

# **PART I - TALK 3**

## **INTRODUCTION INTO ADVANCED CONCEPTS - STOCHASTIC/CONTINUOUS PETRI NETS -**

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Brandenburg University of Technology Cottbus**

# THE FRAMEWORK

**QUALITATIVE**

**STOCHASTIC**

**CONTINUOUS**

**QUALITATIVE**

*time-free*

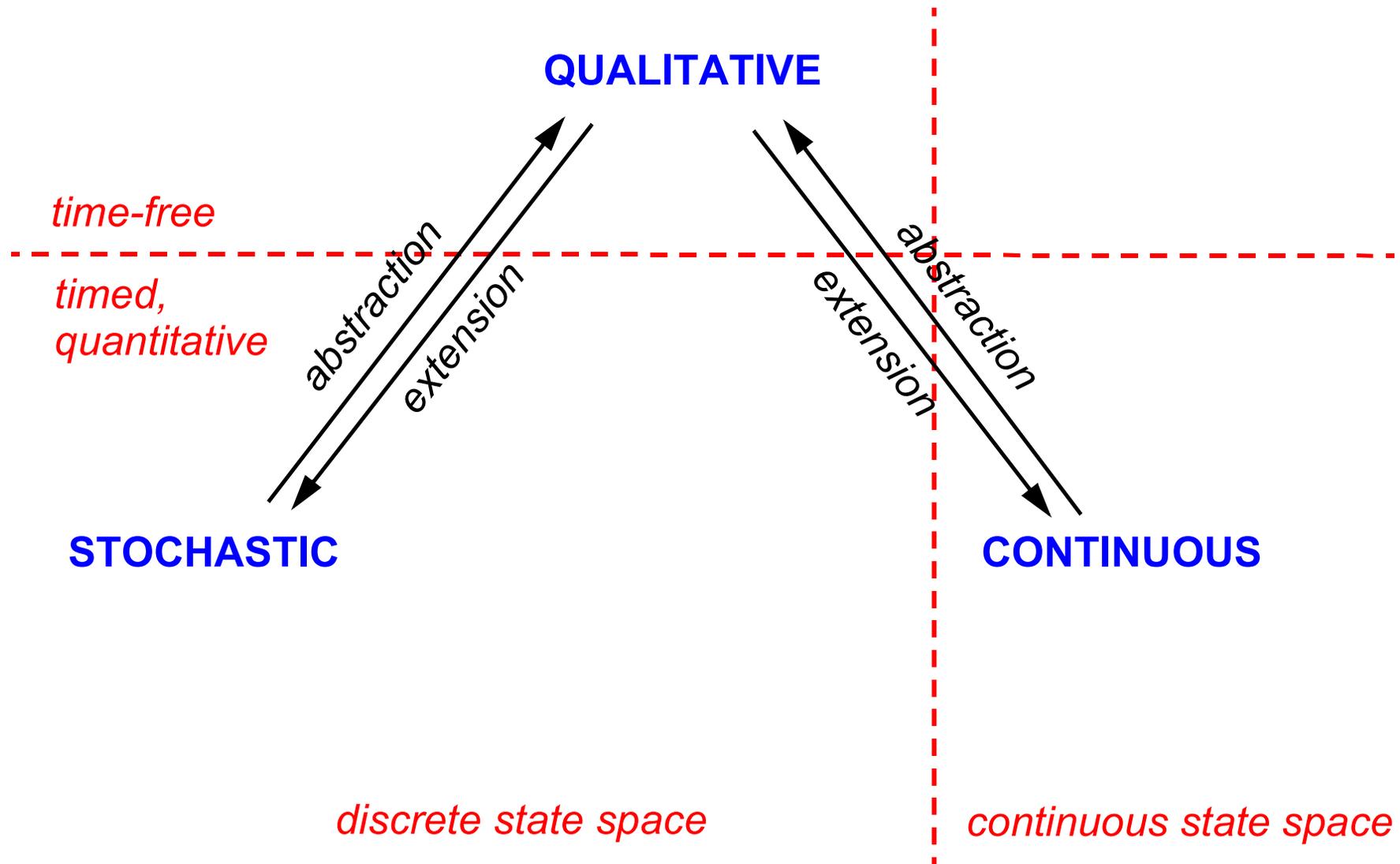
*timed,  
quantitative*

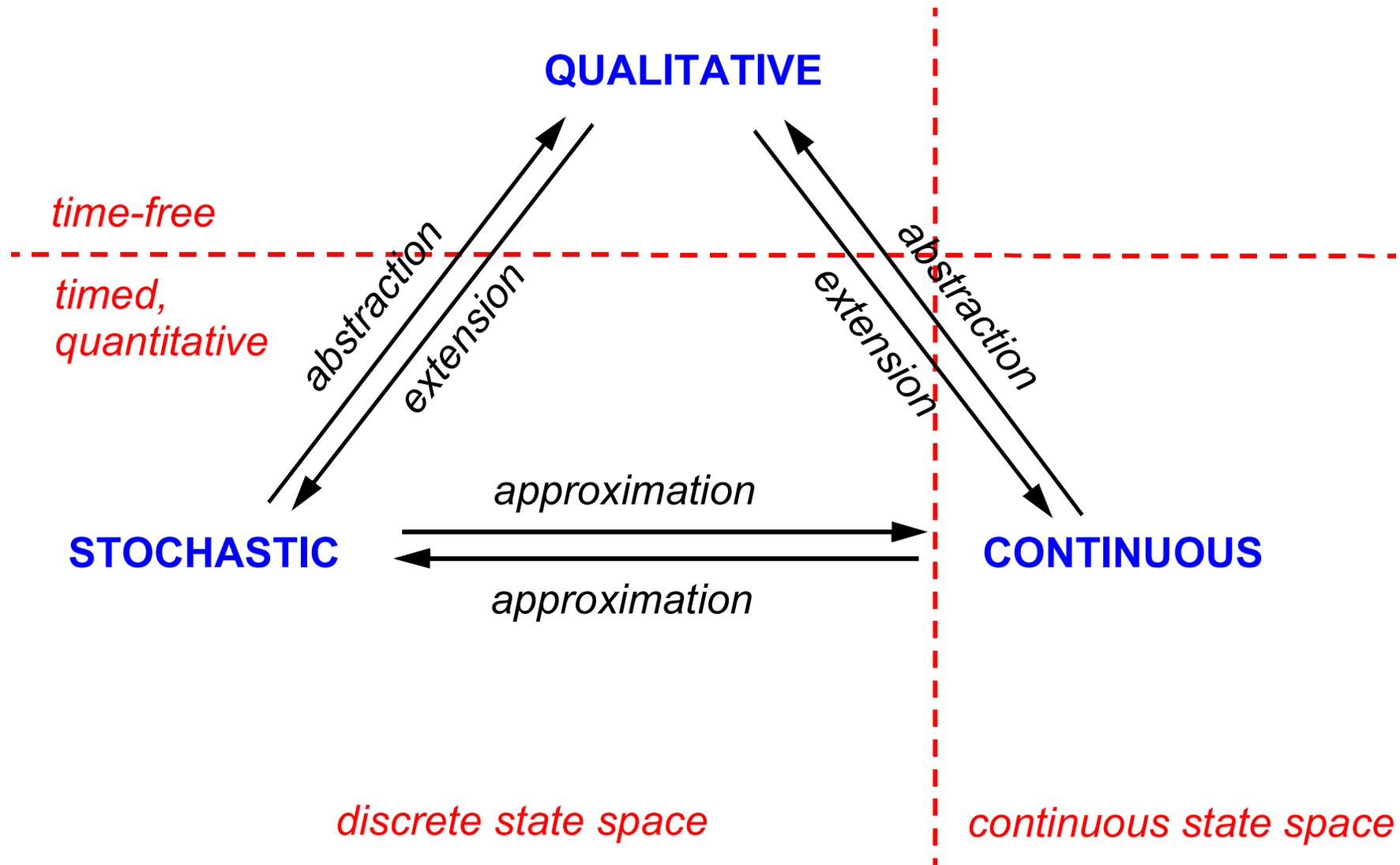
**STOCHASTIC**

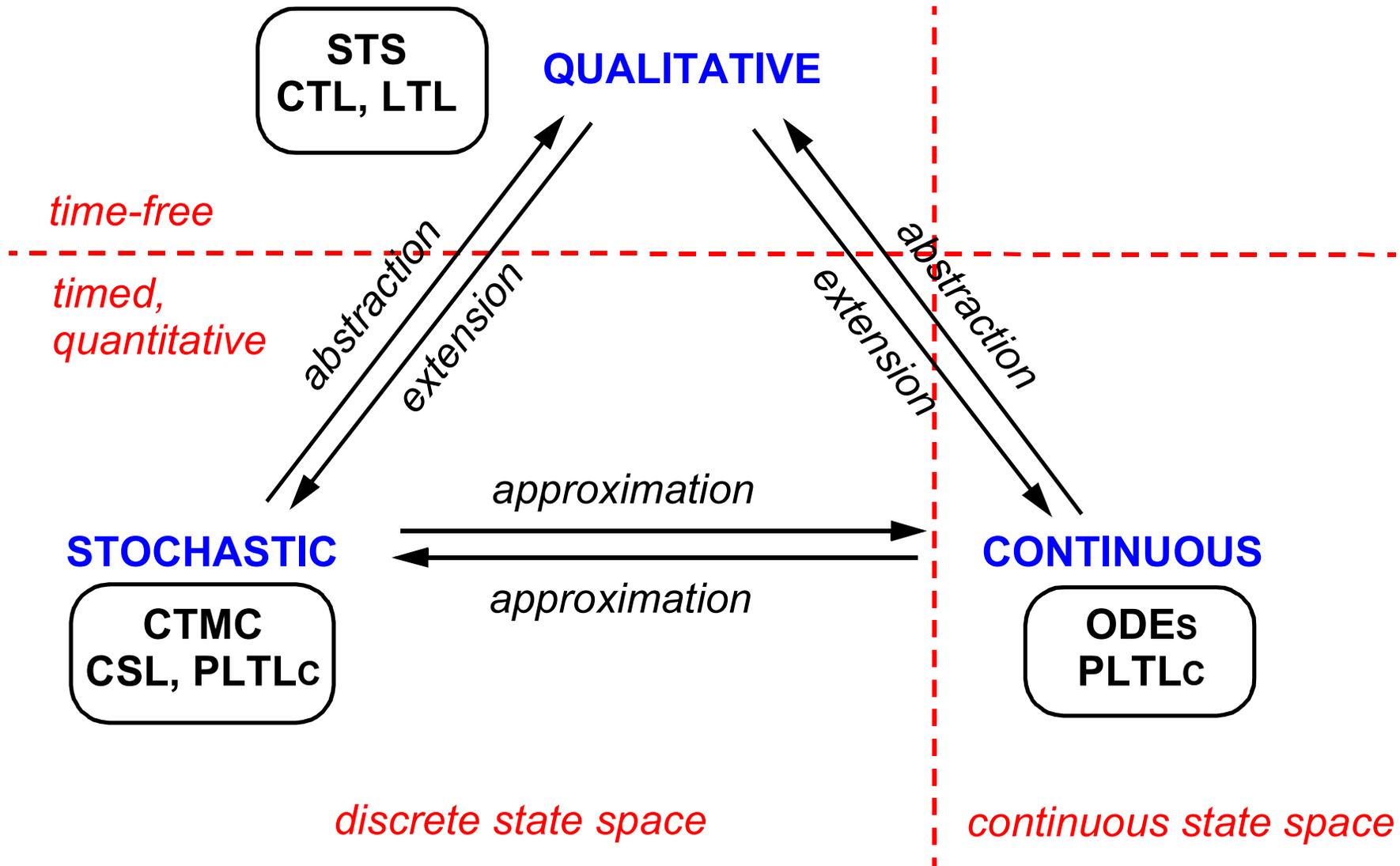
**CONTINUOUS**

*discrete state space*

*continuous state space*

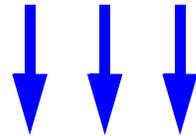






