

Master internship

Exhaustive analysis of the dynamics of Process Hitting through Answer Set Programming

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joint work Olivier ROUX & Morgan MAGNIN

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Context and Aims

MeForBio team:
Algebraic modelling to study
complex dynamical biological systems

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Predicting the **evolutions** of the network.

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Predicting the **evolutions** of the network.

3) What for?

searching of PH **properties** through ASP (Fixed points, reachability).

Plan

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 - Definition
 - Example of an ASP program

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- 5) Conclusion & prospect

Answer Set Programming

ASP:

- Logic program written in language of AnsProlog*
- Form of rules :

$$\begin{aligned}
 & \textit{head} \leftarrow \textit{body}. \\
 L_0 & \leftarrow L_1, \dots, L_m, \textbf{not } L_{m+1}, \dots, \textbf{not } L_n.
 \end{aligned}$$

with each L_j : literal in the sense of classical logic.

Rule's meaning:

If L_1, \dots, L_m are **true** and if L_{m+1}, \dots, L_n are **false**
then L_0 is **true**.

Answer Set Programming

Special types of rules:

- **Constraint** :

$$\leftarrow L_1, \dots, L_m, \text{ not } L_{m+1}, \dots, \text{ not } L_n.$$

- **Fact** :

$$L_0.$$

- **Cardinality** :

$$\min\{L_0, \dots, L_j\} \max \leftarrow L_1, \dots, L_m, \text{ not } L_{m+1}, \dots, \text{ not } L_n.$$

Answer Set Programming

Example:

bird(*X*) ← *lays_egg*(*X*).

mammal(*X*) ← *engender*(*X*).

fly(*X*) ← *bird*(*X*), **not** *mammal*(*X*).

lays_egg(*tweety*).

Answer Set Programming

Example:

$bird(X) \leftarrow lays_egg(X).$

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$lays_egg(tweety).$

Solution:

$bird(tweety) \leftarrow True.$

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$fly(tweety) \leftarrow True.$

Answer: $fly(tweety), bird(tweety).$

The Process Hitting modeling



Sorts: components a, b, z

The Process Hitting modeling



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Processes: local states / levels of expression z_0, z_1, z_2

The Process Hitting modeling

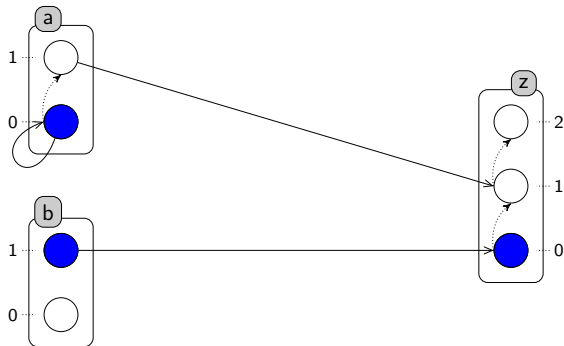


Sorts: components a, b, z

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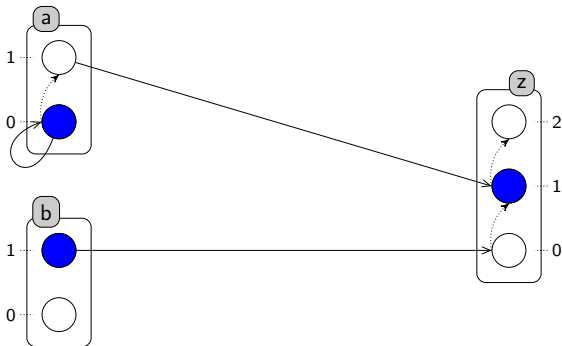
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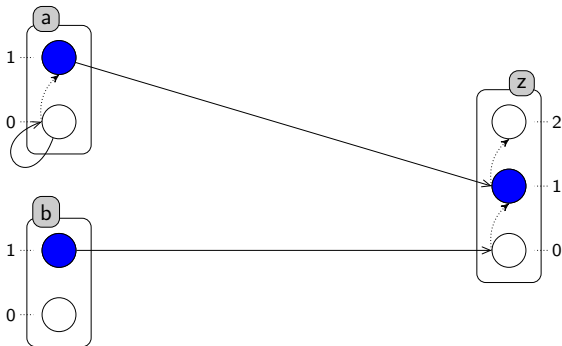
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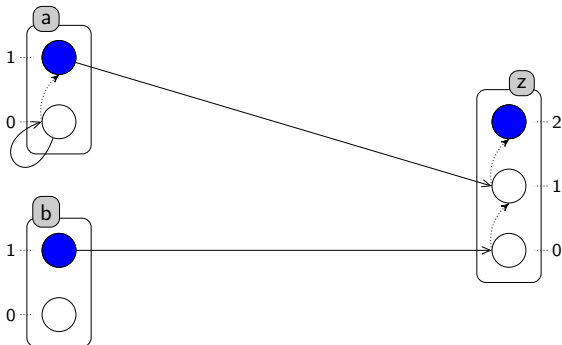
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PH through ASP

