

Challenges with the logical modeling of biological systems: learning and analyzing discrete and hybrid models

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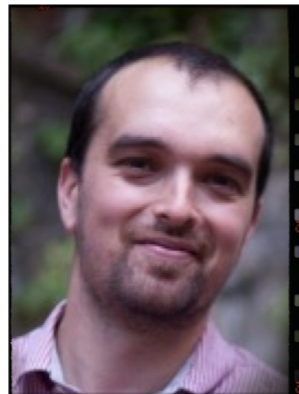
National Institute of Informatics - Japan

Inoue Lab.

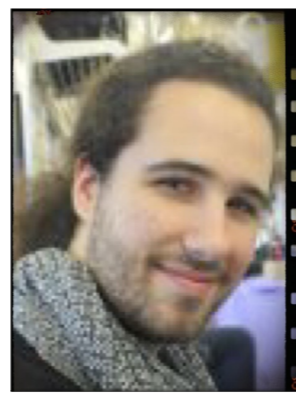
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Olivier
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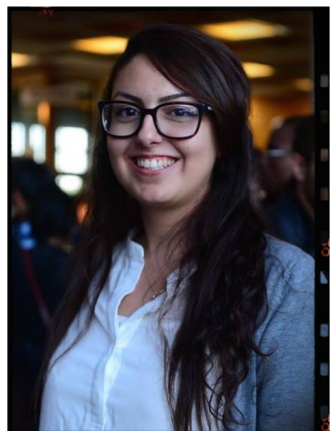
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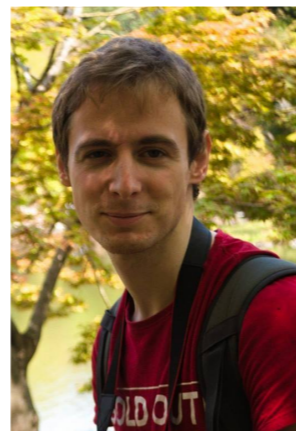


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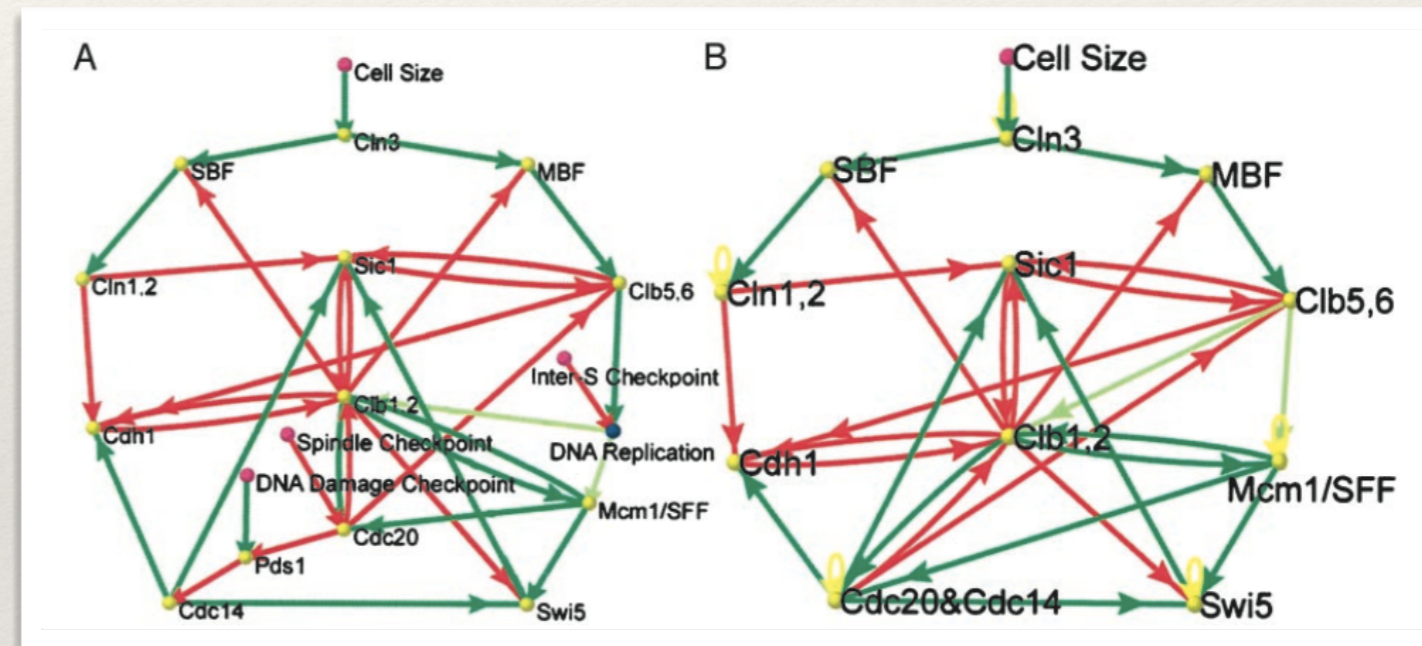
Mitsuhiro
ODAKA

Motivations

- ❖ Goals:
 - ❖ Help to the **diagnosis** and **prevention** of diseases
 - ❖ Develop new **therapies**
- ❖ **Systems biology**
 - ❖ Consider a living organism as a system of **interacting networks**
 - ❖ Link between physiological functions / **dynamics** of the regulations

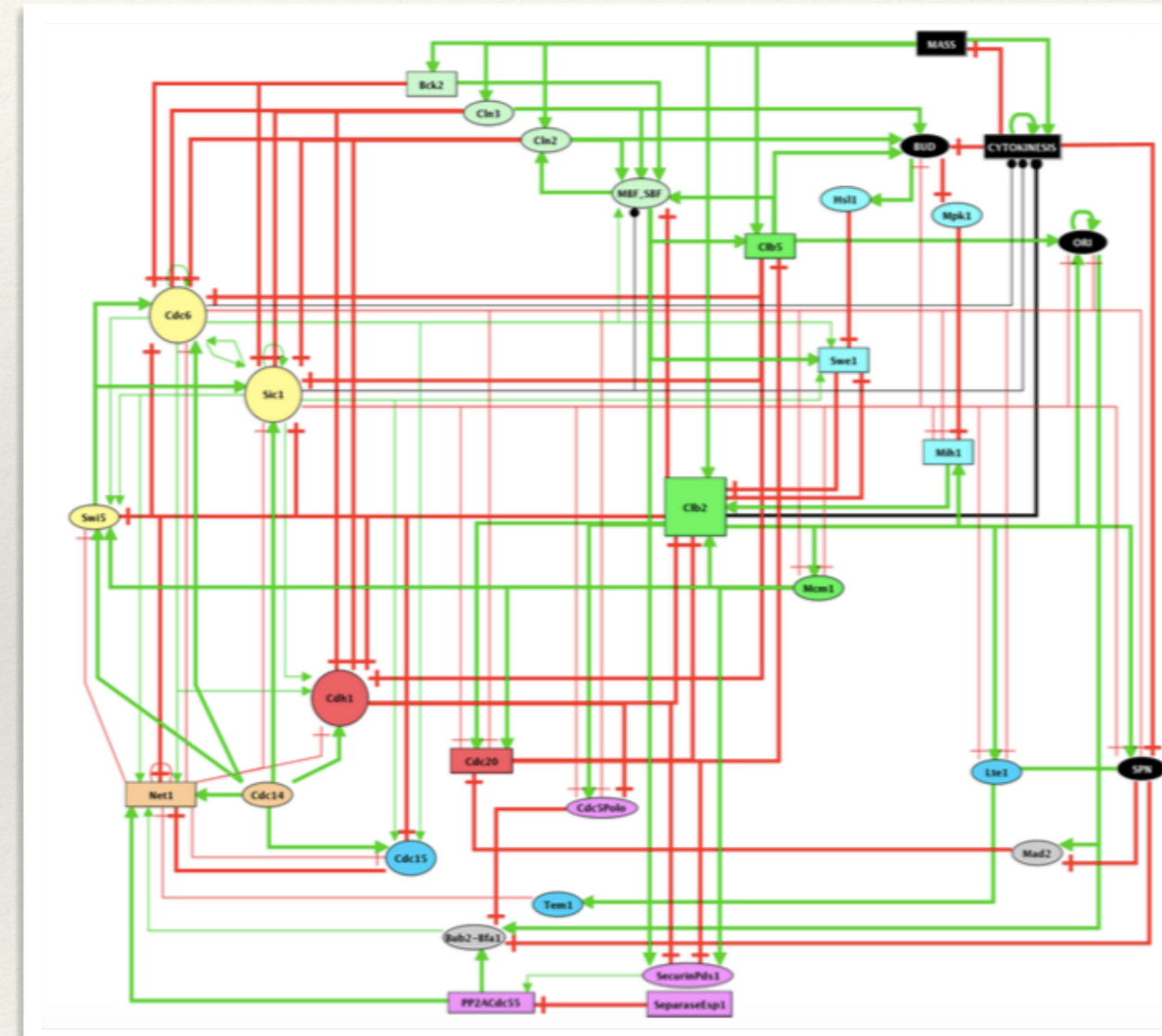


Models: why?



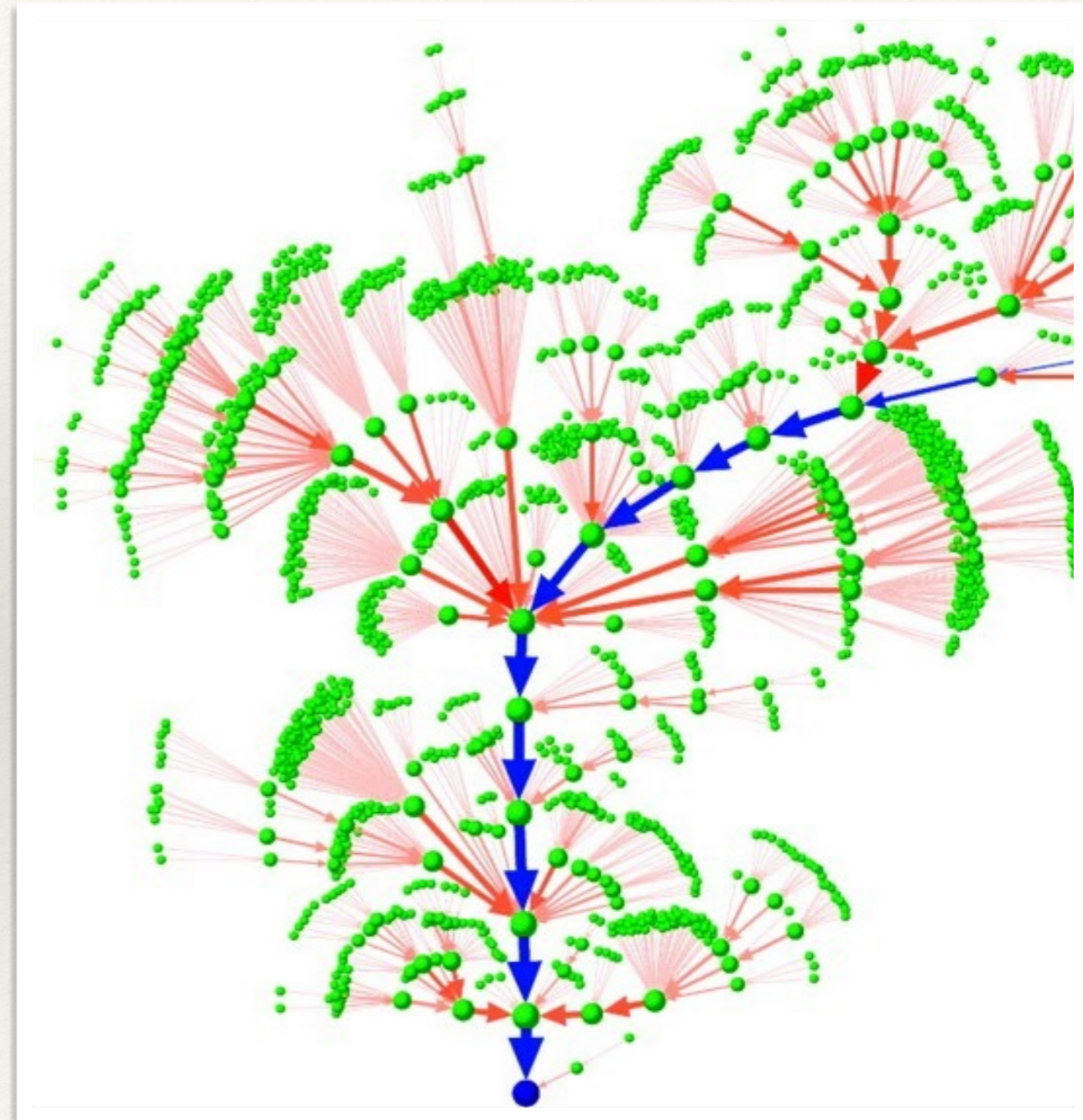
(A) A first model, and (B) a simplified model of the cellular cycle of the budding yeast [LLL+04]

Models: why?



Logical regulatory graph of the budding yeast
[FNL+09]

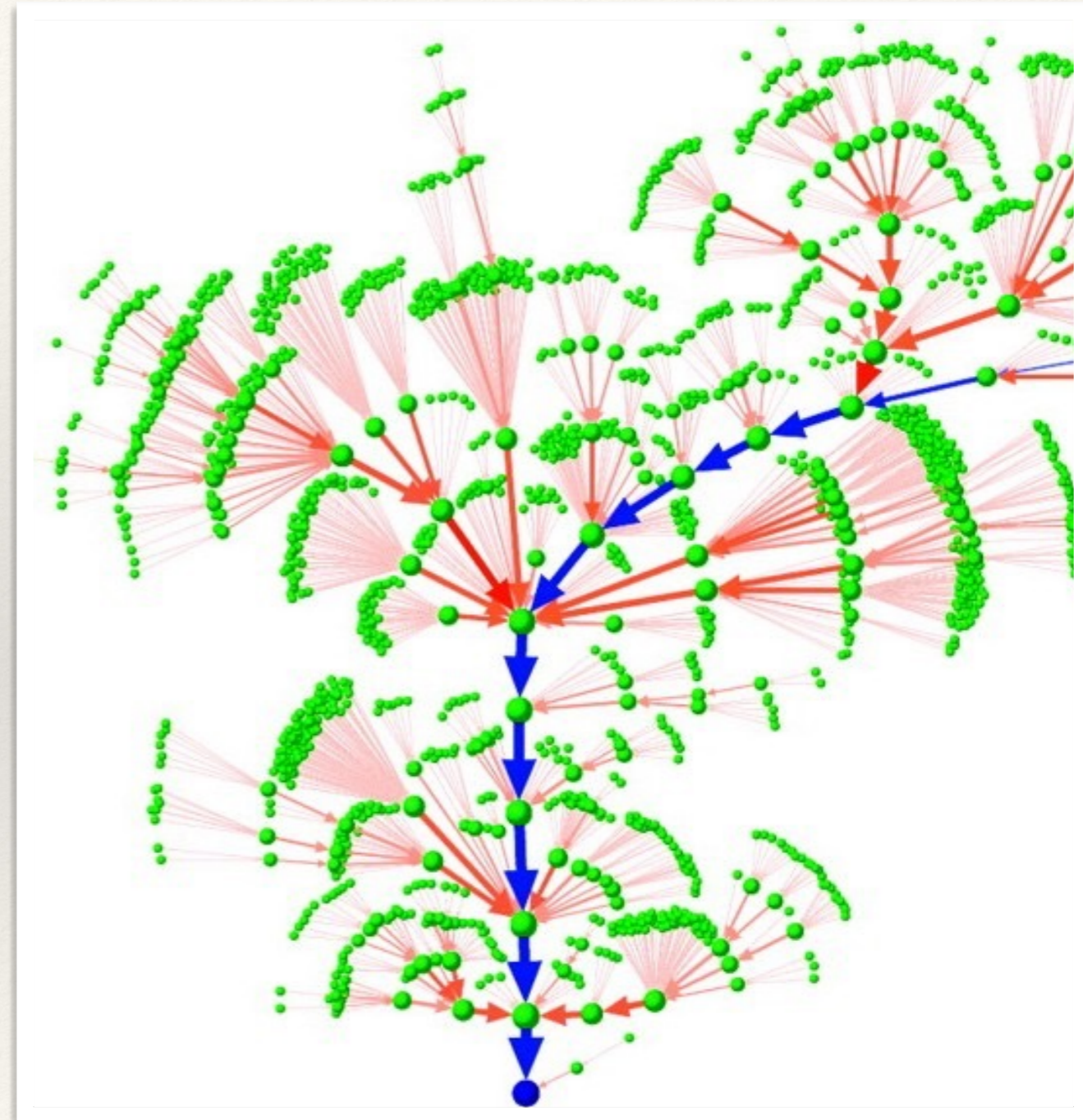
Models: why?



State space of the cellular cycle of the budding yeast
[LLL+04]

Models: why?

- ❖ Objectives:
 - ❖ **Check** whether some expected / unexpected **behaviors** are **possible**
 - ❖ Identify **attractors, oscillations,** etc.
 - ❖ Understand which are the components **at the origin** of some behaviors
- ❖ Problem: how to **circumvent the combinatorial explosion** of the number of behaviors?



State space of the cellular cycle of the budding yeast
[LLL+04]

Modeling the dynamics

Discrete event systems:

$t_1 t_2 t_3 t_1 t_1 t_5 \dots$

- ❖ Modeled through transition systems
- ❖ Capture the **chronology** (the order) of the events

Timed systems:

$(t_1, d_1) (t_2, d_2) (t_3, d_3) (t_1, d_4) (t_1, d_5) (t_5, d_6) \dots$

- ❖ Modeled through timed transition systems
- ❖ Capture the **chronometry** of the events (with their dates)

Scientific challenge

Goals: build predictive dynamic **models** from **time series data**

Our motto: **logical modeling** has its own merits w.r.t. continuous approaches (e.g., ODE)

Underlying questions:

- What are **efficient algorithms** for **discretization** and **learning**?
- What is a *good* predictive model?
- How to validate the **benefits** of our algorithms on "**real-life**" data?

Our scientific challenge

Time series data



Our scientific challenge

Time series data

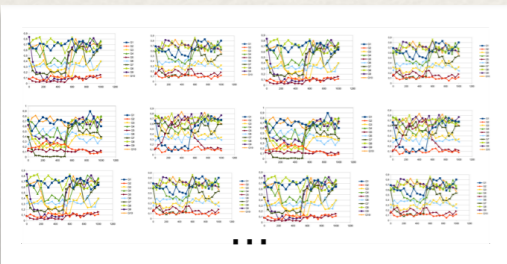


Discretization



Our scientific challenge

Time series data



Discretization

Inductive Learning
[Ino2014]

Our scientific challenge

Time series data



Discretization

Inductive Learning
[Ino2014]

Dynamical models

```

ARNTL(0,T) :- CLOCK(0,T-1).
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Logic programs

Our scientific challenge

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Discretization

Inductive Learning
[Ino2014]

Dynamical models

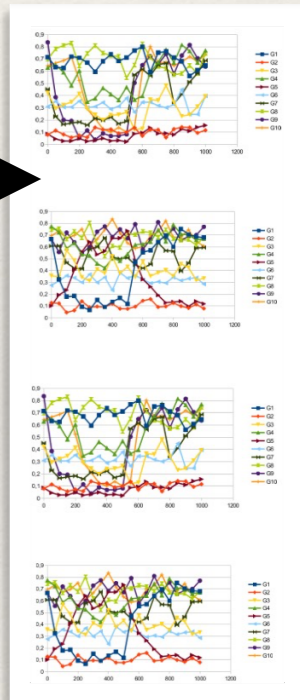
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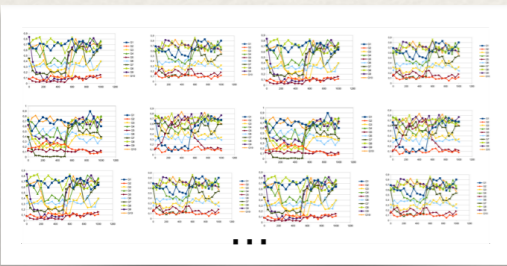
Logic programs

Analysis of
dynamic
behavior



Our scientific challenge

Time series data



Discretization

Inductive Learning
[Ino2014]

Dynamical models

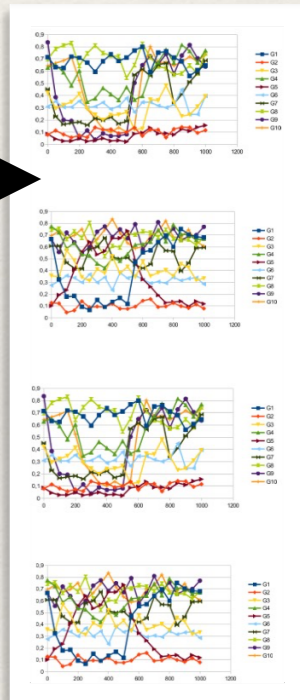
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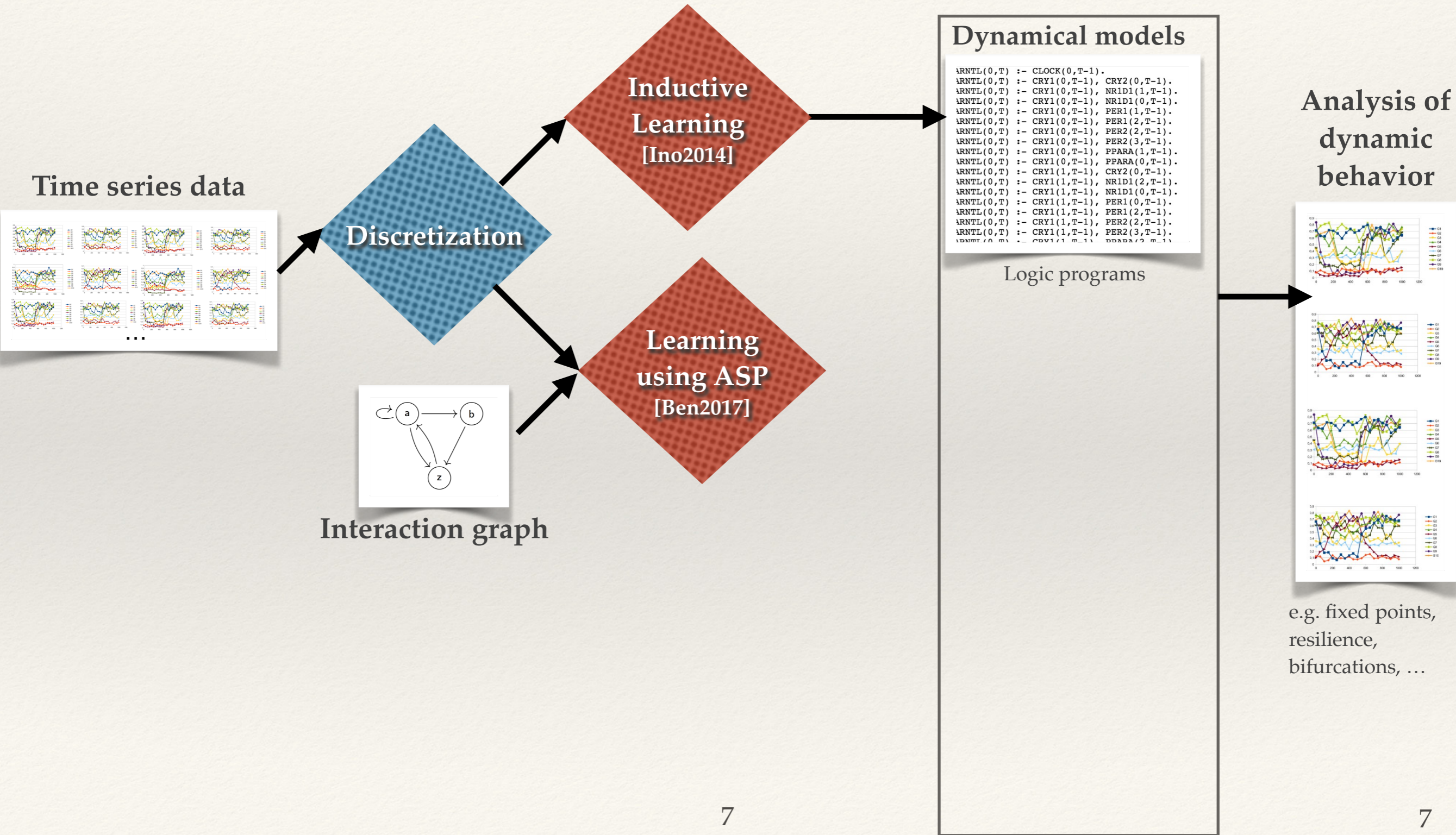
Logic programs

Analysis of
dynamic
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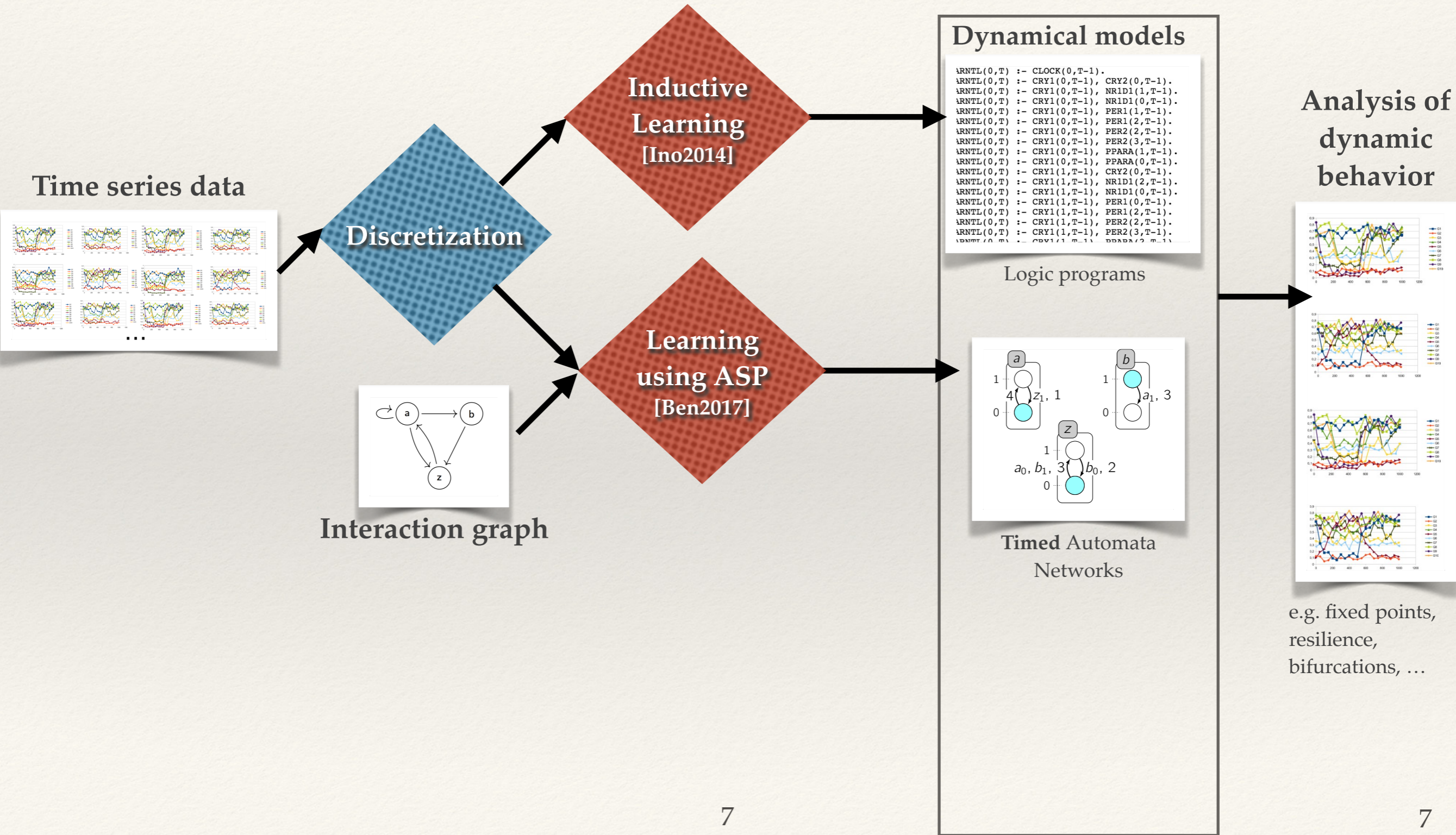


e.g. fixed points,
resilience,
bifurcations, ...

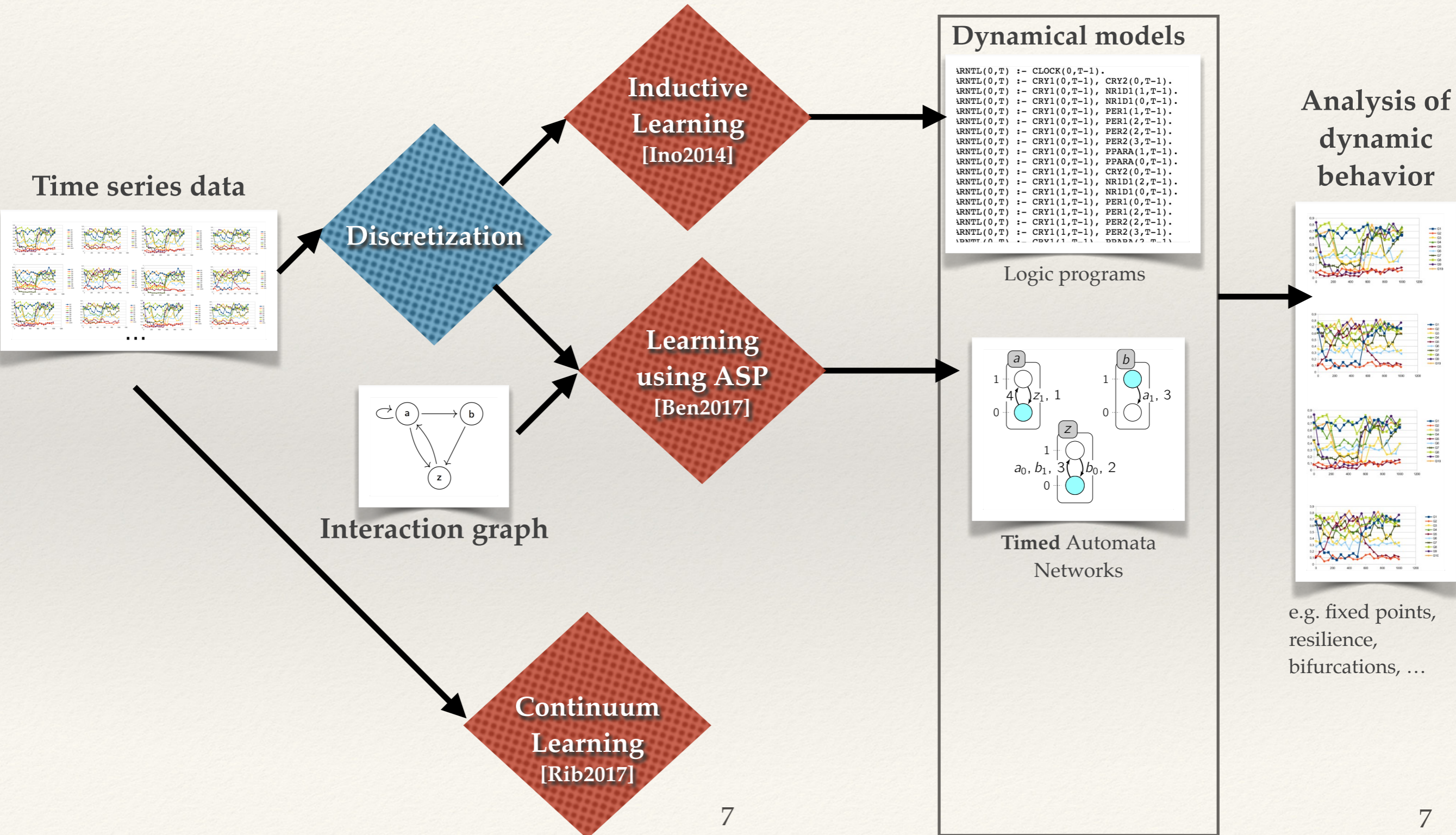
Our scientific challenge



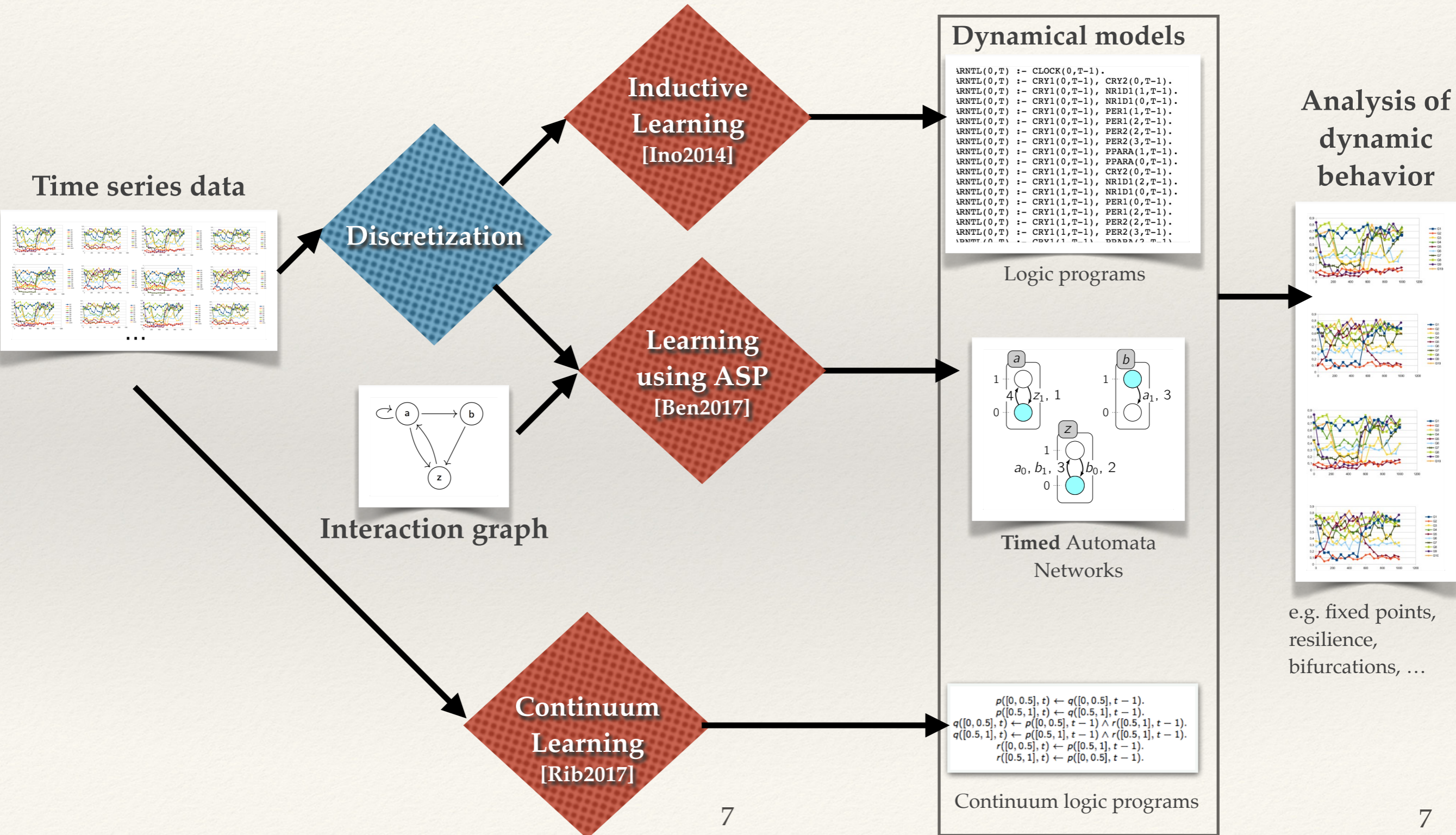
Our scientific challenge



Our scientific challenge

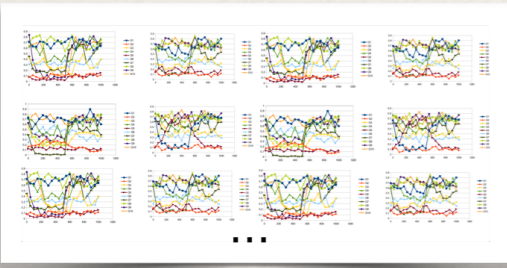


Our scientific challenge

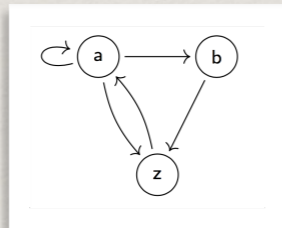


Our scientific challenge

Time series data



Discretization



Interaction graph

Inductive Learning
[Ino2014]

Learning using ASP
[Ben2017]

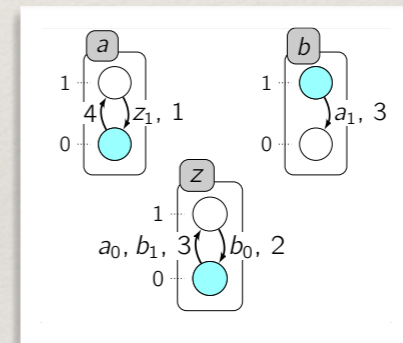
Continuum Learning
[Rib2017]

Dynamical models

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Logic programs



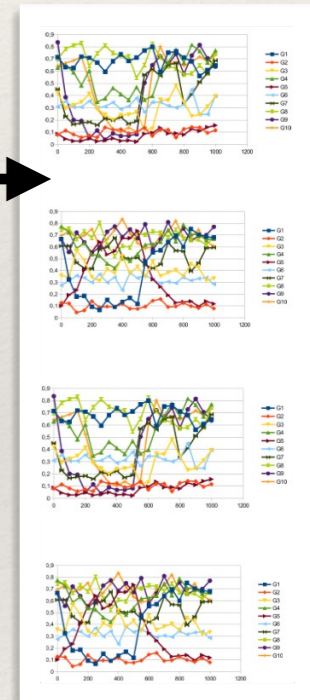
Timed Automata Networks