**3**

**MODELS SHARING STRUCTURE**



**QUANTITATIVE MODEL = QUALITATIVE MODEL  
+  
RATE FUNCTIONS  
(KINETICS)**

❑ **transitions get a stochastic waiting time**

-> *exponential distribution with parameter lambda*

❑ **state-dependent lambda defined by rate function**

-> *any arithmetic function including*

*the transition's pre-places as integer variables and  
user-defined real-valued parameters*

-> *modifier arcs*

-> *popular kinetics:*

*mass-action semantics, level semantics*

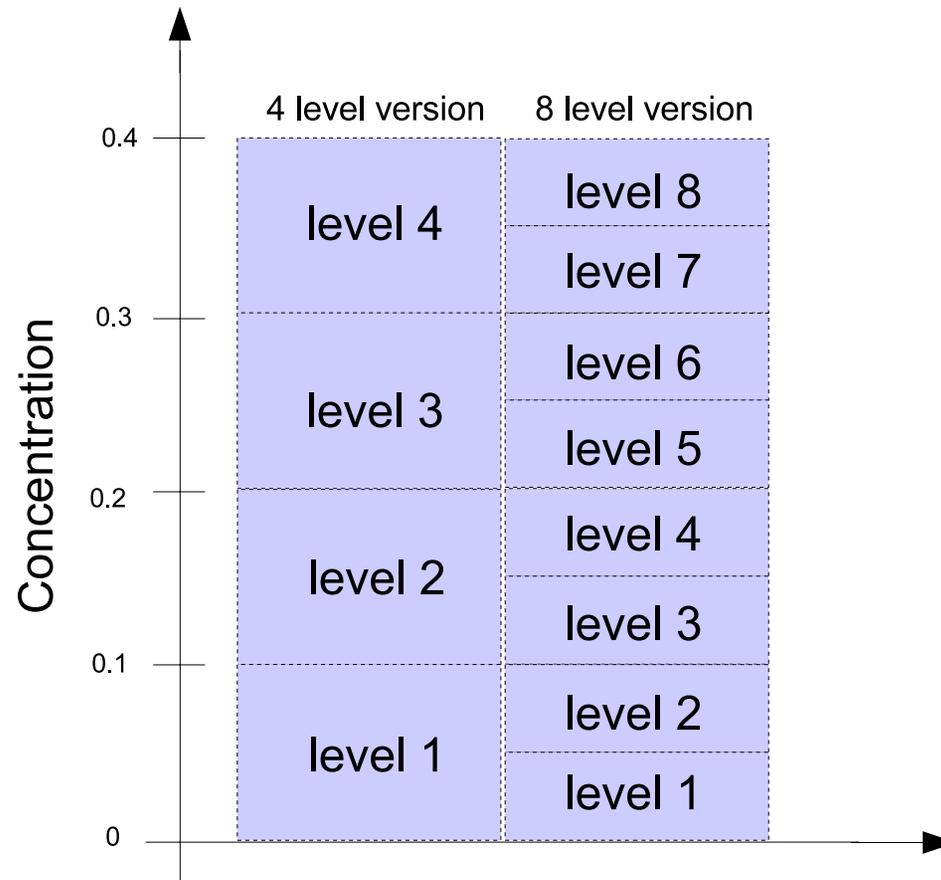
- ❑ **transitions get a stochastic waiting time**
  - > *exponential distribution with parameter lambda*
  
