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# **Two Qualitative Dynamics Semantics for SBGN Process Description Maps**

**Adrien Rougny, Christine Froidevaux and Loic Paulevé** 

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## Context

- □ Larger and larger reaction networks modelling various biological processes (from databases, automatic inference)
- □ Standards to represent reaction networks: e.g. Systems Biology **Graphical Notation Process Description** language (SBGN-PD)
- □ Analysis of the dynamics to understand and control these processes

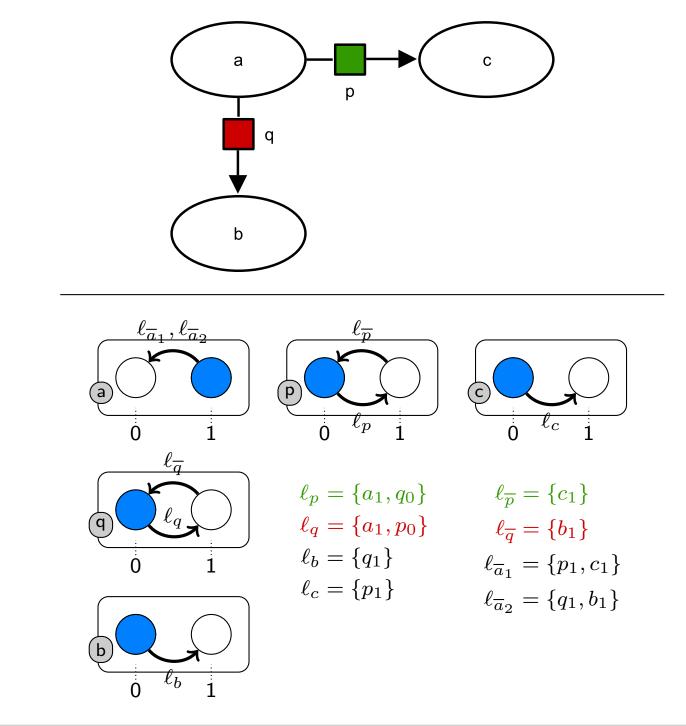
### Motivations

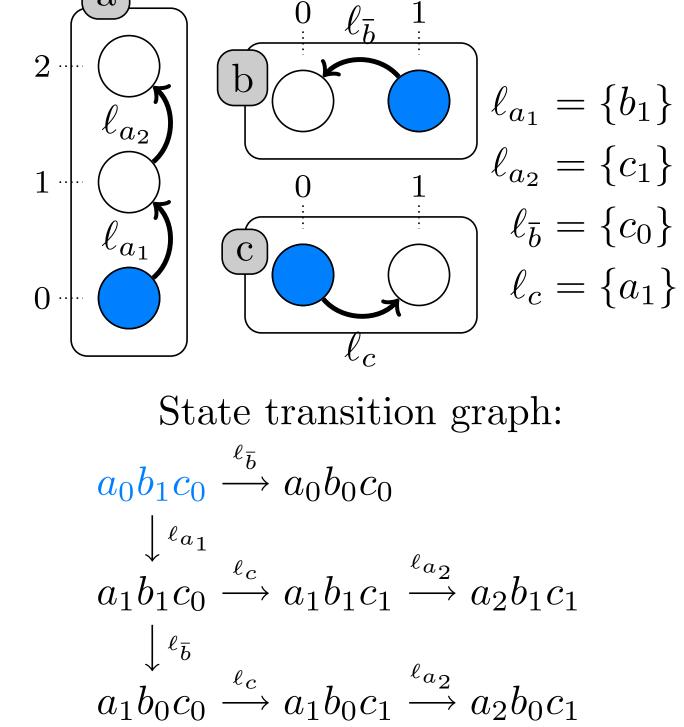
- □ Qualitative semantics allows to capture **important dynamical features** (e.g. attractors, reachability) without numerical parameters □ Model SBGN-PD maps under qualitative semantics
- We propose two semantics formalized by asynchronous automata **networks**: the *general semantics*, together with a refinement called process conflicts, and the stories semantics

