
- Each component has a finite set of **qualitative levels**; e.g.  $\{0, 1, 2\}$ ;
- may be seen as a quantization of the concentration of the component.



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

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

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



[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

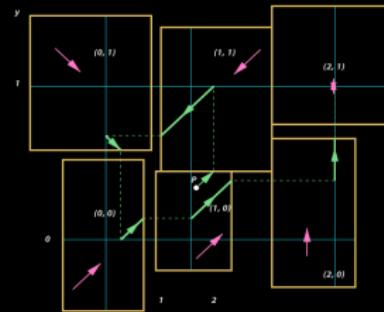
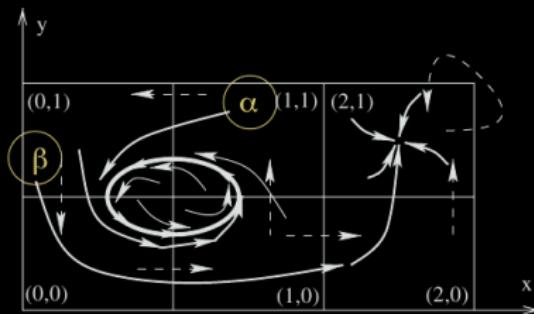
## Introduce delays to actions

### Stochastic Models

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

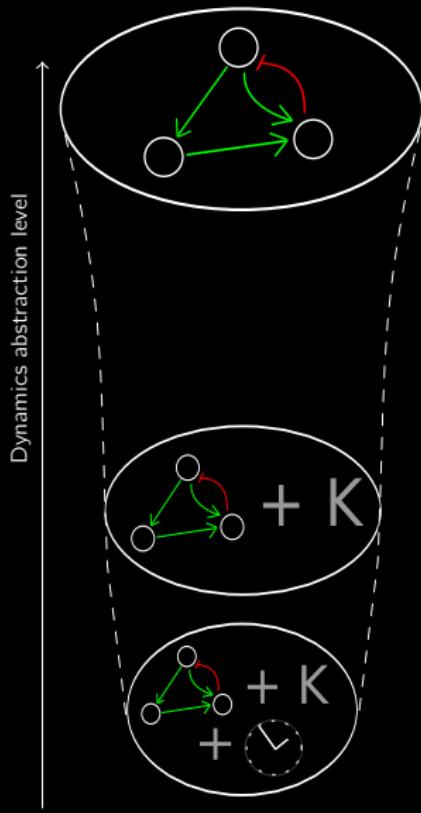
Stochastic Petri Nets /  $\pi$ -calculus, etc.