- ❑ **state-dependent lambda defined by rate function**
  - > *any arithmetic function including  
the transition's pre-places as integer variables and  
user-defined real-valued parameters*
  - > *modifier arcs*
  - > *popular kinetics:  
mass-action semantics, level semantics*
  
- ❑ **semantics: Continuous Time Markov Chain (CTMC)**
  - > *reachability graph + state transition rates*
  
- ❑ **analysis**
  - > *standard Markov analysis techniques: transient, steady state*
  - > *stochastic simulation algorithms (SSA), e.g. Gillespie's SSA*

## □ mass-action semantics

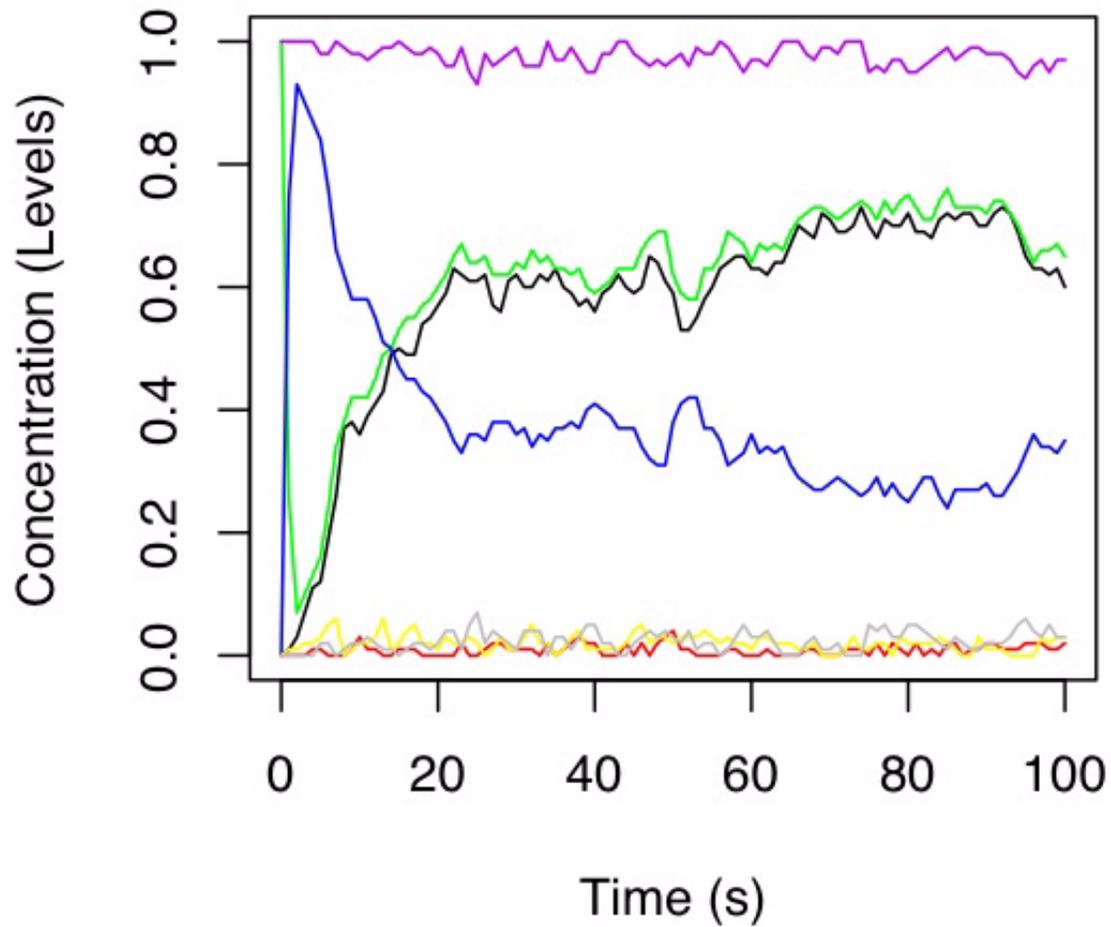
$$h_t := c_t \cdot \prod_{p \in \bullet t} \binom{m(p)}{f(p, t)}$$

## □ level semantics

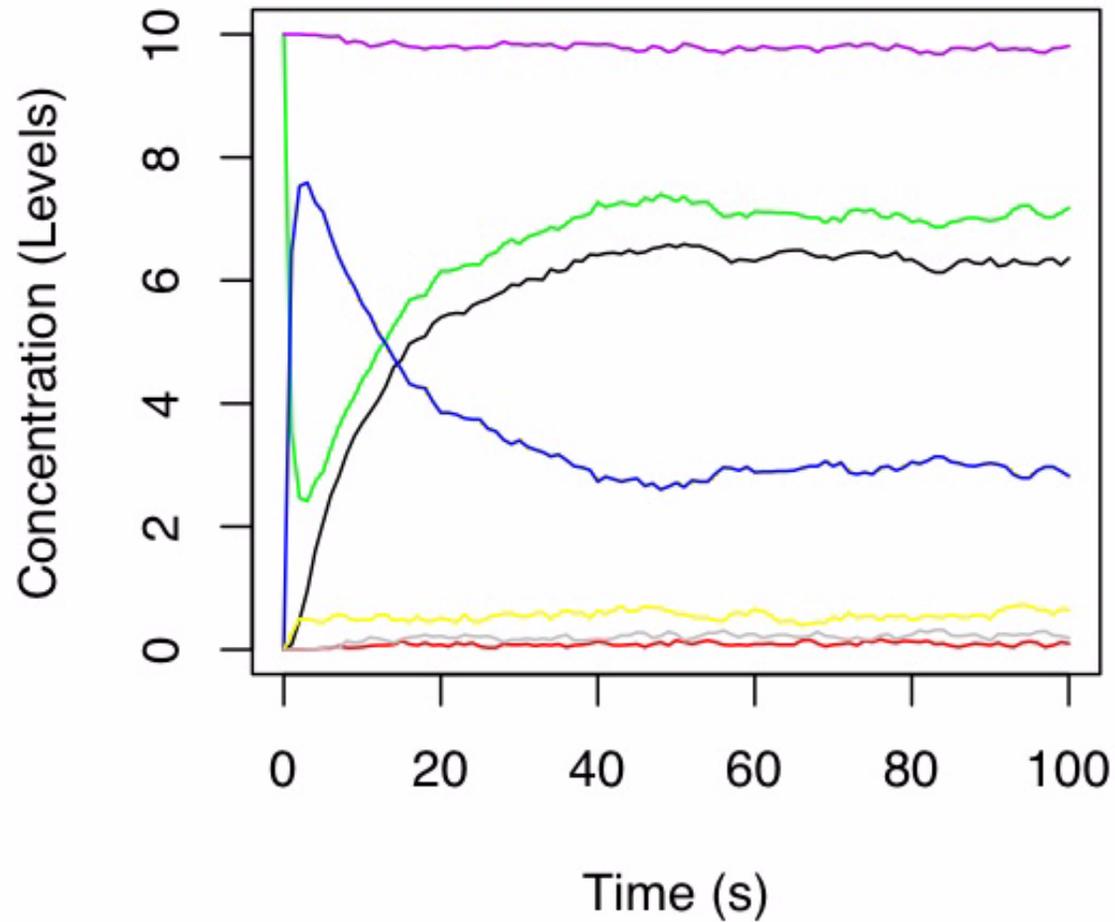
$$h_t := k_t \cdot N \cdot \prod_{p \in \bullet t} \left( \frac{m(p)}{N} \right)$$