### Refinement 1: Process Conflicts

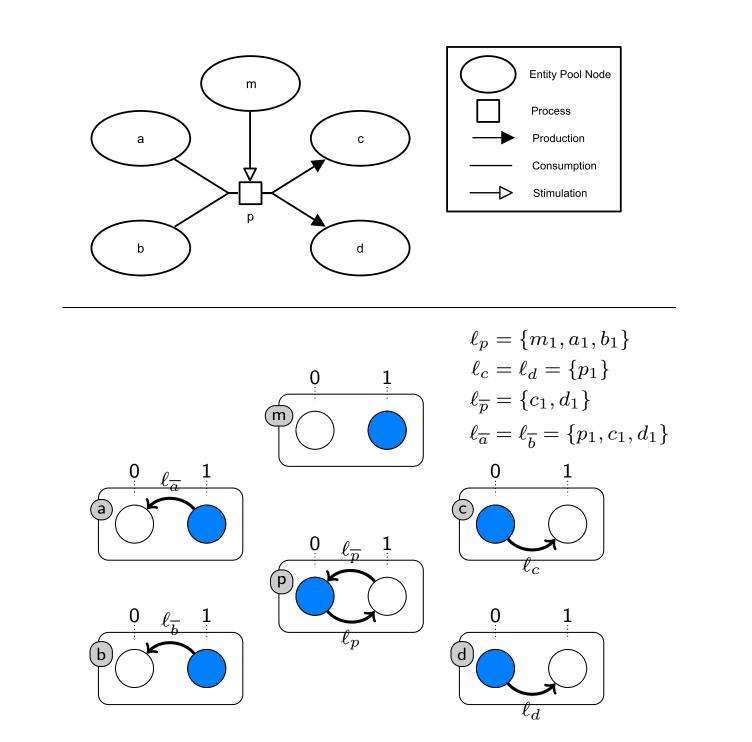
### (a)