### Timed Models

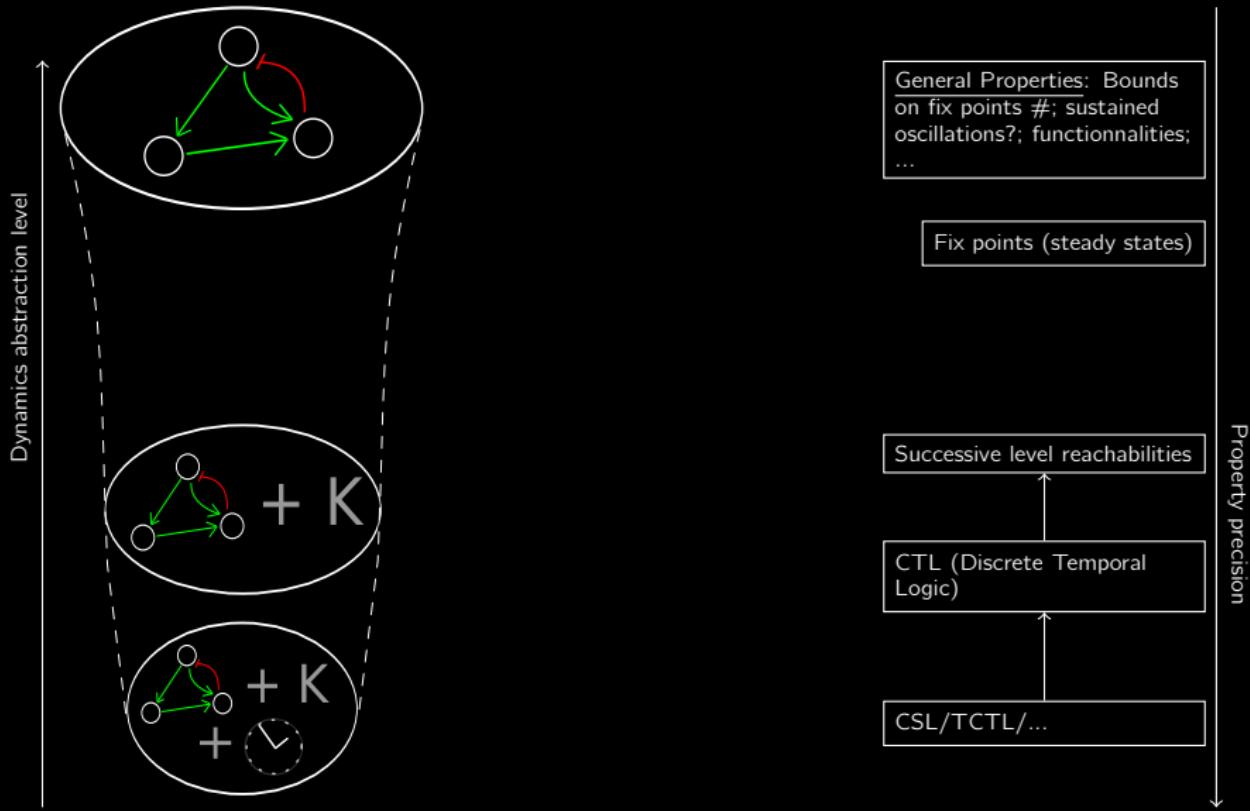


Timed / Hybrid Automata

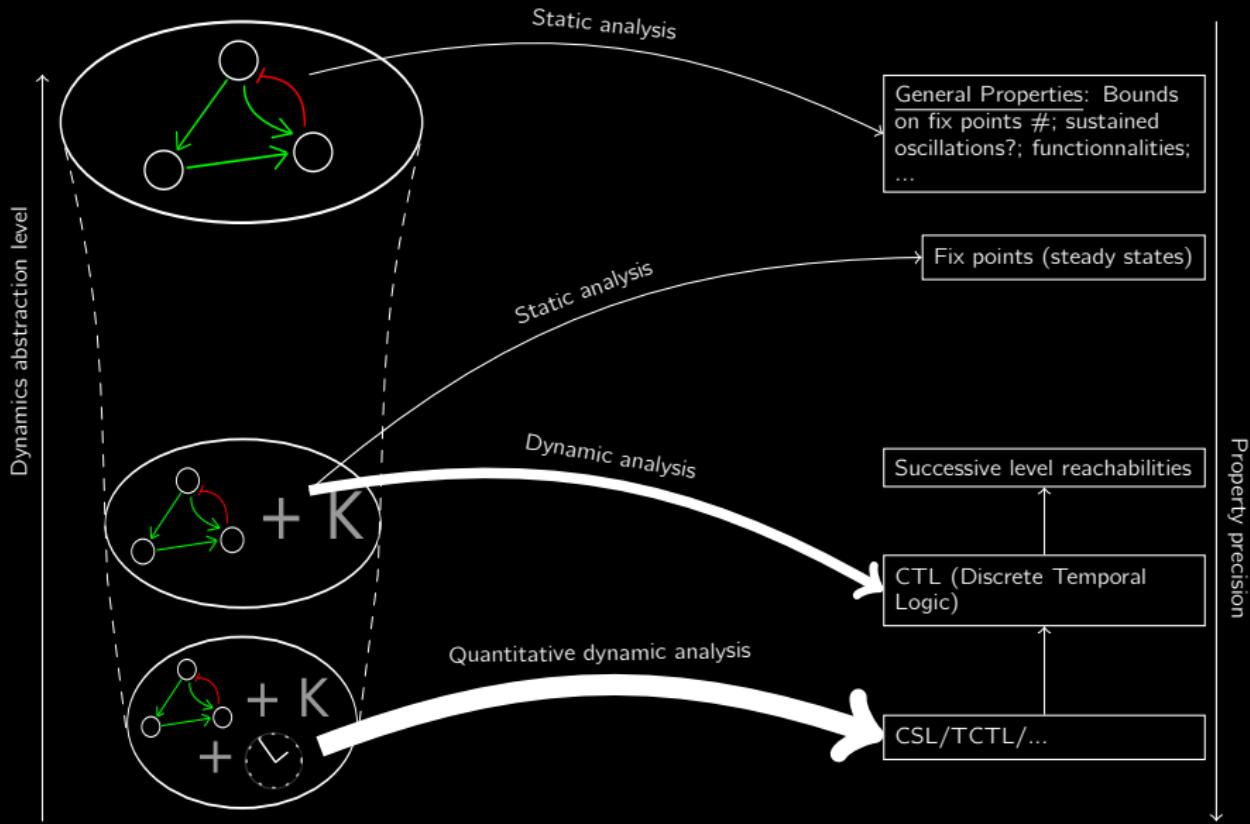
## Summary and Contribution



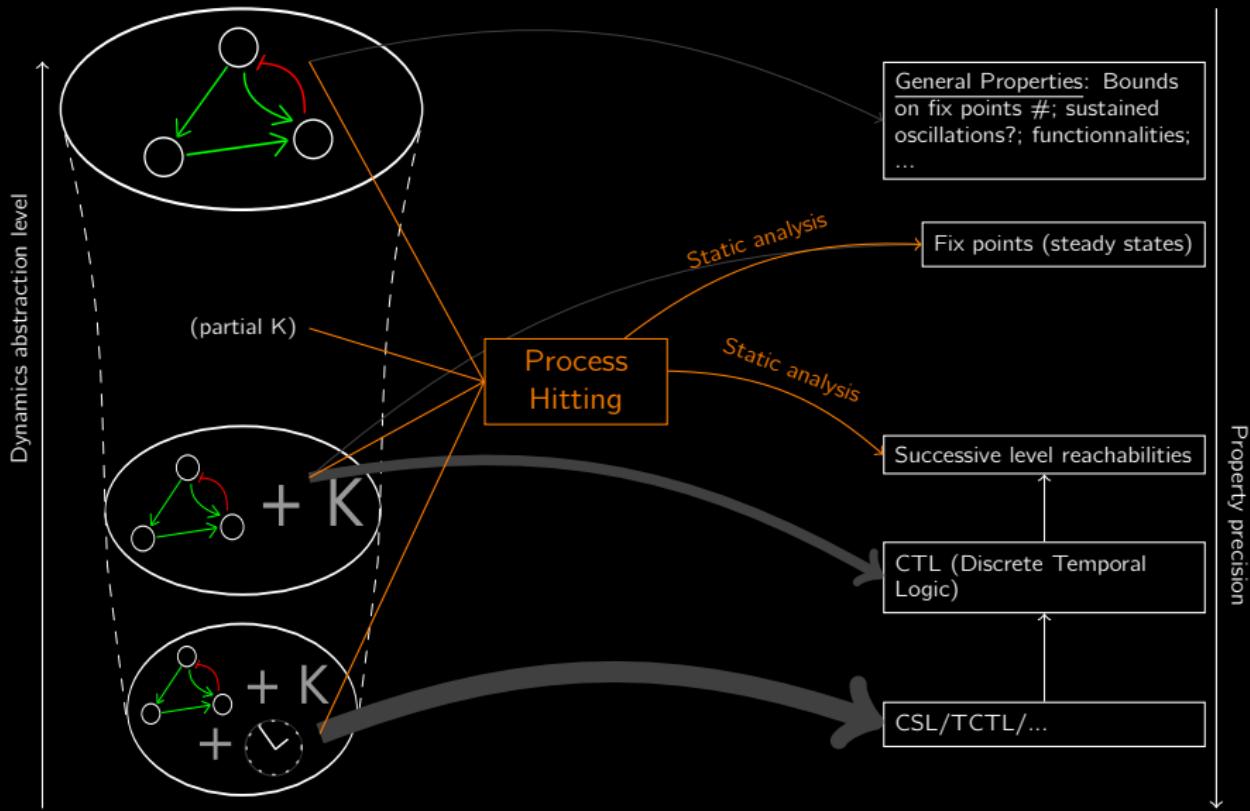
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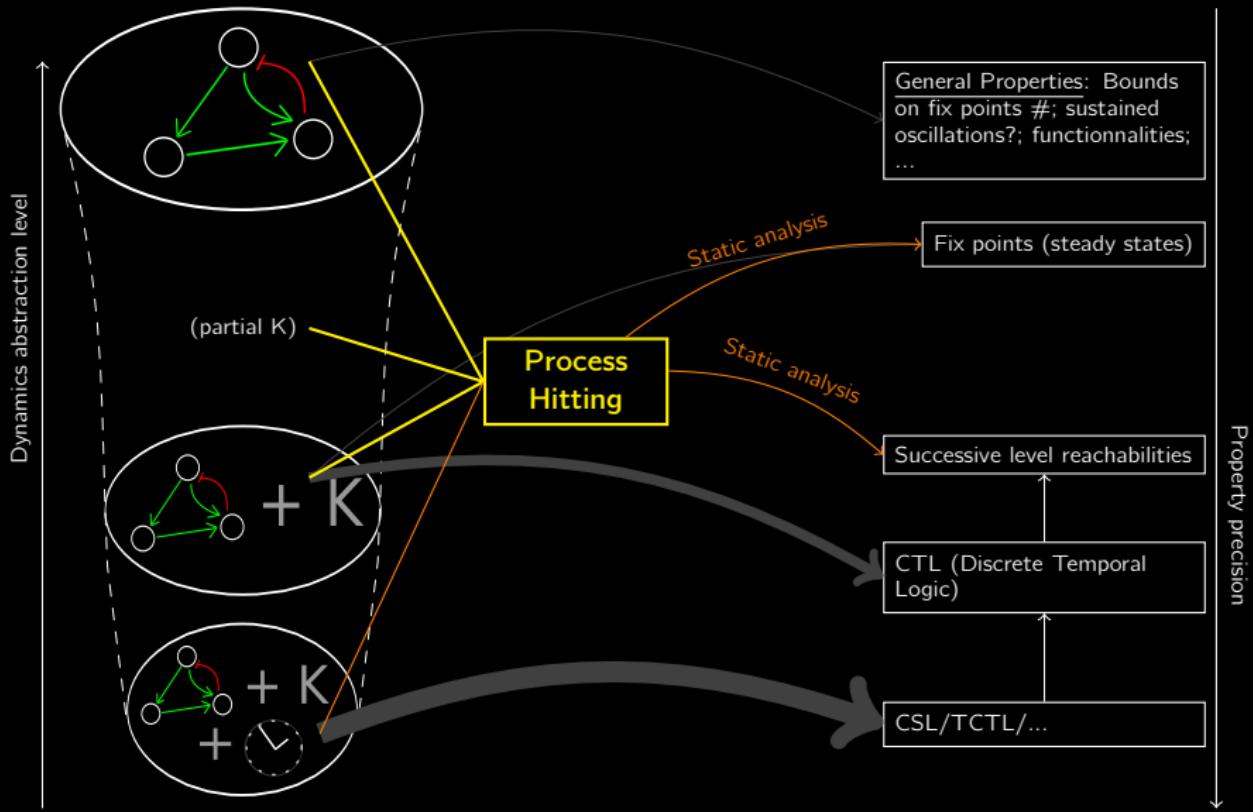
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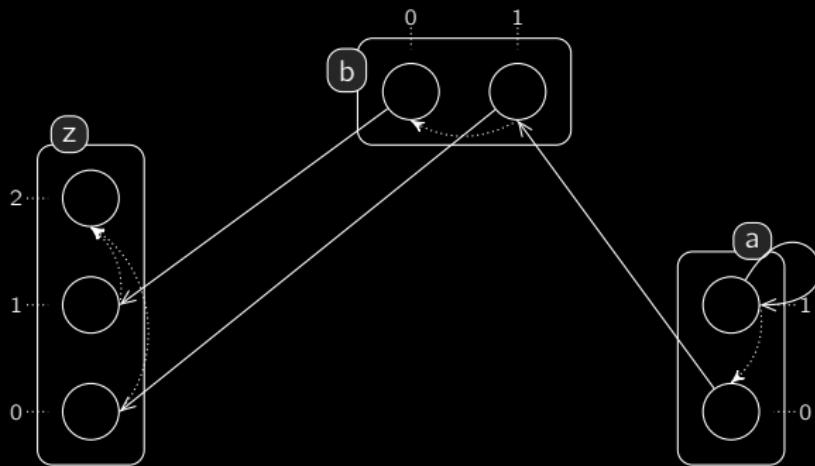


# Outline



# The Process Hitting Framework

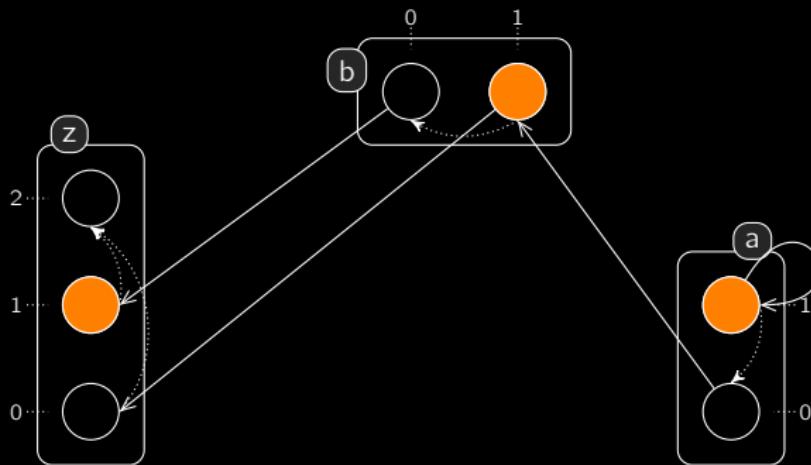
[Paulevé, Magnin, Roux in TCSB 2011]