Network traduction:

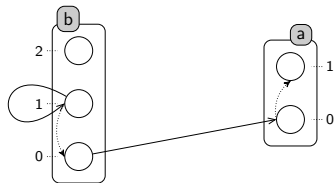
- **Sort:** `sort(A)` .
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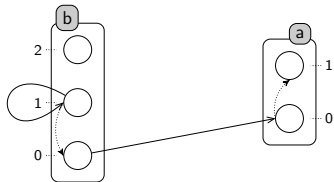


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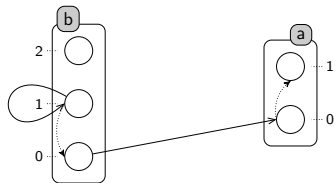
`sort("a").` `sort("b").`

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```

sort("a").  sort("b").
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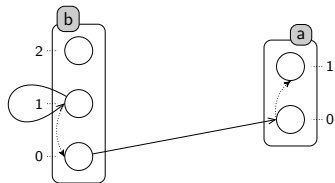
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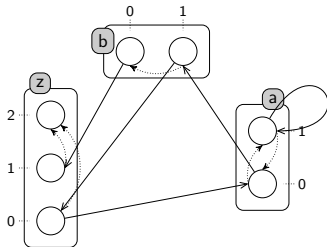
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sort("a").  sort("b").
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action("b",0,"a",0,1).
action("b",1,"b",1,0).

```

Fixed Points

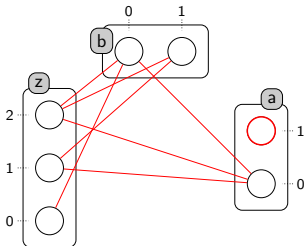
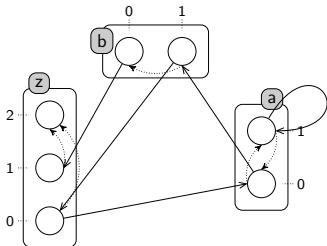
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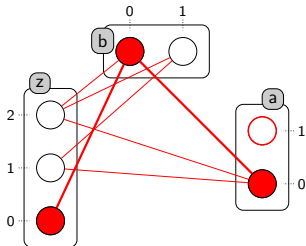
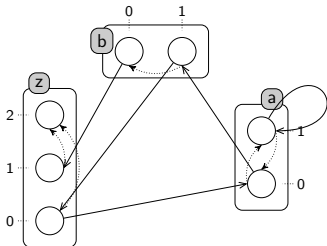
→ Hitless Graph



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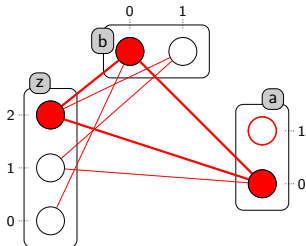
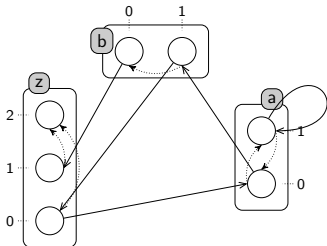
→ Hitless Graph → **n-clics** = fixed points



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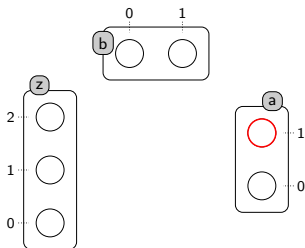
→ Hitless Graph → **n-clics** = fixed points



Implementation of the algorithm (N-Cliques)

Definition of hitless graph :

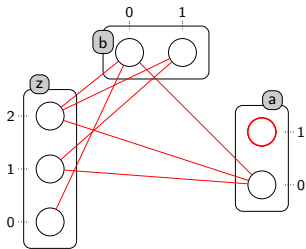
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noAction(A,I,B,J) :- not hit(A,I,B,J), not hit(B,J,A,I), A!=B,
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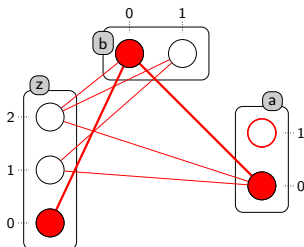