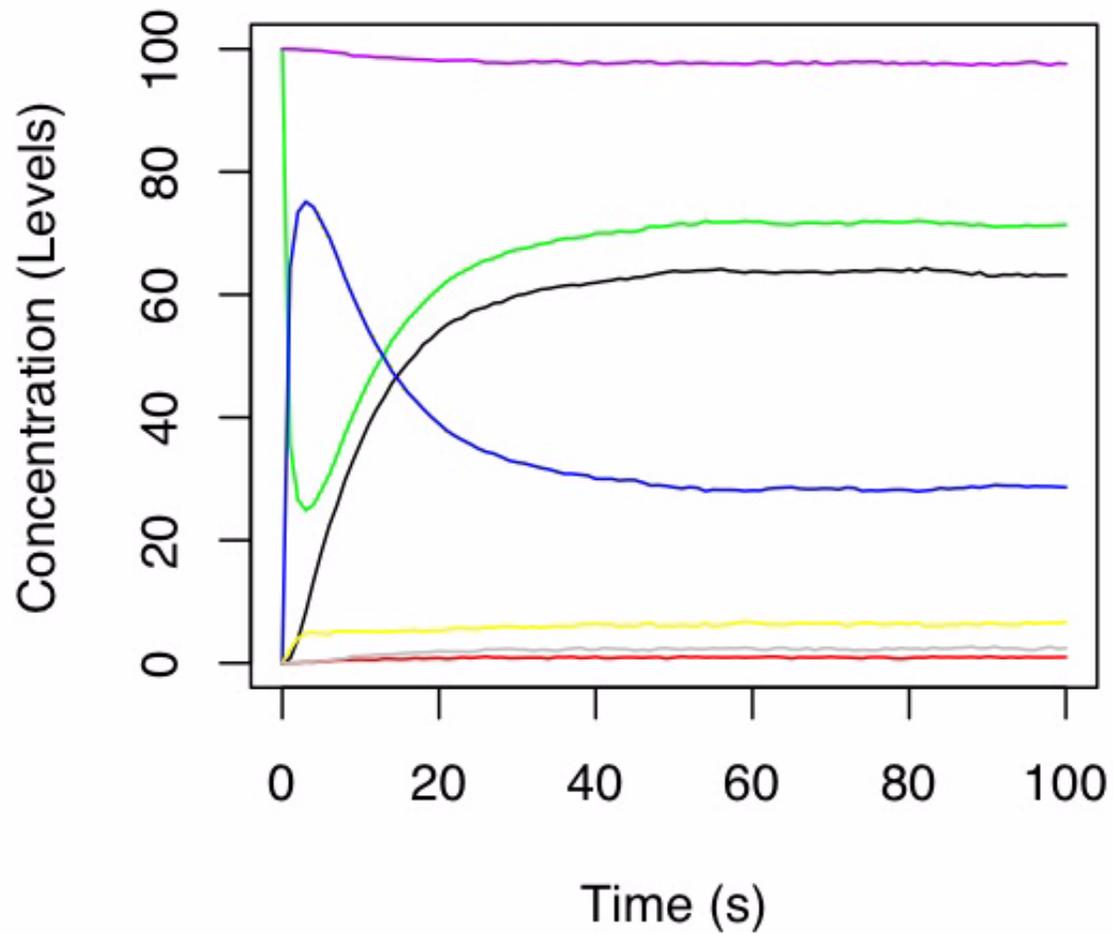
## Stochastic Output – 1 Level



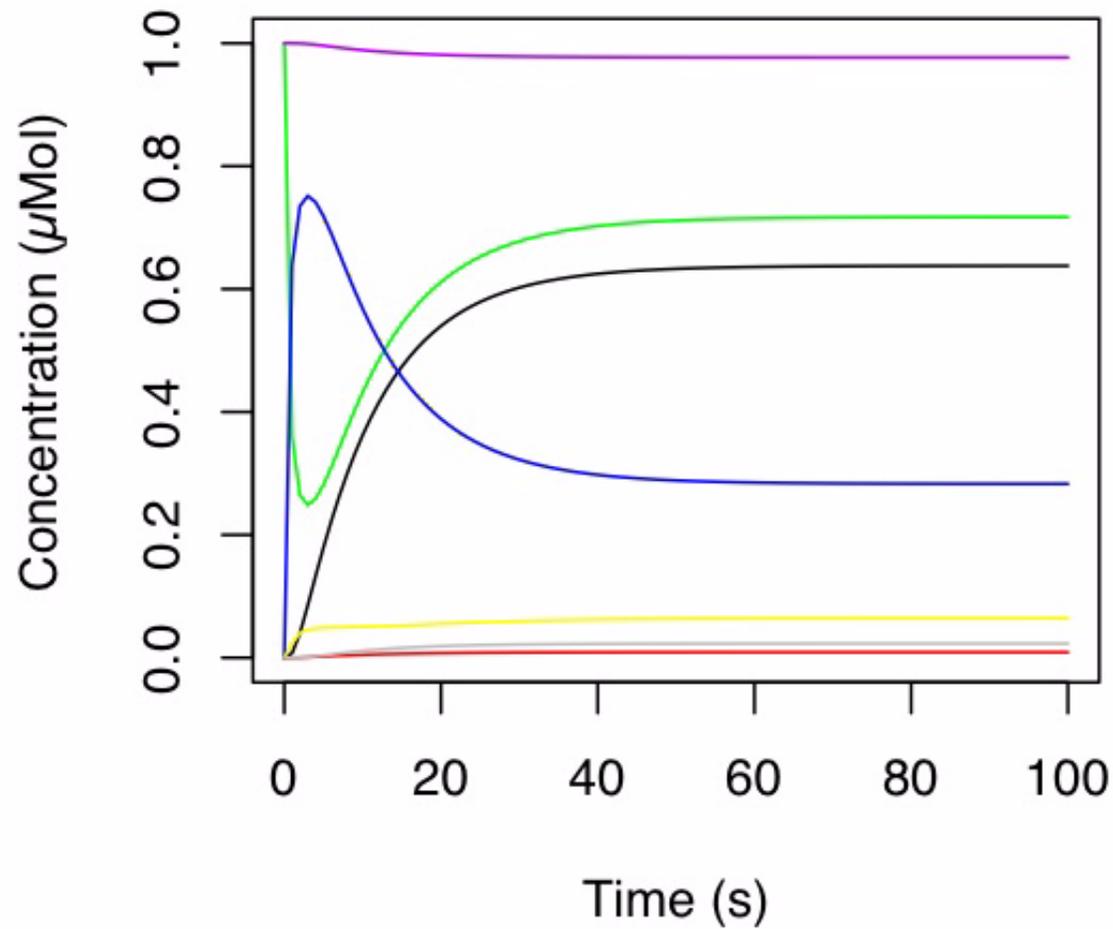
## Stochastic Output – 10 Levels



## Stochastic Output – 100 Levels



## Deterministic Output



❑ **transitions fire continuously**

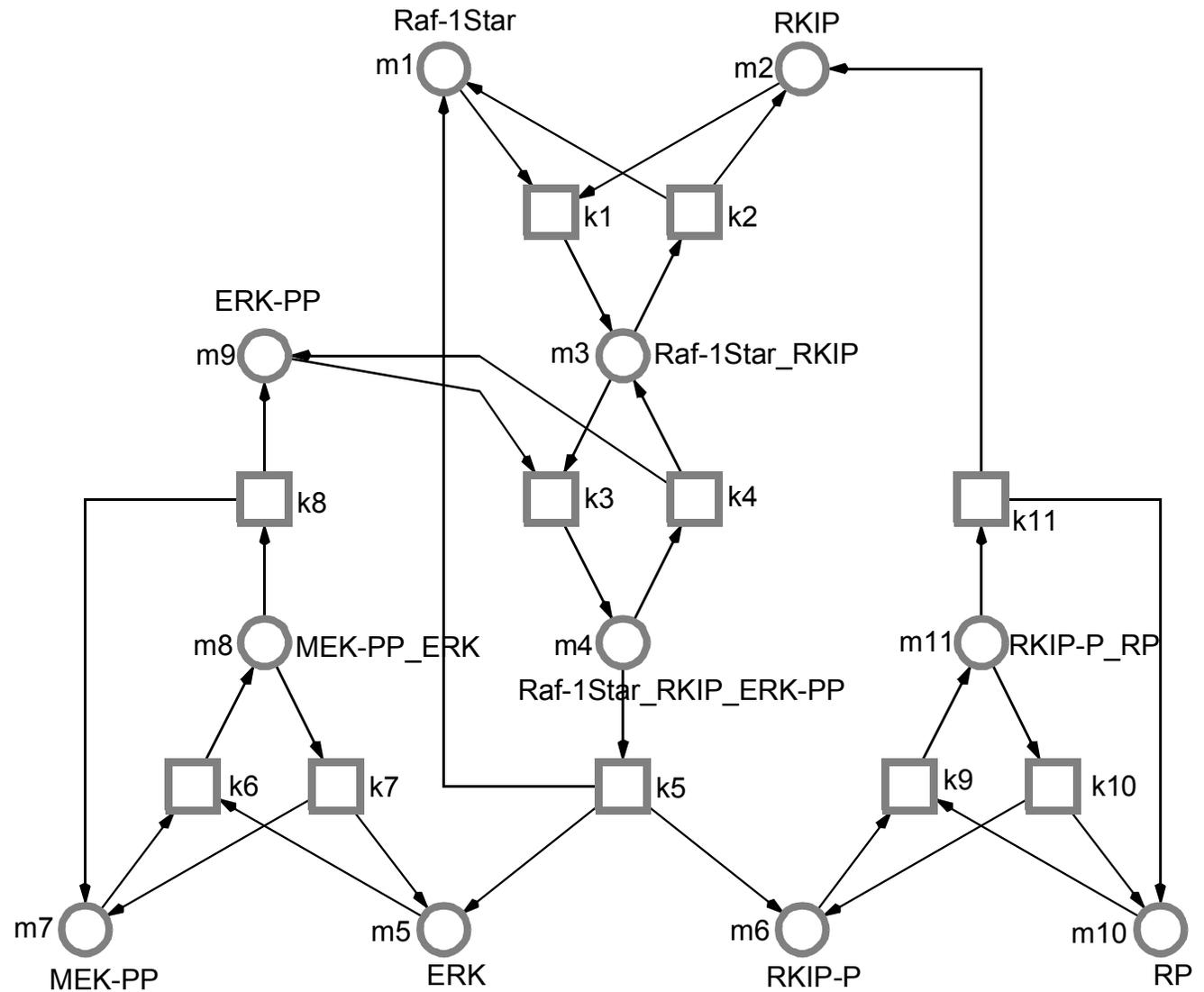
❑ **rate functions**

-> *any arithmetic function including  
the transition's pre-places as real-valued variables and  
user-defined real-valued parameters*