- **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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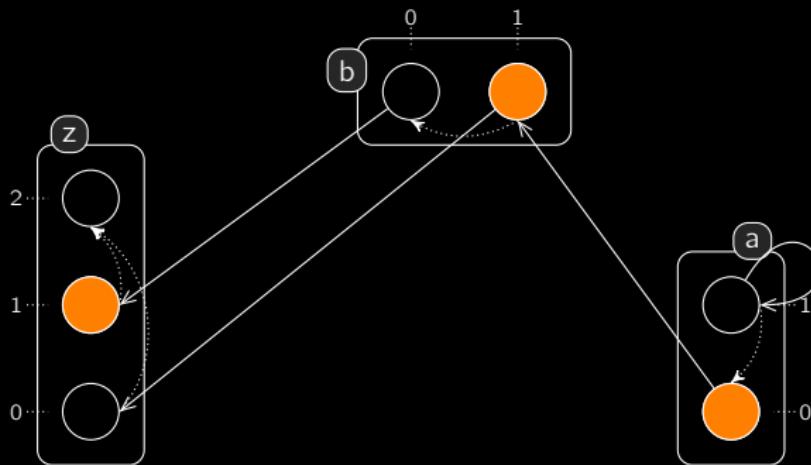
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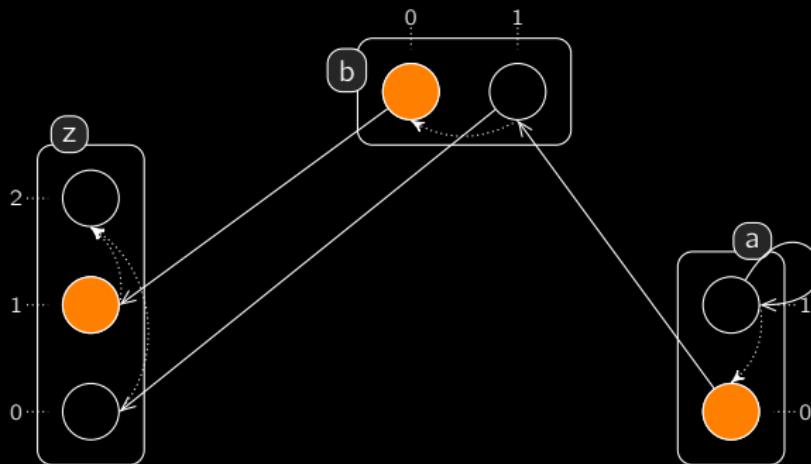
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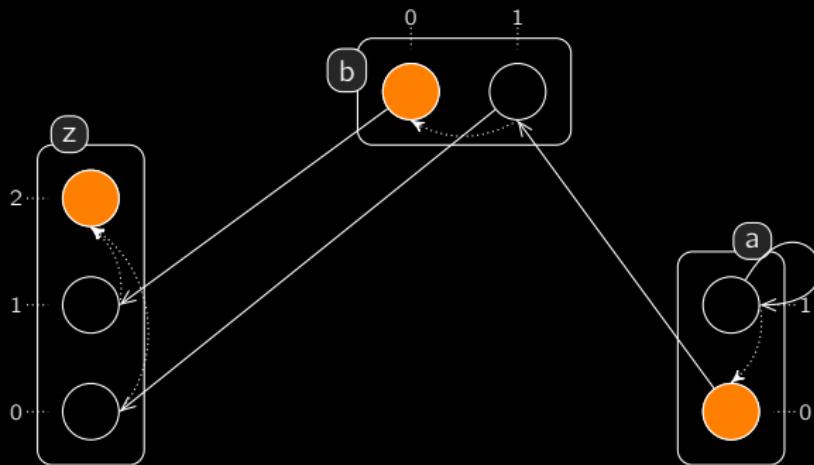
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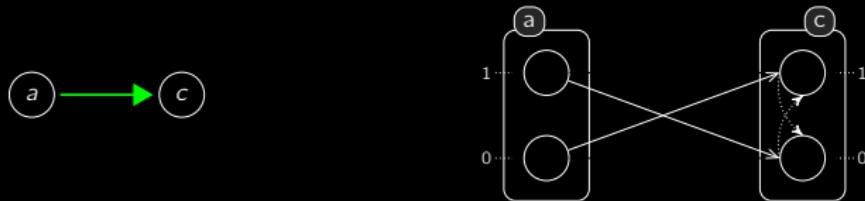
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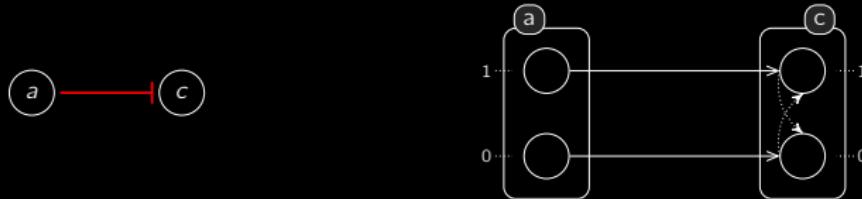
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# From BRNs to Process Hittings



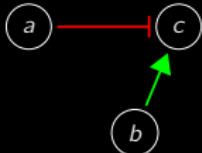
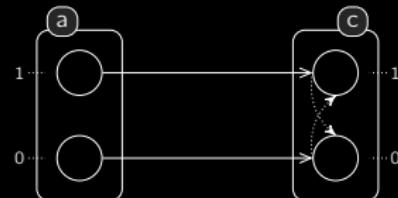
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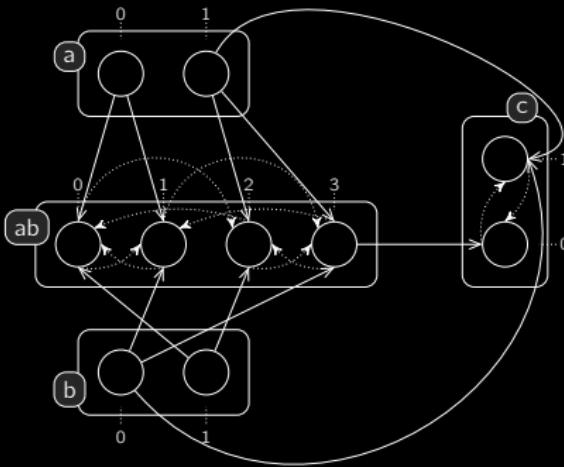


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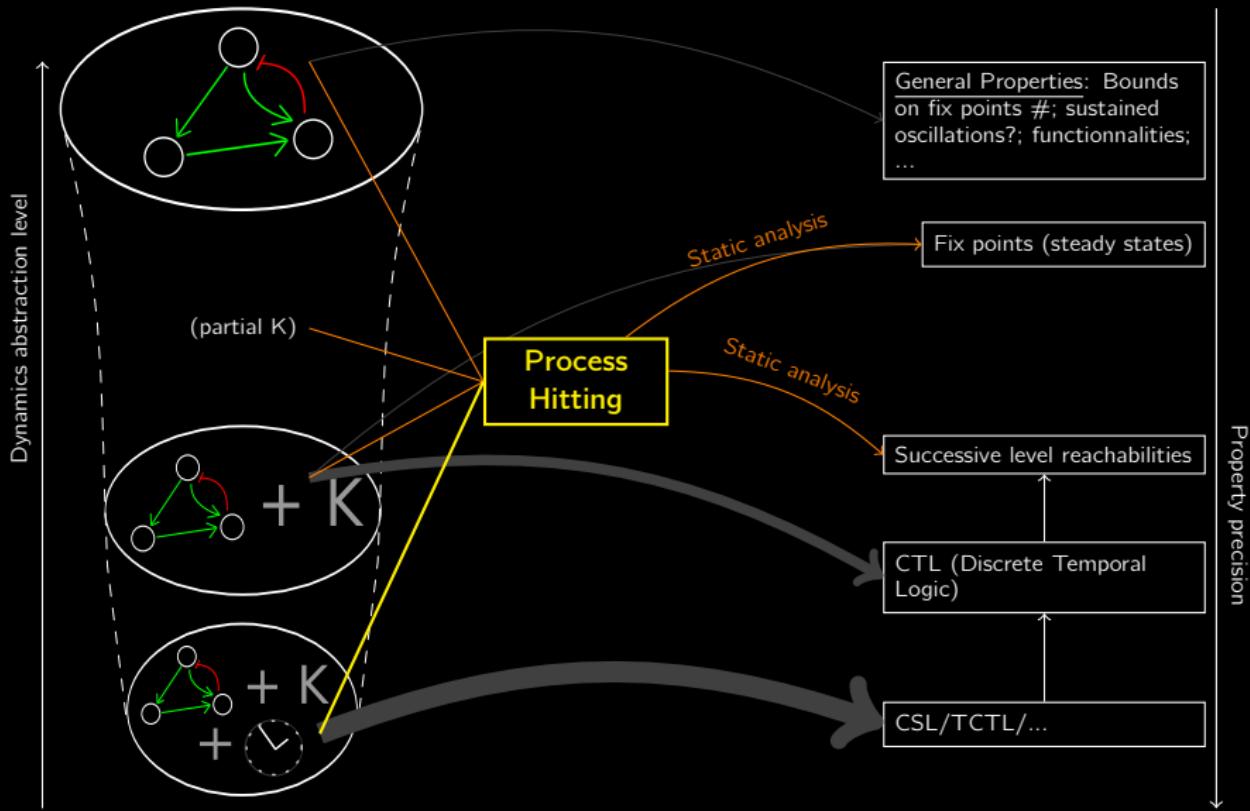


$$c = \neg a \wedge b$$



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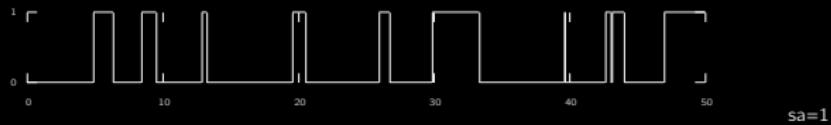
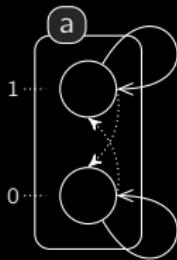
# Outline



## Stochasticity Absorption Factor

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

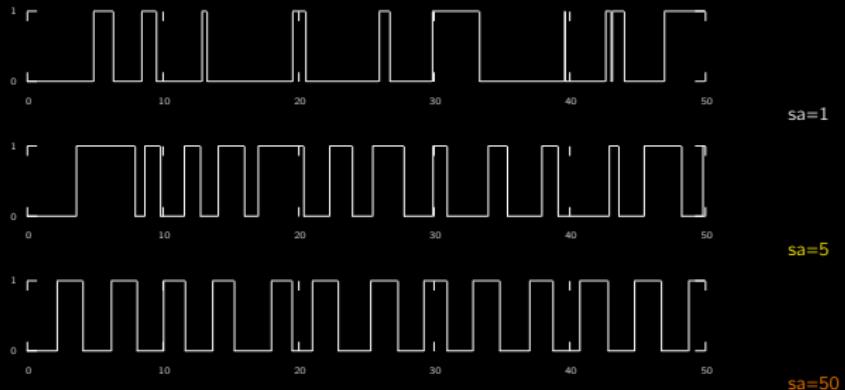
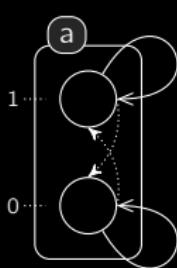
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- (Markov) Exponential distribution: mean  $r^{-1}$ ; variance  $r^{-2}$ .
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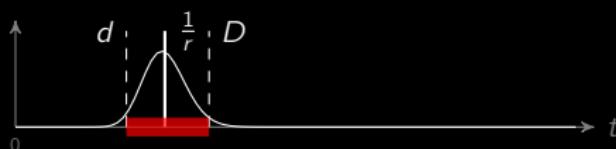
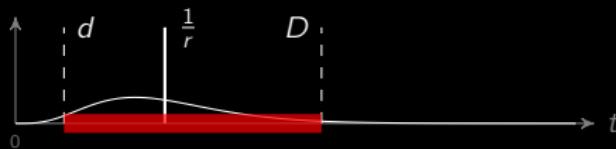
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- At our level of abstraction, we need to tune time features.
- Provide a stochasticity absorption factor  $sa$ :
- duration follows the sum of  $sa$  exponentials of rate  $r.sa$ ;
- mean  $r^{-1}$ ; variance  $r^{-2}sa^{-1}$  (Erlang distribution).