```

p([0, 0.5], t) ← q([0, 0.5], t - 1).
p([0.5, 1], t) ← q([0.5, 1], t - 1).
q([0, 0.5], t) ← p([0, 0.5], t - 1) ∧ r([0.5, 1], t - 1).
q([0.5, 1], t) ← p([0.5, 1], t - 1) ∧ r([0.5, 1], t - 1).
r([0, 0.5], t) ← p([0.5, 1], t - 1).
r([0.5, 1], t) ← p([0, 0.5], t - 1).
    
```

Continuum logic programs

Analysis of dynamic behavior



e.g. fixed points, resilience, bifurcations, ...

Targets

- ❖ Predictive modeling competitions: DREAM Challenges
- ❖ Collaboration with biologists:
 - ❖ Circadian clock (F. Delaunay)
 - ❖ Radiation oncology (F. Paris)

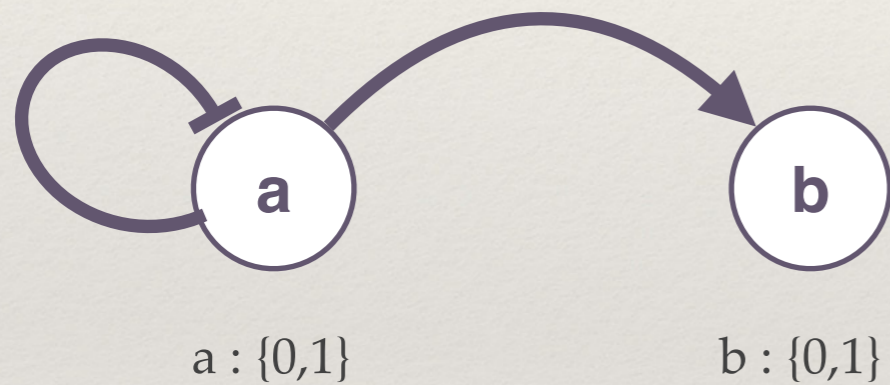
Overview

- ❖ Modeling frameworks for dynamical analysis
- ❖ Analysis of formal models
- ❖ Inference and Learning approaches
- ❖ Application to biological case studies

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- ❖ **Modeling frameworks for dynamical analysis**
- ❖ Analysis of formal models
- ❖ Inference and Learning approaches
- ❖ Application to biological case studies

Principles of abstraction



Example of regulatory Boolean network

Abstract:

- ❖ Components
- ❖ Concentrations
- ❖ Interactions
- ❖ Time
- ❖ Order

Modeling the dynamics

Discrete event systems:

$t_1 t_2 t_3 t_1 t_1 t_5 \dots$

- ❖ Modeled through transition systems
- ❖ Capture the **chronology** (the order) of the events

Automata Networks (a.k.a. Process Hitting), Boolean Networks / Thomas networks, Logic Programs

Timed systems:

$(t_1, d_1) (t_2, d_2) (t_3, d_3) (t_1, d_4) (t_1, d_5) (t_5, d_6) \dots$

- ❖ Modeled through timed transition systems
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Timed Automata Networks

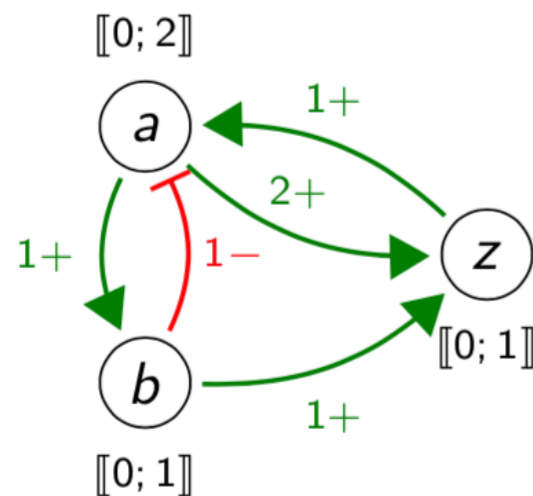
Different modeling frameworks

Discrete Networks / Thomas Modeling

[Kauffman, *Journal of Theoretical Biology*, 1969]

[Thomas, *Journal of Theoretical Biology*, 1973]

- A set of components $N = \{a, b, z\}$
- A discrete domain for each component $\text{dom}(a) = \llbracket 0; 2 \rrbracket$
- Discrete parameters / evolution functions $f^a : \mathcal{S} \rightarrow \text{dom}(a)$
- Signs & thresholds on the edges (redundant) $a \xrightarrow{2+} z$

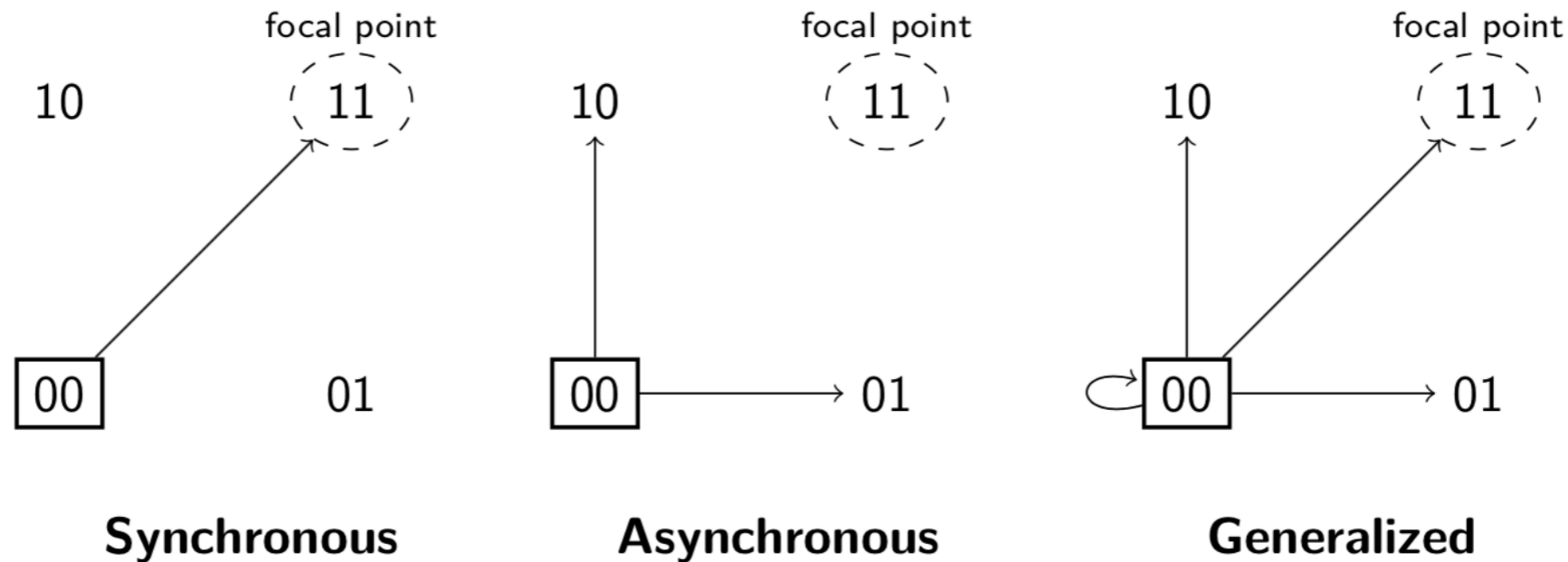


a	f^b	z	b	f^a	a	b	f^z
0	0	0	0	1	0	0	0
1	1	0	1	0	0	1	0
2	1	1	0	1	1	0	0
		1	1	2	1	1	0
					2	0	0
					2	1	1

Semantics = From this information, what are the next possible state(s)?

Choice of an appropriate semantics

GULA: Semantics-Free Learning of a BRN ◦ Discrete Networks



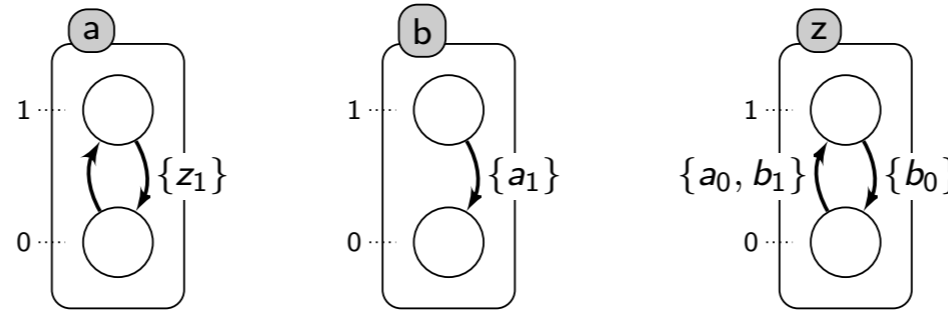
Other semantics:

- Round-robin
- With memory
- With priorities
- ...

Different modeling frameworks

Modeling Delayed Dynamics in Biological Regulatory Networks from Time Series Data

Automata Networks

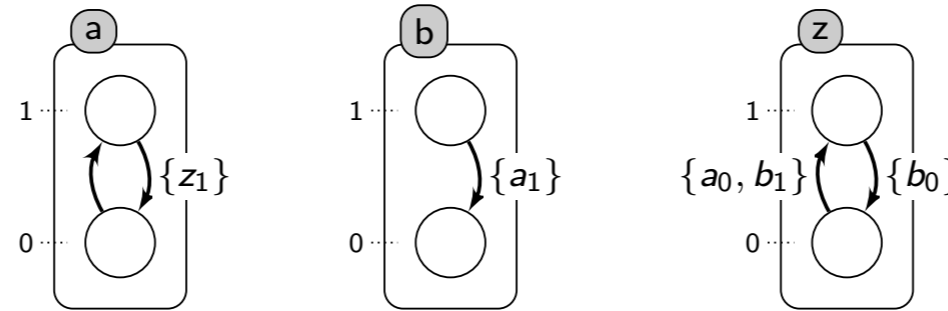


Automata: components a, b, z

Different modeling frameworks

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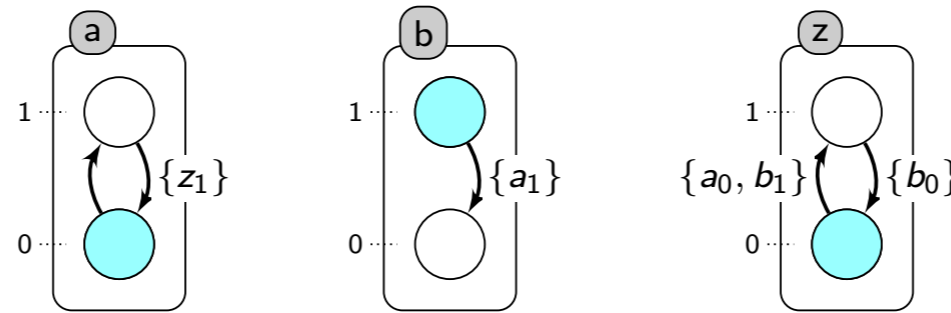
Automata: components a, b, z

Local states: levels of expression $\{a_0, a_1\}$ $\{b_0, b_1\}$ $\{z_0, z_1\}$

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Automata: components a, b, z

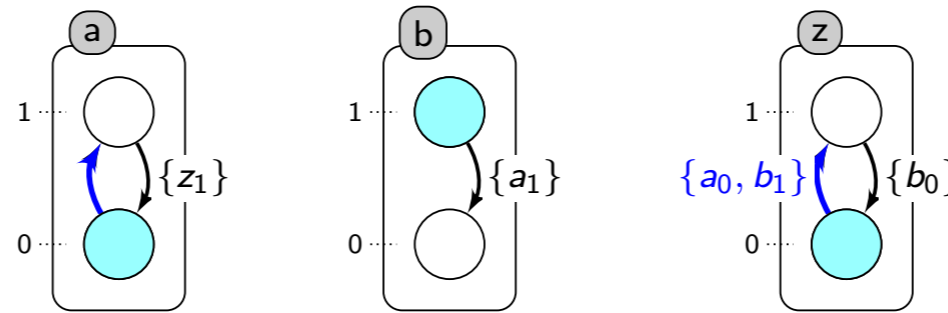
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State: set of active local states $\langle a_0, b_1, z_0 \rangle \sim 010$

Different modeling frameworks

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Local transitions:

$$\tau_1 = a_0 \longrightarrow a_1; \tau_3 = a_1 \xrightarrow{\{z_1\}} a_0; \tau_4 = b_1 \xrightarrow{\{a_1\}} b_0; \tau_2 = z_0 \xrightarrow{\{a_0, b_1\}} z_1; \tau_5 = z_1 \xrightarrow{\{b_0\}} z_0;$$

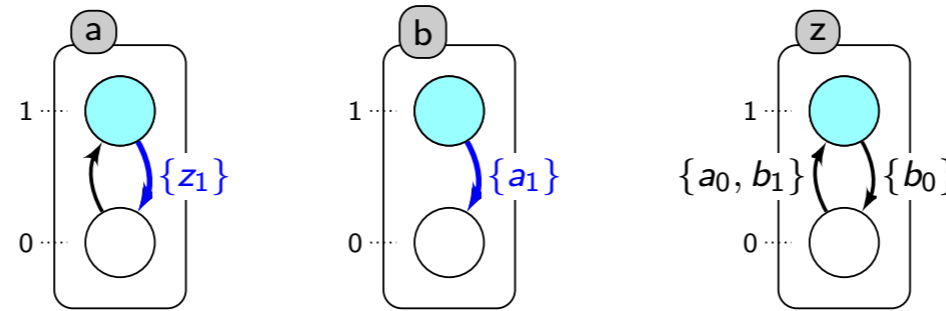
Synchronous : [Kauffman, 1969]

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Different modeling frameworks

Modeling Delayed Dynamics in Biological Regulatory Networks from Time Series Data

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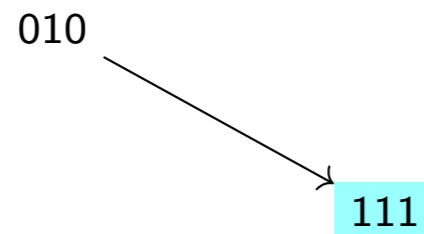
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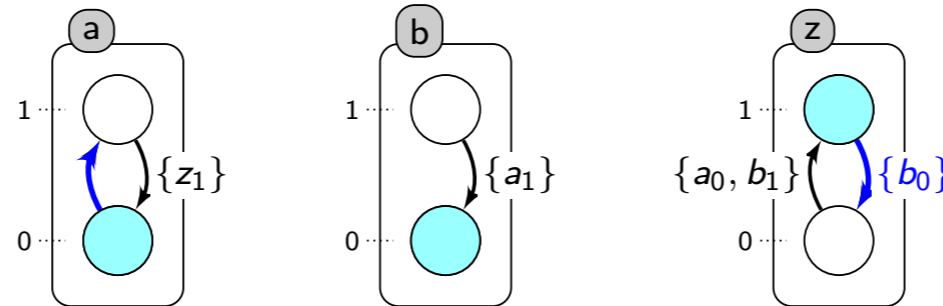
Synchronous : [Kauffman, 1969]



Different modeling frameworks

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Automata Networks



Automata: components a, b, z

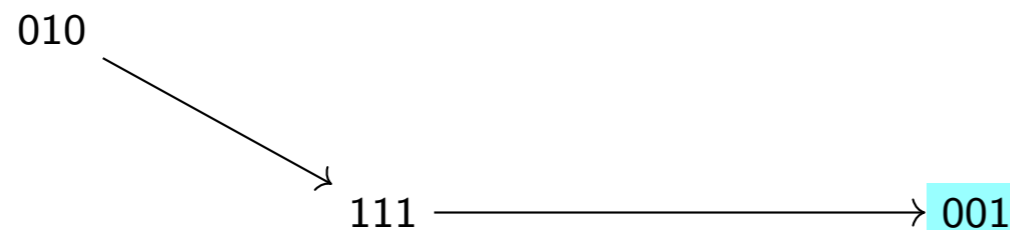
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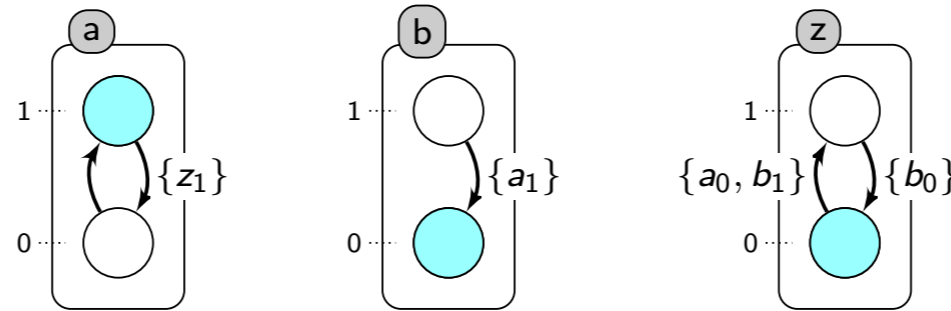
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Different modeling frameworks

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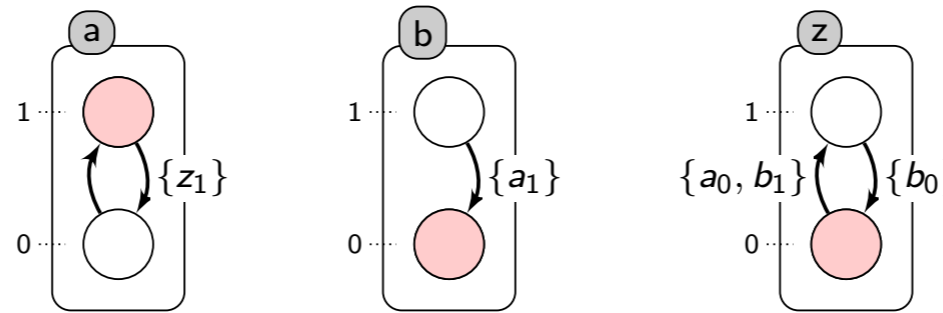
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Automata: components a, b, z

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State: set of active local states $\langle a_1, b_0, z_0 \rangle \sim 100$

Local transitions:

$$\tau_1 = a_0 \longrightarrow a_1; \tau_3 = a_1 \xrightarrow{\{z_1\}} a_0; \tau_4 = b_1 \xrightarrow{\{a_1\}} b_0; \tau_2 = z_0 \xrightarrow{\{a_0, b_1\}} z_1; \tau_5 = z_1 \xrightarrow{\{b_0\}} z_0;$$

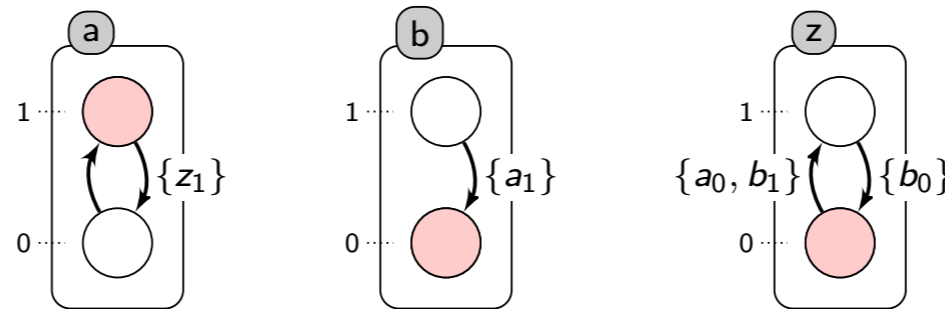
Synchronous : [Kauffman, 1969]



Different modeling frameworks