❑ **real-valued tokens**

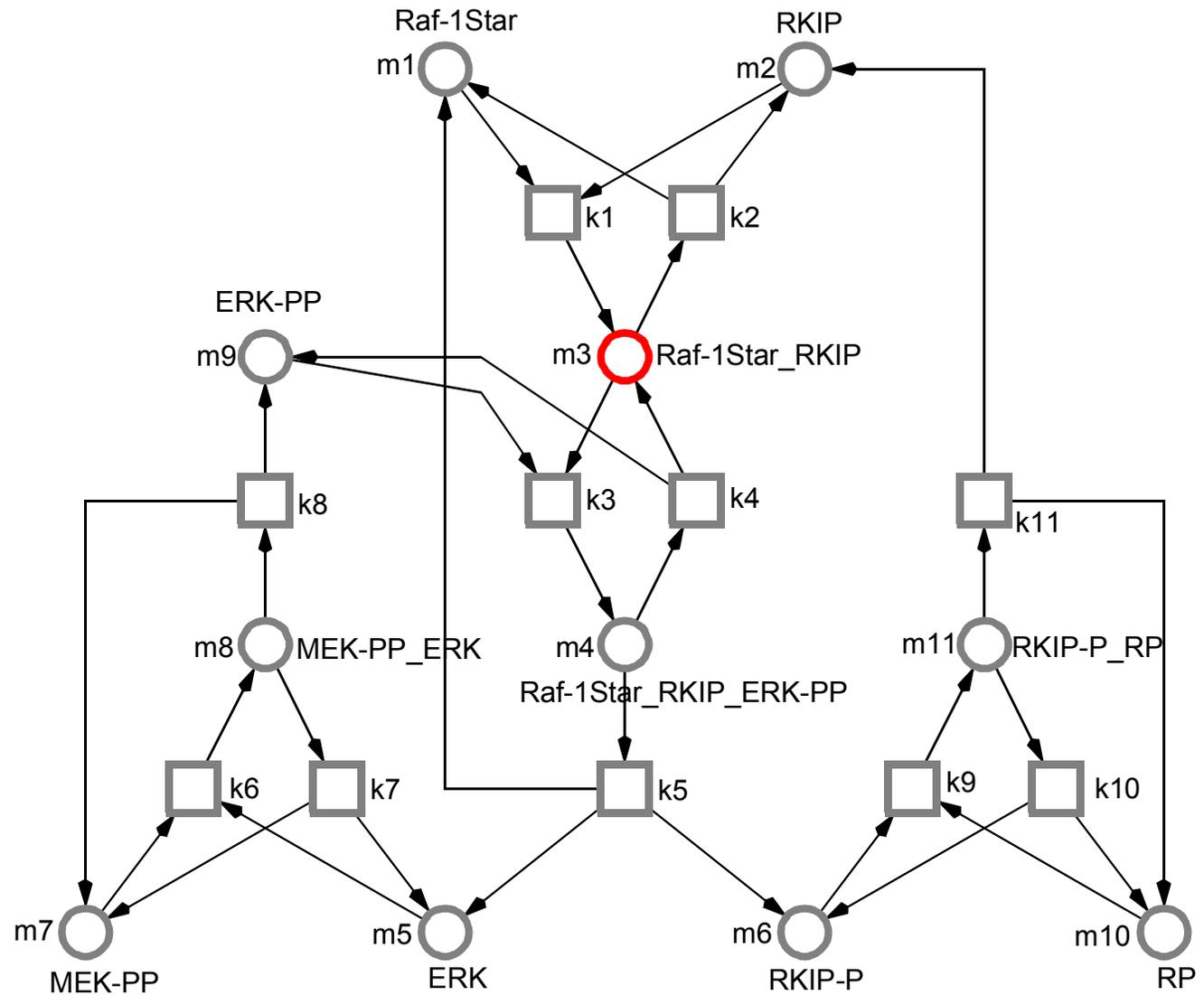
-> *concentrations*

- ❑ **transitions fire continuously**
- ❑ **rate functions**
  - > *any arithmetic function including the transition's pre-places as real-valued variables and user-defined real-valued parameters*
- ❑ **real-valued tokens**
  - > *concentrations*
- ❑ **semantics: set of Ordinary Differential Equations (ODEs)**
  - > *uniquely defined, but not vice versa*
  - > *typically non-linear*
- ❑ **simulation (numerical integration)**
  - > *stiff/unstiff solvers*