## 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  $1 - \alpha$ .
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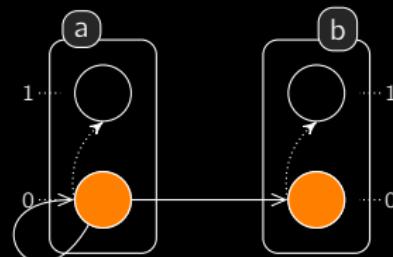
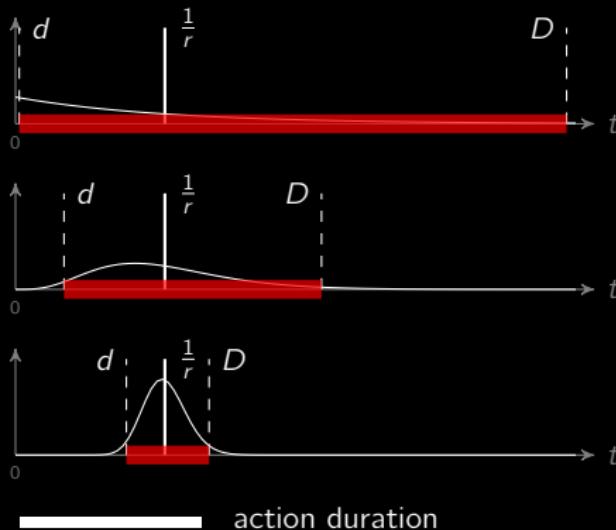


action duration

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

# Simulation and Model Checking

## Stochastic Model Checking

- Translation from the Erlang stochastic  $\pi$ -calculus to PRISM [Paulevé, Magnin, Roux in IEEE TSE, 2010].
- Applies to the Process Hitting as well.
- Not tractable with large stochasticiy 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].
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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

- Static analysis: derive dynamical properties without executing the model.
- Aim at coping with the combinatorial explosion of the dynamics.
- Several approaches: topological analysis, control flow analysis, constraints, abstractions, etc.

### BRNs analysis

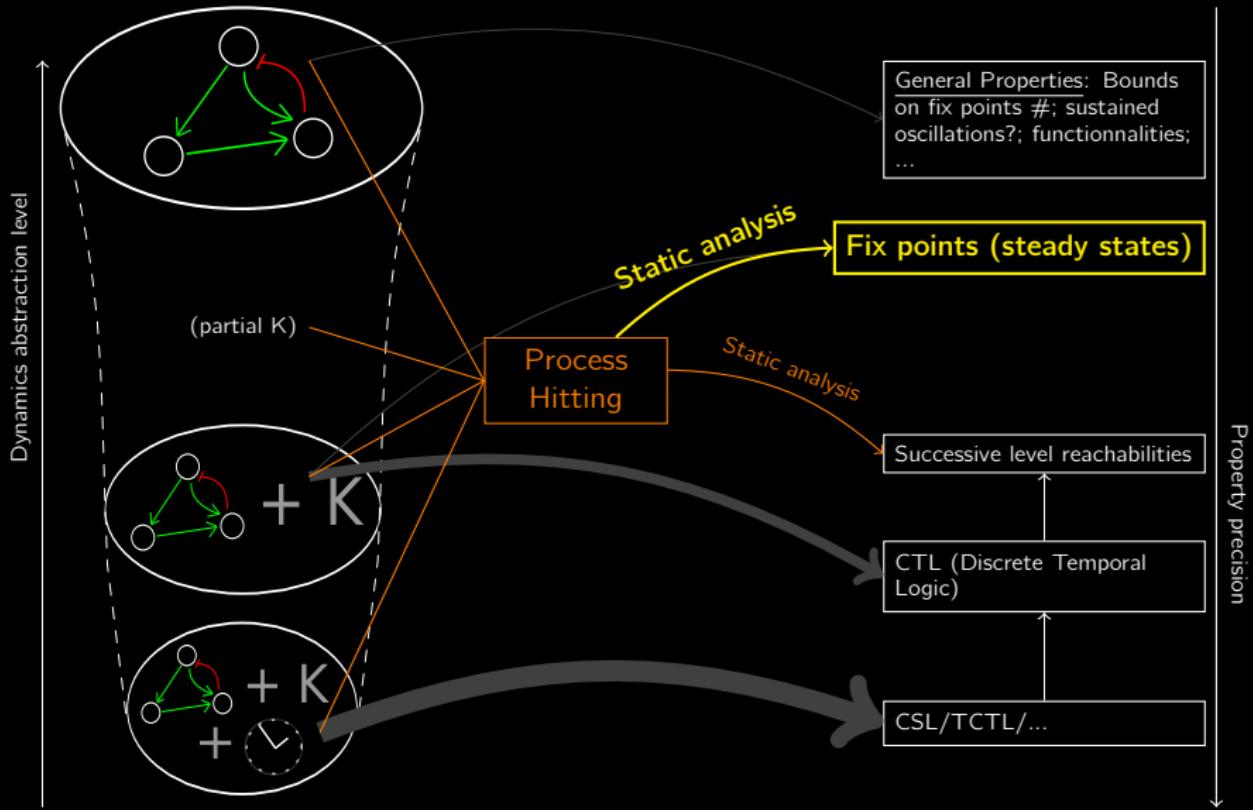
- Topological analyses of the interaction graph: bounds on fix points #; possibility of observing sustained oscillations, etc.
- Decision diagrams to derive fix points (from full BRN specification).

### Our Contributions using Process Hitting

- Topological analysis: complete characterisation of fix points.
- Abstract interpretation: over- and under-approximations of reachability properties.

⇒ brings new insight for deriving efficiently more precise dynamical properties from BRNs.

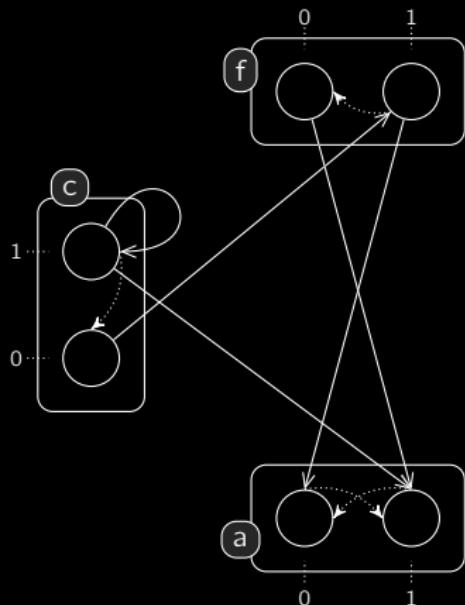
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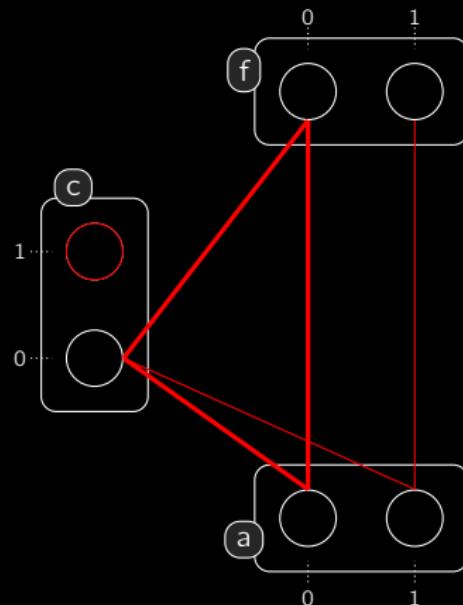
## Fix Points

[Paulevé, Magnin, Roux in TCSB 2011]

## Process Hitting



## Hitless graph

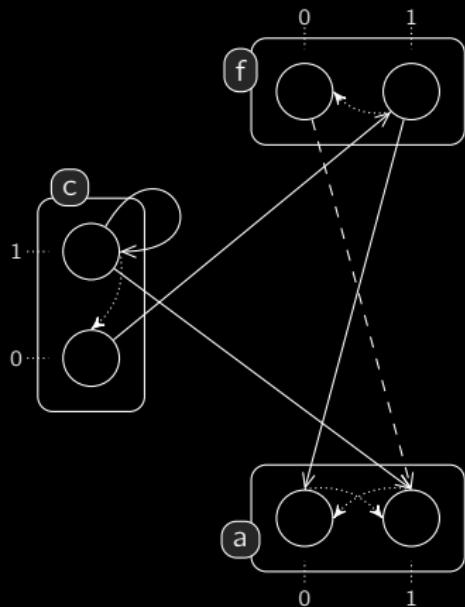


*n*-cliques are fix points

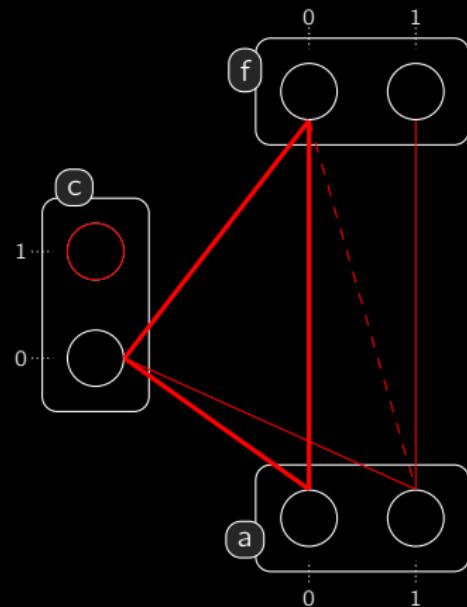
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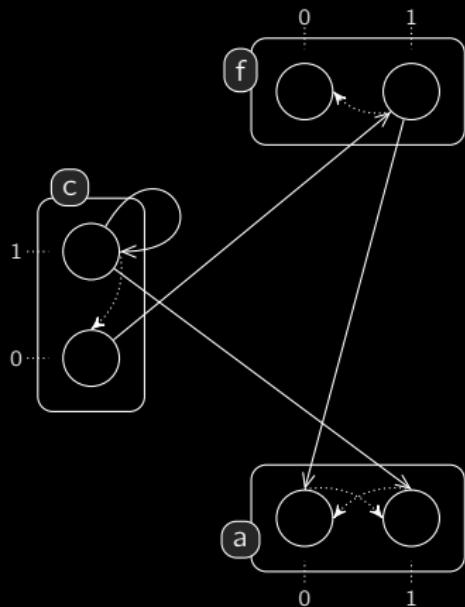