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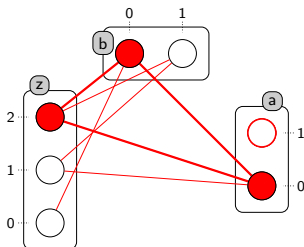
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Find Fixed points :

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noHit(A,I,B,J) :- noAction(A,I,B,J),
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noExistFixPoint :- 0 {noHit(A,I,B,J)} 0, selectProcess(A,I),
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Implementation of the algorithm (N-Cliques)

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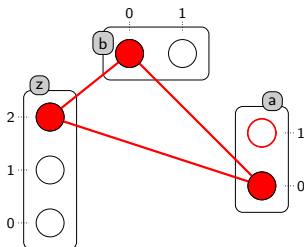
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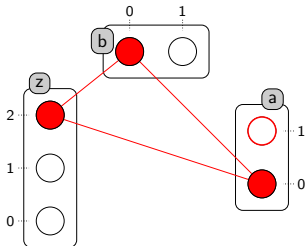


Static analysis

Fixed point through ASP

ASP program result:

Answer 1: `fixProcess(a,0), fixProcess(b,0), fixProcess(z,2).`



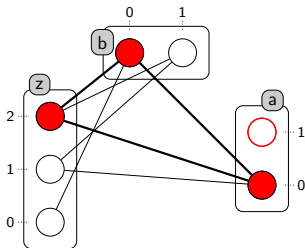
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fixProcess(A,I) :- selectProcess(A,I).
  
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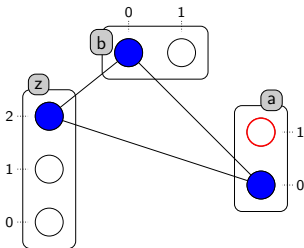


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Static analysis

Fixed Point

Comparison

Model	#sorts	#states	#fix-point	mthd1	mthd2	PINT
mvbrn	3	12	1	0.000s	0.000s	0.006s
ERBB	42	2^{70}	3	0.220s	0.000s	0.017s
tcrsig40	54	2^{73}	1	0.220s	0.020s	0.021s
tcrsig94	133	2^{194}	0	2.540s	0.060s	0.027s
egfr104	193	2^{320}	0	8.220s	0.140s	0.074s

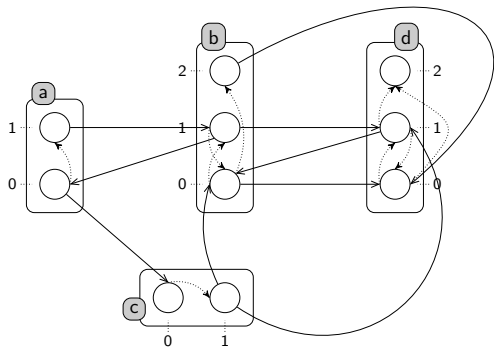
Figure: Execution time of ASP methods and PINT applied for biological networks with a desktop computer (core i5 and 4GB RAM).

PINT : a library developed to parse and study PH models.