### **General Semantics**





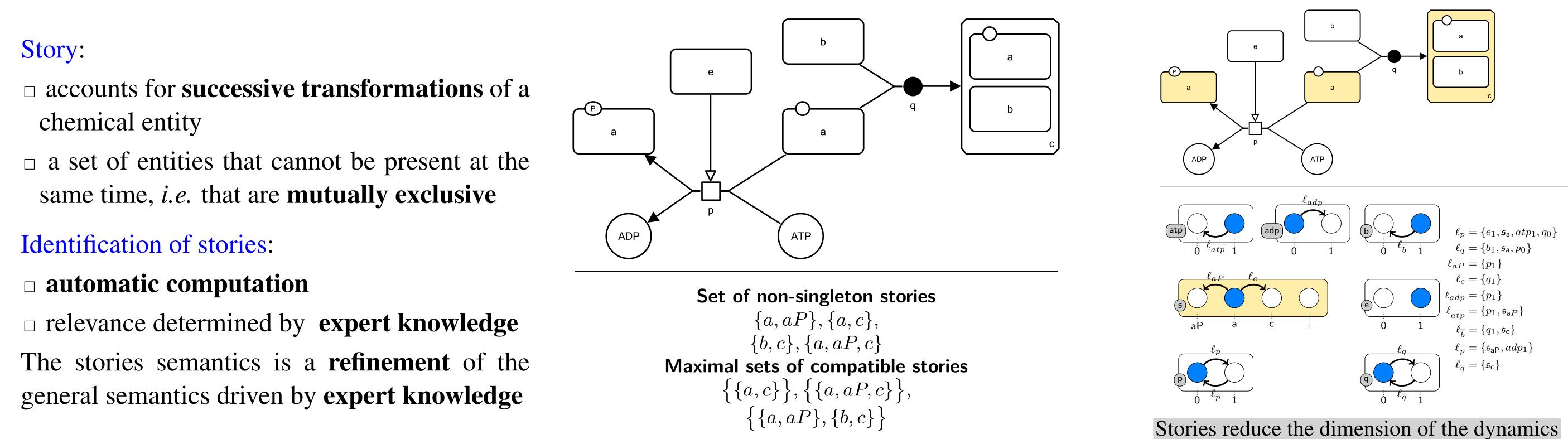
Asynchronous Automata Networks



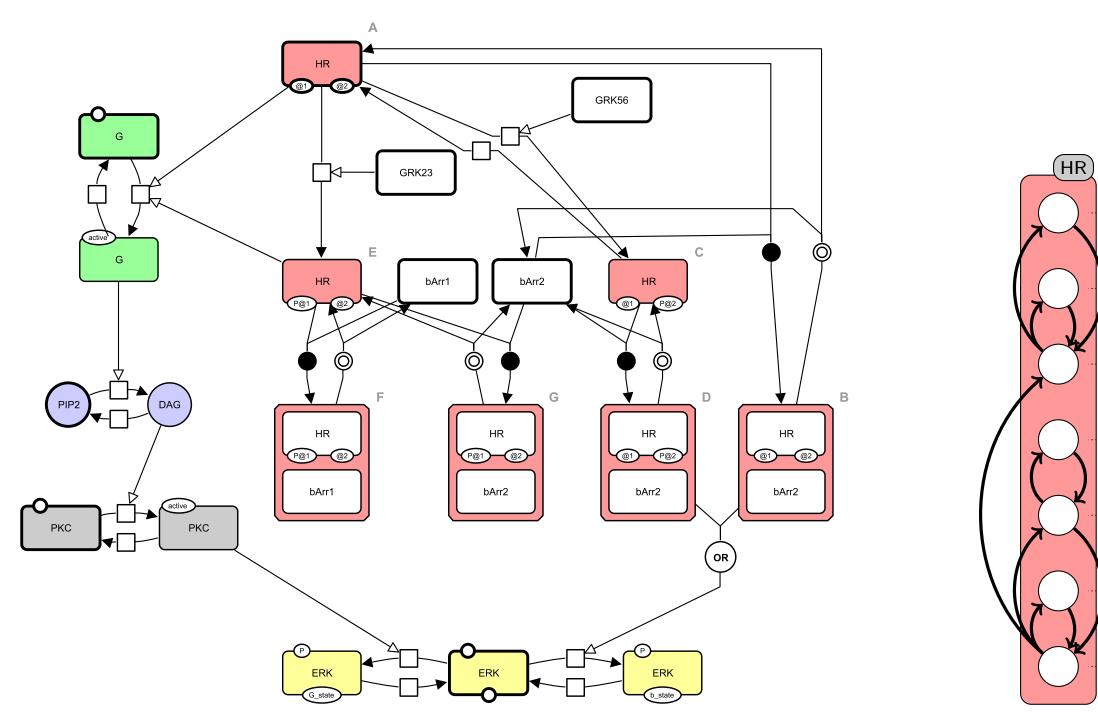
Each chemical entity is modelled by one automaton

Processes p and q cannot occur at the same time

## **Refinement 2: Stories Semantics**



### Example: the $AT_{1A}R$ -induced network



## **Conclusion and Prospects**

- □ **Main features** of SBGN-PD are supported
- $\square$  The stories semantics is a refinement of the general semantics that:
- models several chemical species with a unique variable
- reduces the state space, **increasing the scalability** of the exhaustive computation of the dynamics
- □ The dynamics of SBGN-PD maps modelled under both semantics can be analyzed with state-of-the art tools (PRISM, NuSMV ...)

SBGN-PD map adapted from: D. Heitzler et al., Competing G protein-coupled receptor kinases balance G protein and b-arrestin signaling, MSB, 2012.

semantics	general	general with conflicts	stories
No. of automata	41	41	30
No. of reachable states	$\simeq 10^{11}$	$\simeq 10^{10}$	$\simeq 10^5$

### Ongoing work and prospects:

□ Application to a **large network**, the E2F/RB pathway:

- -153 out of 208 molecules in 28 stories
- -No. of automata reduced from 367 to 242
- Explore the relationship between the stories semantics and **Boolean semantics** applied to **SBGN-AF maps**