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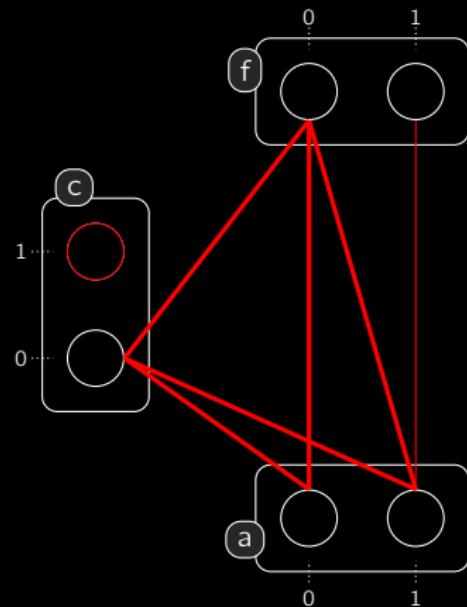
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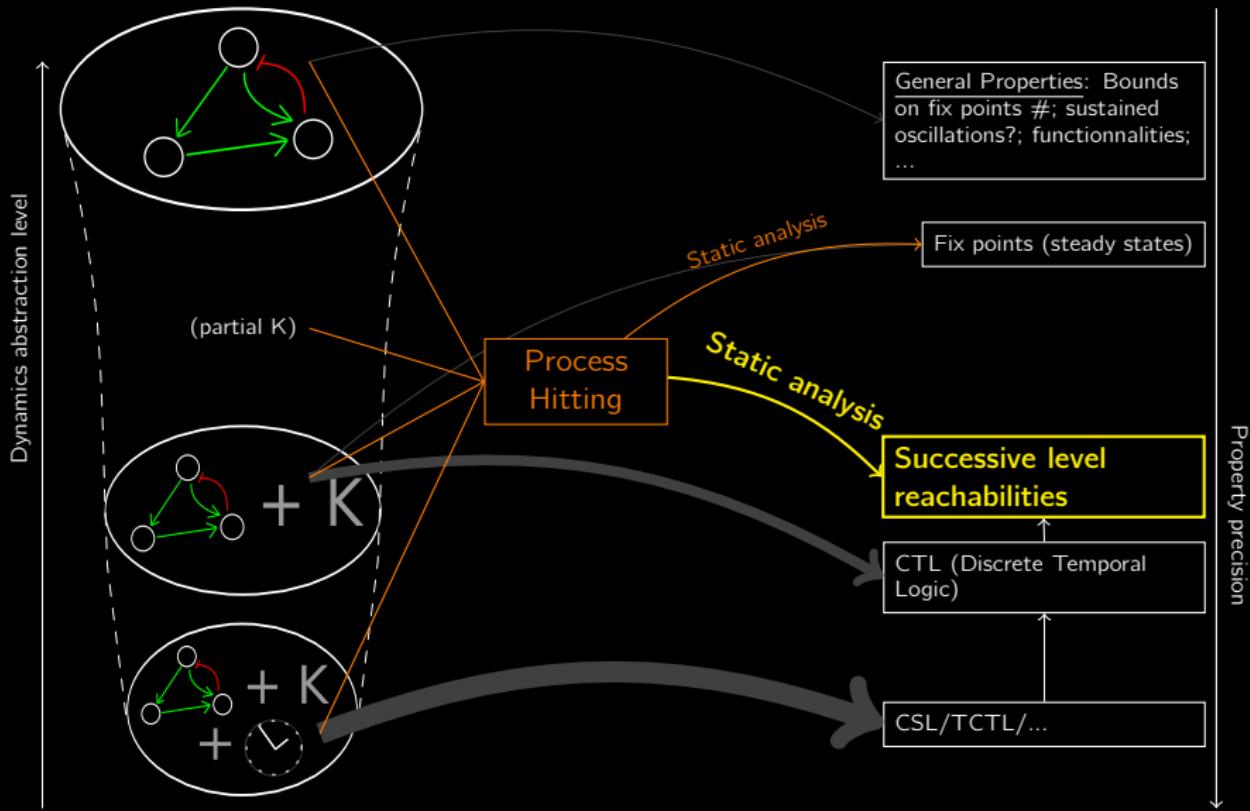


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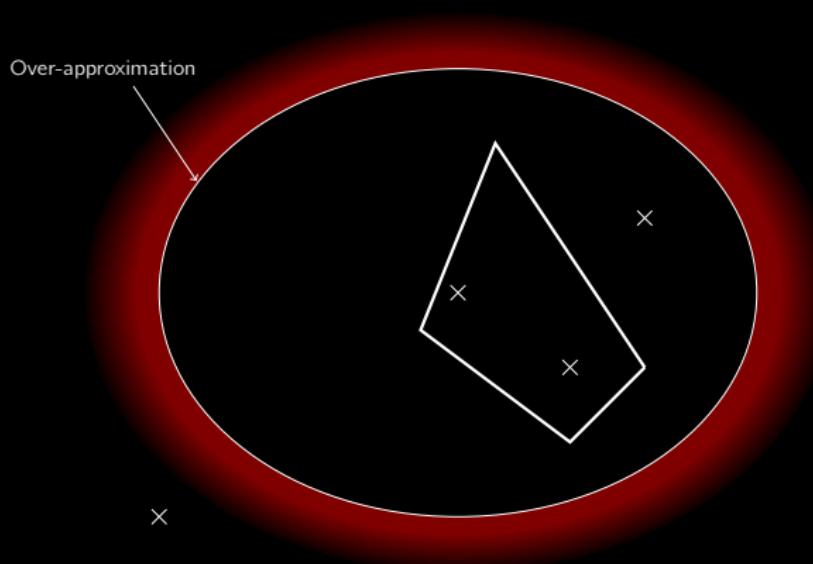


## Approximation of Reachability Properties

- 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.

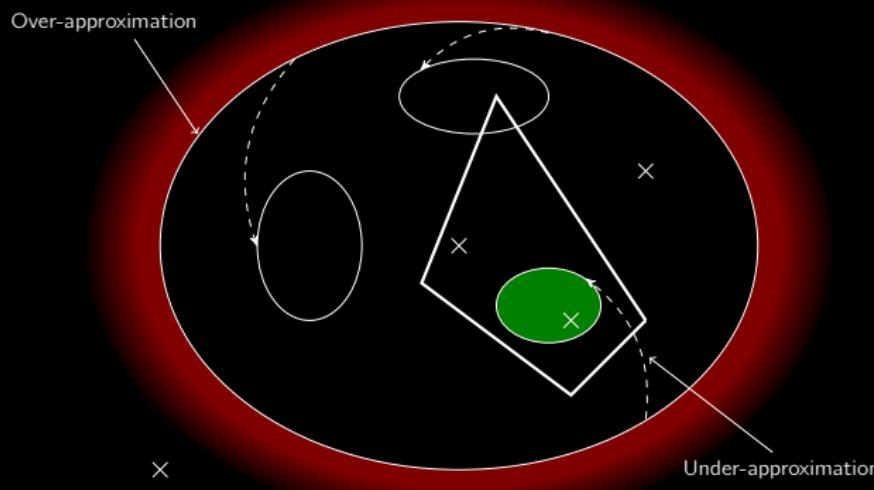
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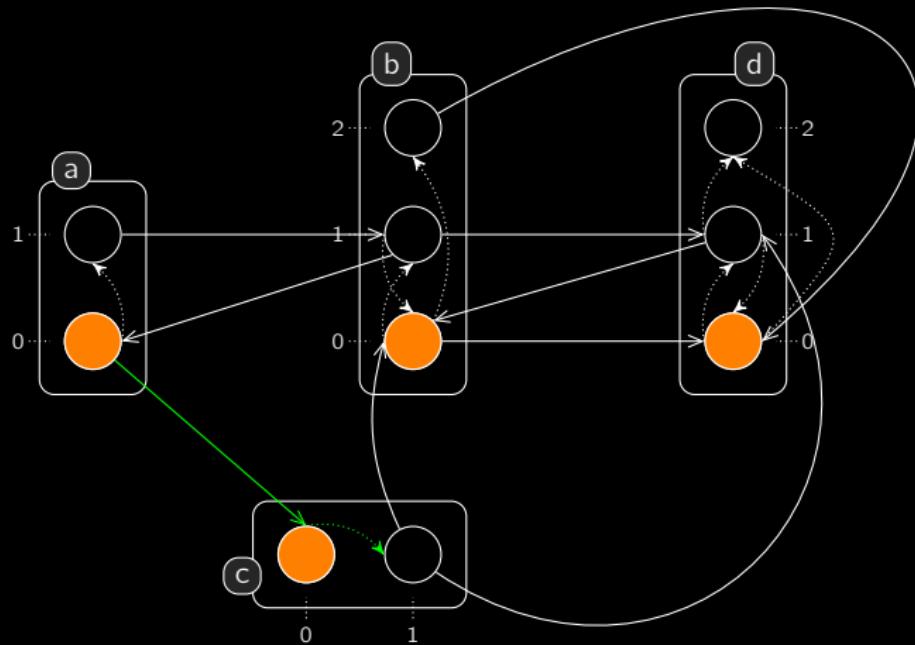