Modeling Delayed Dynamics in Biological Regulatory Networks from Time Series Data

Automata Networks



Automata: components a, b, z

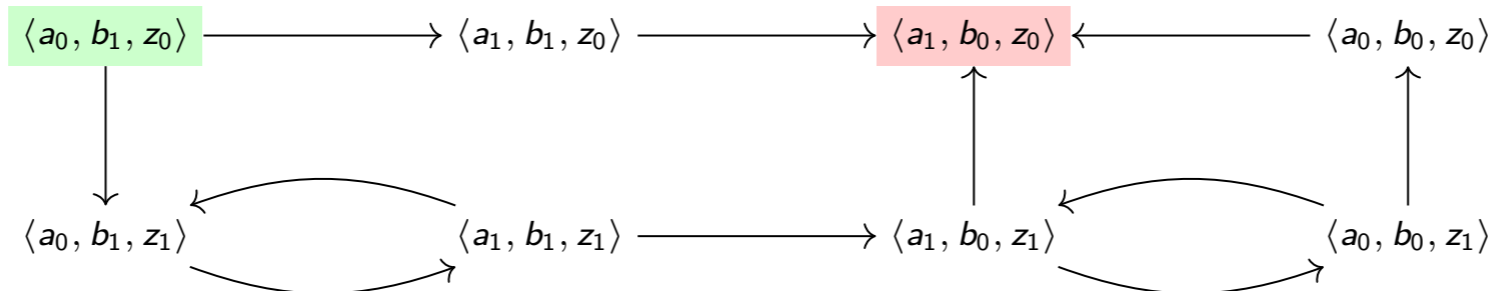
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Asynchrone: [Thomas, 1978]

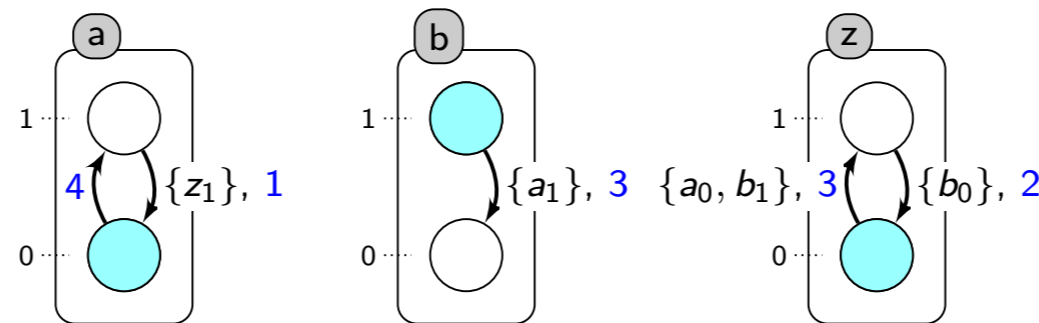


Different modeling frameworks

Modeling Delayed Dynamics in Biological Regulatory Networks from Time Series Data

Timed Automata Networks

Definition



Refinement of th AN dynamics by delays integration : $\delta \in \mathbb{N}$

each delay δ is **specific** of a local transition τ

and δ represents **the necessary time duration** of the local transition τ to occur.

Example:

$b_1 \xrightarrow[3]{\{a_1\}} b_0$: **3** represents the time it takes for **changing** b from the local level 1 to 0.

⇒ How to calculate the states graph of **the refined dynamics** of a T-AN ?

⇒ What would be **the semantics of the dynamics** of a T-AN that shows this refinement?

Logic programs

$$\underbrace{x_0^{val_0}(t)}_{\text{head}} \leftarrow \underbrace{x_1^{val_1}(t-1) \wedge x_2^{val_2}(t-1) \wedge \dots \wedge x_n^{val_n}(t-1)}_{\text{body}}.$$

Where:

$x_0, x_1, x_2, \dots, x_n:$

Variables

$val_0, val_1, val_2, \dots, val_n:$

Values

$t:$

Time step

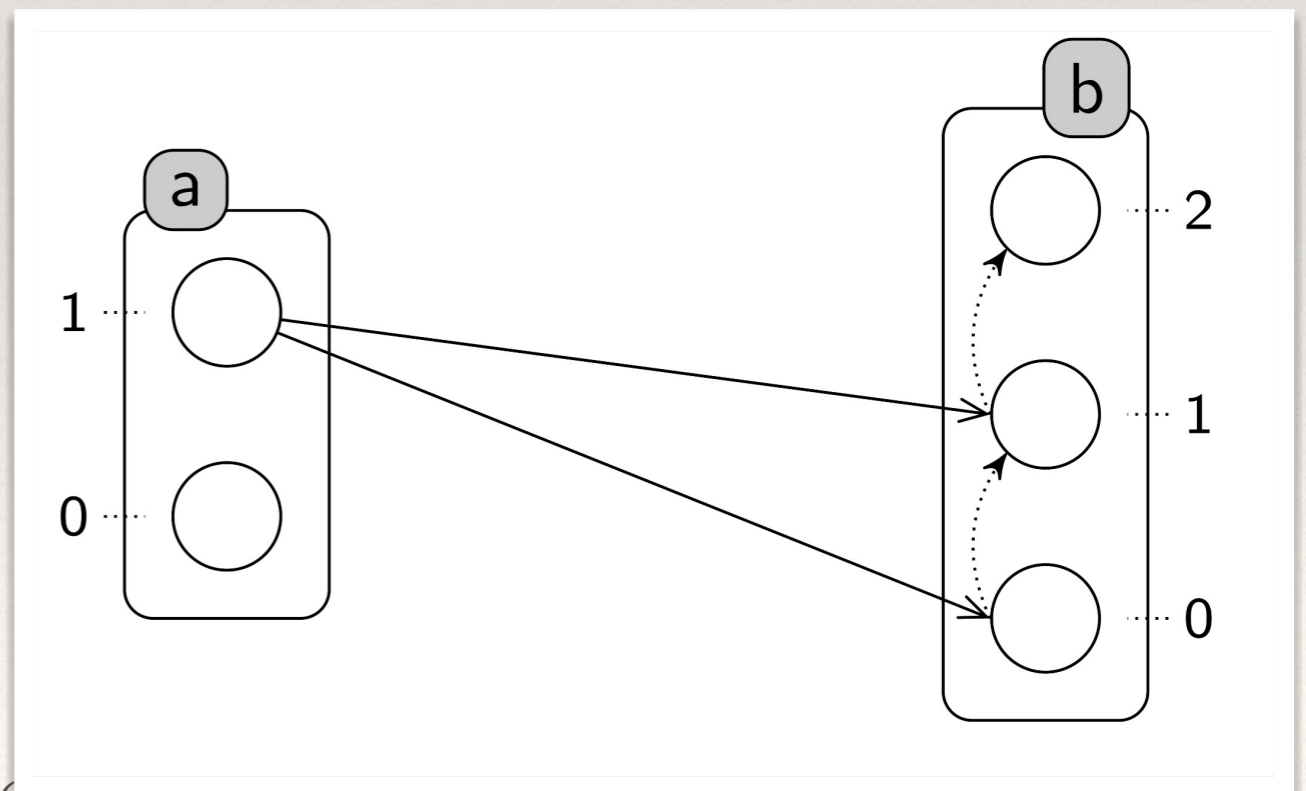
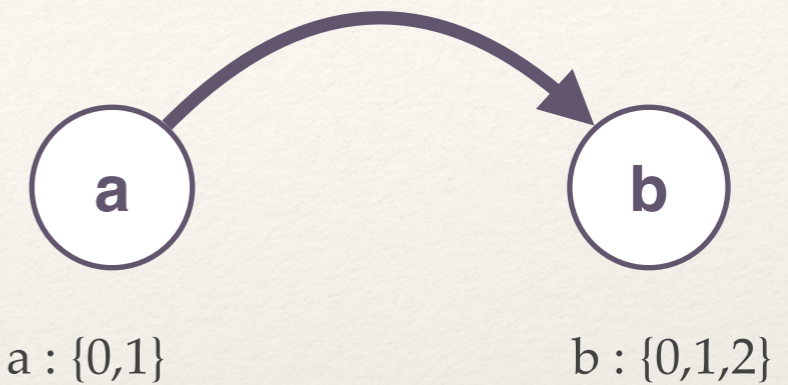
\leftarrow

Reverse implication

All atoms in the body are in conjunction

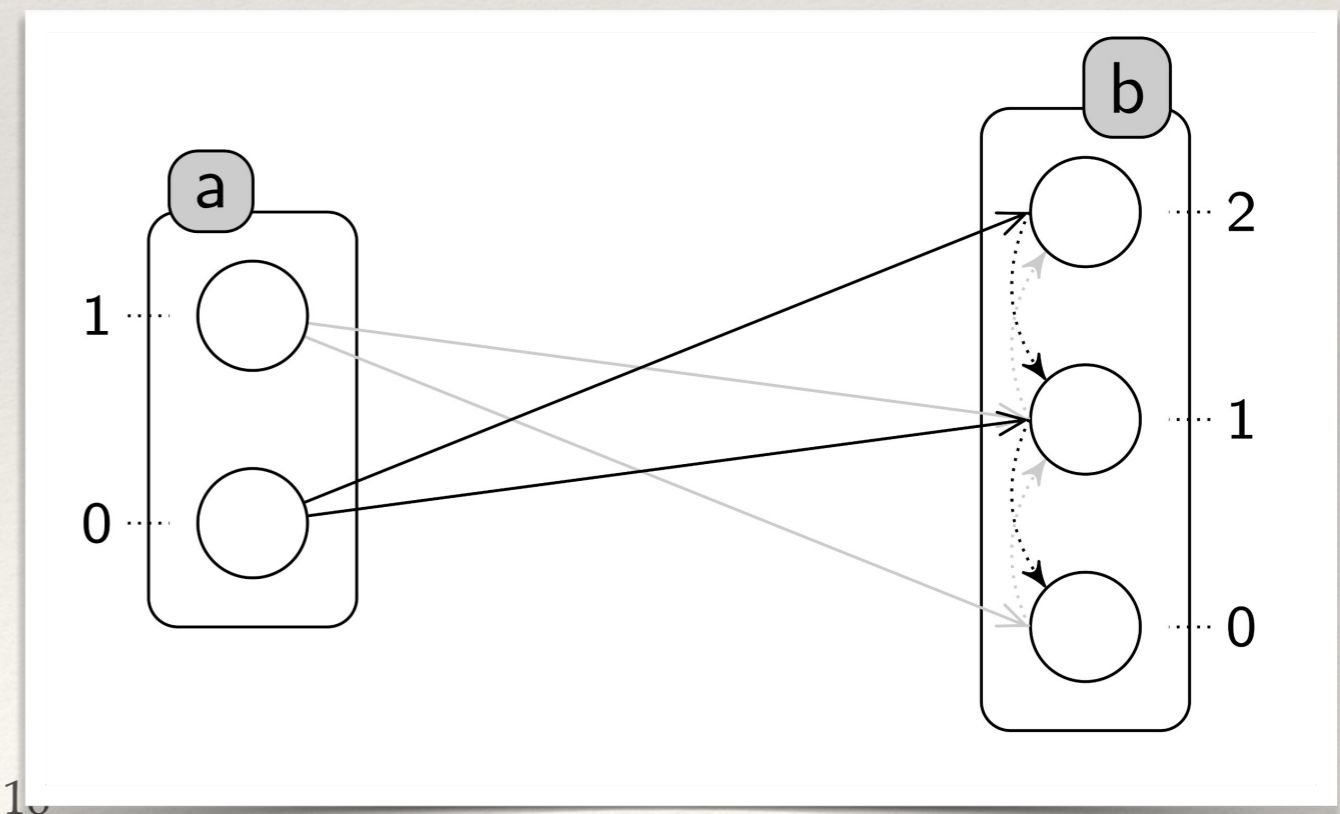
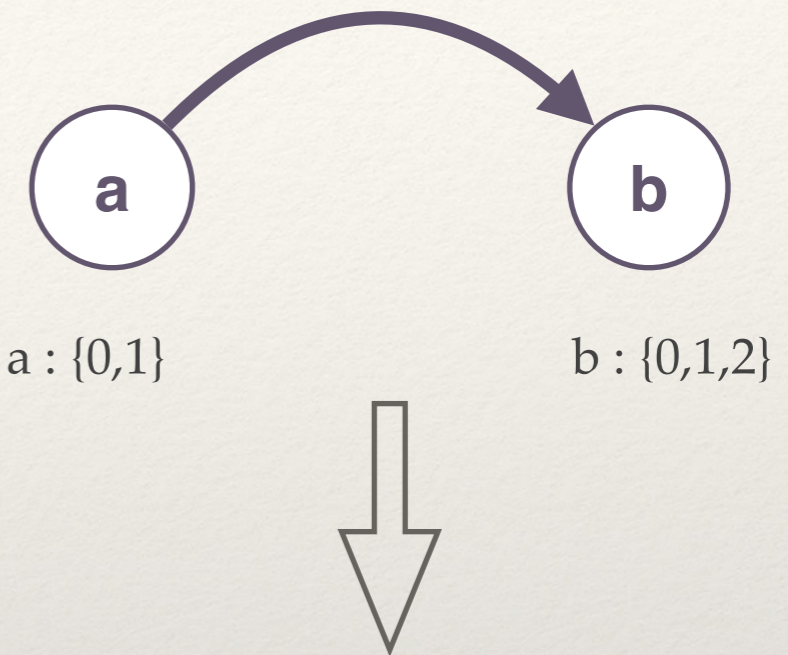
Hypotheses for translation from biology

- ❖ Discretization of the **concentration (expression)** of a component: Boolean or multi-valued
- ❖ Discretization of **time**: event-bases
- ❖ **Unitarian** dynamics
- ❖ Semantics: **non-deterministic asynchronous**



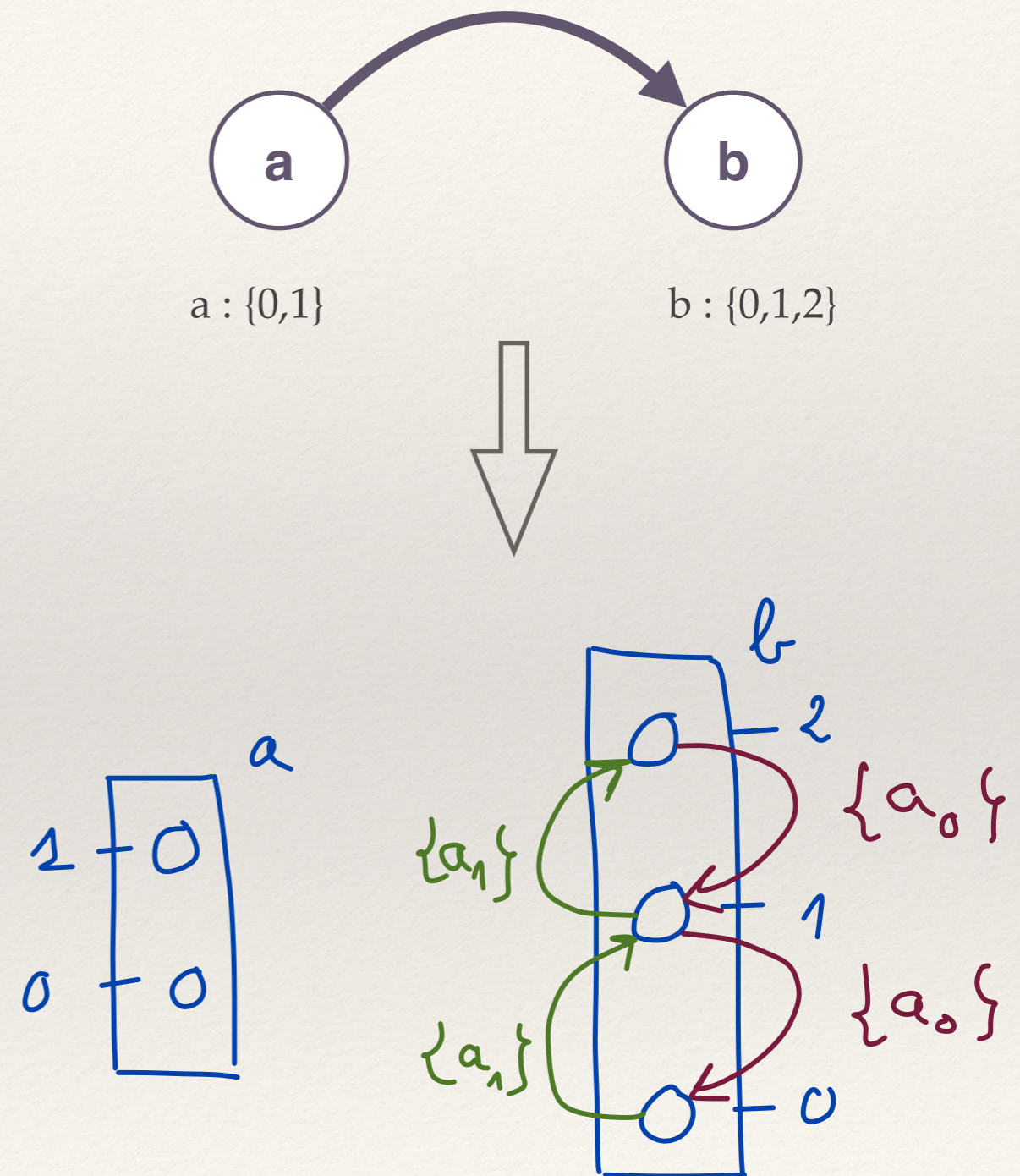
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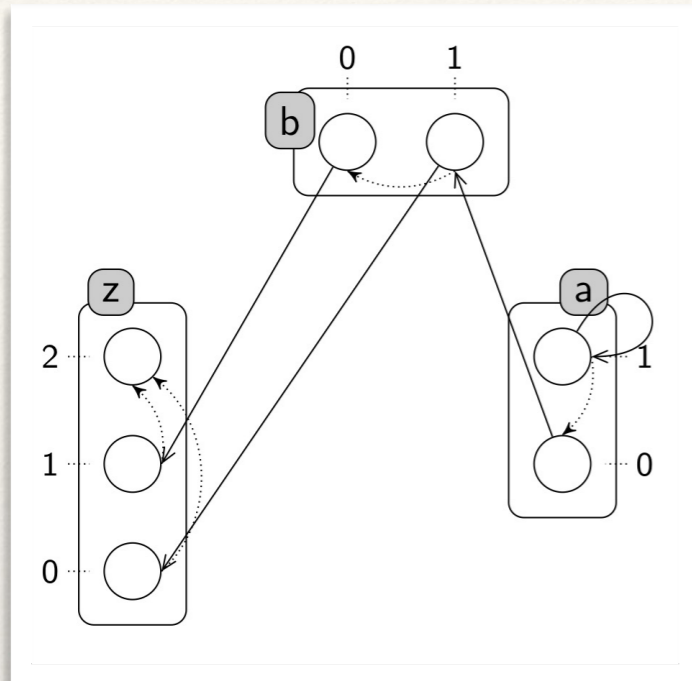


Overview

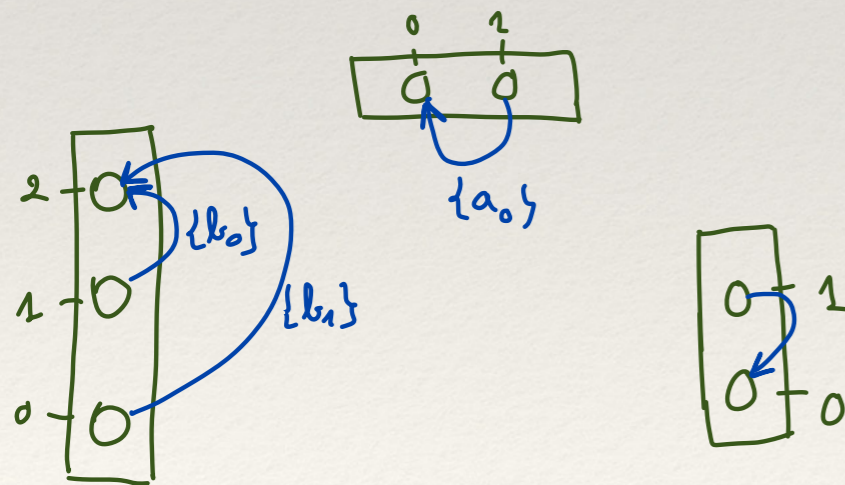
- ❖ Modeling frameworks for dynamical analysis
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- ❖ Inference and Learning approaches
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An example of static analysis

[PMR11a,PCF+14]



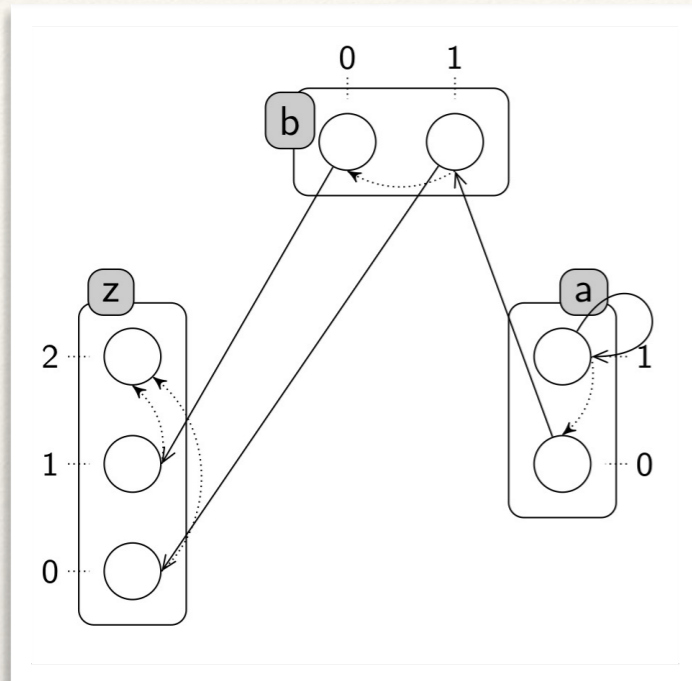
Process Hitting framework



Asynchronous automata network

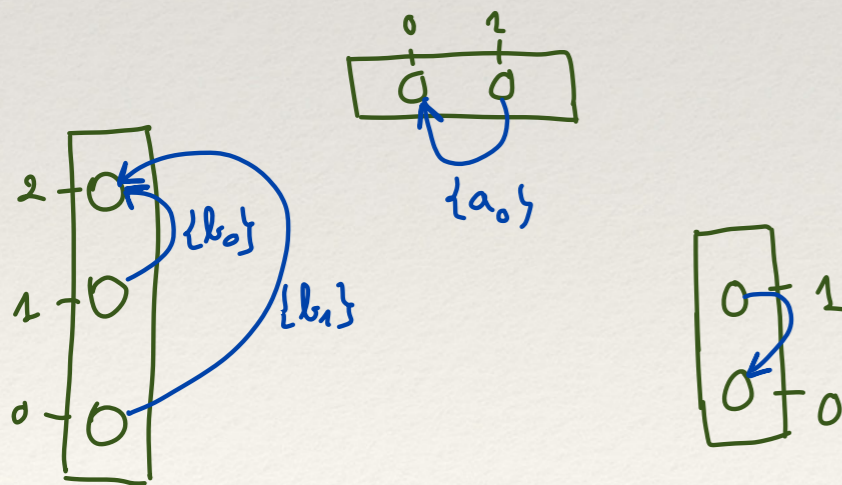
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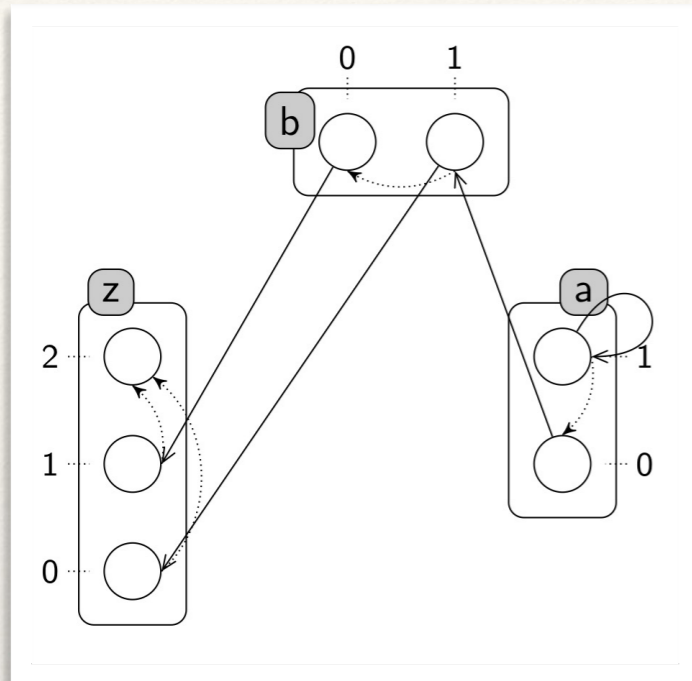
What are the fixed points?



Asynchronous automata network

An example of static analysis

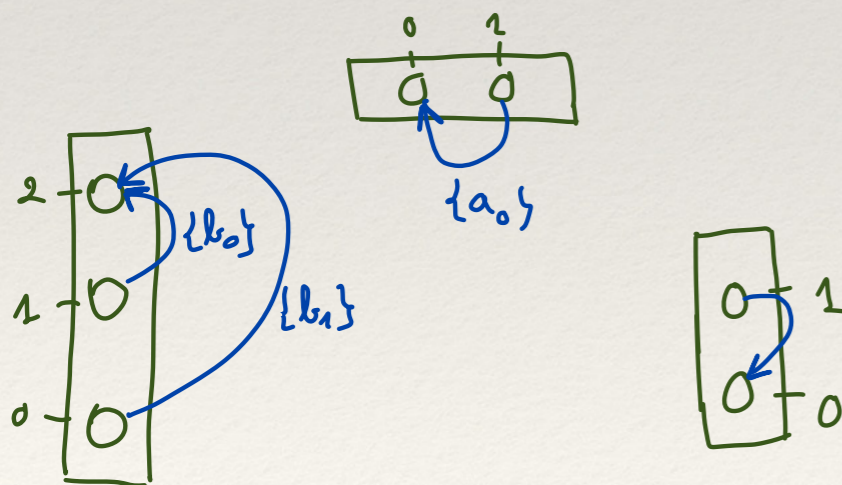
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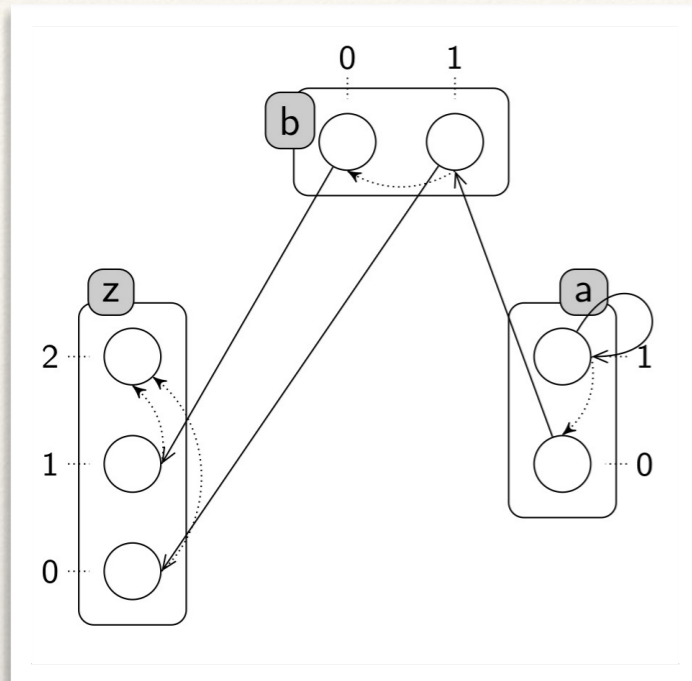
- ❖ Naive method: compute all possible scenarios



Asynchronous automata network

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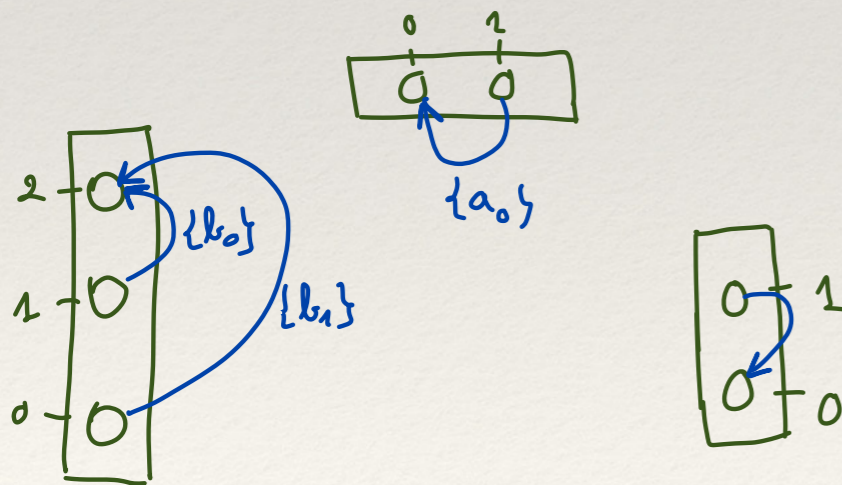
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Process Hitting framework

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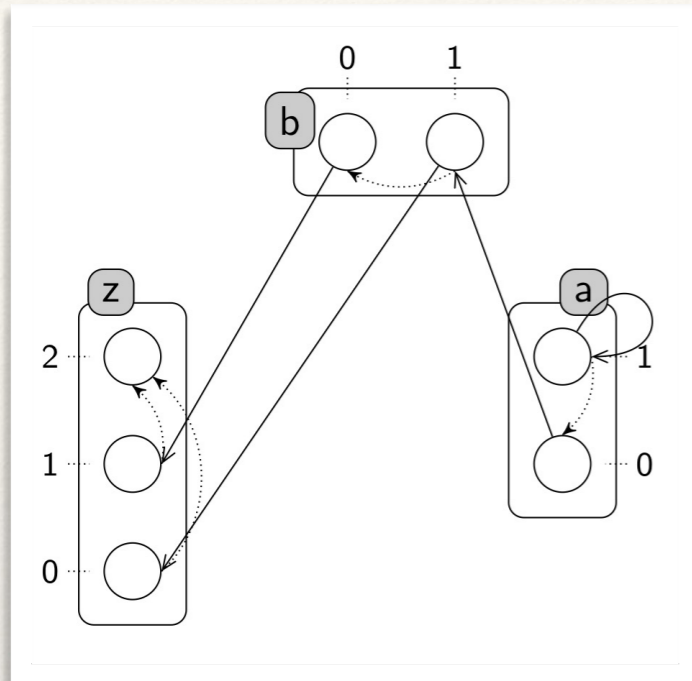
- ❖ Our proposition: static analysis



Asynchronous automata network

An example of static analysis

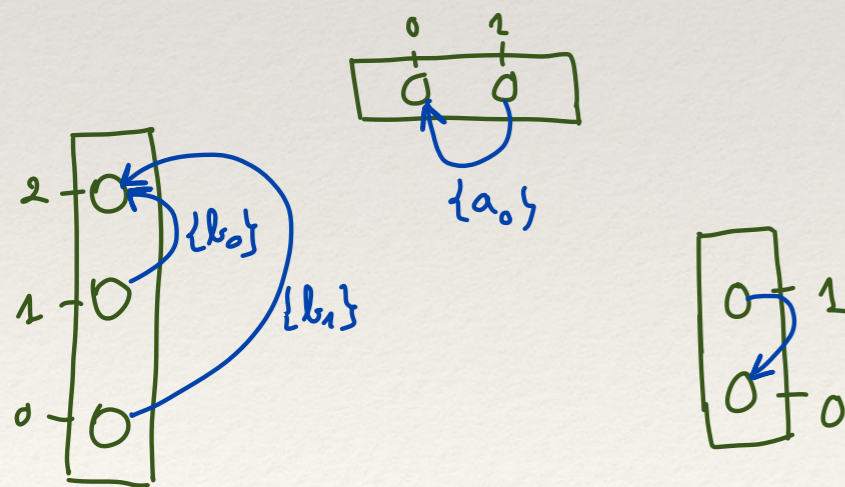
[PMR11a,PCF+14]



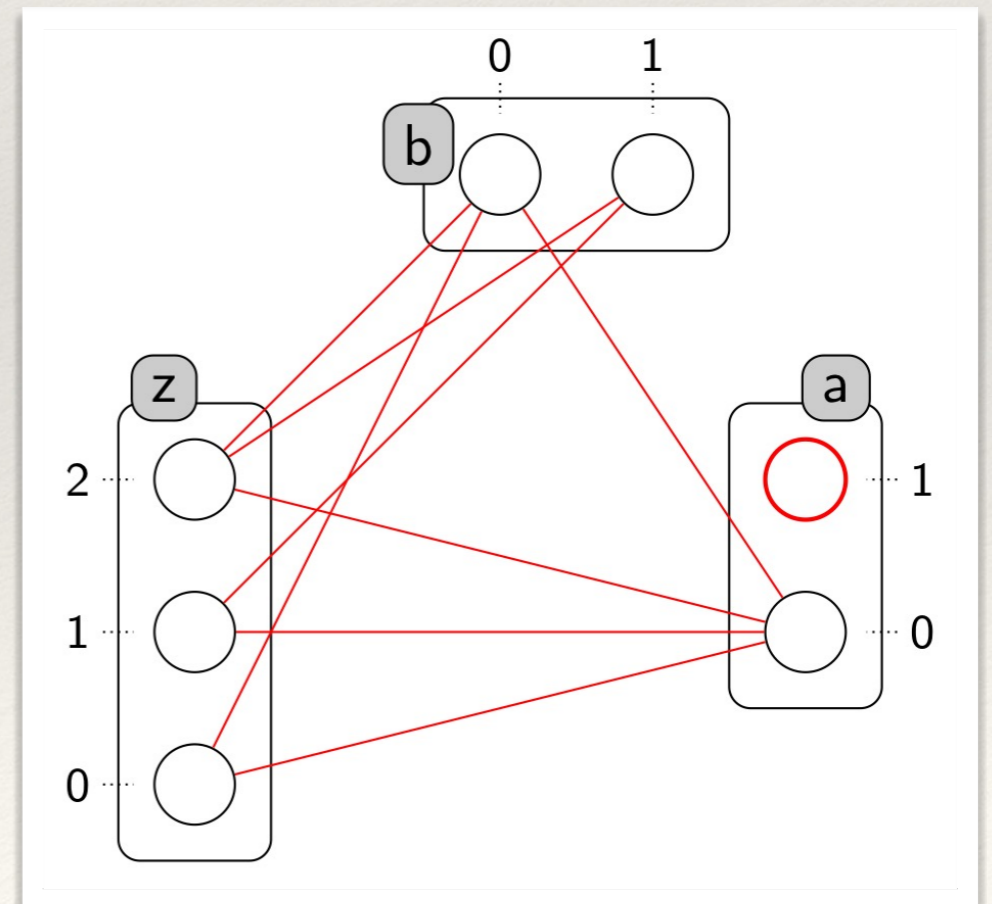
Process Hitting framework

What are the fixed points?

- ❖ Our proposition: static analysis
- ❖ Compute hitless graph



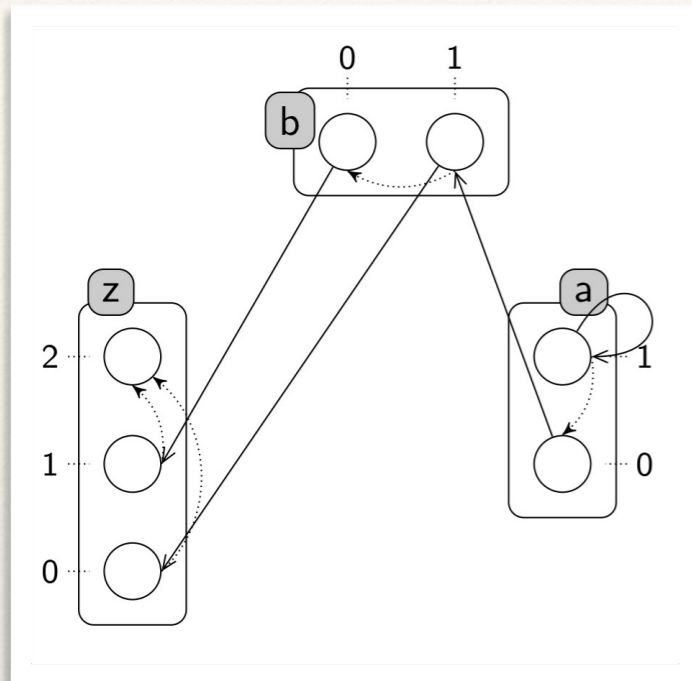
Asynchronous automata network



Hitless graph

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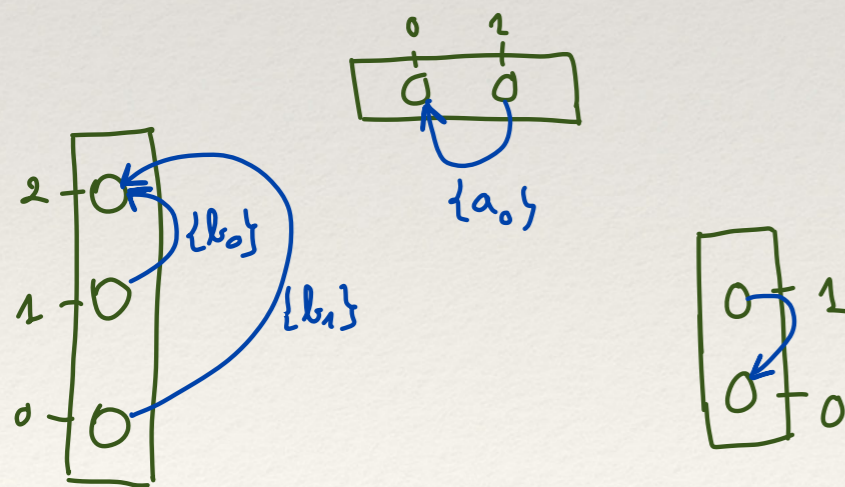
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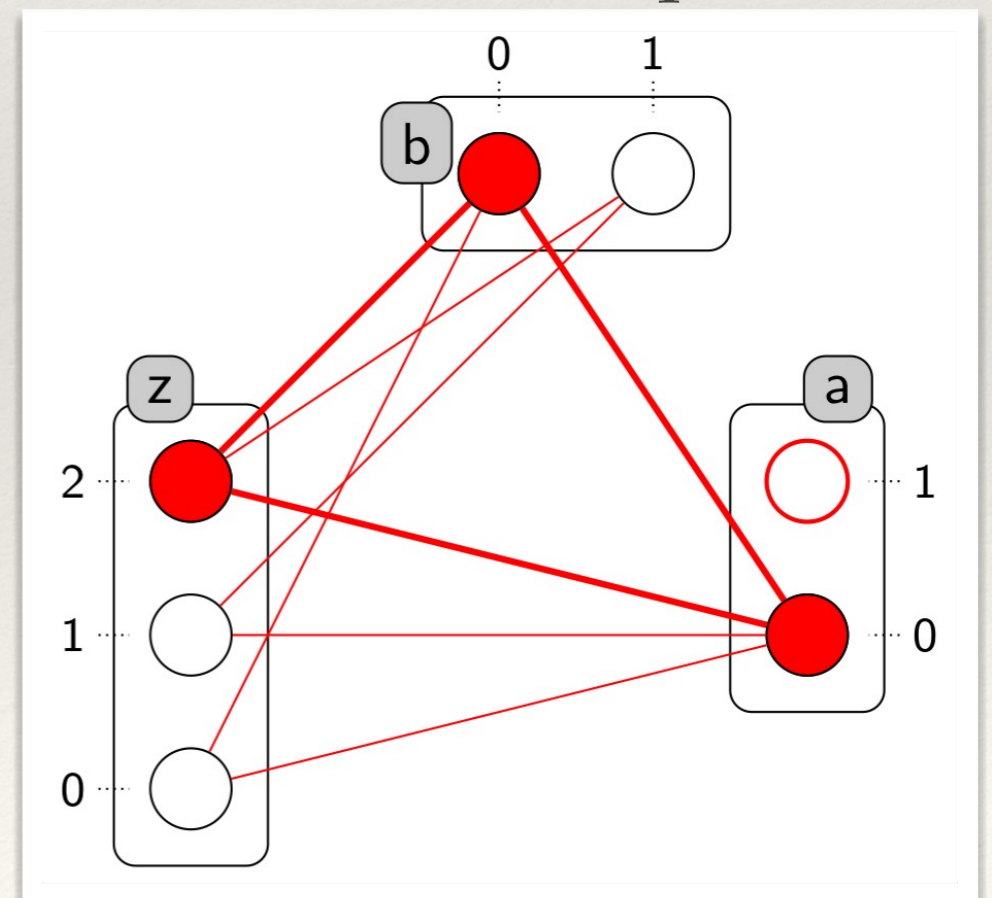
Process Hitting framework

What are the fixed points?

- ❖ Our proposition: static analysis
- ❖ Compute hitless graph
- ❖ Enumerate its n -cliques



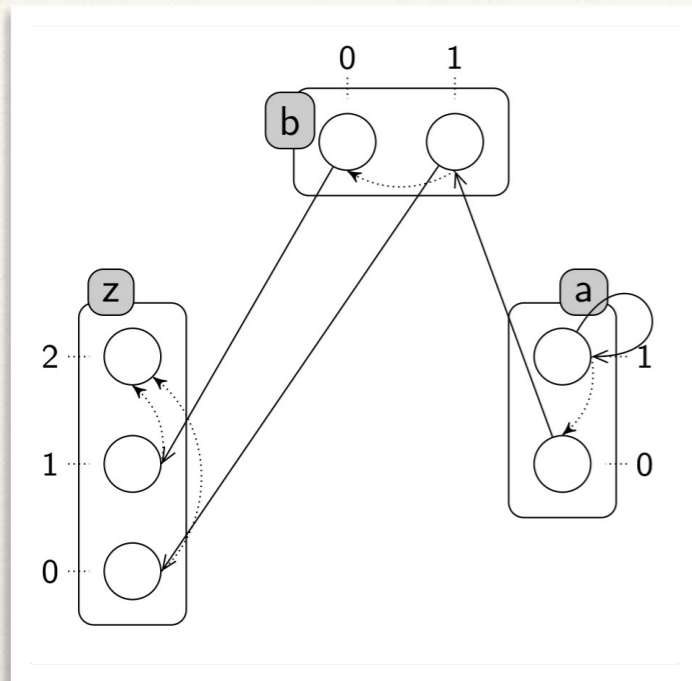
Asynchronous automata network



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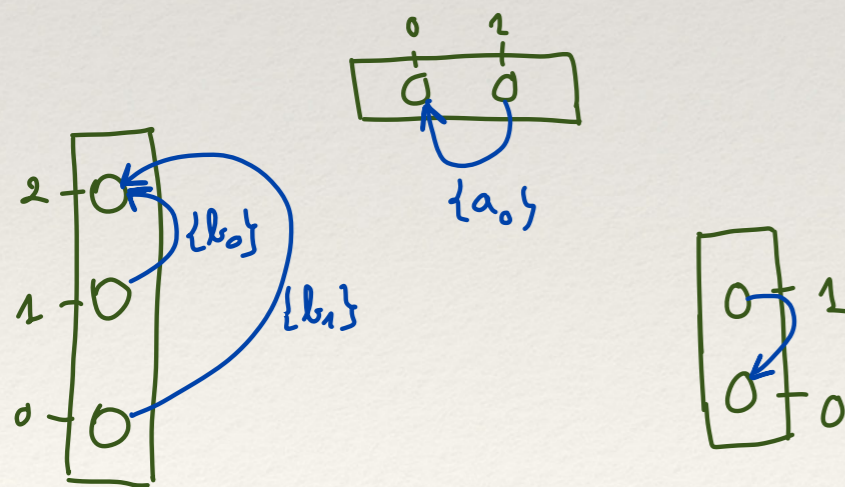
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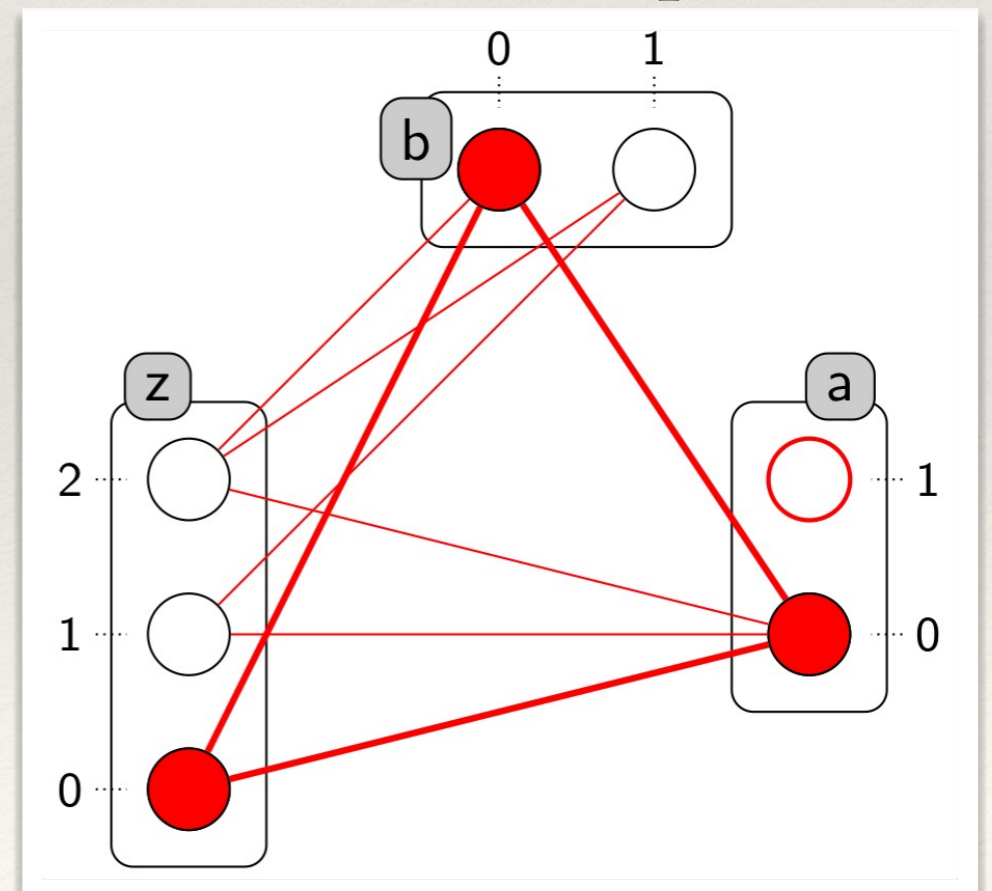
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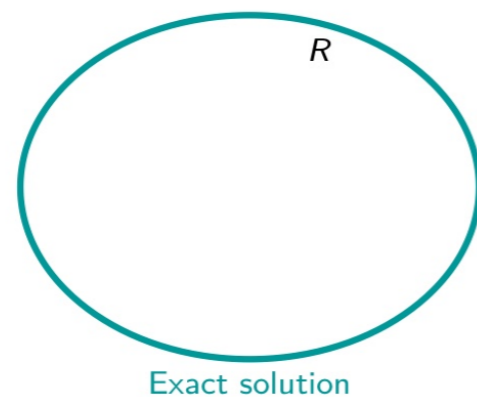
Hitless graph

Reachability analysis [PMR12]

Goal: check properties of the following type

« From an initial state S_0 , can we reach a state S_n where a_i is active? »

Our proposition: **under-approximation** P and **over-approximation** Q of the dynamics s.t. $P \Rightarrow R \Rightarrow Q$

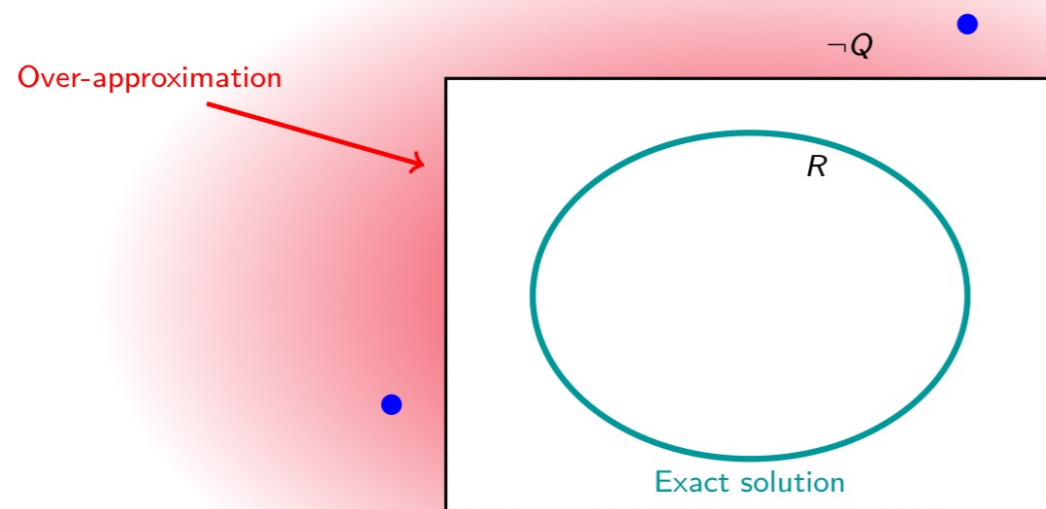


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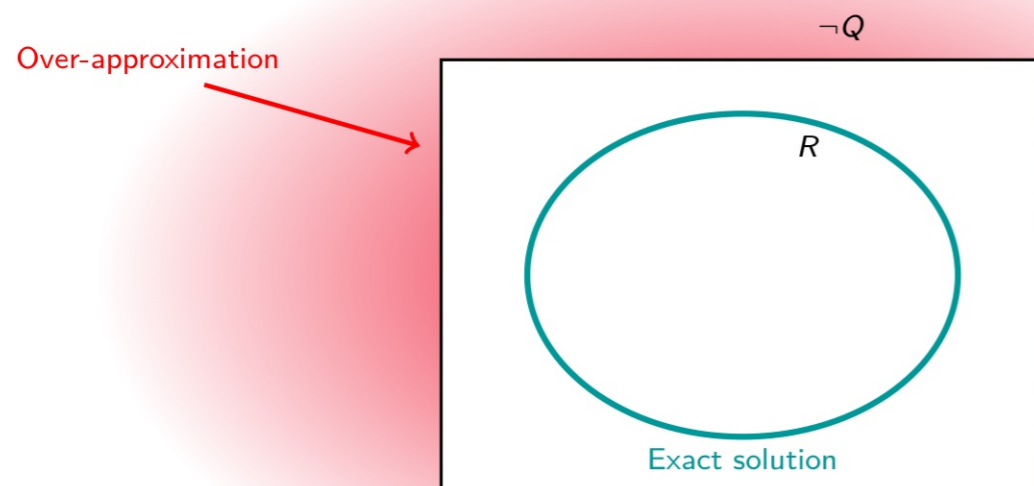


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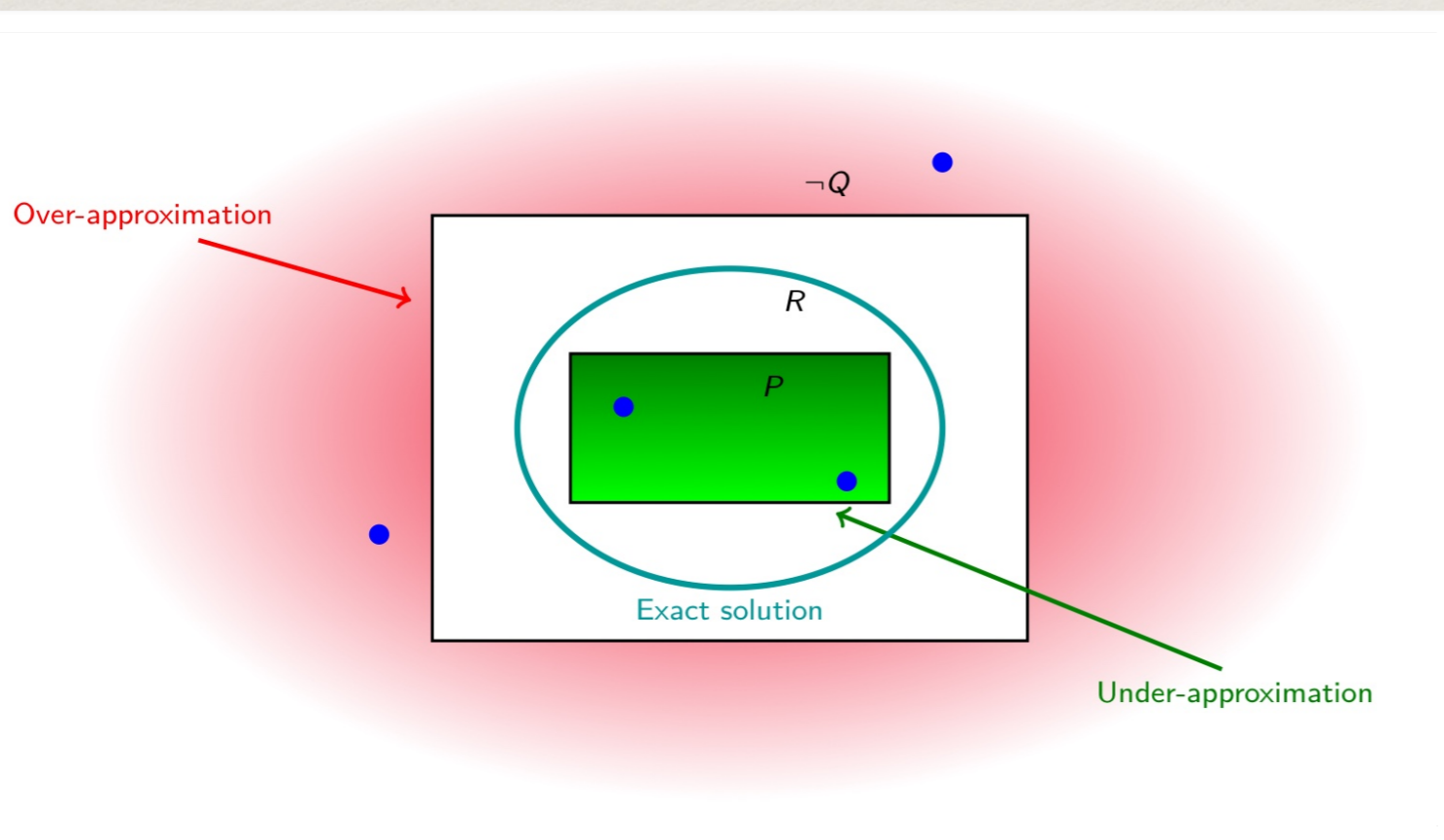


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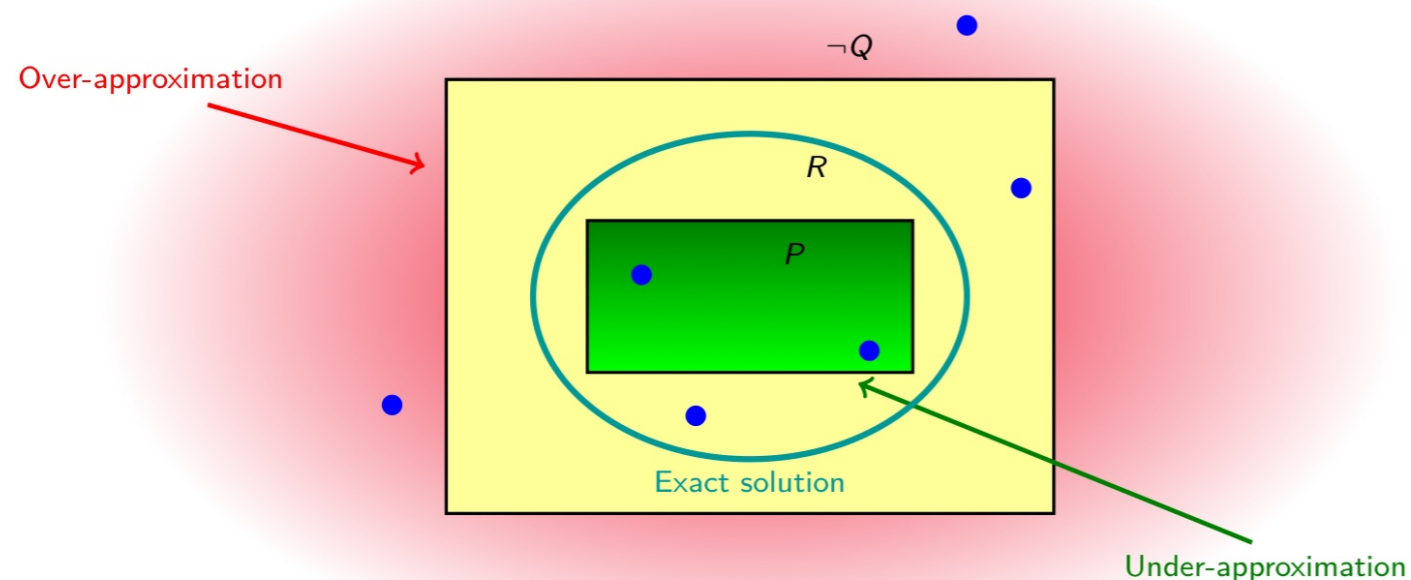


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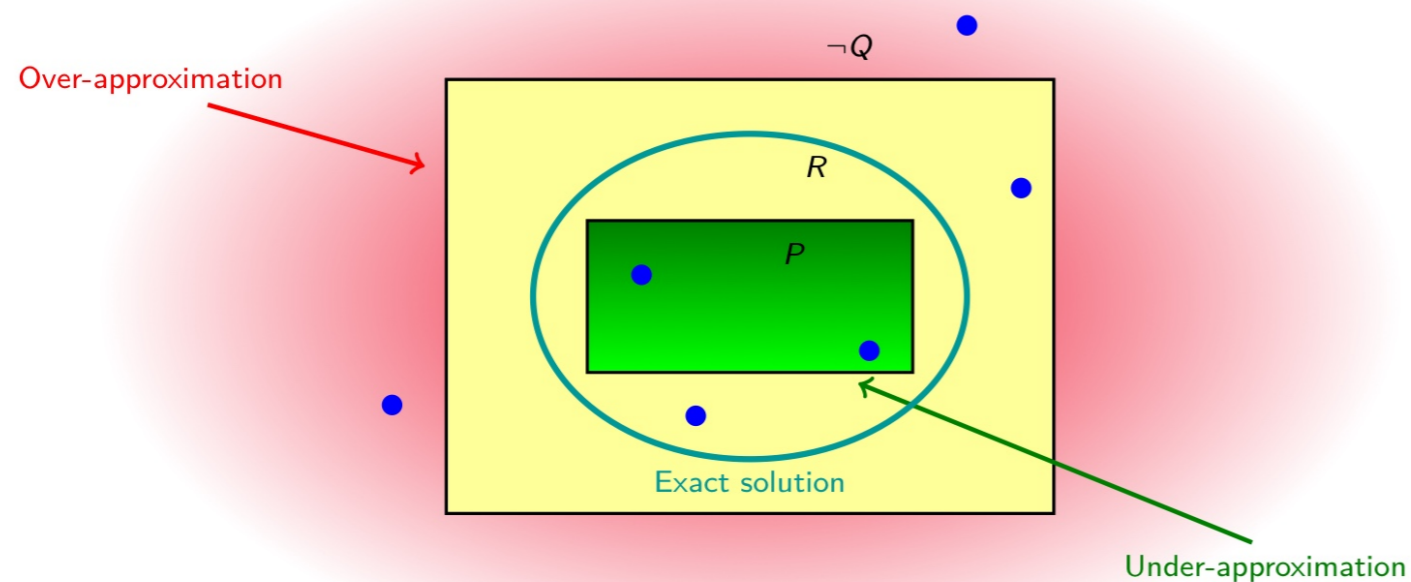


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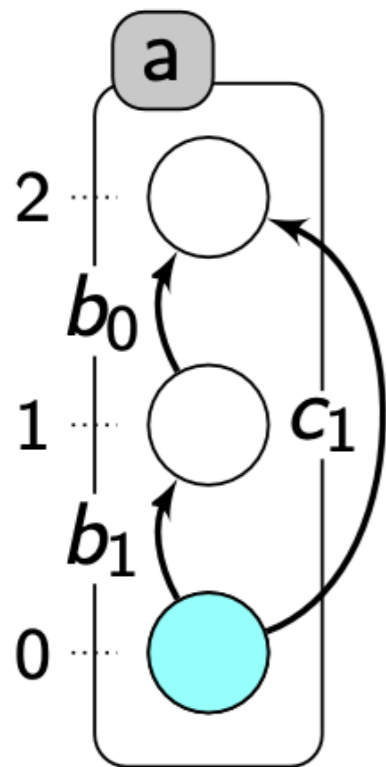
Polynomial in the number of automata

Exponential in the number of local states of each automaton (usually very low, max. 4)