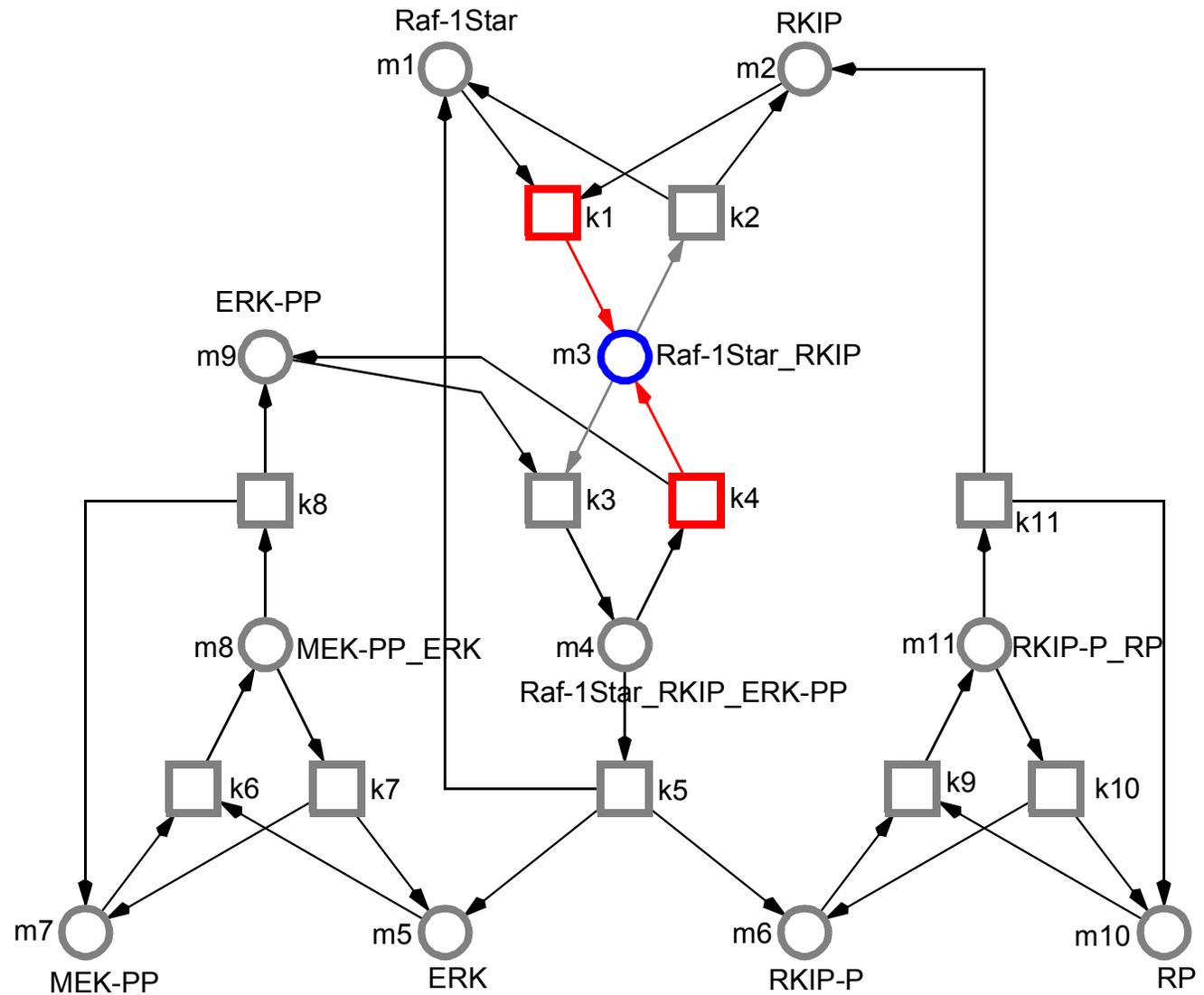
# CONTINUOUS PETRI NET DEFINES ODES

$$\frac{dm_3}{dt} =$$



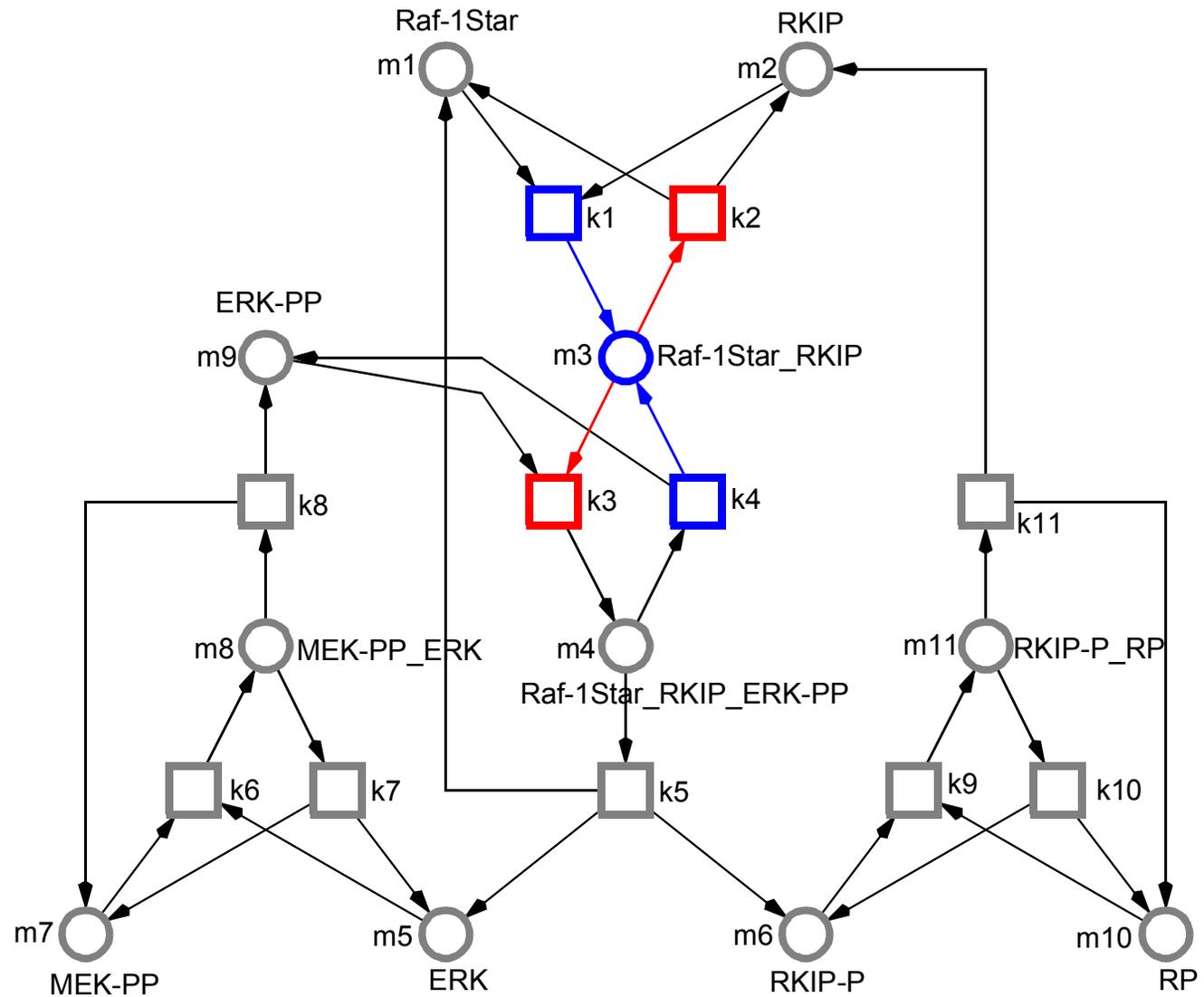
# CONTINUOUS PETRI NET DEFINES ODES

$$\frac{dm_3}{dt} = +r_1 + r_4$$



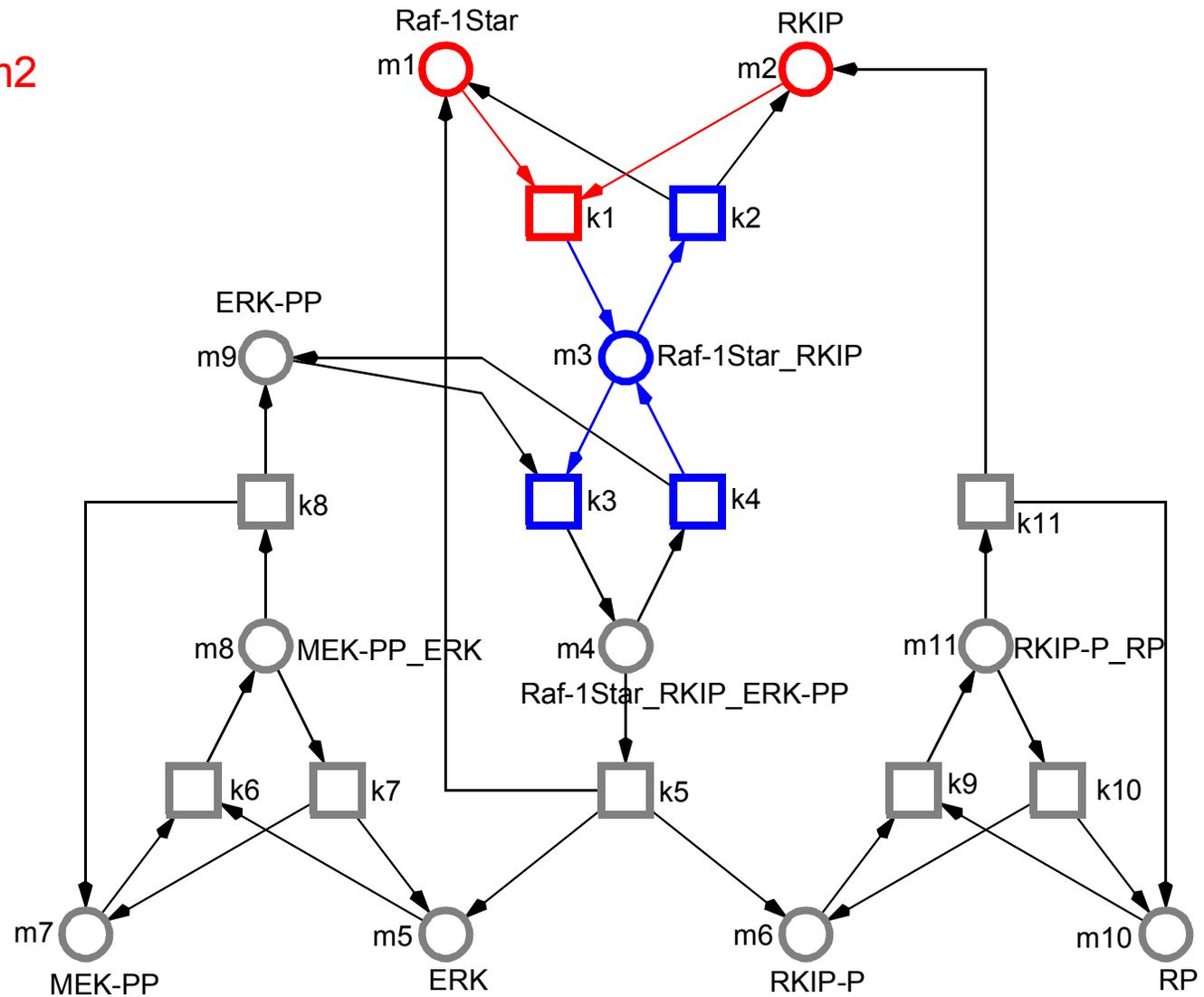
# CONTINUOUS PETRI NET DEFINES ODES

$$\frac{dm_3}{dt} = + r_1 + r_4 - r_2 - r_3$$