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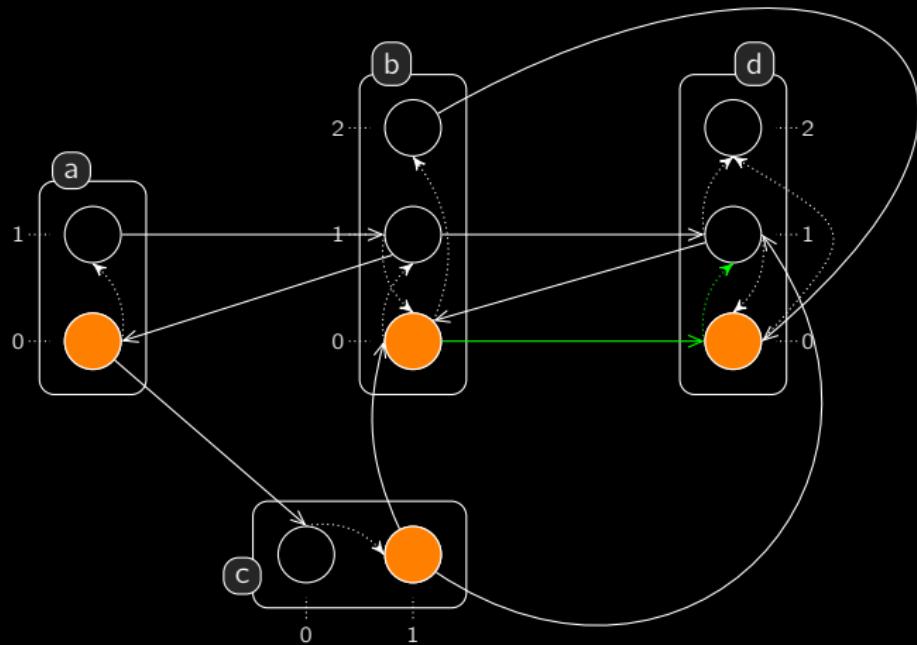
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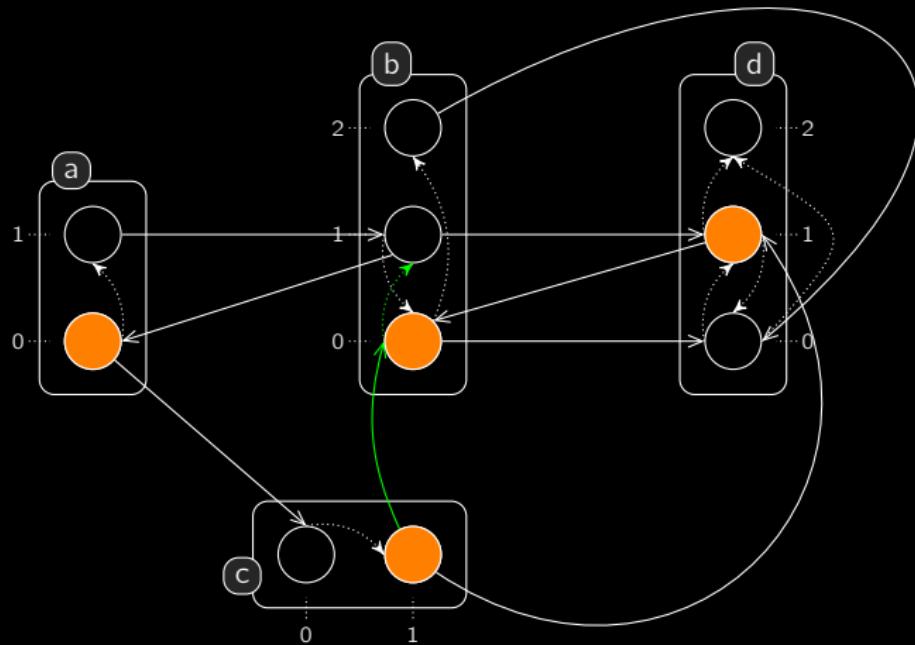
## Scenarios


$$a_0 \rightarrow c_0 \uparrow c_1 :: b_0 \rightarrow d_0 \uparrow d_1 :: c_1 \rightarrow b_0 \uparrow b_1 :: b_1 \rightarrow d_1 \uparrow d_2$$

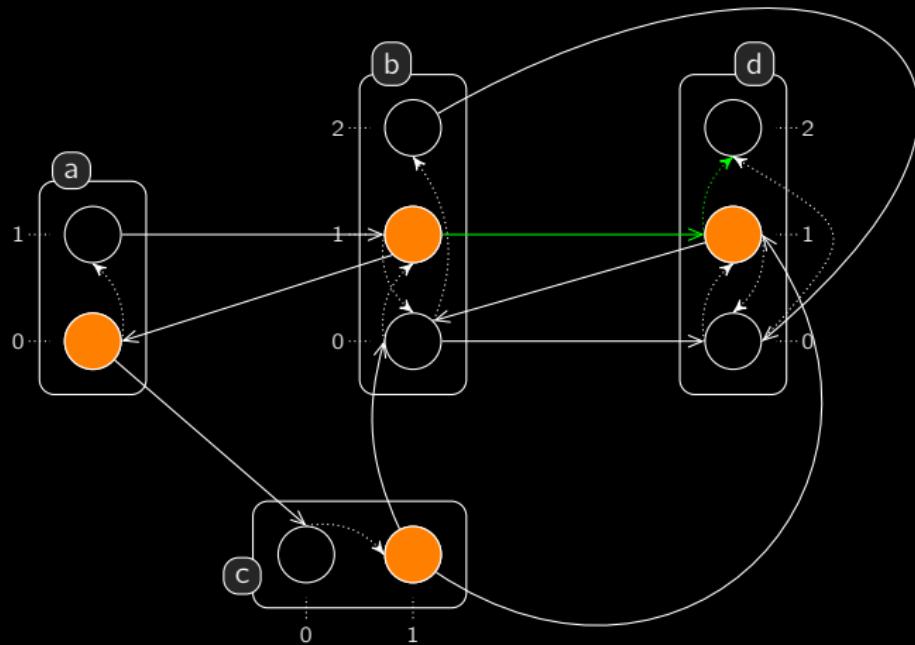
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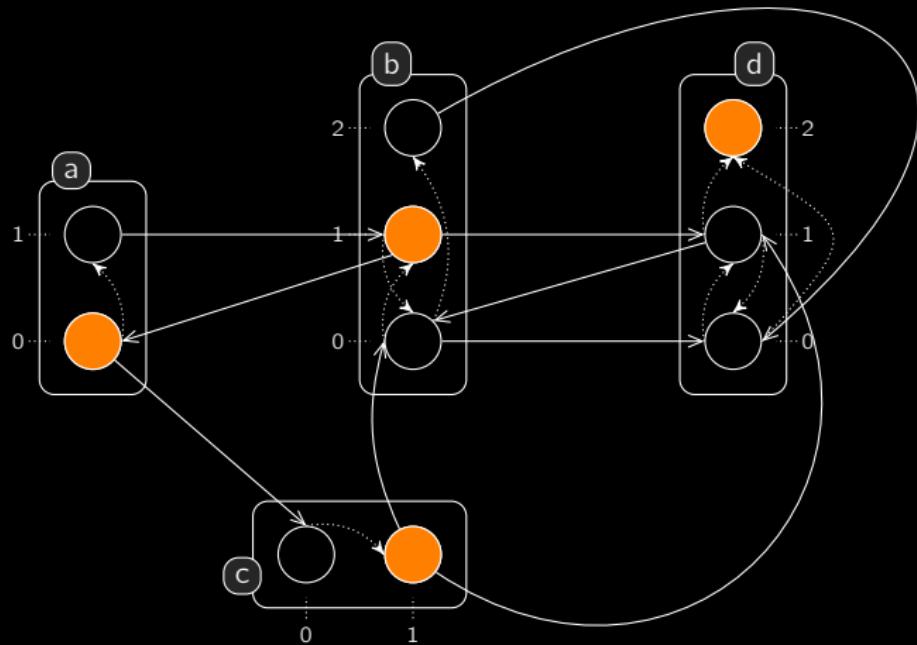
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## Abstract Interpretation of Scenarios

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

Scenarios – Successively playable actions.

- E.g.  $\delta = a_0 \rightarrow c_0 \uparrow c_1 :: b_0 \rightarrow d_0 \uparrow d_1 :: c_1 \rightarrow b_0 \uparrow b_1 :: b_1 \rightarrow d_1 \uparrow d_2$ .

Context — For each sort, subset of initial processes.

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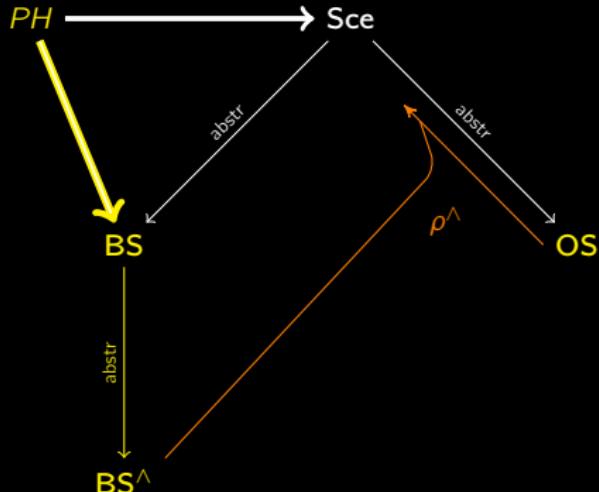
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- E.g.  $\varsigma = \langle a_0, b_0, b_2, c_0, d_0 \rangle$ .