Reachability analysis [PMR12]

- ❖ Idea: resolve the reachability of a local state of automata

Example : reachability from a_0 to a_2 : (objective $a_0 \rightsquigarrow a_2$)



- 2 local solutions :

- $a_0 \xrightarrow{c_1} a_2$ (c_1 required)

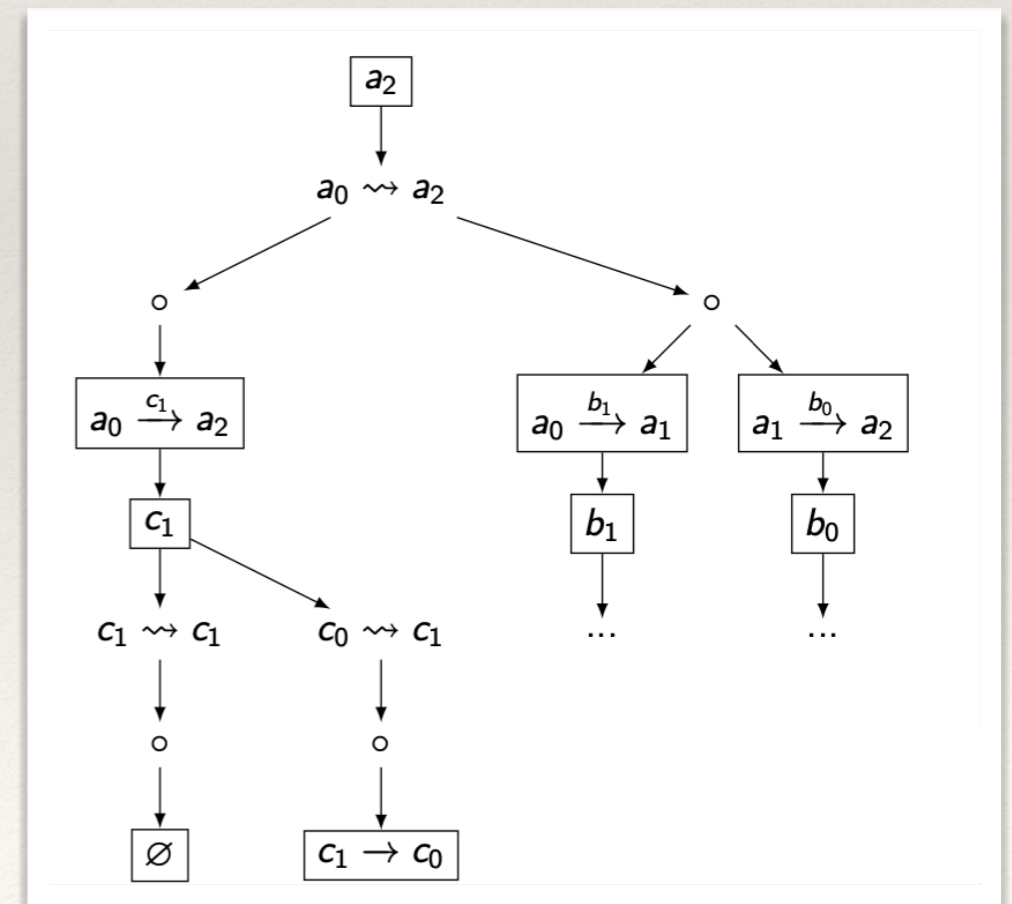
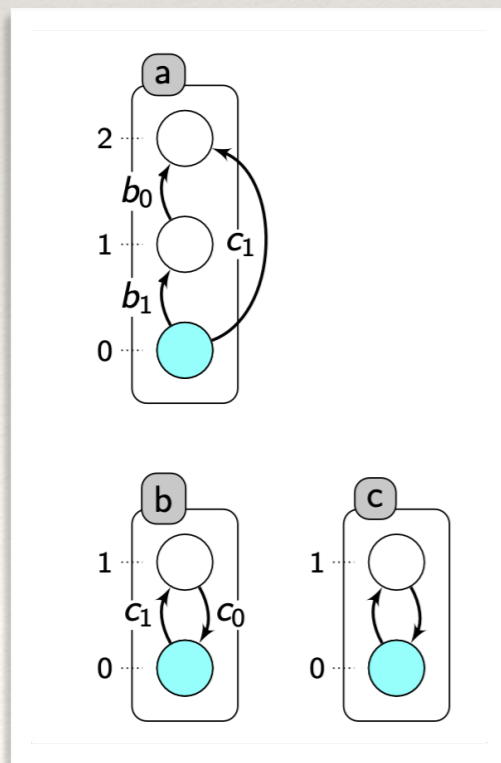
- $a_0 \xrightarrow{b_1} a_1$; $a_1 \xrightarrow{b_0} a_2$ (b_0 and b_1 required)

Reachability analysis [PMR12]

- ❖ Idea: resolve the reachability of a local state of automata
- ❖ **Over-approximation: abstraction of the order** in which the required processes are necessary
- ❖ **Under-approximation:** all required processes must be activated in all possible orders

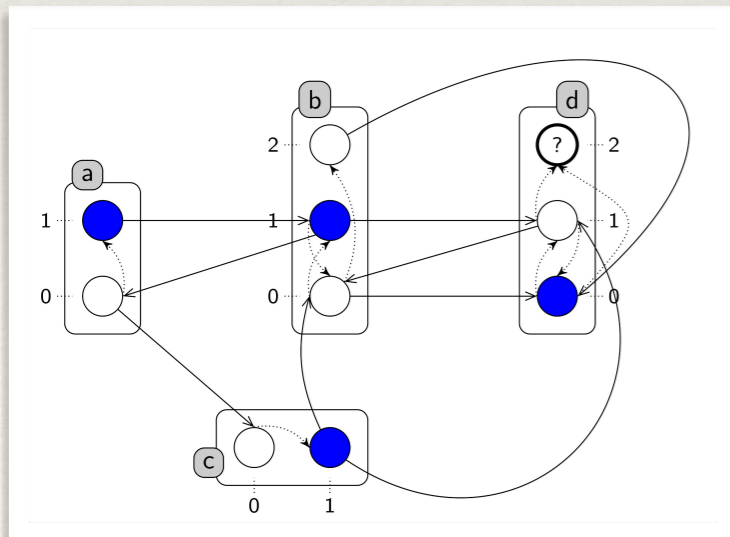
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- ❖ Construction of **local causality graph**



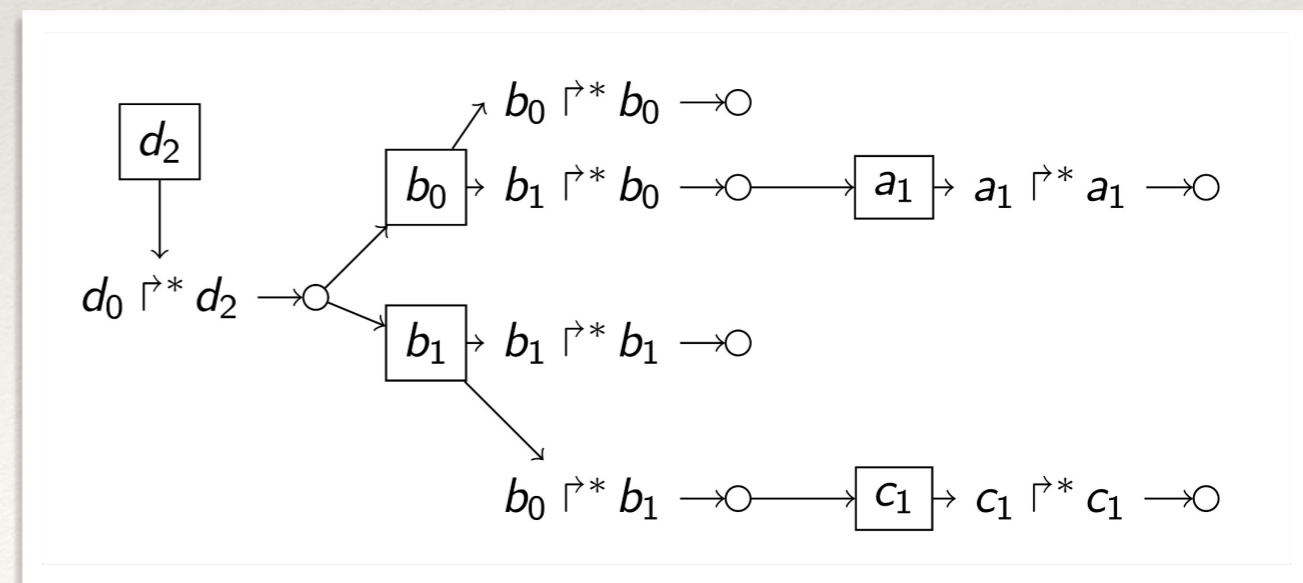
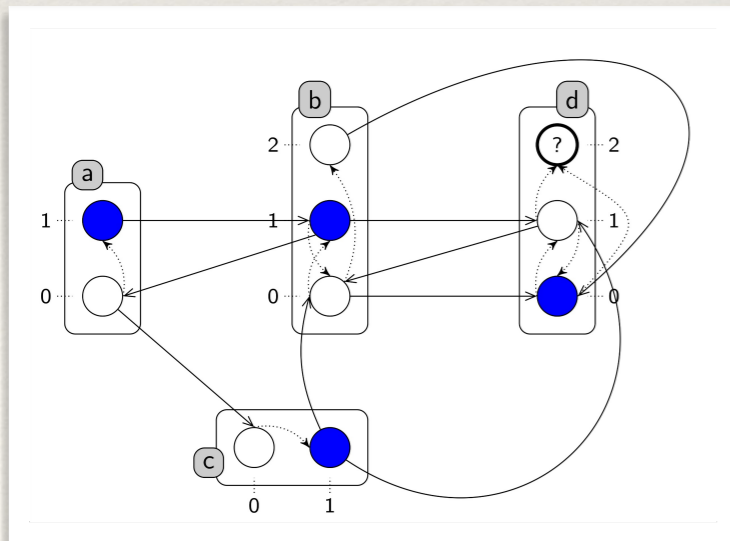
Implementation of Local Causality Graph [PMR12]

- ❖ **Computation** of the local causality graph:
 - ❖ **Polynomial** in the number of automata
 - ❖ **Exponential** in the number of local states of each automaton (usually low, max. 4)
- ❖ **Analysis** of the graph (sufficient condition): **polynomial** in the size of the abstract graph
- ❖ **Enumeration** of the subsets of solutions (if needed): **exponential** in the size of the abstract graph



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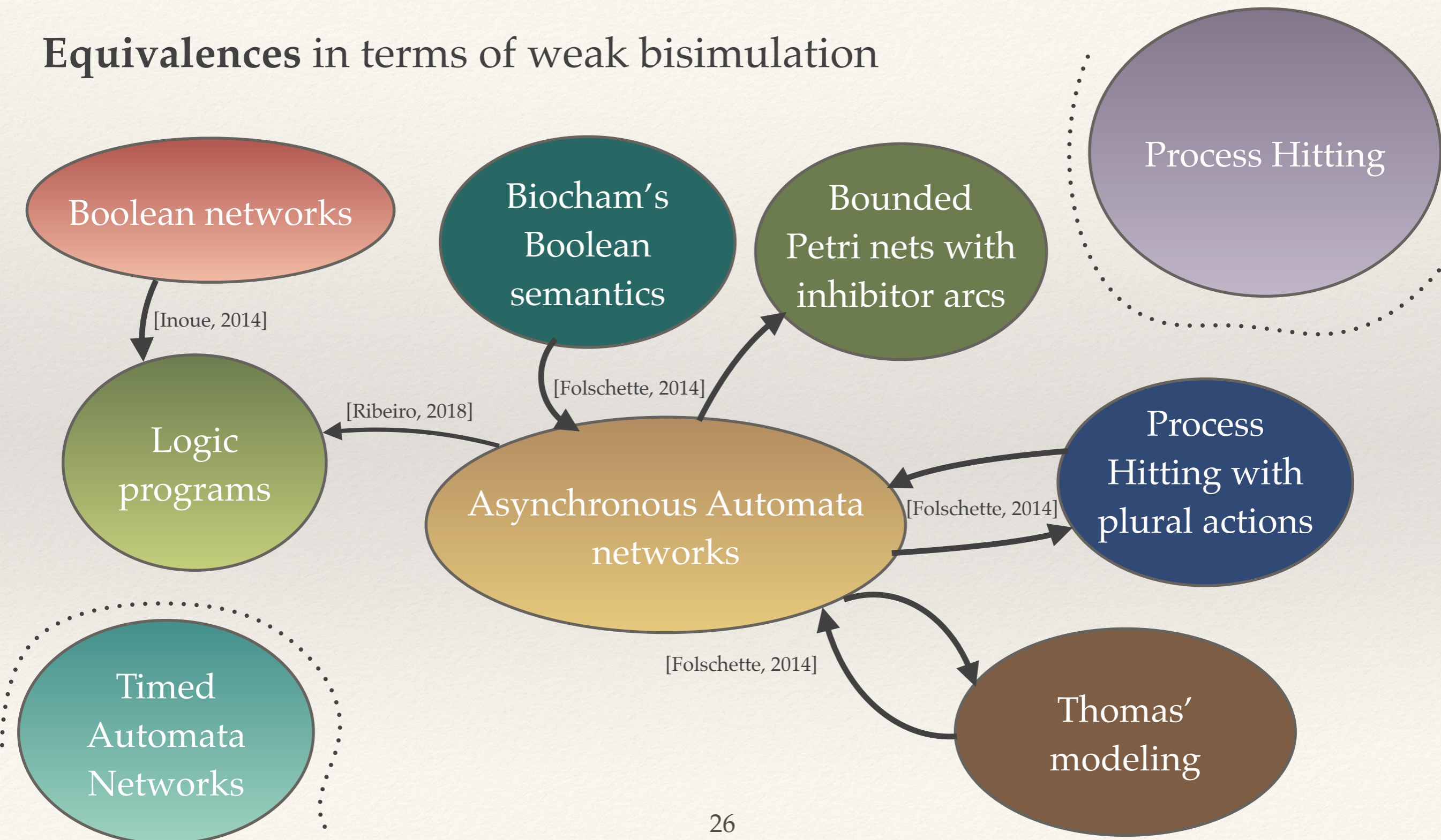


Overview

- ❖ Modeling frameworks for dynamical analysis
- ❖ Analysis of formal models
- ❖ **Inference and Learning approaches**
- ❖ Application to biological case studies

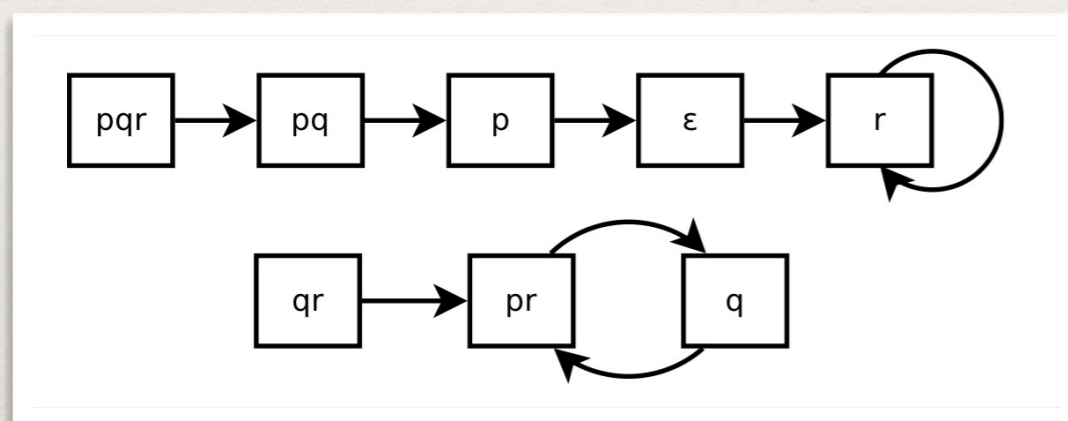
Correspondance between formalisms

Equivalences in terms of weak bisimulation



Learn systems dynamics

Principle of the LFIT framework [IRS14]: learn a **logic program** by observing the behavior of the system expressed as a succession **states-transitions**.



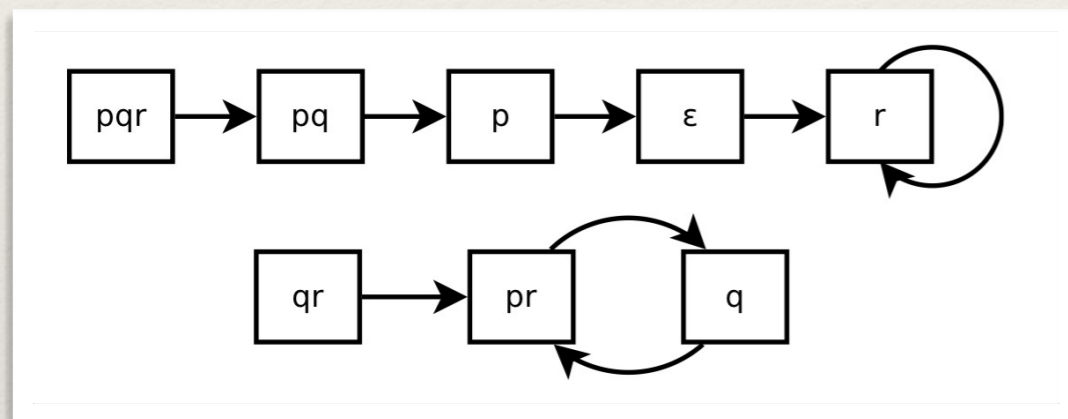
$$\begin{aligned}
 p(t+1) &\leftarrow q(t). \\
 q(t+1) &\leftarrow p(t) \wedge r(t). \\
 r(t+1) &\leftarrow \neg p(t).
 \end{aligned}$$

Input: observation of the behavior of the system

Output: logic program

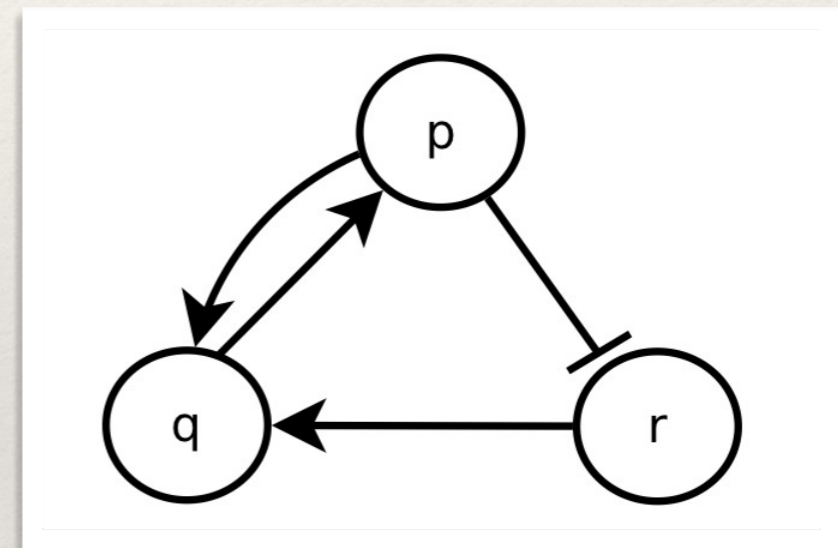
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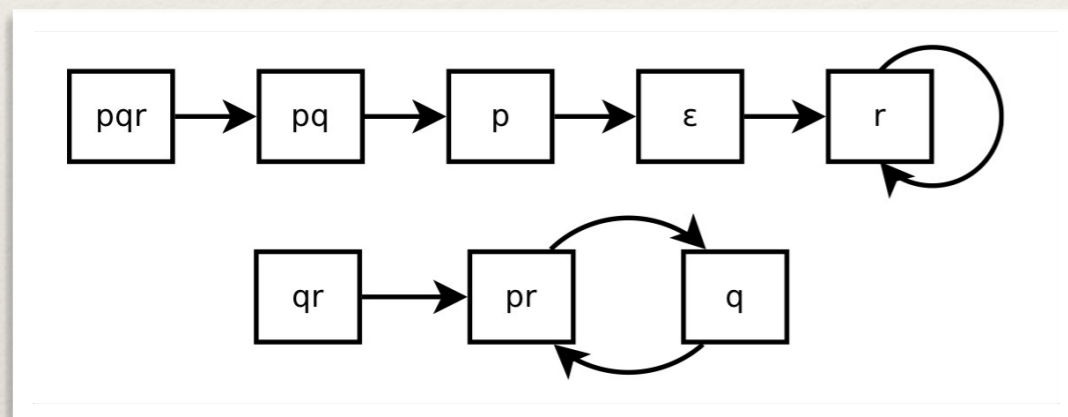
LFIT



Output: Boolean network

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LFIT

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Boolean variables

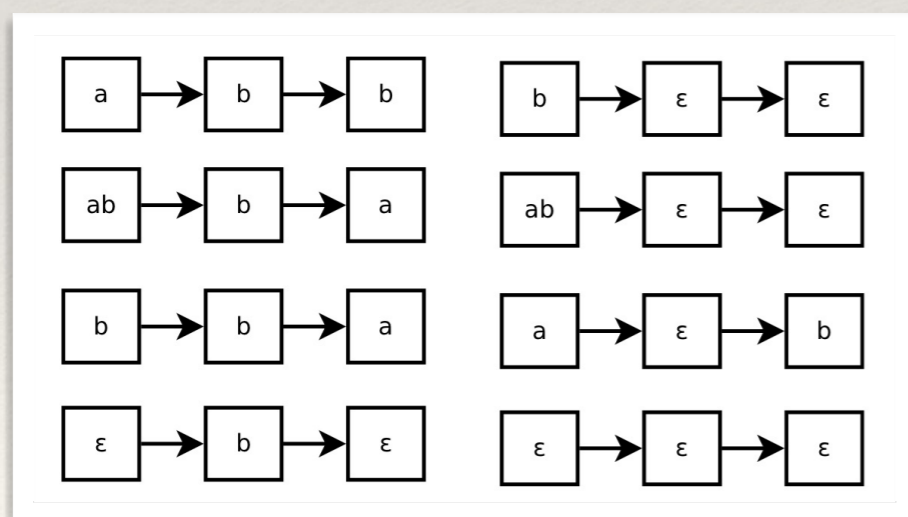
Output: logic program

Synchronous and **deterministic** dynamics

Apply LFIT to biological systems

[RMI+15a,RMI+15b]

- ❖ Successive extensions :
 - ❖ **Delayed influences** (models with memory)
 - ❖ **Multi-valued variables**
 - ❖ **Asynchronous dynamics**
 - ❖ **Non-deterministic semantics**



$a(t) \leftarrow b(t-1), b(t-2)$
 $b(t) \leftarrow a(t-2), \neg b(t-2)$

Input: observation of the system's behavior

Output: logic program

A wide range of possible models (and inference algorithms)

<https://github.com/Tony-sama/pylfit>

Values	Boolean	Multi-valued	Captured symbolically (e.g. intervals)
Time	Chronological: Event-driven model	Chronometrical: Discrete time	Chronometrical: Dense time
Semantics of discrete transitions	Synchronous	Asynchronous	Generalized
Determinism of discrete transitions	Deterministic	Non-deterministic	Probabilistic
Markov property	With memory	Memoryless	

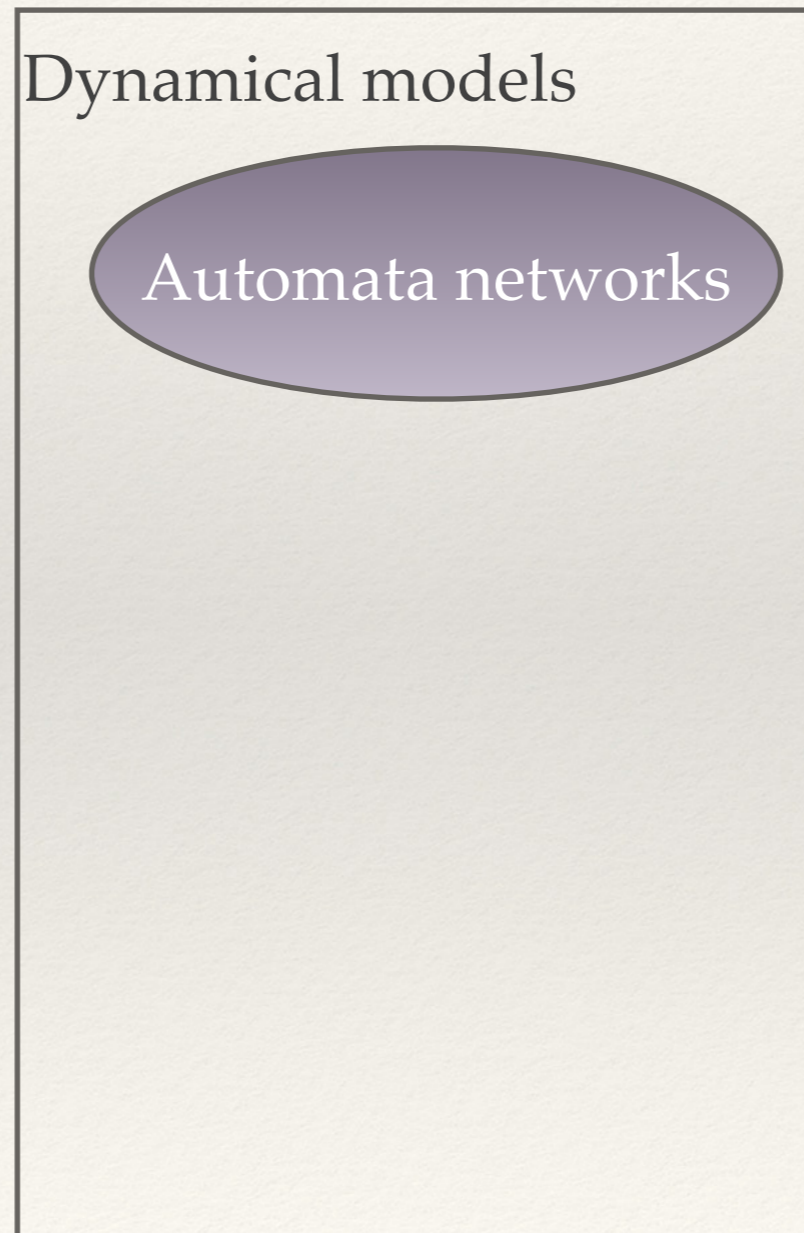
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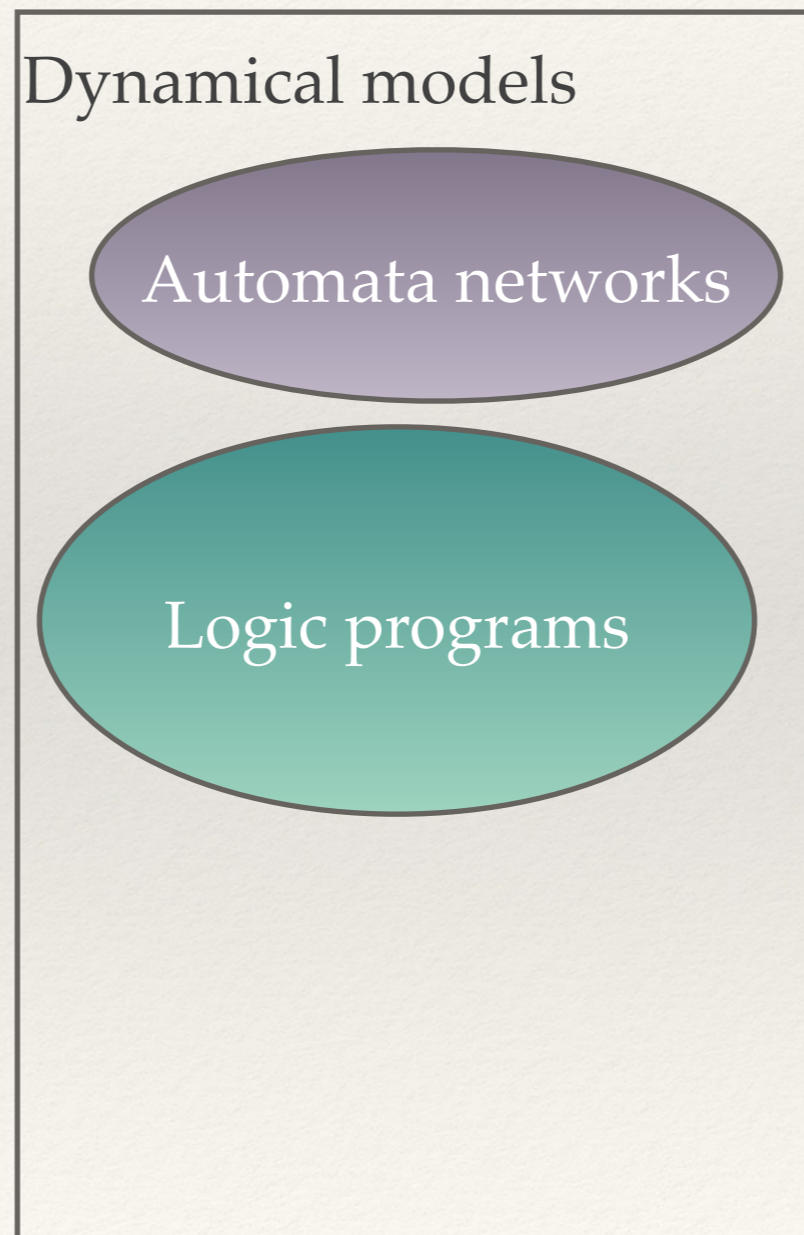
Synthesis of our contributions

Dynamical models

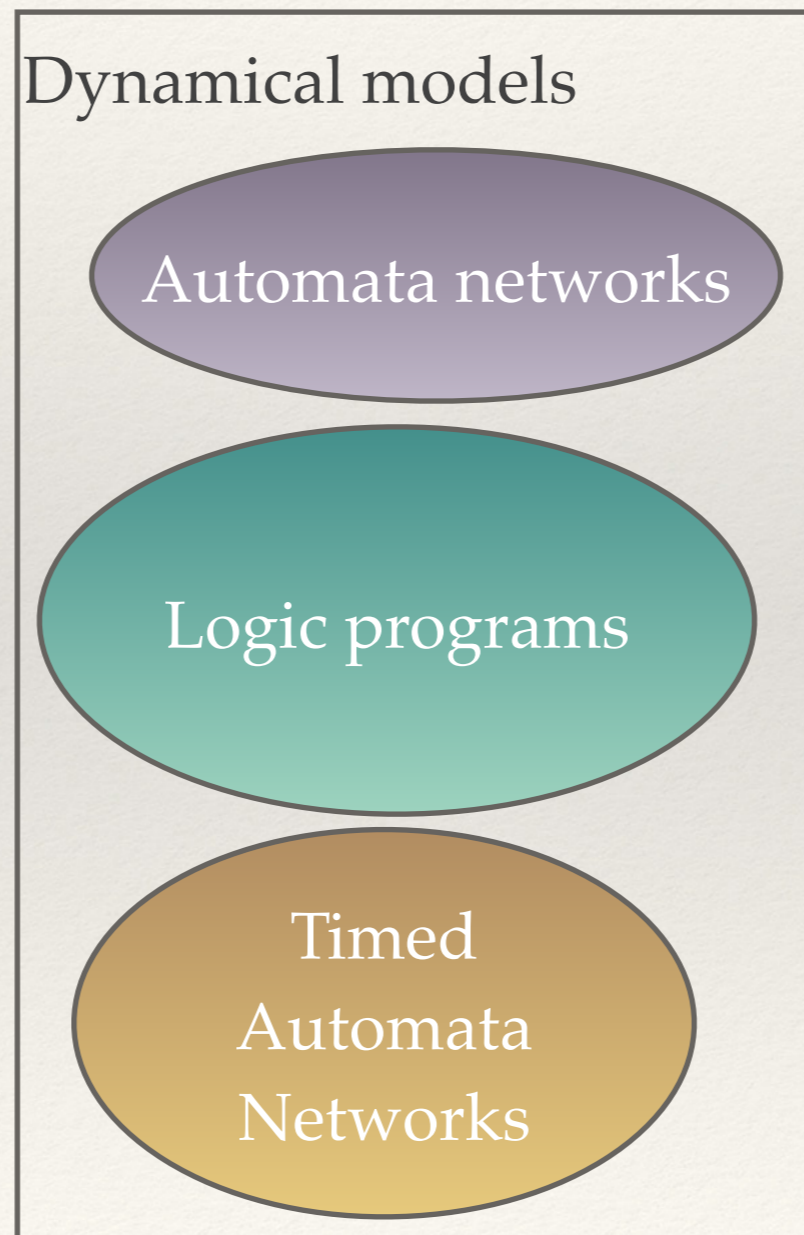
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Question?

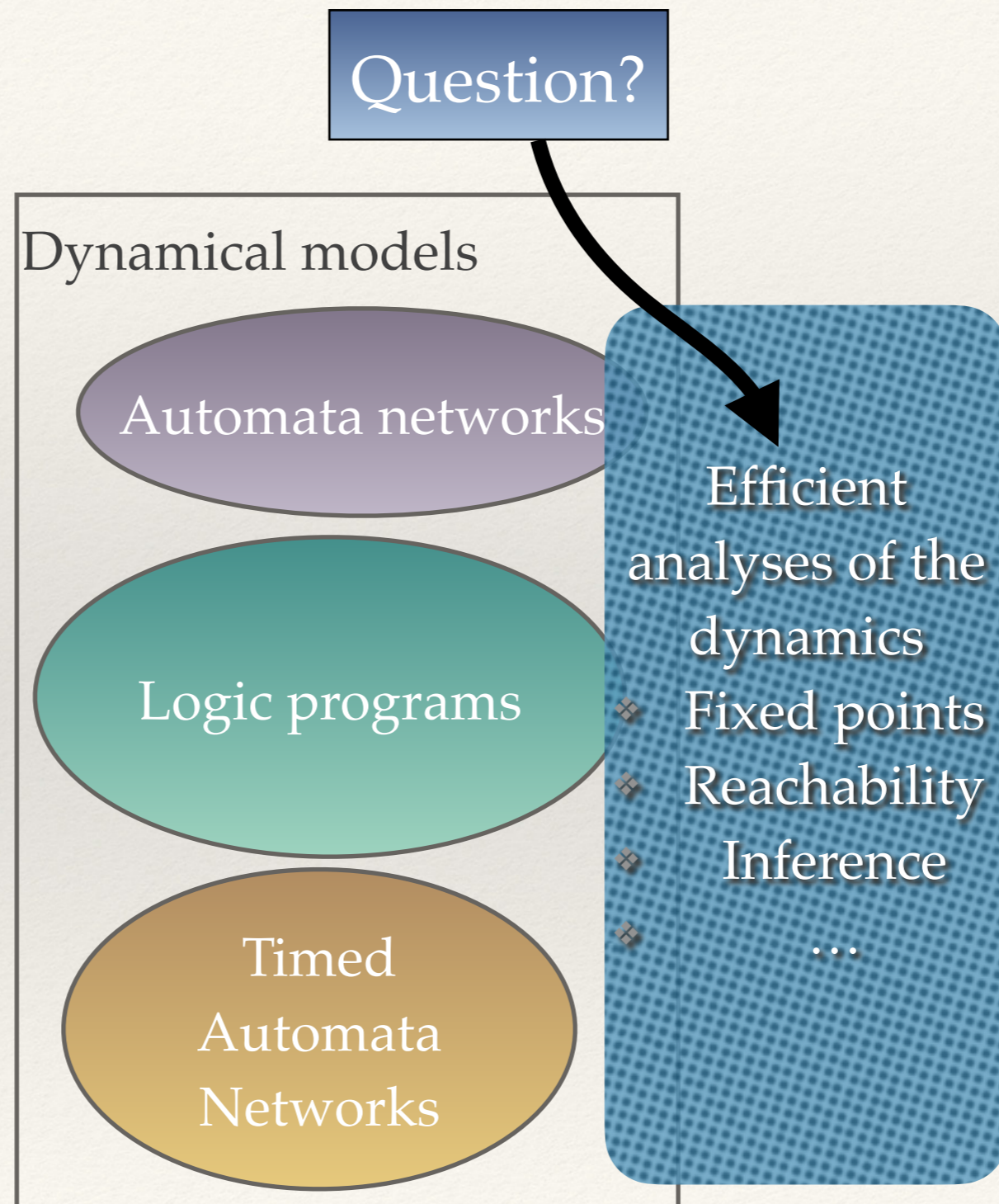
Dynamical models

Automata networks

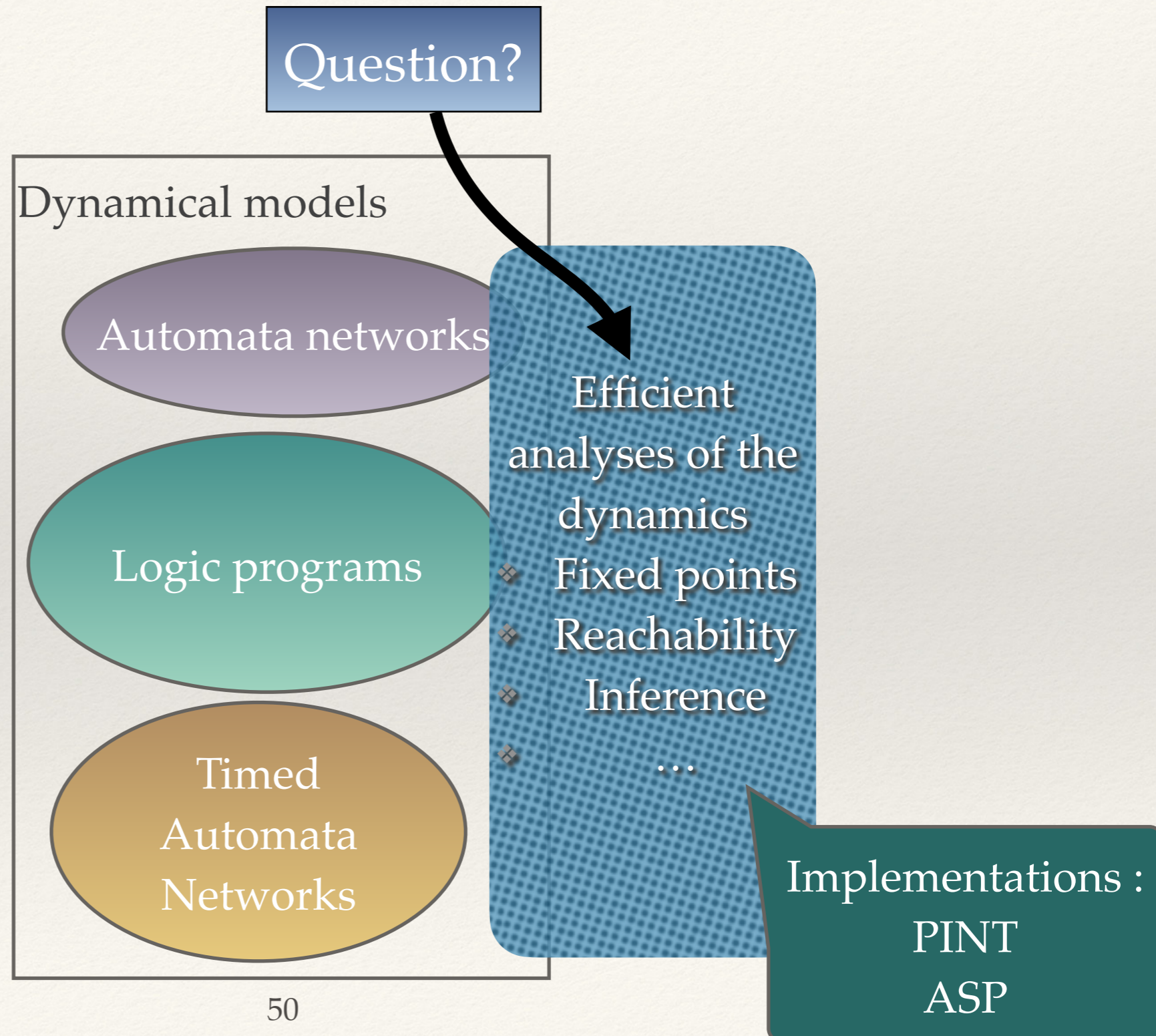
Logic programs

Timed
Automata
Networks

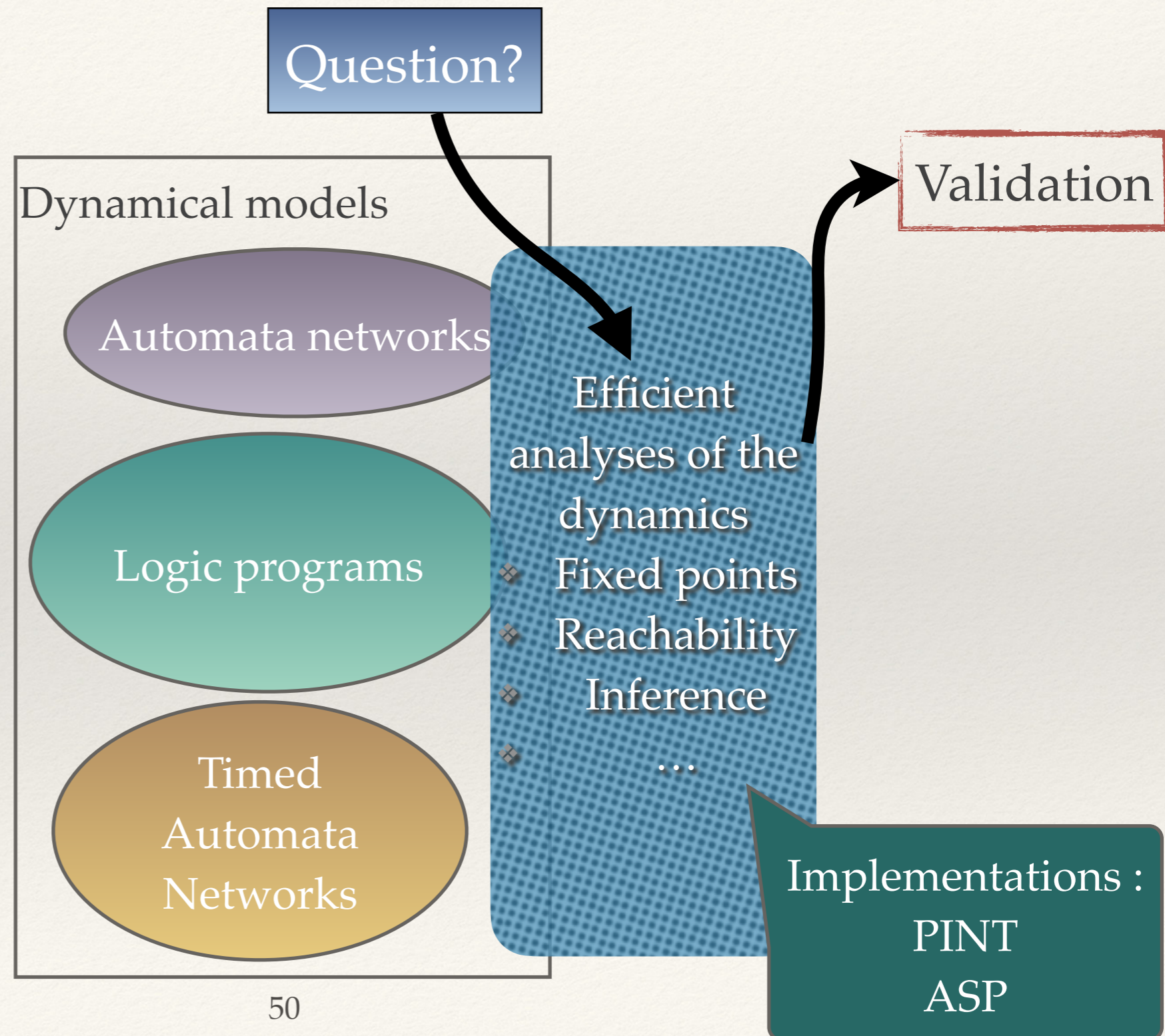
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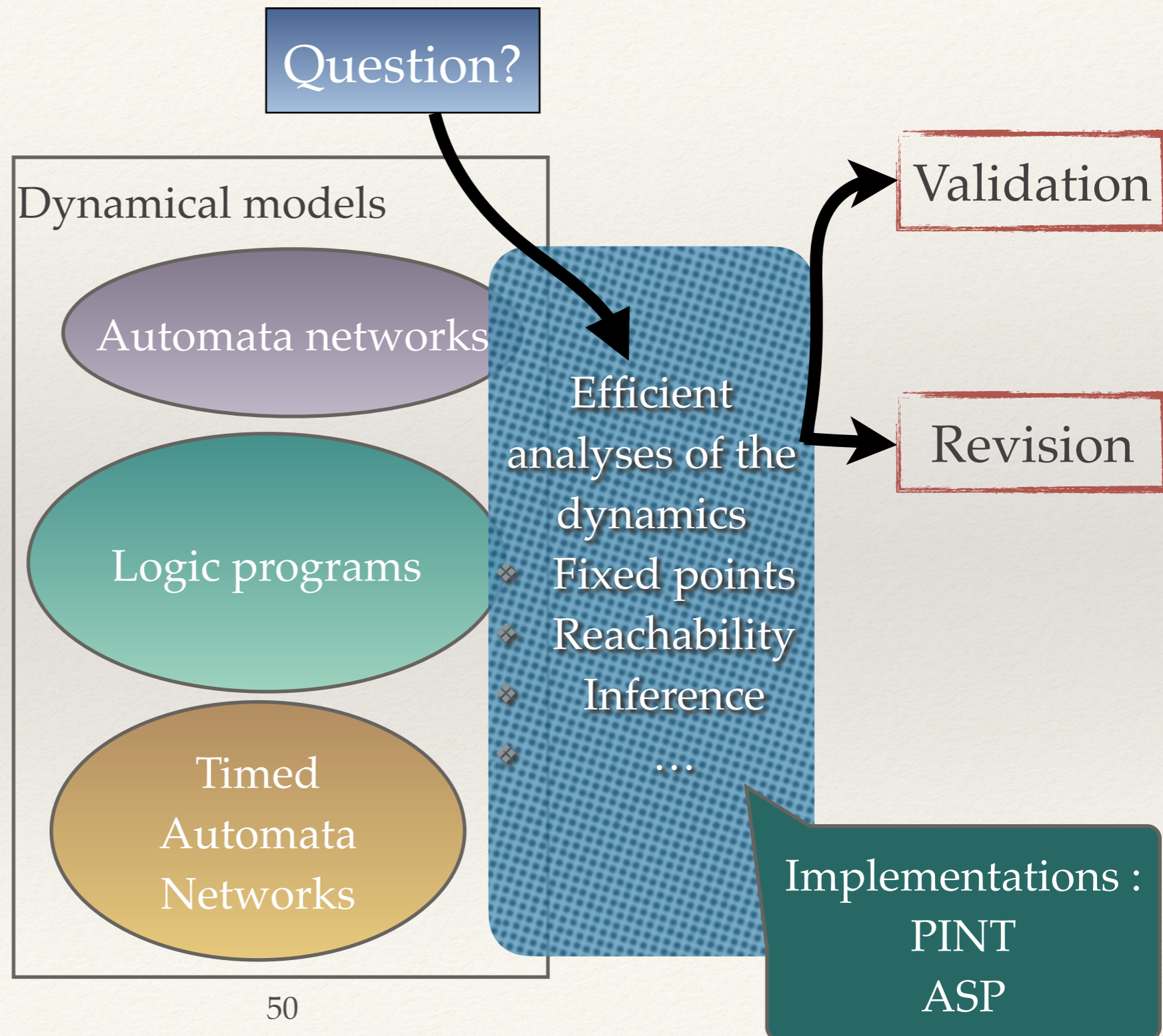
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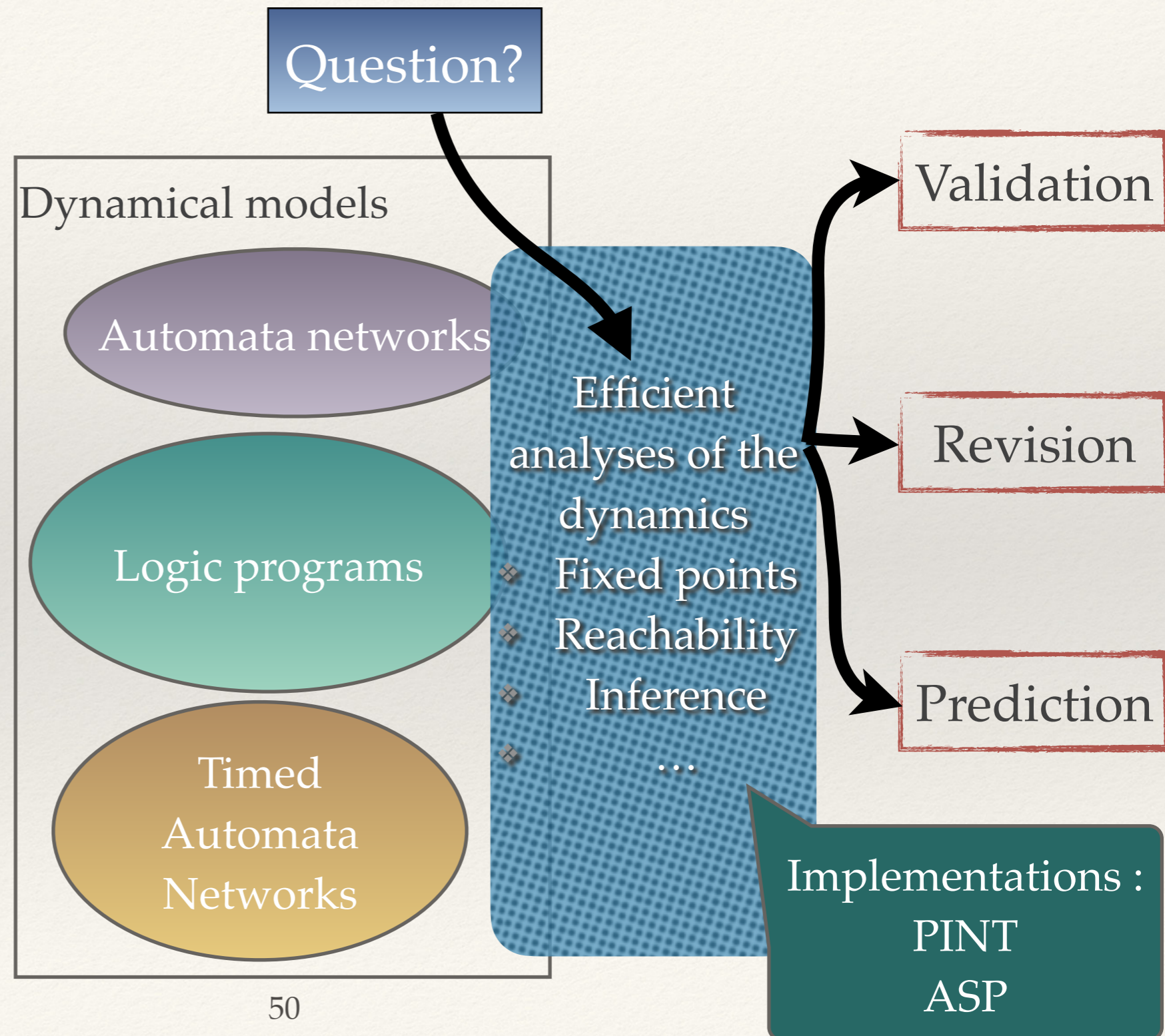
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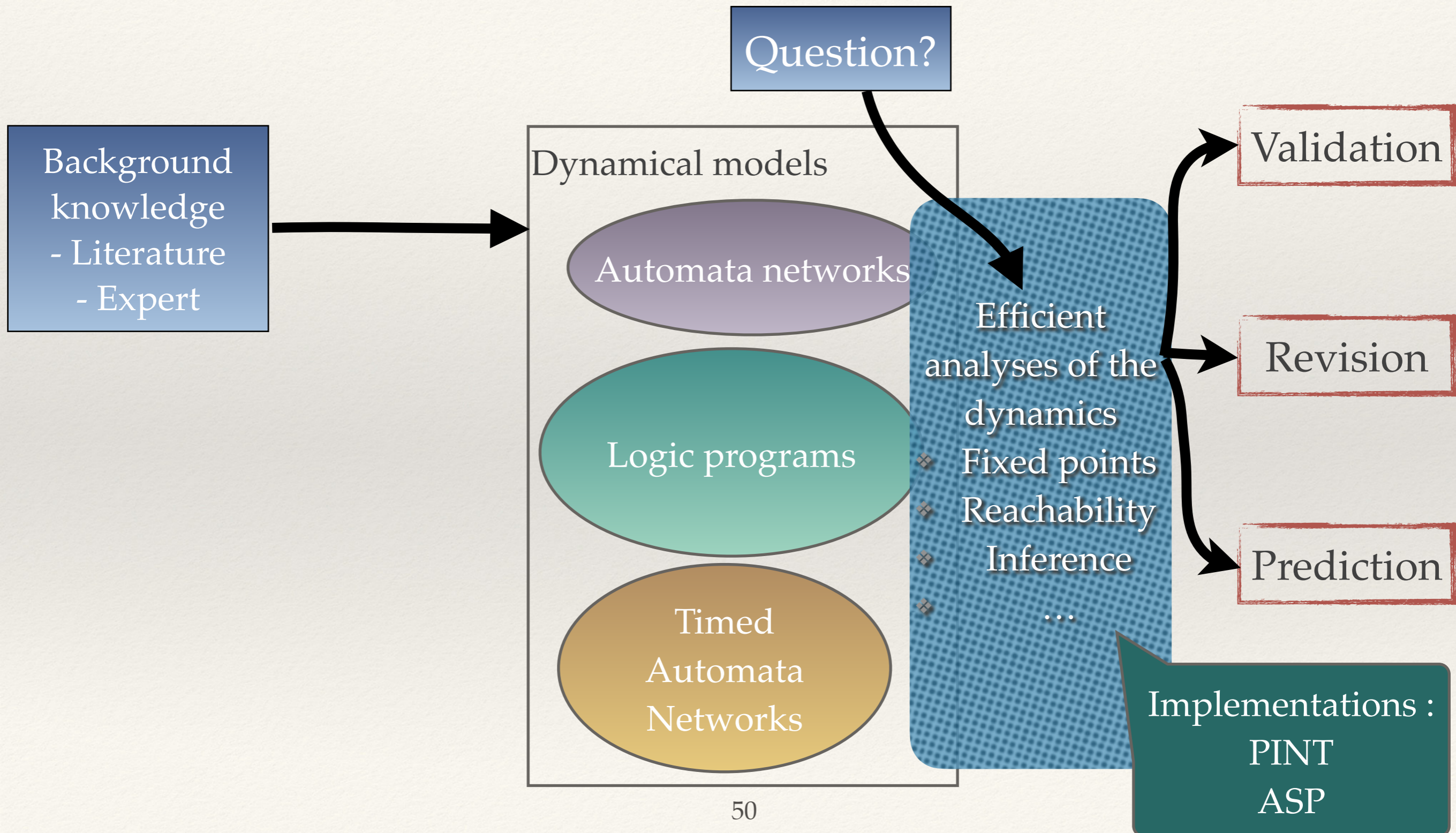
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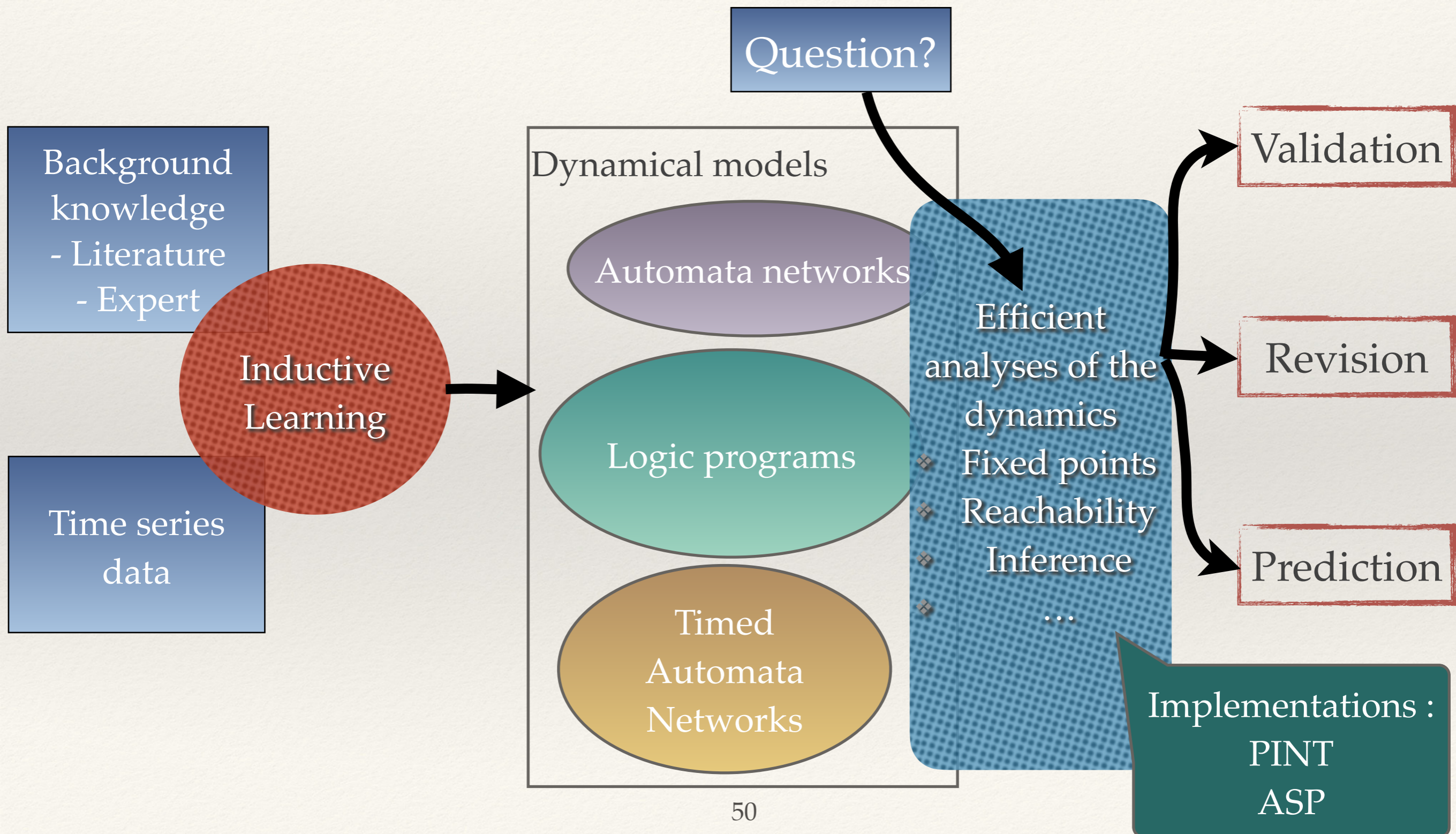
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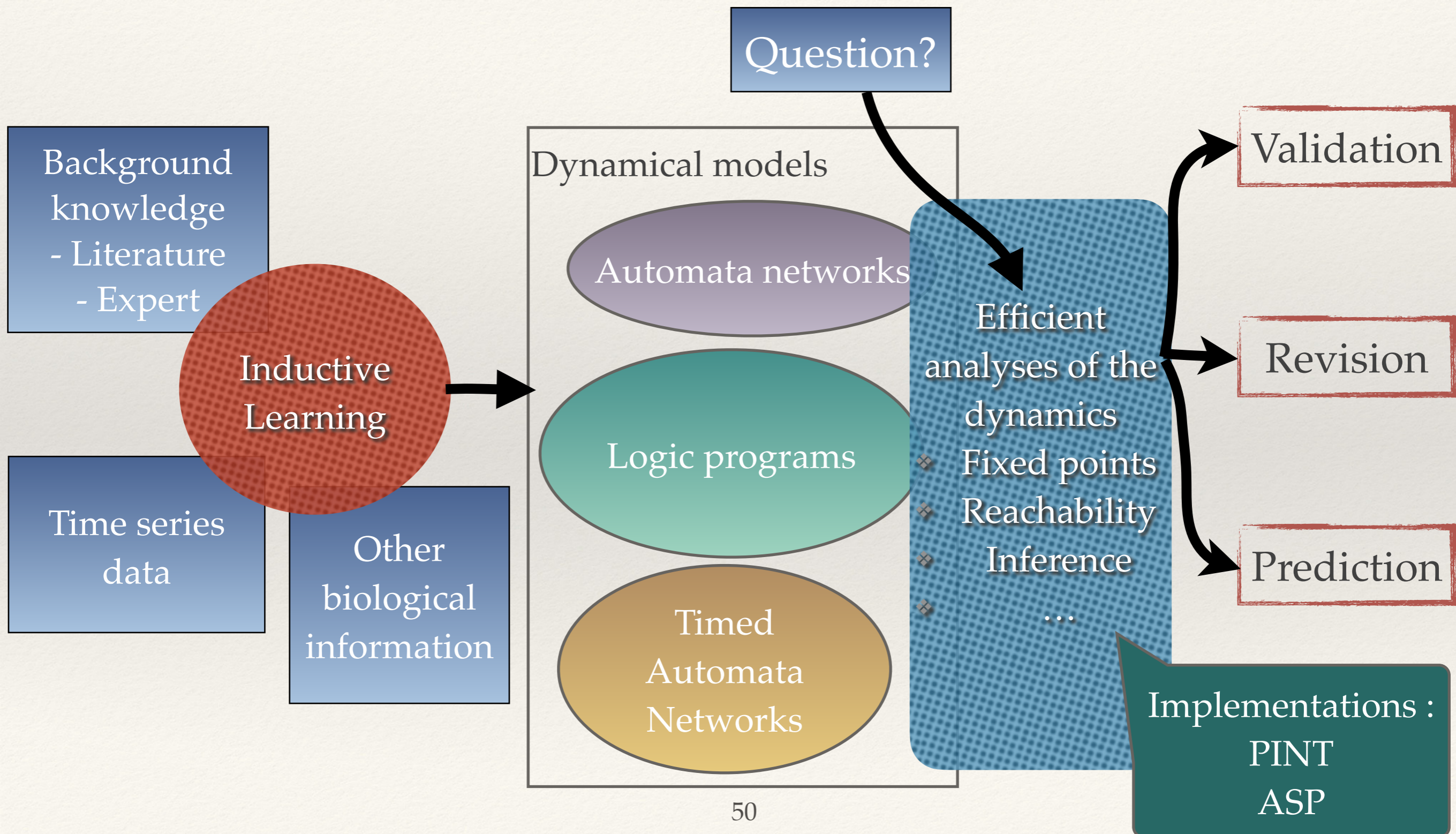
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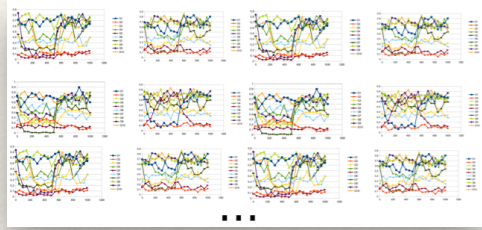
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Methodological goals

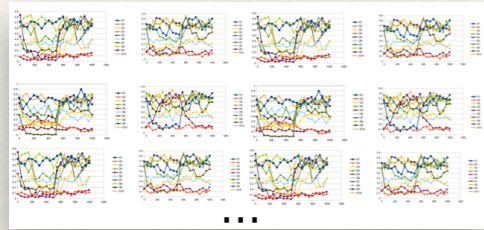
Methodological goals

Time series data



Methodological goals

Time series data

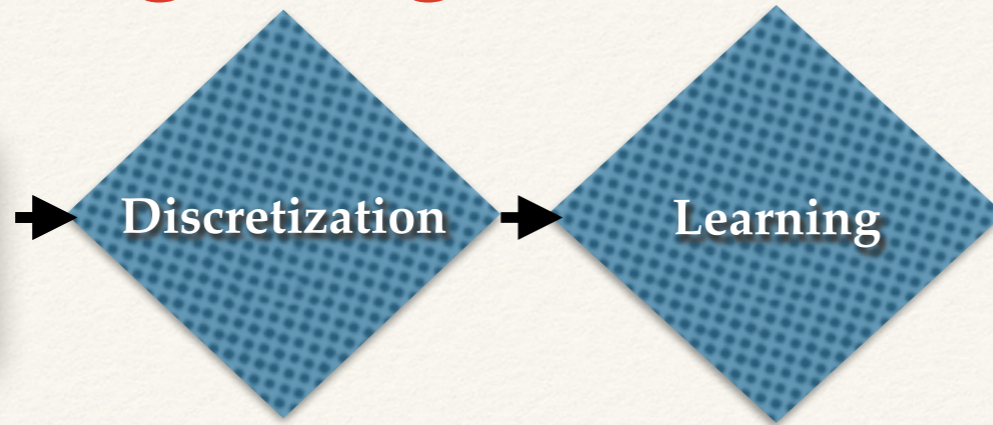
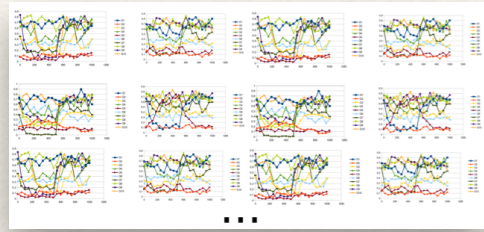


Discretization



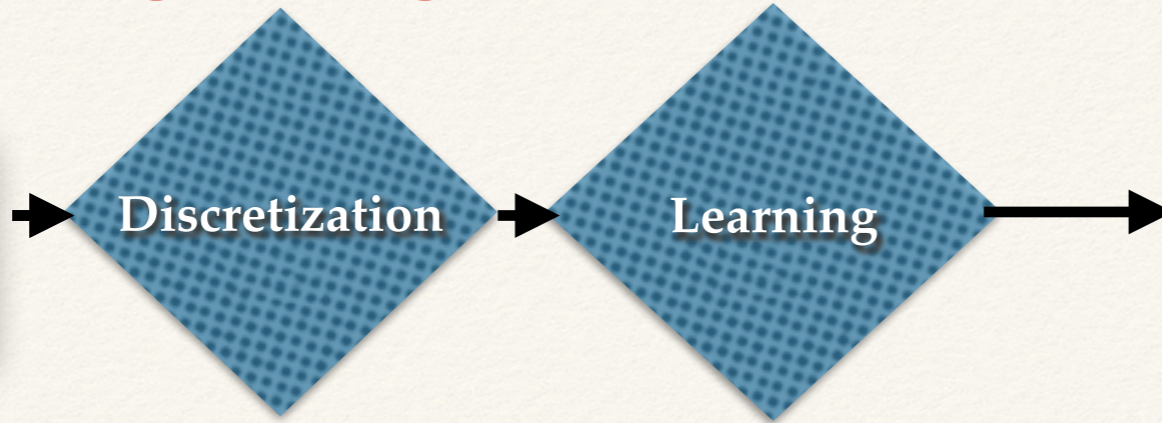
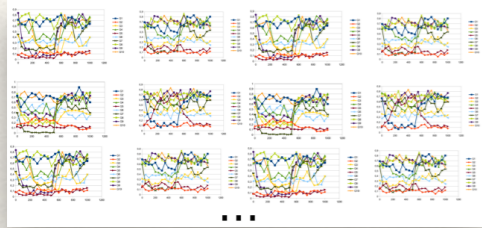
Methodological goals

Time series data



Methodological goals

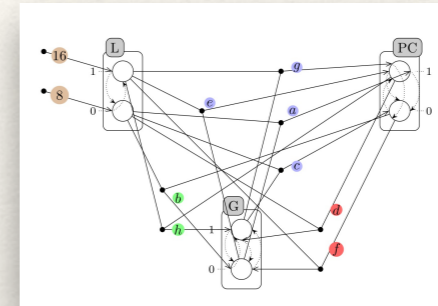
Time series data



Dynamical models