Dynamic analysis

Reachability

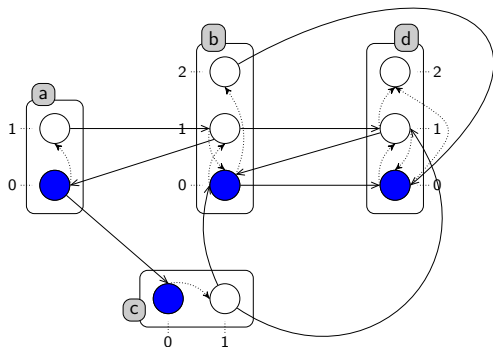
Reachability of processes:



Dynamic analysis

Reachability

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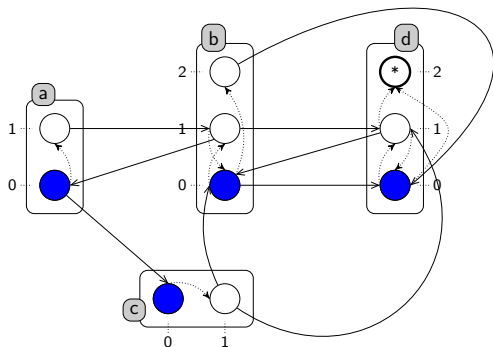
- Initial context

$\langle a_0, b_0, c_0, z_0 \rangle$

Dynamic analysis

Reachability

Reachability of processes:



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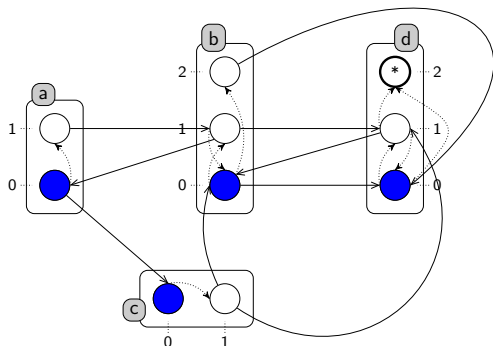
- Objectives

$[\uparrow d_2]$

Dynamic analysis

Reachability

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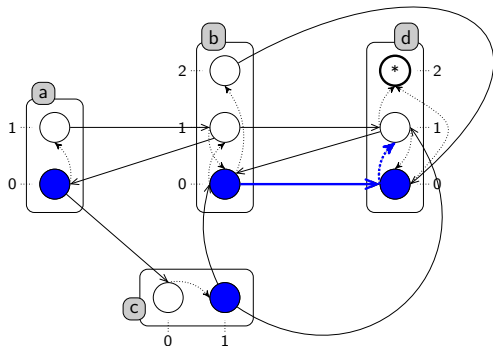
→ Concretization of the objective = scenario

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

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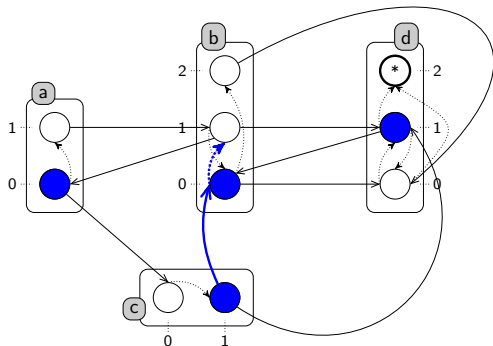
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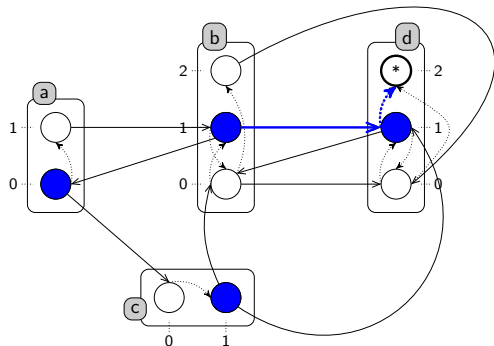
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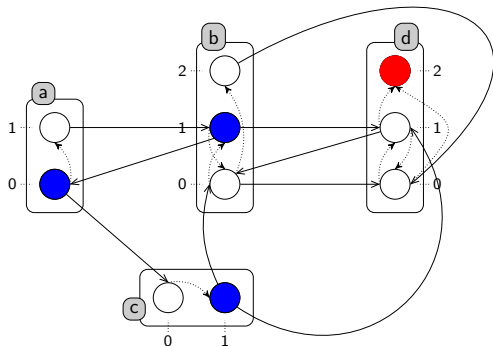
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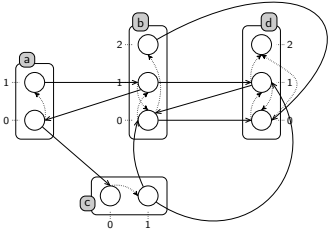
→ Concretization of the objective = scenario

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

Dynamic analysis

Evolution through ASP

Network evolution through ASP



Dynamic analysis

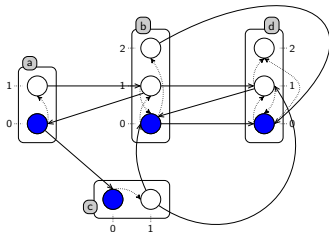
Evolution through ASP

Network evolution through ASP

Initializing :

`init(activeProcess("a",0)).`

avec a: sorte, 0: indice du processus



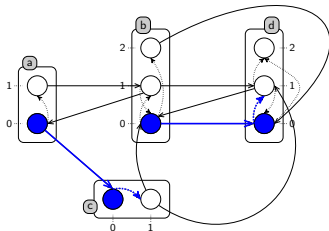
Dynamic analysis

Evolution through ASP

Network evolution through ASP

Playable actions at step T :