Overall approach

- 2 complementary abstractions;
- Bounce Sequences BS;
- Objective Sequences OS;
- Concretization:  
 $\gamma_\varsigma : OS \mapsto \wp(Sce)$ ;
- Refinements:  
 $\rho : OS \mapsto \wp(OS)$ ;
- $\gamma_\varsigma(\omega) = \gamma_\varsigma(\rho(\omega))$ .



# Two Complementary Abstractions

$$a_0 \rightarrow c_0 \uparrow c_1 :: b_0 \rightarrow d_0 \uparrow d_1 :: c_1 \rightarrow b_0 \uparrow b_1 :: b_1 \rightarrow d_1 \uparrow d_2$$

## Abstraction by Objective Sequences

- $c_0 \uparrow^* c_1 :: d_0 \uparrow^* d_1 :: b_0 \uparrow^* b_1 :: d_1 \uparrow^* d_2;$

## Two Complementary Abstractions

$$a_0 \rightarrow c_0 \uparrow c_1 :: b_0 \rightarrow \textcolor{blue}{d}_0 \uparrow d_1 :: c_1 \rightarrow \textcolor{blue}{b}_0 \uparrow \textcolor{blue}{b}_1 :: b_1 \rightarrow d_1 \uparrow \textcolor{blue}{d}_2$$

### Abstraction by Objective Sequences

- $c_0 \uparrow^* c_1 :: d_0 \uparrow^* d_1 :: b_0 \uparrow^* b_1 :: d_1 \uparrow^* d_2$ ;
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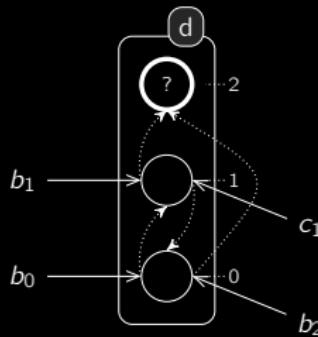
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E.g.:  $b_0 \rightarrow d_0 \uparrow d_1 :: b_1 \rightarrow d_1 \uparrow d_2$  ( $d_0 \uparrow^* d_2$ )

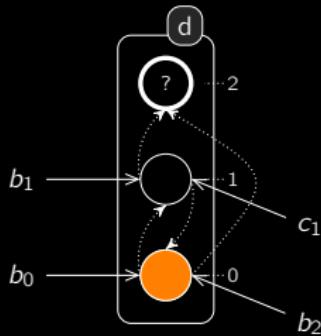
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E.g.:  $b_0 \rightarrow d_0 \uparrow d_1 :: b_1 \rightarrow d_1 \uparrow d_2$  ( $d_0 \uparrow^* d_2$ )

⇒ can be computed off-line:

- $\mathbf{BS}(d_0 \uparrow^* d_2) = \{b_0 \rightarrow d_0 \uparrow d_1 :: b_1 \rightarrow d_1 \uparrow d_2, b_2 \rightarrow d_0 \uparrow d_2\};$
- $\mathbf{BS}^\wedge(d_0 \uparrow^* d_2) = \{\{b_0, b_1\}, \{b_2\}\}.$

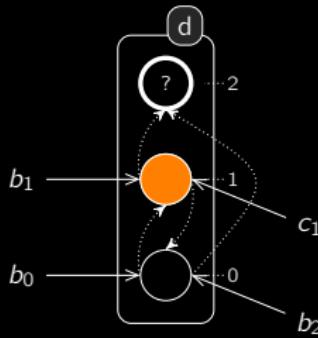
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## Objective Sequence Refinements

$$\gamma_\varsigma(\omega) = \{\delta \in \mathbf{Sce} \mid \omega \text{ abstracts } \delta \wedge \text{support}(\delta) \subseteq \varsigma\}.$$

Idea: the more details we know, the better  $\gamma_\varsigma(\omega)$  should be understood.

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**Objective Refinement by  $\mathbf{BS}^\wedge$ :**  $\rho^\wedge$

$\mathbf{Obj} \times \wp(\mathbf{BS}^\wedge)$	$\wp(\mathbf{OS})$
$d_0 \xrightarrow{*} d_2$	$\star \xrightarrow{*} \textcolor{blue}{b}_0 :: b_0 \xrightarrow{*} \textcolor{blue}{b}_1 :: d_0 \xrightarrow{*} d_2,$
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$\{\{b_0, b_1\}, \{b_2\}\}$	$\star \xrightarrow{*} \textcolor{blue}{b}_2 :: d_0 \xrightarrow{*} d_2$
$\gamma_\varsigma(d_0 \xrightarrow{*} d_2)$	$= \gamma_\varsigma(\rho^\wedge(d_0 \xrightarrow{*} d_2, \mathbf{BS}^\wedge(d_0 \xrightarrow{*} d_2)))$

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Generalization to  $\mathbf{OS}$  refinements:  $\tilde{\rho}$

$\mathbf{OS} \times \wp(\mathbf{BS}^\wedge)$	$\wp(\mathbf{OS})$
$\omega, \mathbf{BS}^\wedge$	interleave $\binom{\omega'}{\omega_{1..n-1}} :: \omega_{n.. \omega }$ where $n \in \mathbb{I}^\omega$ and $\omega' :: \omega_n \in \rho^\wedge(\omega_n, \mathbf{BS}^\wedge(\omega_n))$
$\gamma_\varsigma(\omega)$	$= \gamma_\varsigma(\tilde{\rho}(\omega, \mathbf{BS}^\wedge))$

## Objective Sequence Concretizability

a.k.a. Process Reachability Problem

$$\gamma_s(a_i \uparrow^* a_j :: \dots :: z_k \uparrow^* z_l) \neq \emptyset?$$

CTL:  $EF(s[a] = a_j \wedge EF(\dots \wedge EF(s'[z] = z_l) \dots)).$

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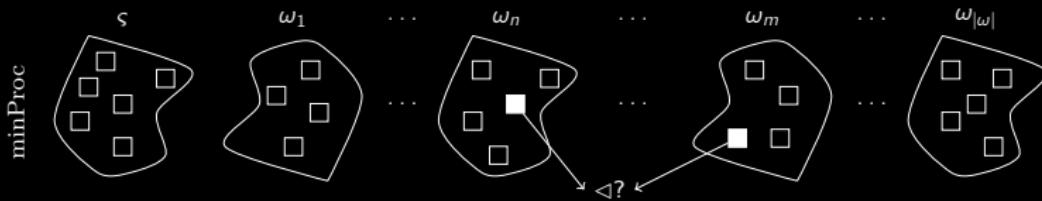
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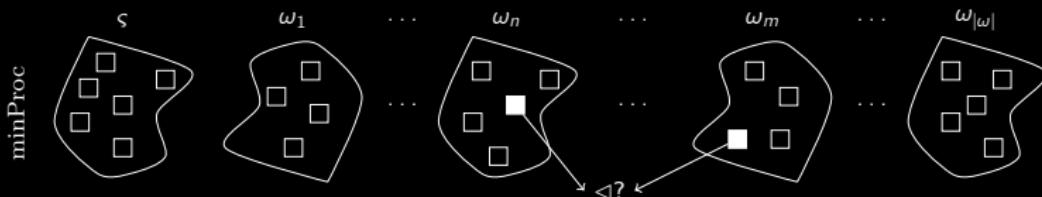
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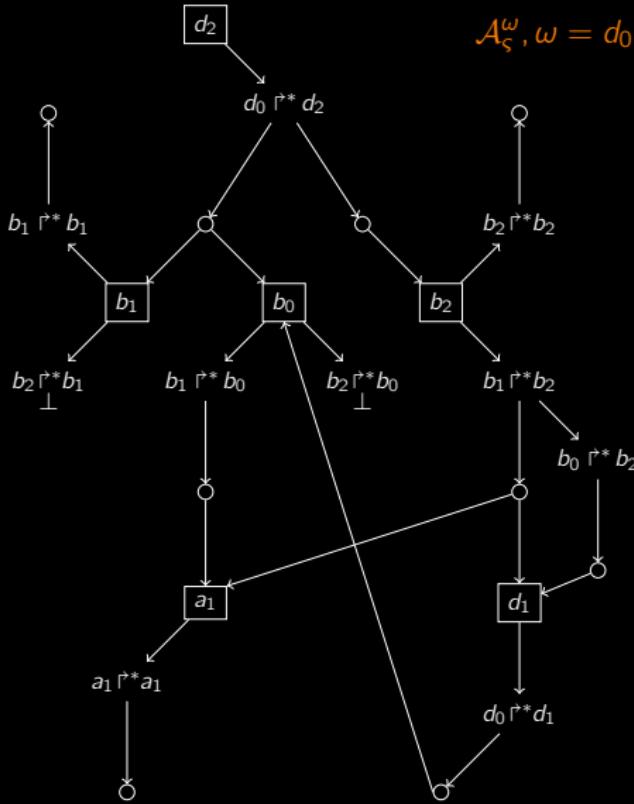
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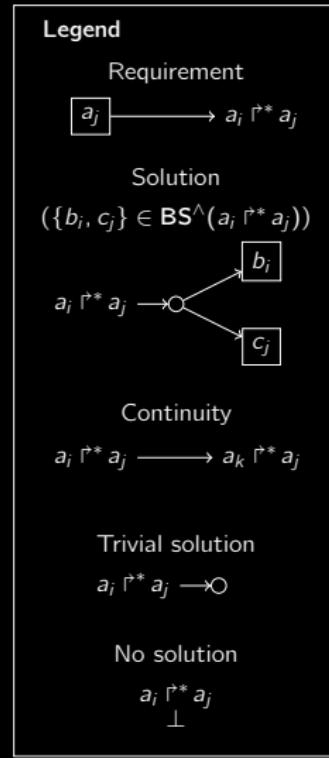
## Under-approximation

- Sufficient condition: impose a particular structure of scenarios.