```
ARNTL(0,T) :- CLOCK(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(0,T-1).
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ARNTL(0,T) :- CRY1(0,T-1), PER1(2,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER2(2,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER2(3,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PPARA(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PPARA(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), CRY2(0,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), PPARA(2,T-1).
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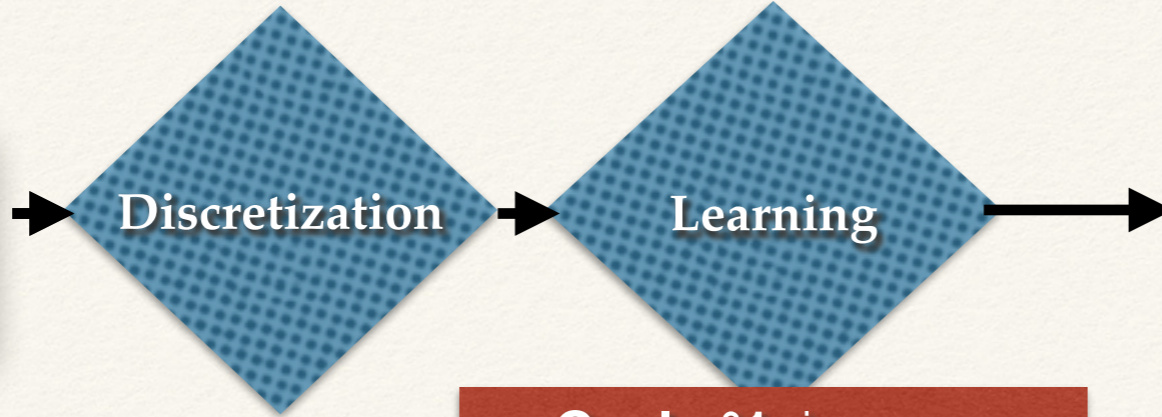
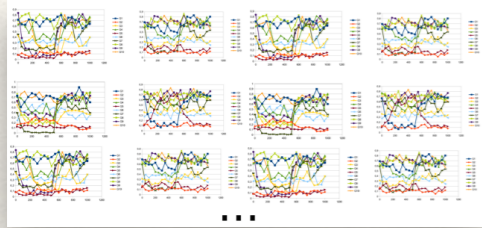
Logic programs