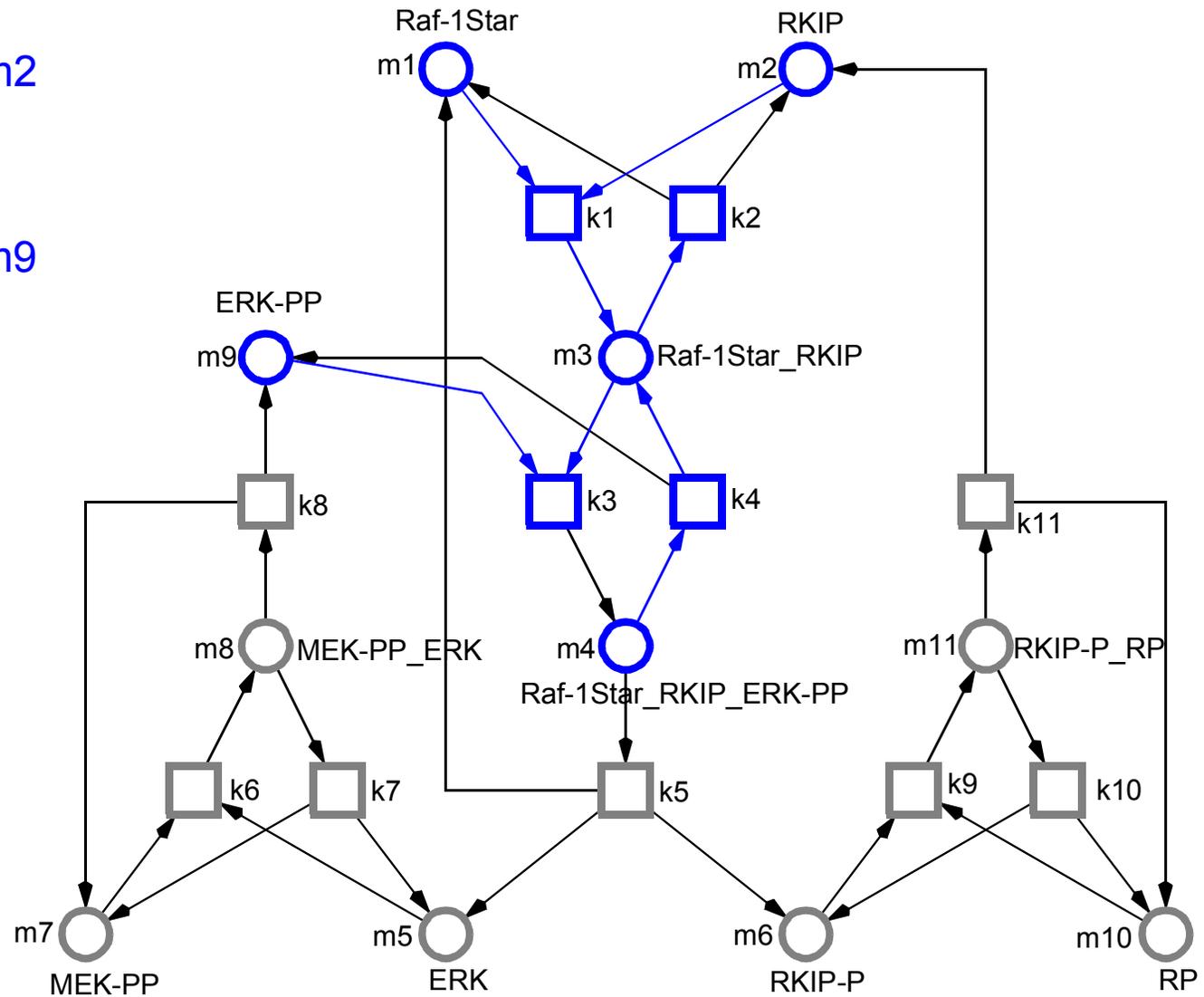
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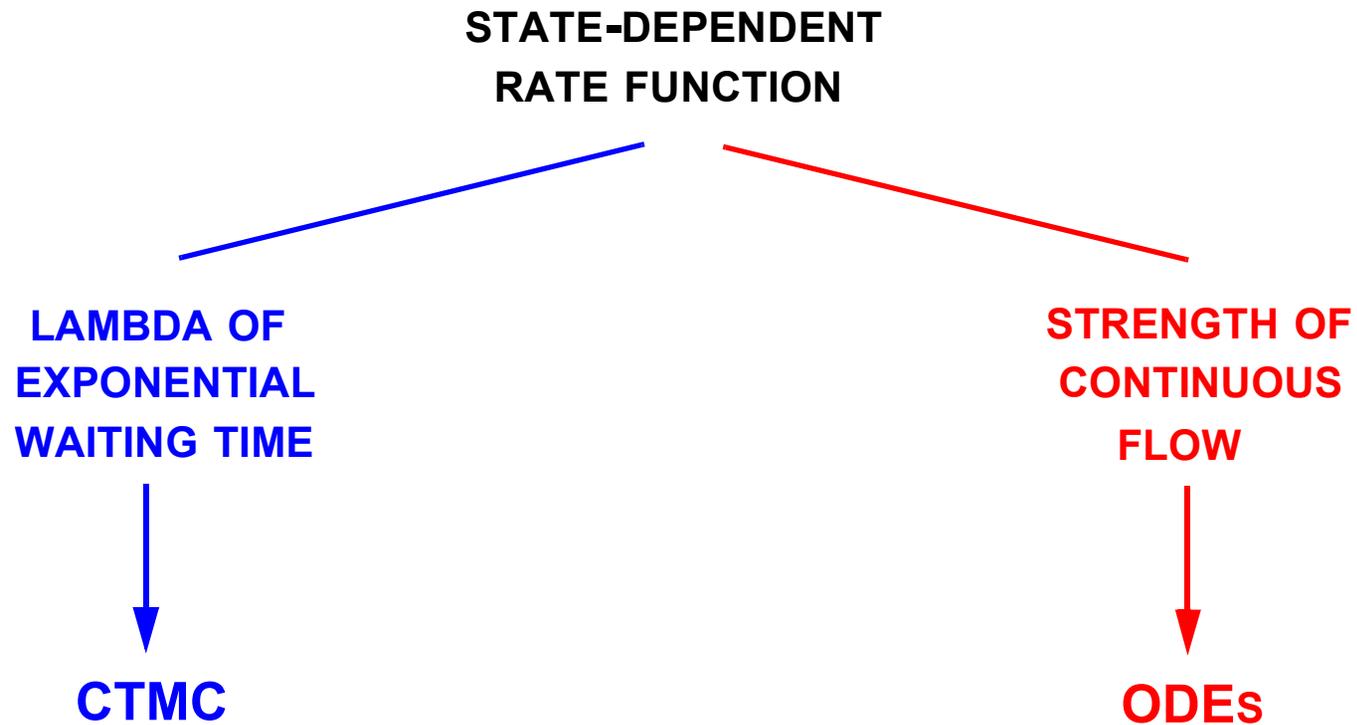
$$\frac{dm_3}{dt} = + k_1 * m_1 * m_2 + r_4 - r_2 - r_3$$



# CONTINUOUS PETRI NET DEFINES ODES

$$\frac{dm_3}{dt} = + k_1 * m_1 * m_2 + k_4 * m_4 - k_2 * m_3 - k_3 * m_3 * m_9$$