# Implementation of Over-approximations



$$\mathcal{A}_\varsigma^\omega, \omega = d_0 \cap^* d_2, \varsigma = \langle a_1, b_1, b_2, c_1, d_0 \rangle$$



# Complexities

## Data structures

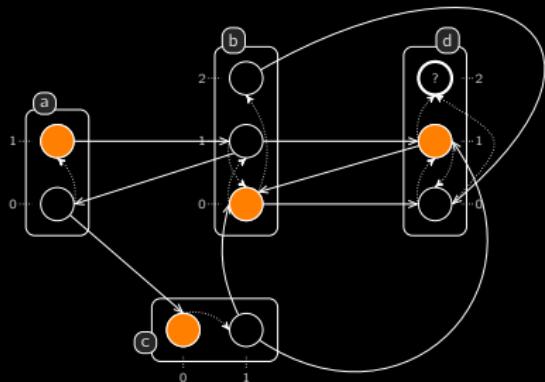
- Computing  $\text{BS}$  is exponential in the number of processes in a single sort.
- Computing  $\text{BS}^\wedge$  is faster, but still exponential.
- The size of  $\mathcal{A}_\varsigma^\omega$  is  $\approx$  polynomial in the number of processes.

## Analyses

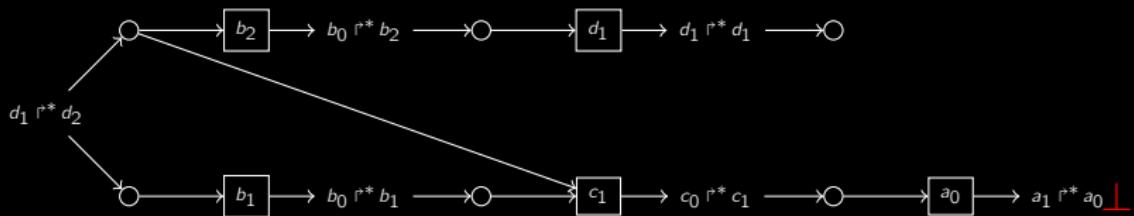
- Over-approximations are polynomial in the size of  $\mathcal{A}_\varsigma^\omega$ .
- Under-approximation is polynomial in the size of  $\mathcal{A}_\varsigma^\omega$ ; and can be applied a number of times exponential in the number of solutions of a single objective.

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

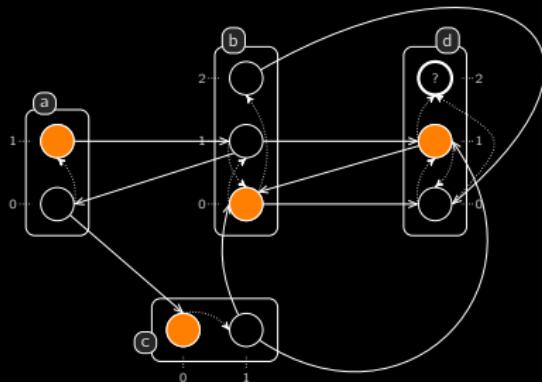
## Un-ordered Over-approximation



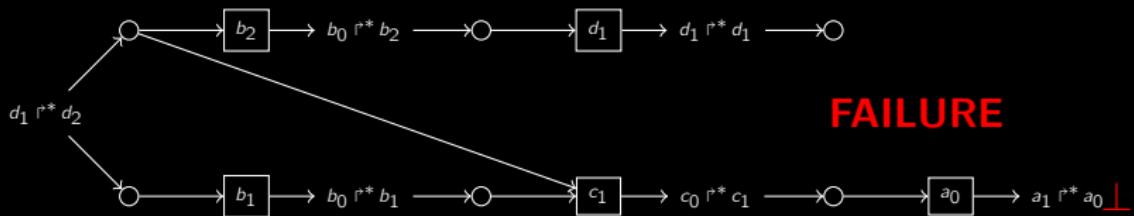
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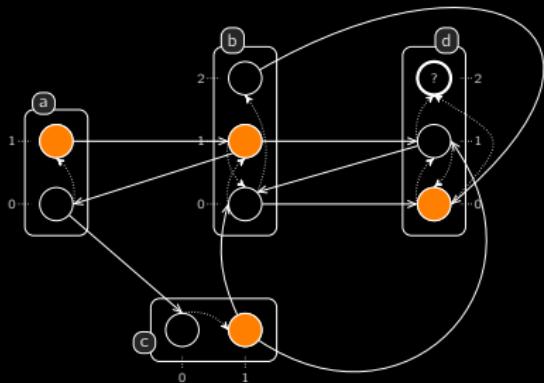
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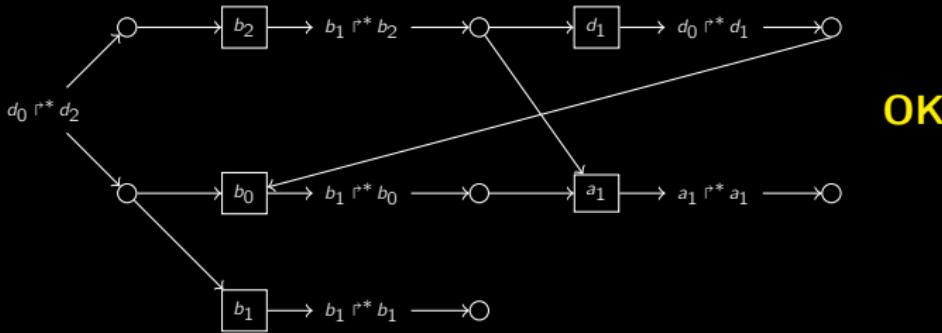
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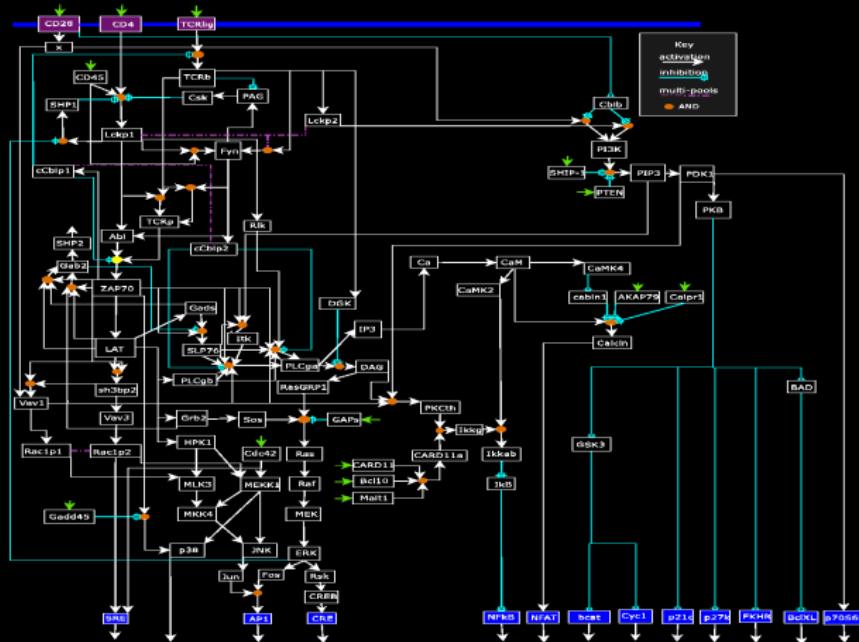


## Outline

- 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

# T-Cell Receptor Signalling Pathway

(94 components)



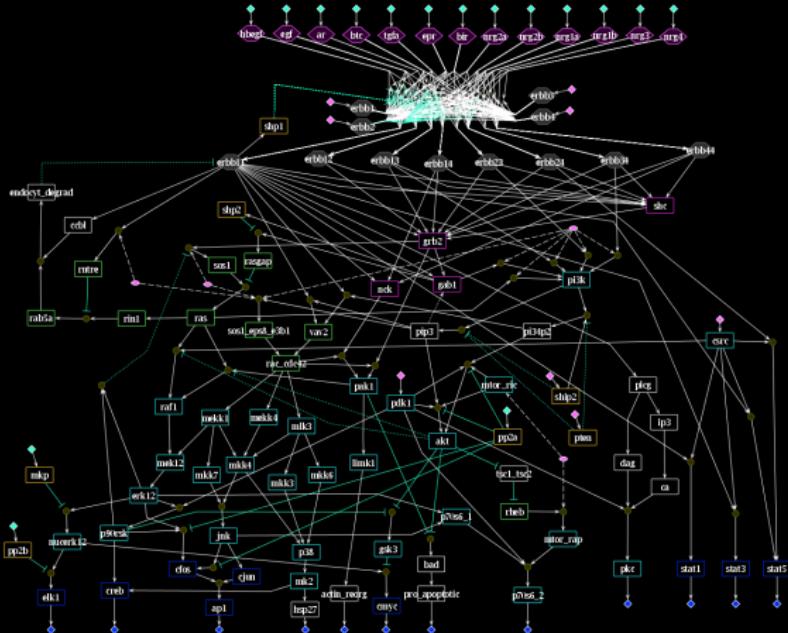
[Saez-Rodriguez, et al.  
in PLoS Comput Biol,  
07]

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

Reachability analysis always conclusive; around **0.01s** (compared to *libddd*: out of memory). [<http://ddd.lip6.fr>]

## 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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## Conclusion

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- *Elementary* framework for **dynamical complex systems**;
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- Generic tuning of **time features within stochastic models** (simulation + standard model checking).
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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

Derive more properties

- Characterisation of **attractors**;
- Verification for **sustained oscillations**;
- etc.

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- Extension to the **Process Hitting with Priorities** (e.g., two classes of actions: instantaneous and non-instantaneous);
- Idea: **detect key concurrencies** leading to the satisfaction or non-satisfaction of a properties.

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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.