Automata networks

Methodological goals

Time series data

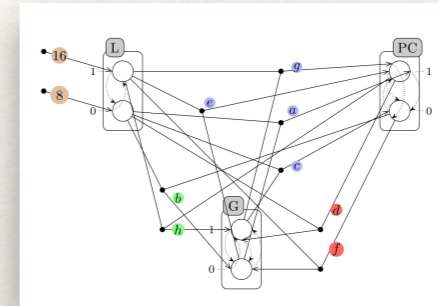


Goal n°1: improve learning by addressing noisy, incomplete and/or conflicting data

Dynamical models

```
ARNTL(0,T) :- CLOCK(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(0,T-1).
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ARNTL(0,T) :- CRY1(0,T-1), PER1(2,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER2(2,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER2(3,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), NR1D1(2,T-1).
ARNTL(0,T) :- CRY1(1,T-1), NR1D1(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PER1(0,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), PPARA(2,T-1).
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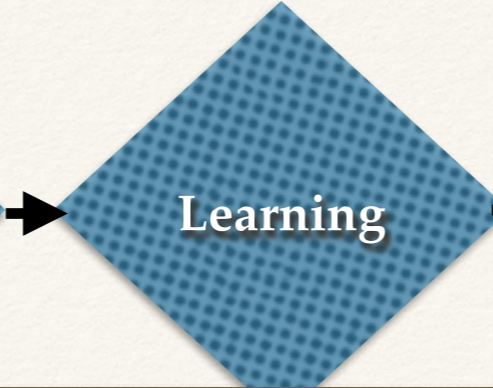
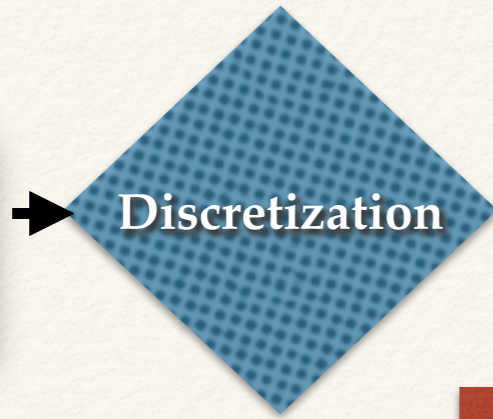
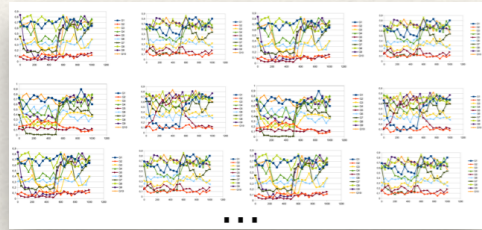
Logic programs



Automata networks

Methodological goals

Time series data

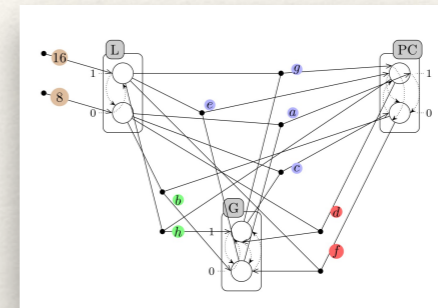


Goal n°1: improve learning by addressing noisy, incomplete and/or conflicting data

Dynamical models

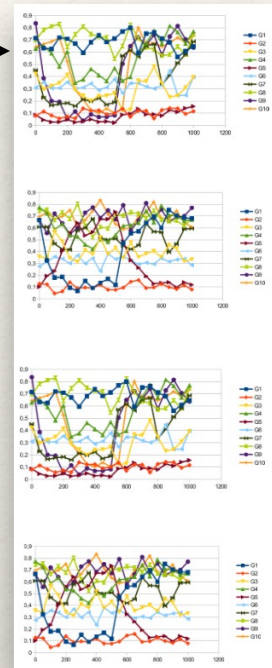
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ARNTL(0,T) :- CLOCK(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER2(2,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER2(3,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PPARA(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PPARA(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), NR1D1(2,T-1).
ARNTL(0,T) :- CRY1(1,T-1), NR1D1(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PER1(0,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), PER2(3,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PPARA(2,T-1).
```