```
playableAction(A,I,B,J,K,T) :- action(A,I,B,J,K),
    instate(activeProcess(A,I),T),
    instate(activeProcess(B,J),T), time(T).
```



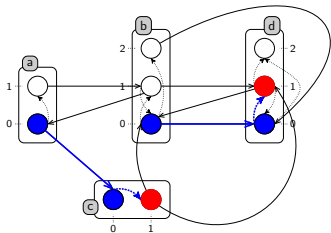
Dynamic analysis

Evolution through ASP

Network evolution through ASP

Change active processes :

```
{activeFromTo(B,J,K,T)} :- playableAction(A,I,B,J,K,T),
                             J!=K, time(T).
                             :- 2{ activeFromTo(B,J,K,T)}, time(T).
```



Dynamic analysis

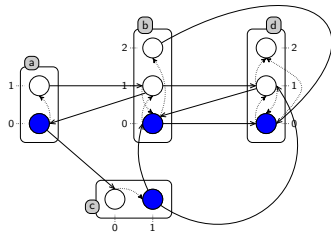
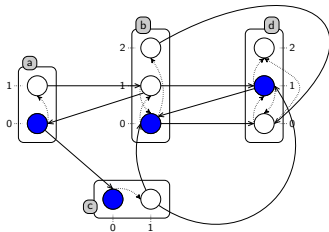
Evolution through ASP

Network evolution through ASP

Active processes at next step (T+1) :

```

instate(activeProcess(B,K),T+1) :- activeFromTo(B,J,K,T), time(T).
instate(activeProcess(A,I),T+1) :- instate(activeProcess(A,I),T),
                                     activeFromTo(B,J,K,T), A!=B, time(T).
  
```



Dynamic analysis

Evolution through ASP

Network evolution through ASP

```
time(0..N).
```

Results ($N = 3$) :

```
Answer 1:  activeFromTo("d",0,1,0) activeFromTo("c",0,1,1)
          actifFromTo("b",0,1,2).
```

```
Answer 2:  activeFromTo("d",0,1,0) activeFromTo("b",0,2,1)
```

```
Answer 3:  activeFromTo("c",0,1,0) activeFromTo("d",0,1,1)
          activeFromTo("d",1,0,2) activeFromTo("b",0,1,3)
```

```
...
```

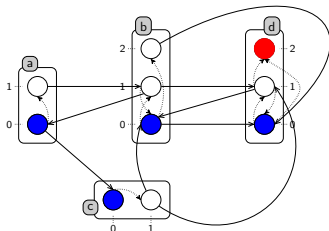
```
Answer 29: activeFromTo("c",0,1,0) activeFromTo("b",0,1,1)
          activeFromTo("a",0,1,2)
```

Dynamic analysis

Reachability through ASP

Success reachability through ASP:

`goal(activeProcess("d",2)).`

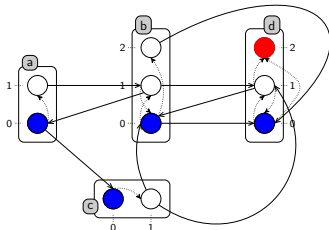


Dynamic analysis

Reachability through ASP

Success reachability through ASP:

```
goal(activeProcess("d",2)).
satisfiable(F,T) :- goal(F), instate(F,T).
:- not satisfiableTot.
```

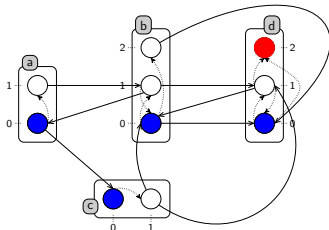


Dynamic analysis

Reachability through ASP

Success reachability through ASP:

```
goal(activeProcess("d",2)).
satisfiable(F,T) :- goal(F), instate(F,T).
:- not satisfiableTot.
time(0..N).
```

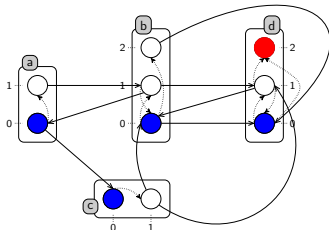


Dynamic analysis

Reachability through ASP

Success reachability through ASP:

```
goal(activeProcess("d",2)).
satisfiable(F,T) :- goal(F), instate(F,T).
:- not satisfiableTot.
time(0..N).
predict N -> Inconvenient
```



Dynamic analysis

Reachability through ASP

Results for ($N = 2$) :

UNSATISFIABLE

Results for ($N = 3$) :

Answer 1: `activeFromTo(c,0,1,0), activeFromTo(d,0,1,1),
activeFromTo(b,0,1,2), activeFromTo(d,1,2,3).`

Answer 2: `activeFromTo("d",0,1,0) activeFromTo("c",0,1,1)
activeFromTo("b",0,1,2) activeFromTo("d",1,2,3)`

Dynamic analysis

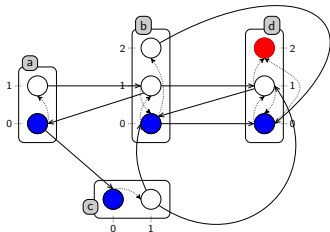
Reachability through ASP

Success reachability through ASP iterative:

```

goal(activeProcess("d",2)).
#base
instate(F,0) :- init(F).
#cumulative t
playableAction(A, I, B, J, K,t), activeFromTo(B, J, K,t),
instate(activeProcess(A, I),t + 1)...
#volatile t
notSatisfaisable(t) :- goal(F), not instate(F,t).
:- notSatisfaisable(t).

```



Dynamic analysis

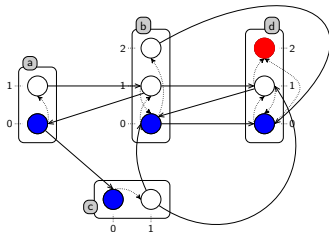
Reachability through ASP

Success reachability through ASP iterative:

Results:

Answer 1: `activeFromTo(c,0,1,0)`, `activeFromTo(d,0,1,1)`,
`activeFromTo(b,0,1,2)`, `activeFromTo(d,1,2,3)`.

Answer 2: `activeFromTo("d",0,1,0)` `activeFromTo("c",0,1,1)`
`activeFromTo("b",0,1,2)` `activeFromTo("d",1,2,3)`



Dynamic analysis

Reachability through ASP

Comparison:

Initializing biological models components and the objectives.

Model	#sorts	#states	#steps	ASP	ASP _i	PINT
Exemple	4	36	4	0.000s	0.000s	0.000s
ERBB	42	2^{70}	18	10.620s	5.020s	0.022s
tcrsig40	54	2^{73}	26	156.500s	127.250s	0.012s

Figure: Excecution time of ASP methods (CLINGO et ICLINGO) and PINT applied for biological networks with a desktop computer (core i5 and 4GB RAM)

Dynamic analysis

Reachability through ASP

Comparison:

Method of Rocca et al.:

- ASP
- CTL properties with model cheking (AF, EF, AG...)
- Transitions graph

Dynamic analysis

Reachability through ASP

Comparison:

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Comparaison of the property **EF**

$$\text{prop} = \mathbf{EF} (I_0, \text{goal})$$

Dynamic analysis

Reachability through ASP

Comparaison:

Example: Tail resorption of tadpole :
12 sorts, 42 process, 139 actions and 524.288 states.

$$\text{prop} = \text{EF}(I_0, \text{goal})$$

Dynamic analysis

Reachability through ASP

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Network traduction :

- Transition graph: 3min6s
- Process Hitting : 0.346s

Dynamic analysis

Reachability through ASP

Comparaison:

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Network traduction :

- Transition graph: 3min6s
- Process Hitting : 0.346s

Property verification :

- Rocca et al. method : 7min17s
- our itirative method : 1.9s

Conclusion & Prospects

- New dynamic analysis of Process Hitting models:
 - Fixed point
 - Network evolution
 - Reachability
- Prospects:
 - Adaptation on other models (PN, model of Thomas...)
 - Eliminating cycles
 - Search attractors
 - Reverse reachability (*goal* \rightarrow I_0 ?)

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Thanks for your attention