Logic programs



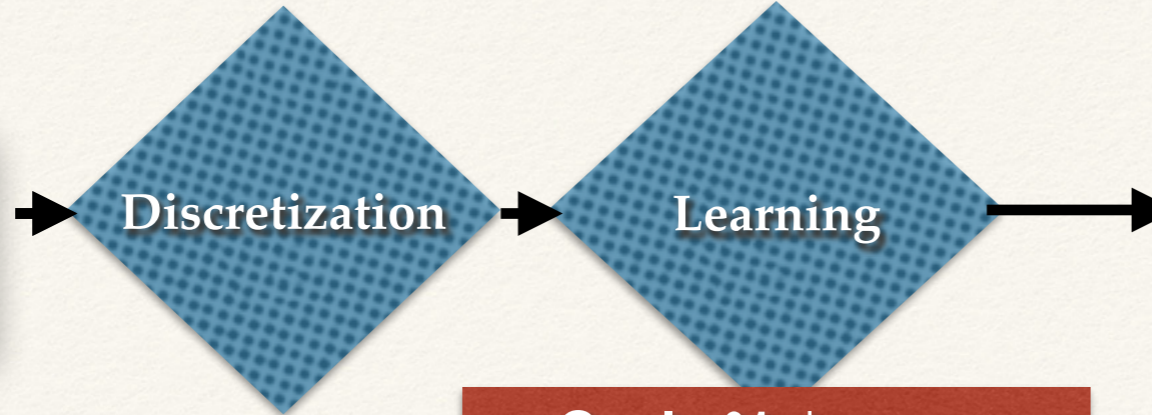
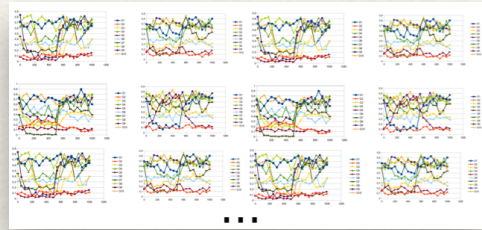
Automata networks

Analysis of dynamic behaviors



Methodological goals

Time series data

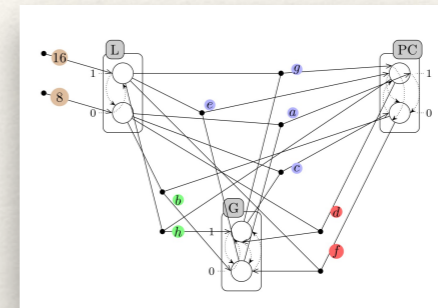


Goal n°1: improve learning by addressing noisy, incomplete and/or conflicting data

Dynamical models

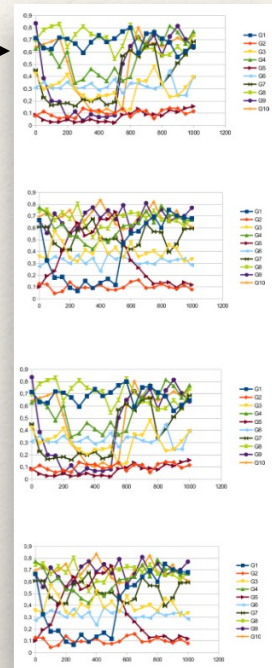
```
ARNTL(0,T) :- CLOCK(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER2(2,T-1).
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ARNTL(0,T) :- CRY1(0,T-1), PPARA(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), NR1D1(2,T-1).
ARNTL(0,T) :- CRY1(1,T-1), NR1D1(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PER1(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PER1(2,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PER2(2,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PER2(3,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PPARA(2,T-1).
```

Logic programs



Automata networks

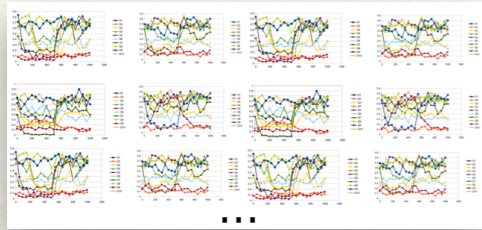
Analysis of dynamic behaviors



Background knowledge on the dynamics

Methodological goals

Time series data



Discretization

Learning

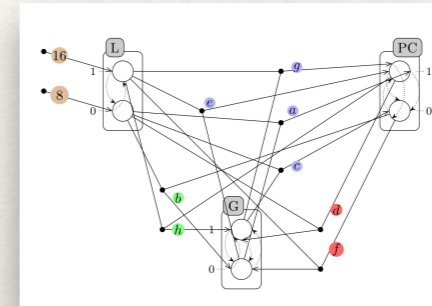
Goal n°1: improve learning by addressing noisy, incomplete and/or conflicting data

Dynamical models

```

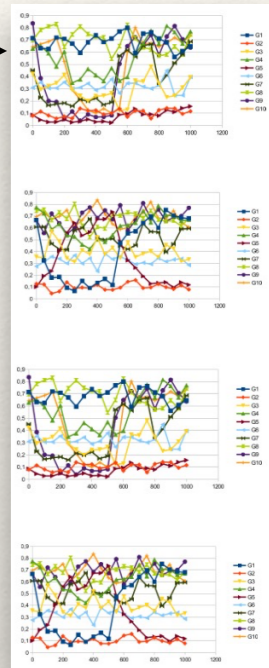
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ARNTL(0,T) :- CRY1(0,T-1), PER2(3,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PPARA(1,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), PER1(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PER1(2,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), PPARA(2,T-1).
    
```

Logic programs



Automata networks

Analysis of dynamic behaviors



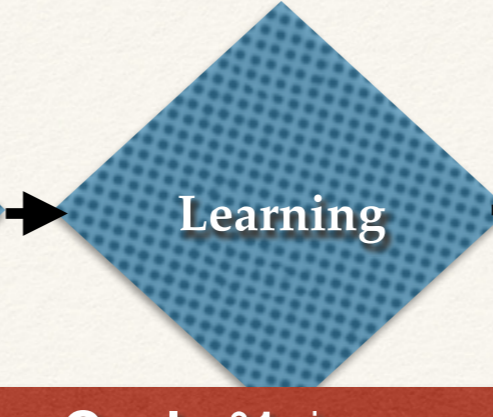
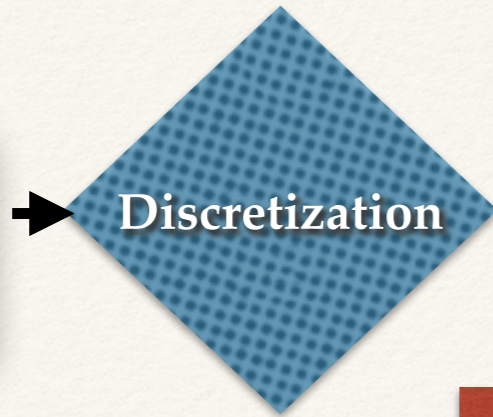
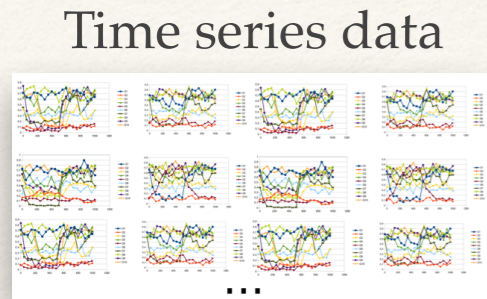
Background knowledge on the dynamics

Formalization

Expected properties in temporal logics

$$(M(p_{G0}) = 1 \wedge M(p_{PC0}) = 1) \rightsquigarrow_{[0,70,1]} (M(p_{G1}) = 1)$$

Methodological goals



Goal n°1: improve learning by addressing noisy, incomplete and/or conflicting data

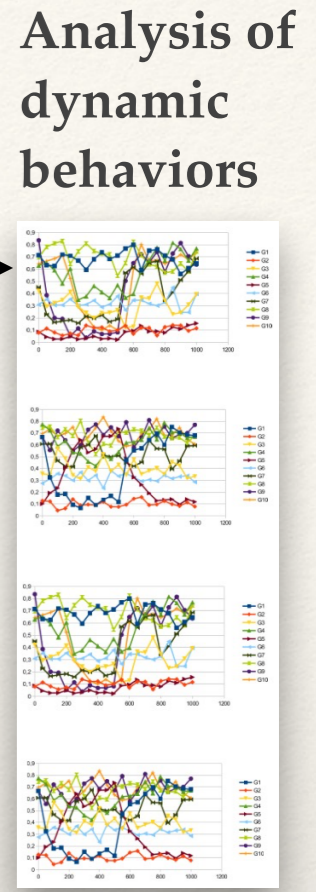
Dynamical models

```

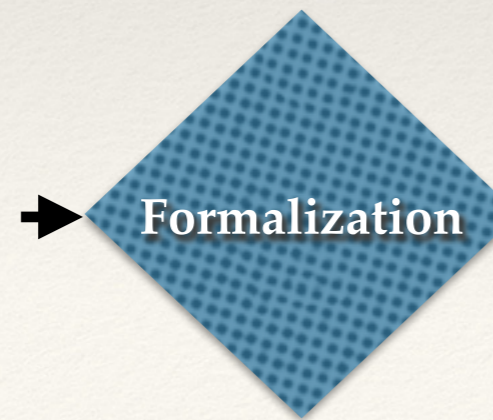
ARNTL(0,T) :- CLOCK(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(1,T-1), NR1D1(2,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), PPARA(2,T-1).
    
```

Logic programs

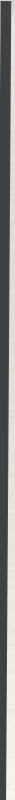
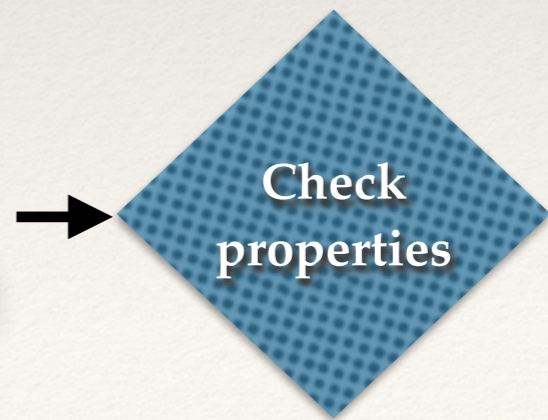
Automata networks



Background knowledge on the dynamics

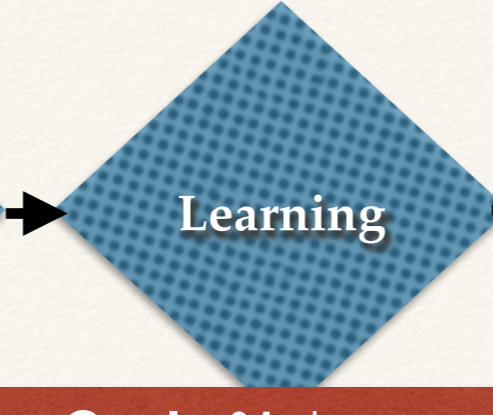
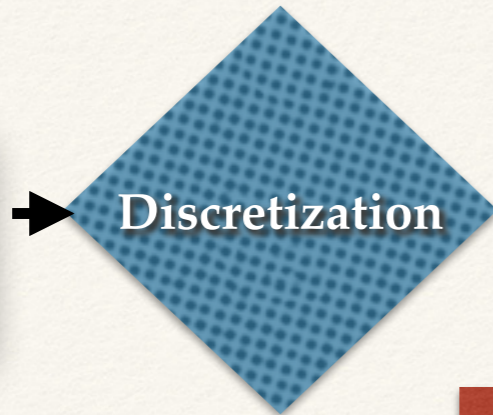
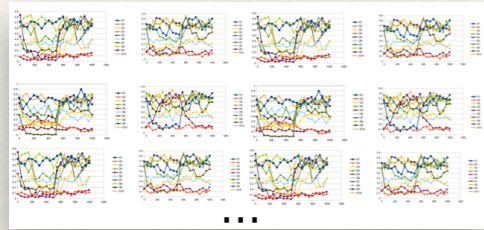


Expected properties in temporal logics

$$(M(p_{G0}) = 1 \wedge M(p_{PC0}) = 1) \rightsquigarrow_{[0, \tau_0, 1]} (M(p_{G1}) = 1)$$


Methodological goals

Time series data



Goal n°1: improve learning by addressing noisy, incomplete and/or conflicting data

Dynamical models

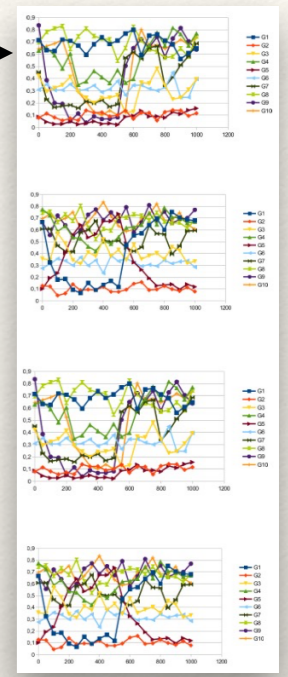
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ARNTL(0,T) :- CLOCK(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), PER1(1,T-1).
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```

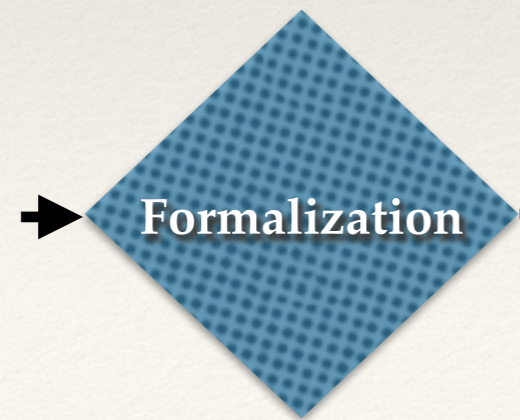
Logic programs

Automata networks

Analysis of dynamic behaviors

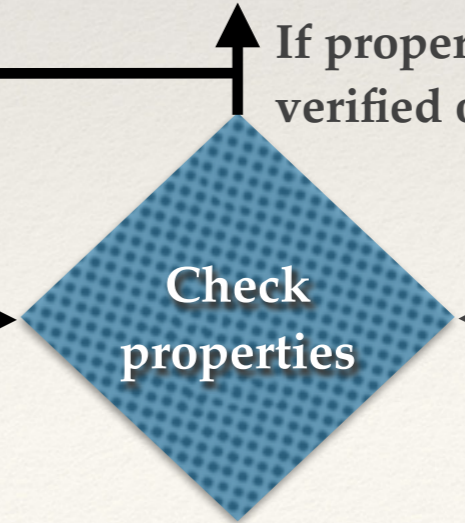


Background knowledge on the dynamics

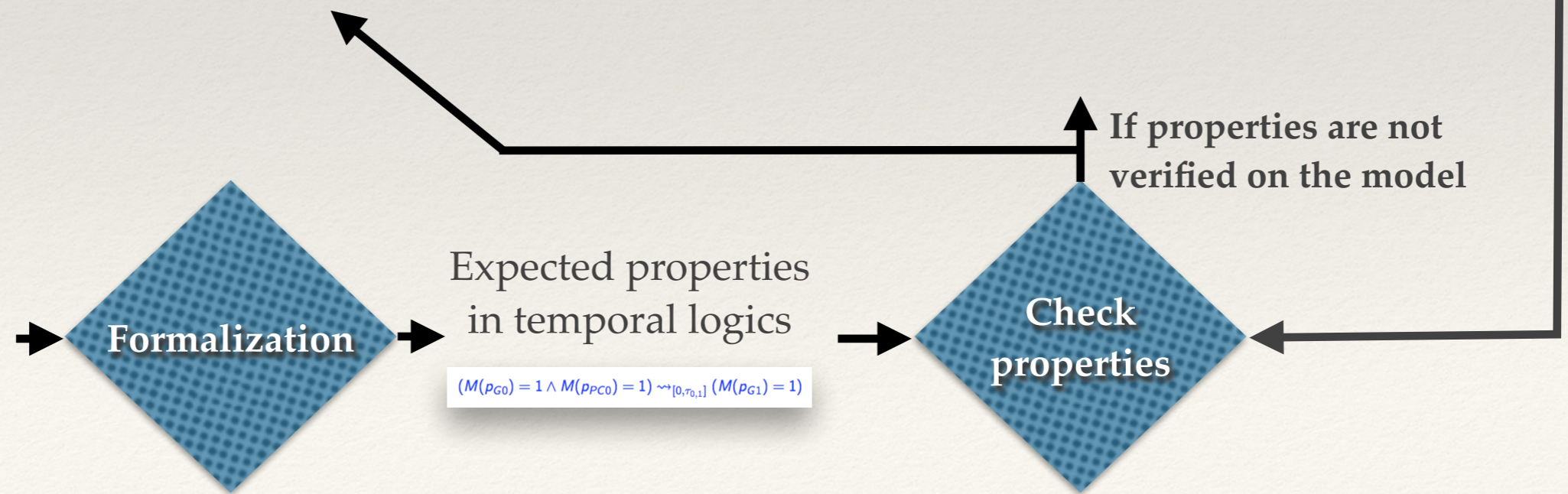


Expected properties in temporal logics

$$(M(p_{G0}) = 1 \wedge M(p_{PC0}) = 1) \rightsquigarrow_{[0,70,1]} (M(p_{G1}) = 1)$$

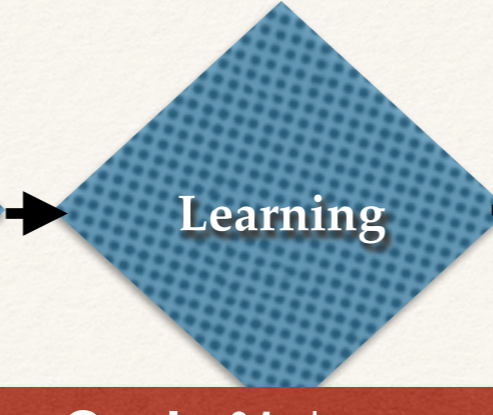
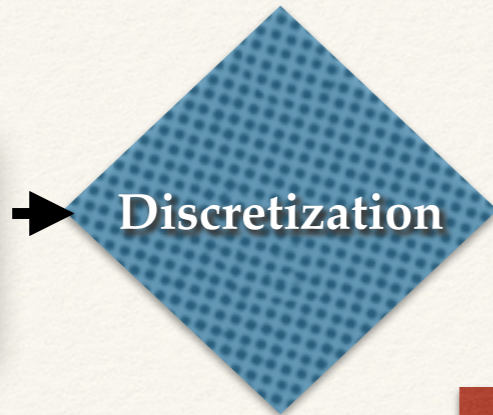
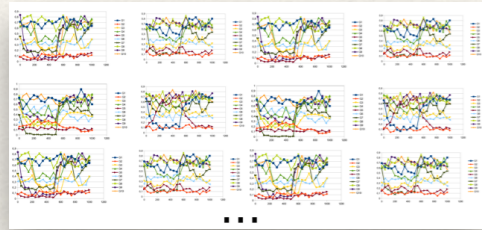


If properties are not verified on the model



Methodological goals

Time series data



Goal n°1: improve learning by addressing noisy, incomplete and/or conflicting data

Dynamical models

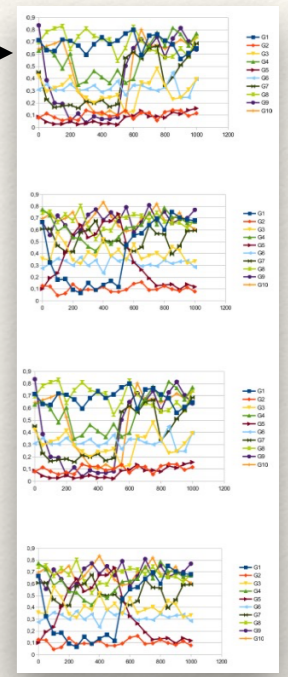
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ARNTL(0,T) :- CLOCK(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), CRY2(0,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
ARNTL(0,T) :- CRY1(0,T-1), NR1D1(0,T-1).
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ARNTL(0,T) :- CRY1(1,T-1), PER2(3,T-1).
ARNTL(0,T) :- CRY1(1,T-1), PPARA(2,T-1).
    
```

Logic programs

Automata networks

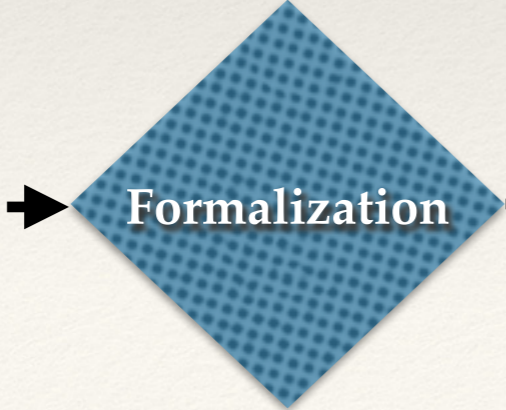
Analysis of dynamic behaviors



Goal n°2: automatically propose modifications to control the model w.r.t. properties

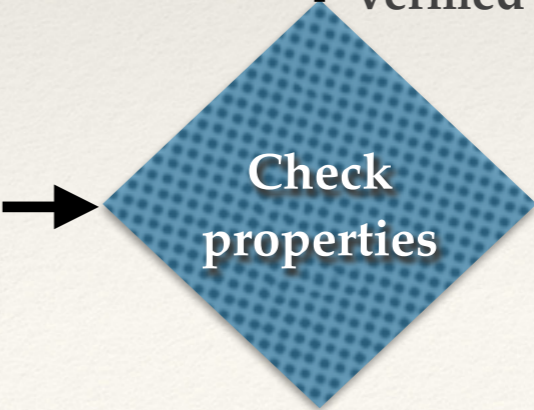


Background knowledge on the dynamics

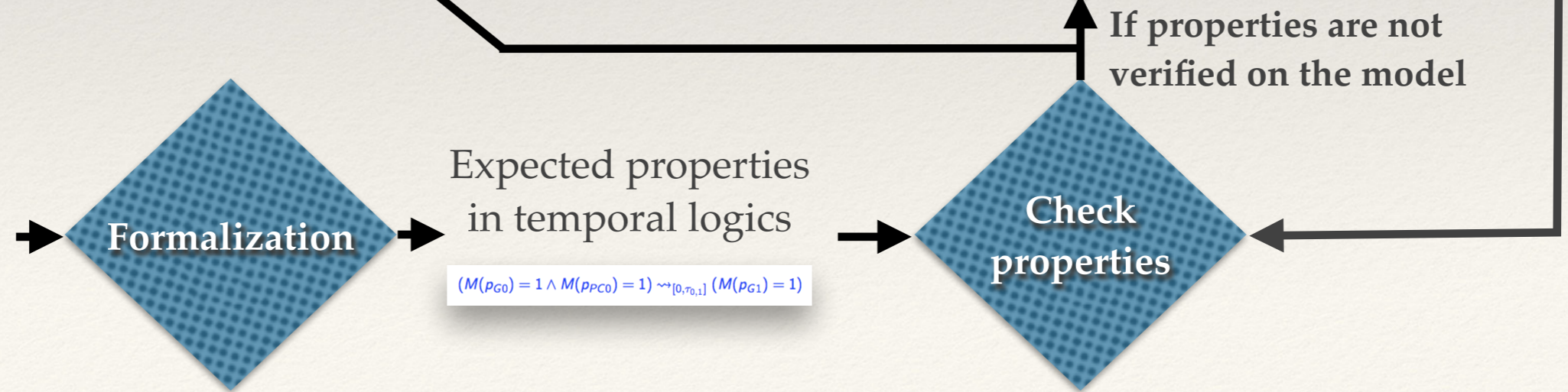


Expected properties in temporal logics

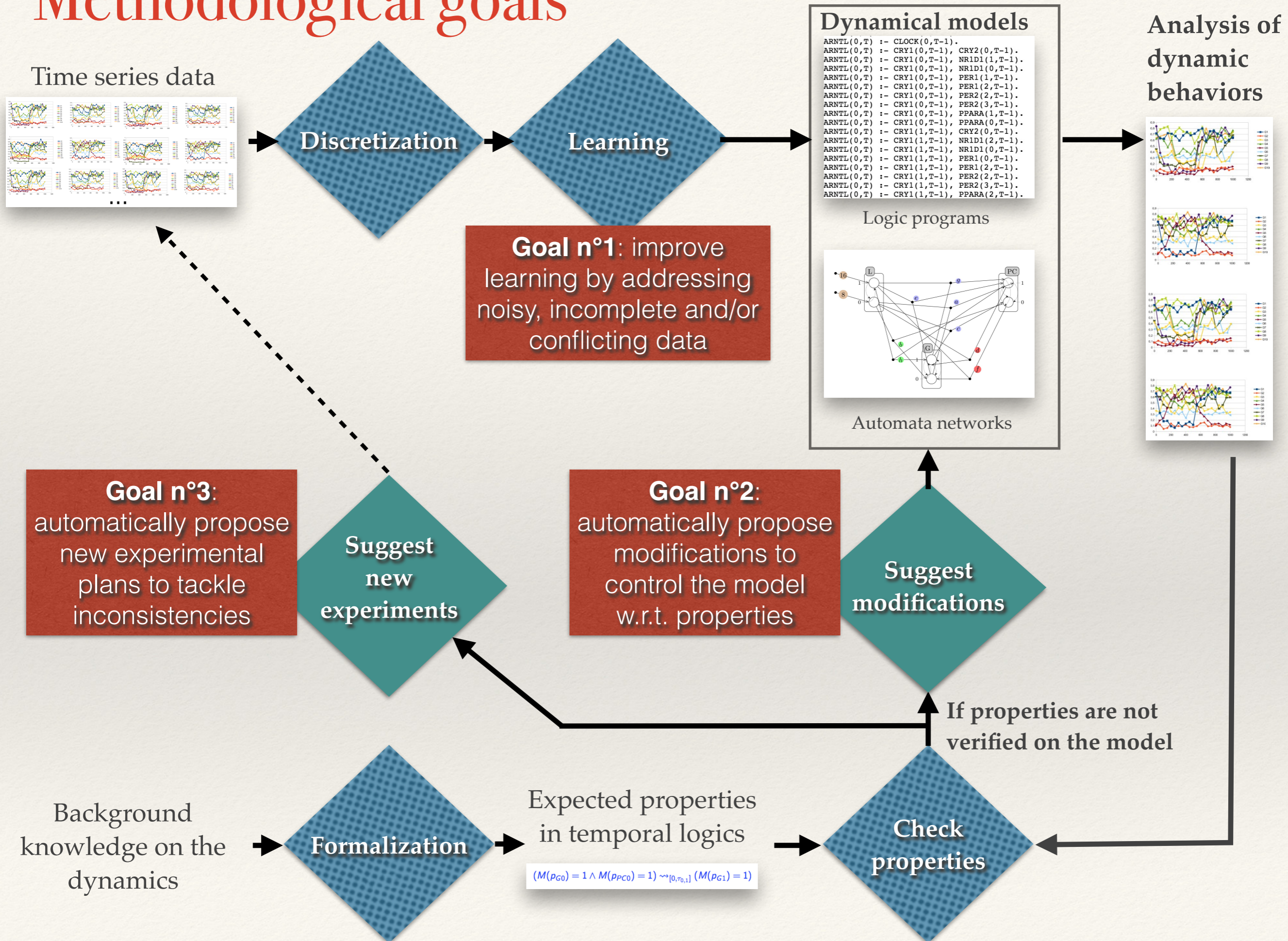
$$(M(p_{G0}) = 1 \wedge M(p_{PC0}) = 1) \rightsquigarrow_{[0,70,1]} (M(p_{G1}) = 1)$$



If properties are not verified on the model

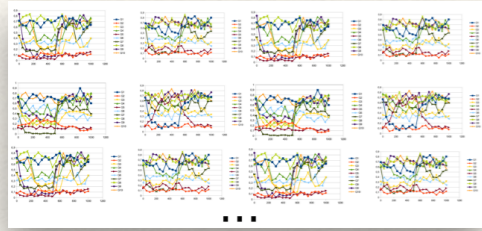


Methodological goals



Current works

Time series data



Discretization

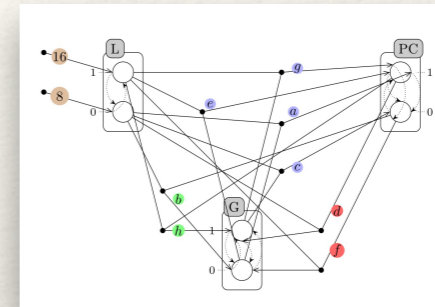
Learning

Dynamical models

```

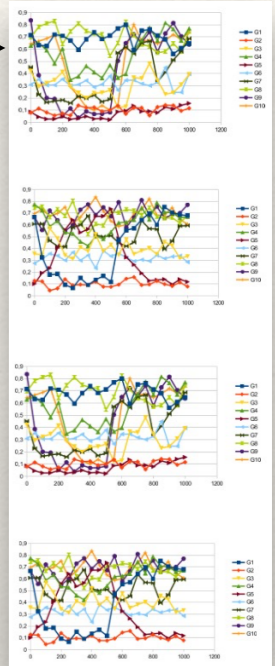
ARNTL(0,T) :- CLOCK(0,T-1).
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```

Logic programs



Automata networks

Analysis of dynamic behaviors



Suggest new experiments

Suggest modifications

If properties are not verified on the model

Additional (new) knowledge on the dynamics

Formalization

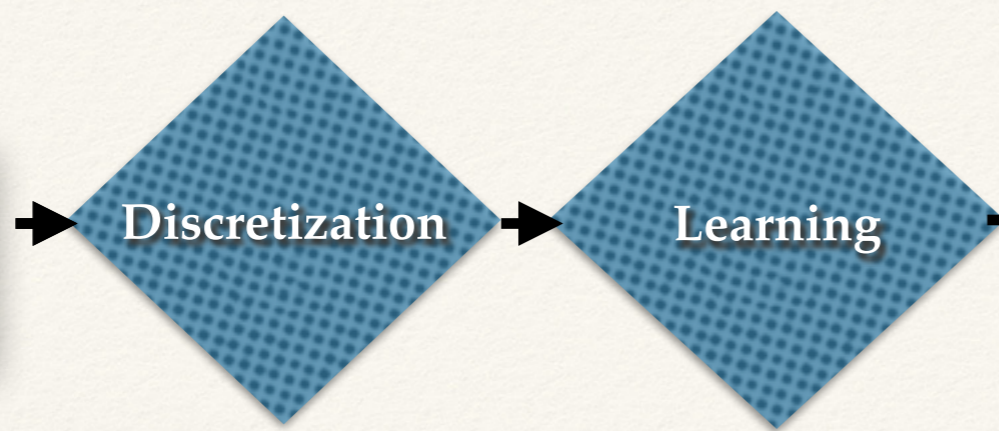
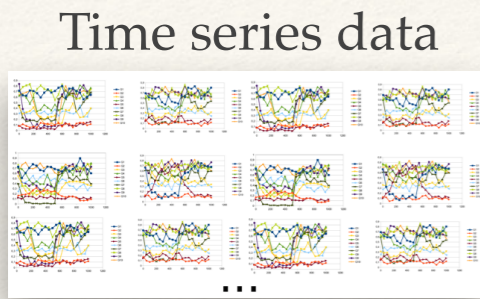
Properties in temporal logics

$$(M(p_{G0}) = 1 \wedge M(p_{PC0}) = 1) \rightsquigarrow_{[0, \tau_0, 1]} (M(p_{G1}) = 1)$$

Check properties

Current works

Extension of LFIT to generalized semantics



Dynamical models

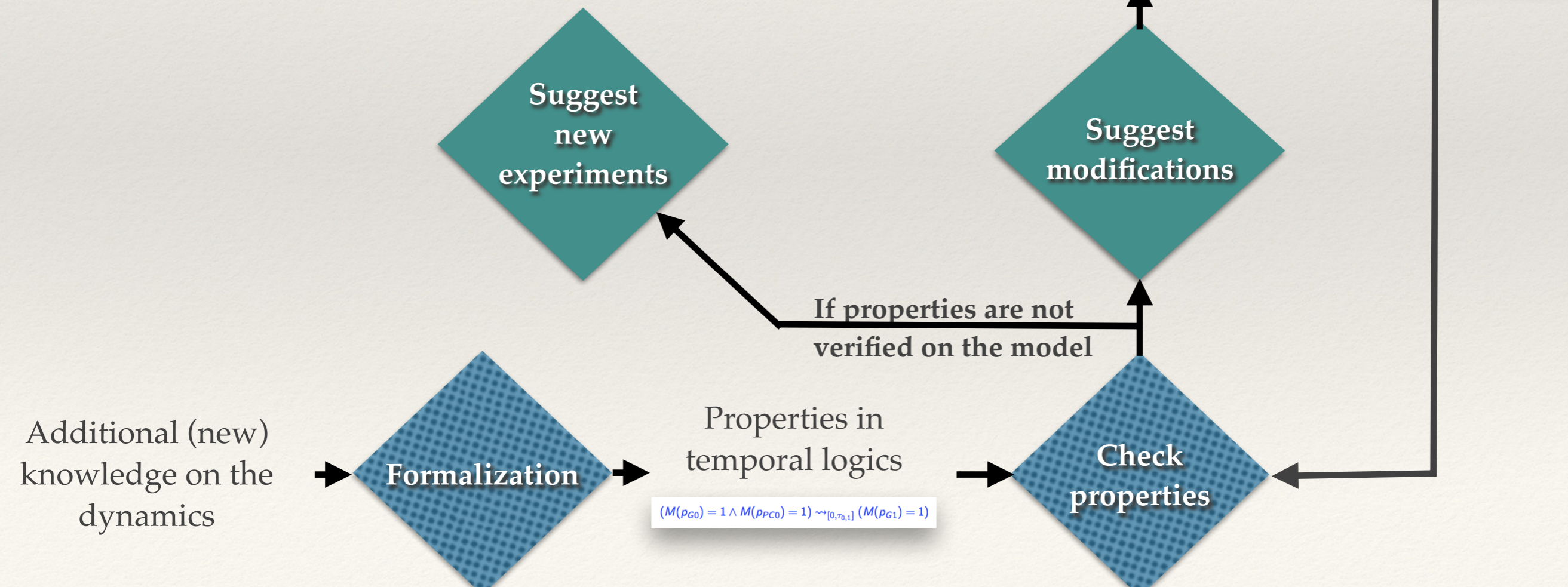
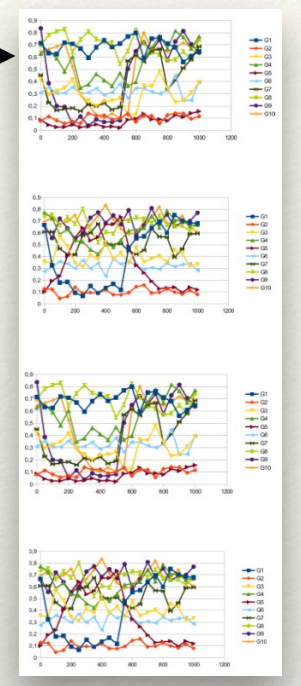
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ARNTL(0,T) :- CRY1(0,T-1), NR1D1(1,T-1).
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Logic programs

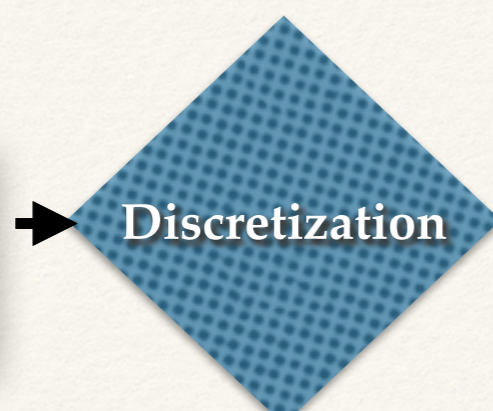
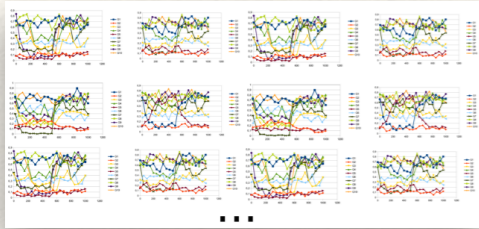
Automata networks

Analysis of dynamic behaviors

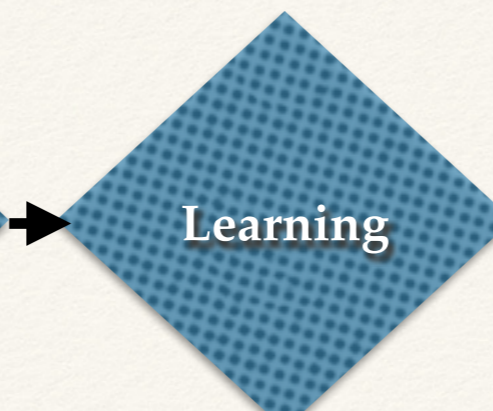


Current works

Time series data



Extension of LFIT to generalized semantics

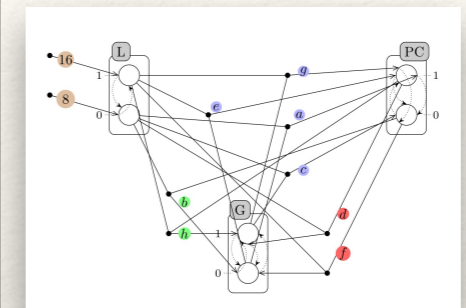


Dynamical models

```

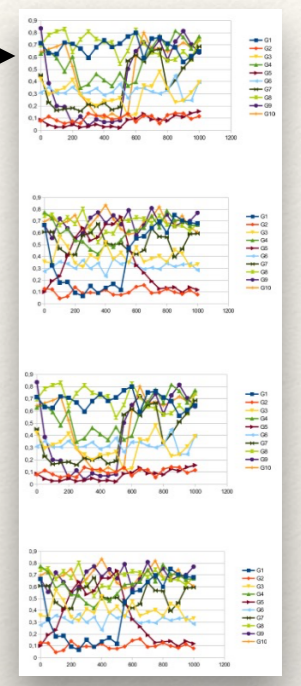
ARNTL(0,T) :- CLOCK(0,T-1).
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```

Logic programs



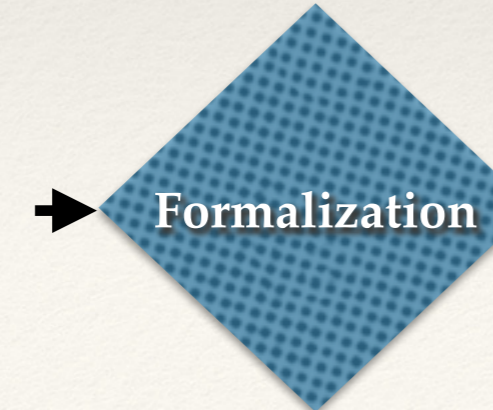
Automata networks

Analysis of dynamic behaviors



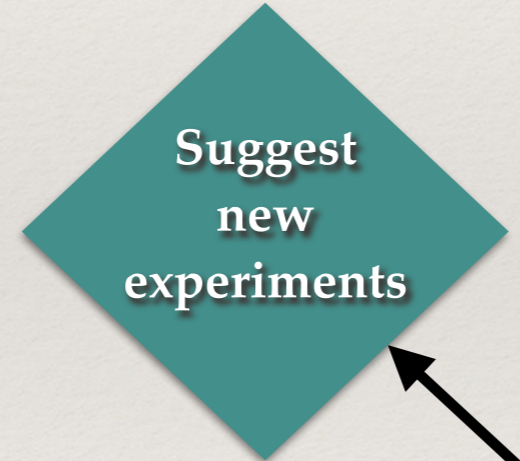
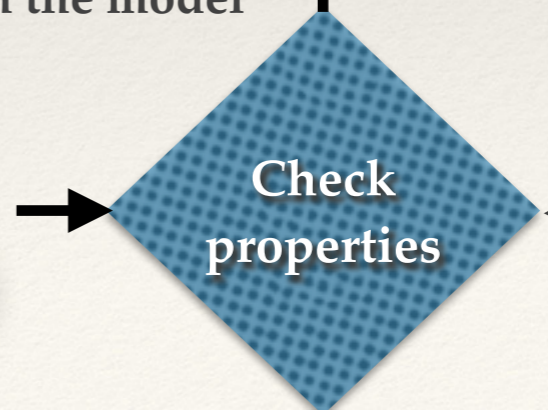
Heuristics to improve reachability analysis

Additional (new) knowledge on the dynamics



Properties in temporal logics

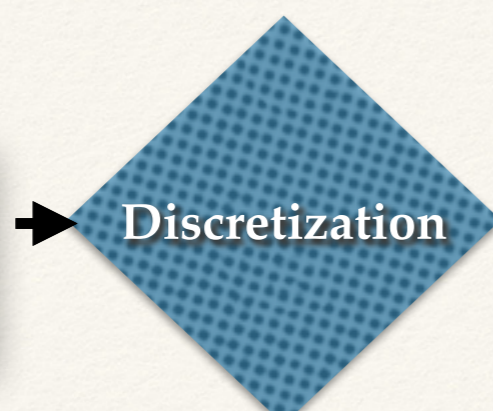
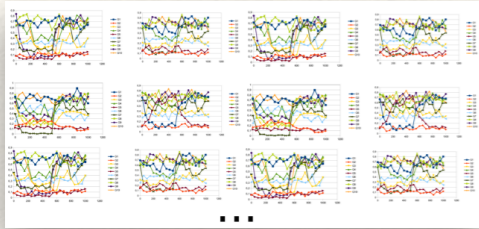
$$(M(p_{G0}) = 1 \wedge M(p_{PC0}) = 1) \rightsquigarrow_{[0, \tau_0, 1]} (M(p_{G1}) = 1)$$



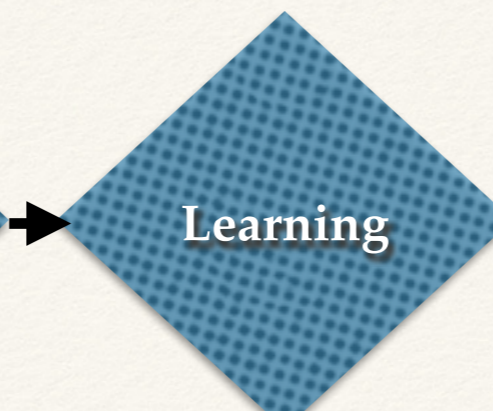
If properties are not verified on the model

Current works

Time series data



Extension of LFIT to generalized semantics

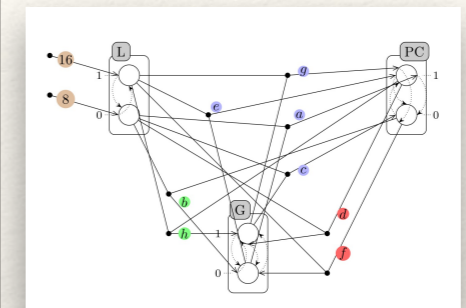


Dynamical models

```

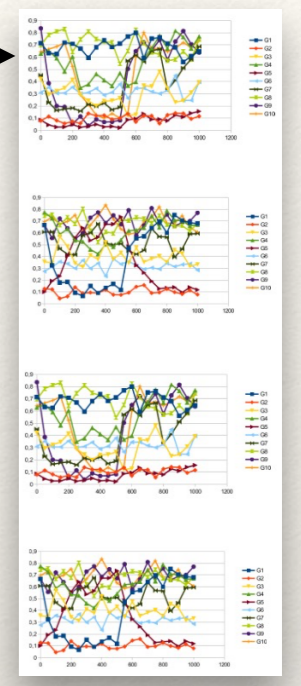
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Logic programs



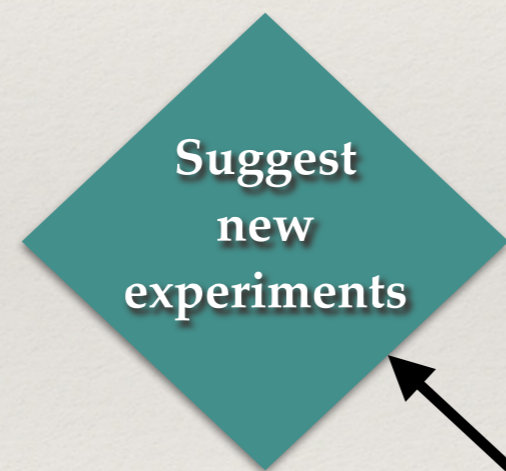
Automata networks

Analysis of dynamic behaviors



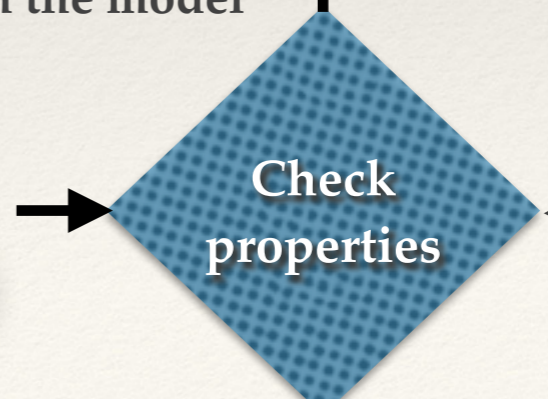
Heuristics to improve reachability analysis

Use of (i) ASP (ii) SAT-solvers to analyze reachability



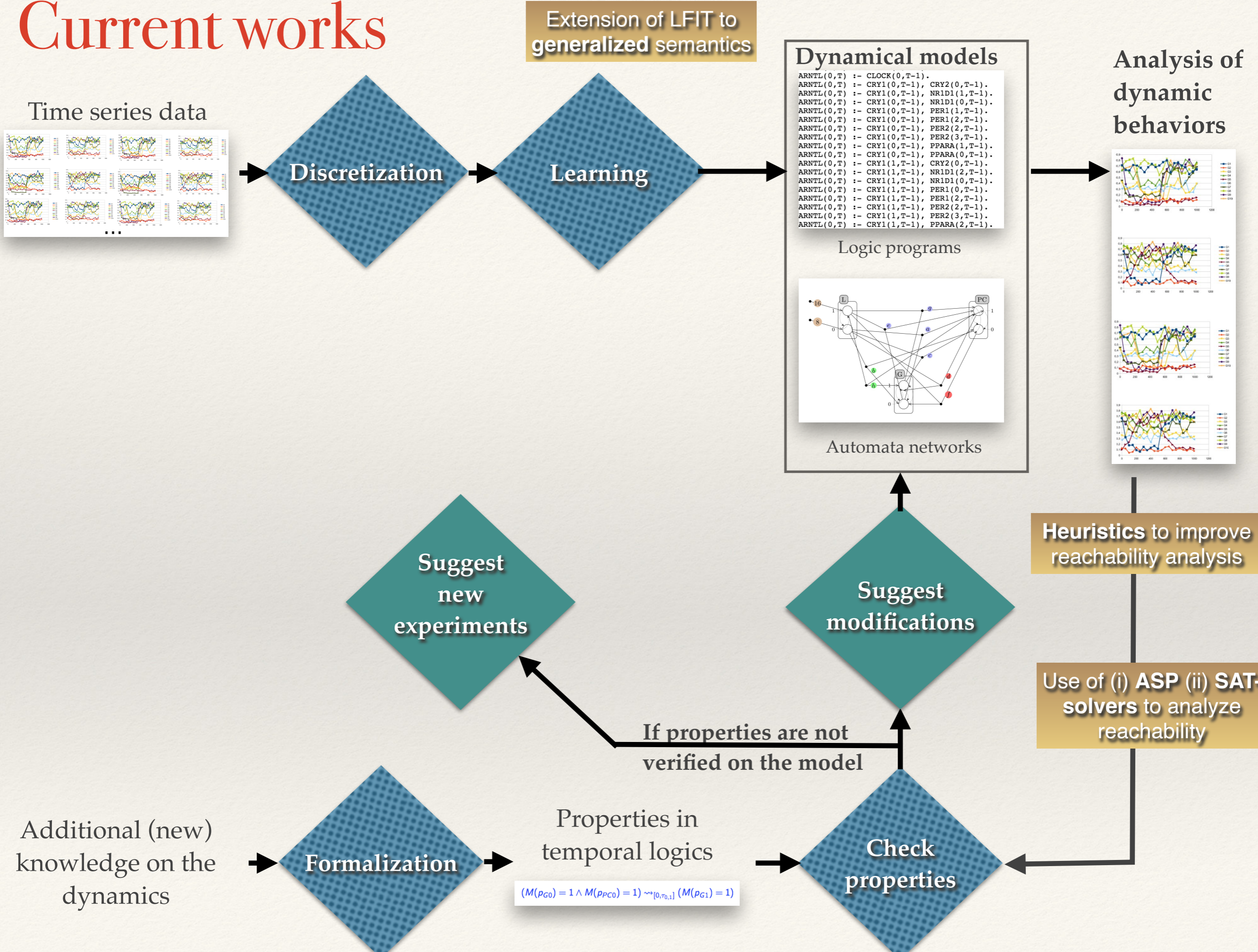
Properties in temporal logics

$$(M(p_{G0}) = 1 \wedge M(p_{PC0}) = 1) \rightsquigarrow_{[0, \tau_0, 1]} (M(p_{G1}) = 1)$$

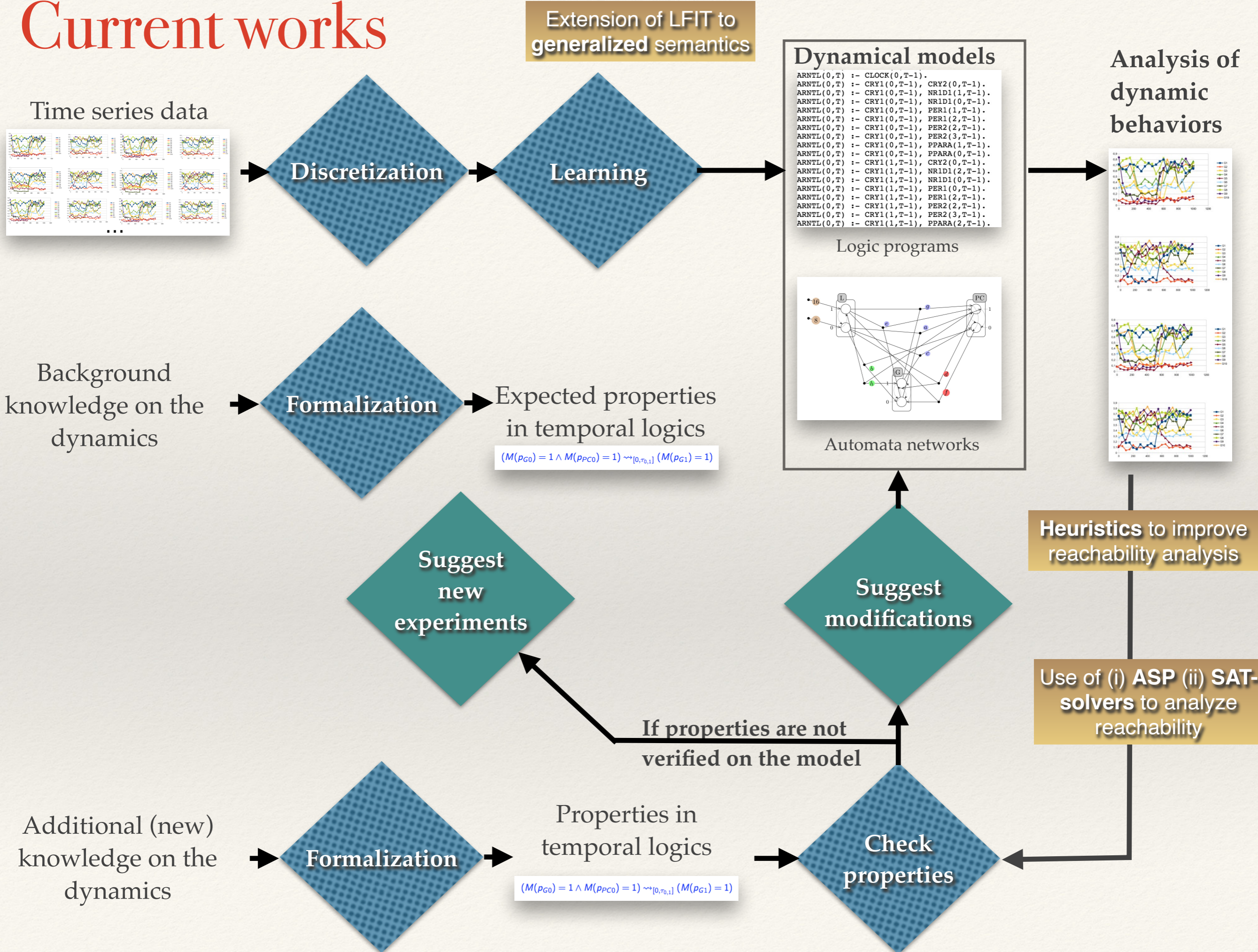


Additional (new) knowledge on the dynamics

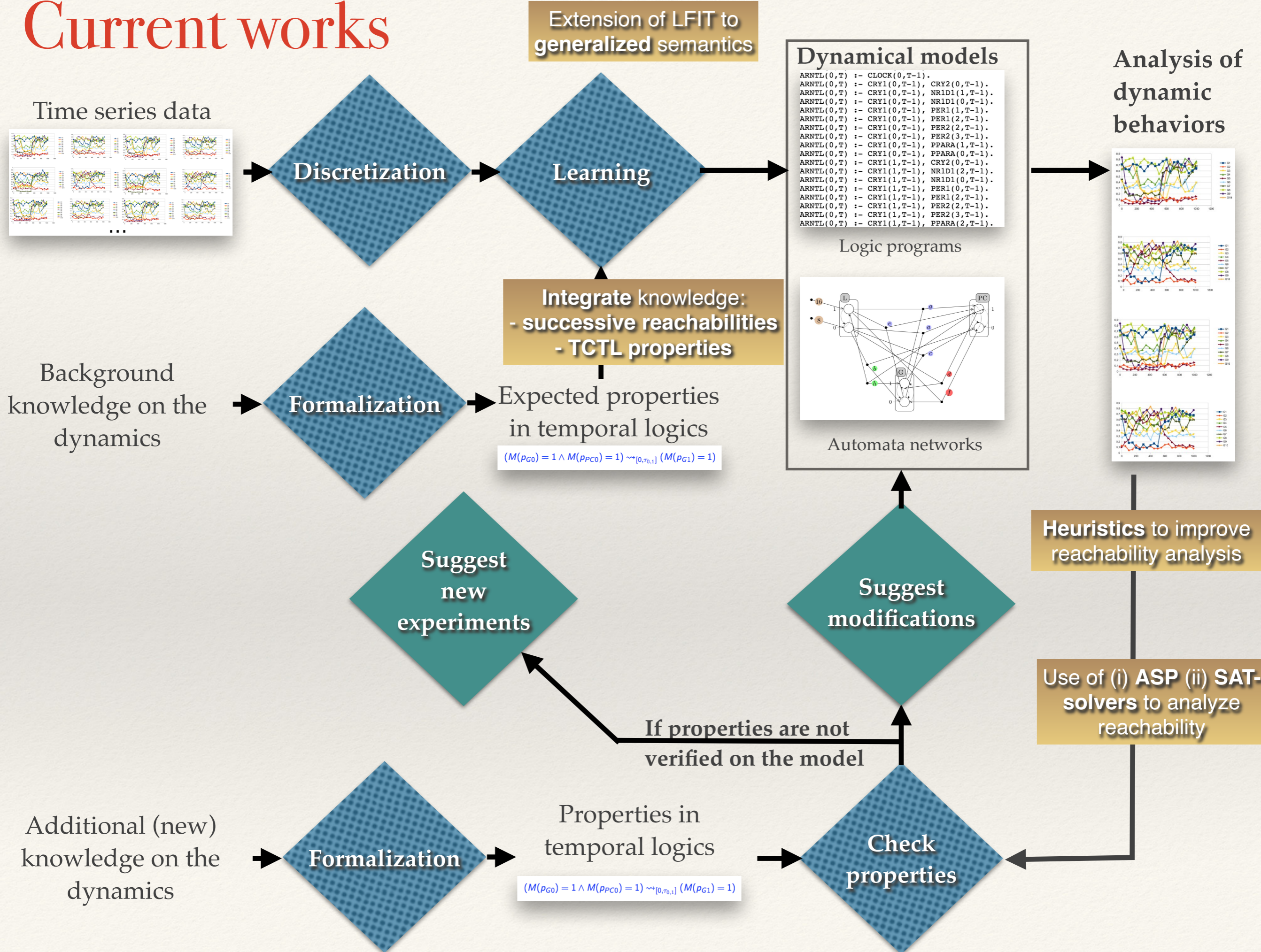
If properties are not verified on the model



Current works



Current works

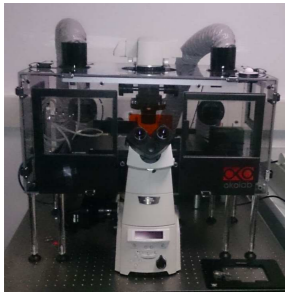


With biologists working on radiotherapy

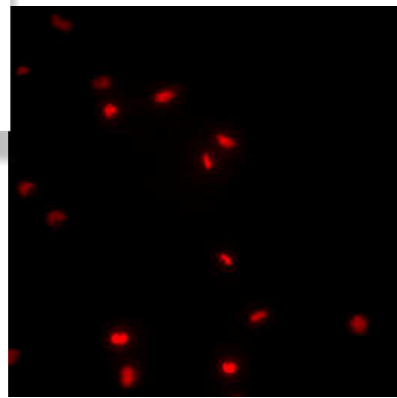
Team of François Paris
(INSERM Nantes)

- ❖ Goals:
 - ❖ Gain **better knowledge** of the radiobiology in function of the radiation therapy (RT) schemes & time
 - ❖ **Assess *in silico*** the effects of new RT protocols, reducing the need of *in vivo* experiments and helping oncologists on their clinical decision allowing personalized treatments

Time lapse



1 image every
10 minutes



1 RT = 1 timelapse

Learning

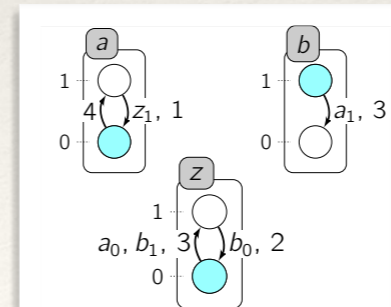
Each cell has a
finite number
of states (<10)

Dynamical models

```

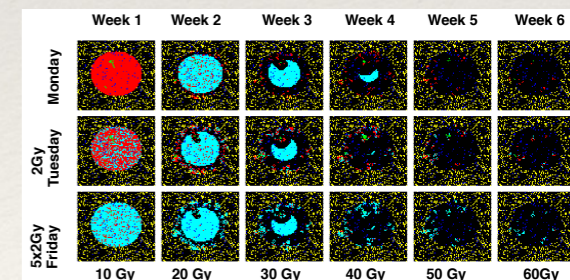
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```

Logic program



Automata networks

Simulation of
novel radiation
protocols

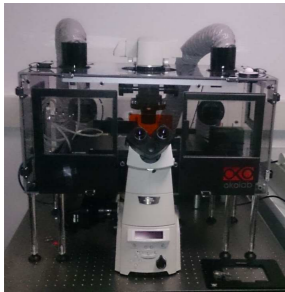


With biologists working on radiotherapy

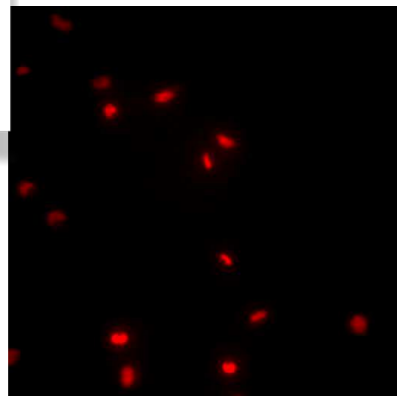
Team of François Paris
(INSERM Nantes)

- ❖ New challenges w.r.t. LFIT:
 - ❖ Model cell **division** (*mitosis*)
 - ❖ Model **spatial interactions** between cells

Time lapse



1 image every
10 minutes



1 RT = 1 timelapse

Learning

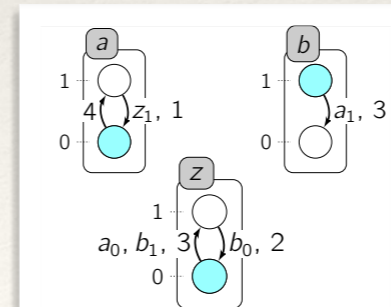
Each cell has a
finite number
of states (<10)

Dynamical models

```

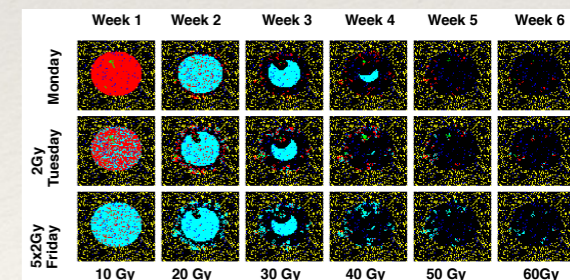
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Logic program



Automata networks

Simulation of
novel radiation
protocols



Références

- ❖ [BFR+15] Abdallah, E. B., Folschette, M., Roux, O., & Magnin, M. (2015, November). Exhaustive analysis of dynamical properties of Biological Regulatory Networks with Answer Set Programming. In *Bioinformatics and Biomedicine (BIBM), 2015 IEEE International Conference on* (pp. 281-285). IEEE.
- ❖ [FPI+12] Folschette, M., Paulevé, L., Inoue, K., Magnin, M., & Roux, O. (2012). Concretizing the process hitting into biological regulatory networks. In *Computational methods in systems biology* (pp. 166-186). Springer Berlin Heidelberg.
- ❖ [FMP+13] Folschette, M., Paulevé, L., Magnin, M., & Roux, O. (2013). Under-approximation of reachability in multivalued asynchronous networks. *Electronic Notes in Theoretical Computer Science*, 299, 33-51.
- ❖ [FPI+15] Folschette, M., Paulevé, L., Inoue, K., Magnin, M., & Roux, O. (2015). Identification of biological regulatory networks from Process Hitting models. *Theoretical Computer Science*, 568, 49-71.
- ❖ [FPM+15] Folschette, M., Paulevé, L., Magnin, M., & Roux, O. (2015). Sufficient conditions for reachability in automata networks with priorities. *Theoretical Computer Science*, 608, 66-83.

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- ❖ [PMR11a] Paulevé, L., Magnin, M., & Roux, O. (2011). Refining dynamics of gene regulatory networks in a stochastic π -calculus framework. In *Transactions on computational systems biology xiii* (pp. 171-191). Springer Berlin Heidelberg.
- ❖ [PMR11b] Paulevé, L., Magnin, M., & Roux, O. (2011). Tuning temporal features within the stochastic π -calculus. *Software Engineering, IEEE Transactions on*, 37(6), 858-871.
- ❖ [PMR12] Paulevé, L., Magnin, M., & Roux, O. (2012). Static analysis of biological regulatory networks dynamics using abstract interpretation. *Mathematical Structures in Computer Science*, 22(04), 651-685.
- ❖ [PCF+14] Paulevé, L., Chancellor, C., Folschette, M., Magnin, M., & Roux, O. (2014). Analyzing large network dynamics with process hitting. *Logical Modeling of Biological Systems*, 125-166.
- ❖ [RMI+15a] Ribeiro, T., Magnin, M., Inoue, K., & Sakama, C. (2015). Learning delayed influences of biological systems. *Frontiers in bioengineering and biotechnology*, 2.
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