-> supported by, e.g., COPASI, Dizzy, ..., Snoopy

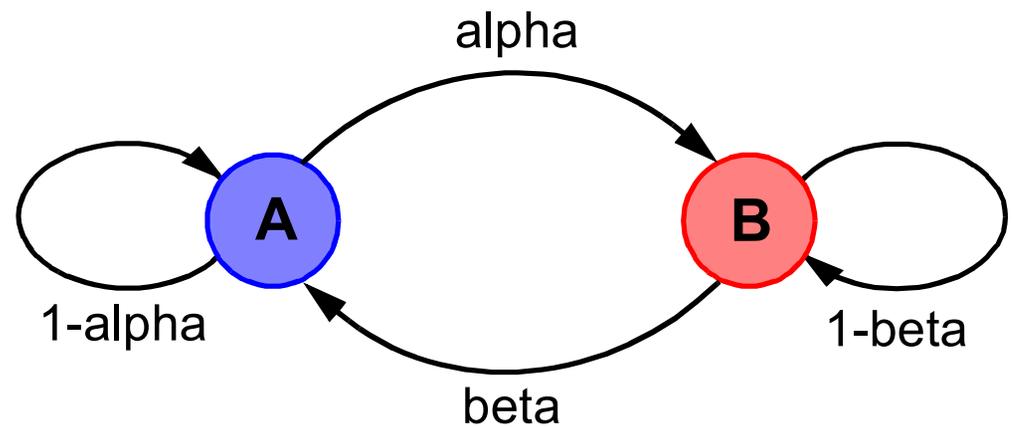
# CASE STUDY - MULTICELL COLONIES

❑ two cell types: phenotype A and B

❑ cell divide

-> cell division may involve mutation of the offspring

-> parent cell keeps its phenotype



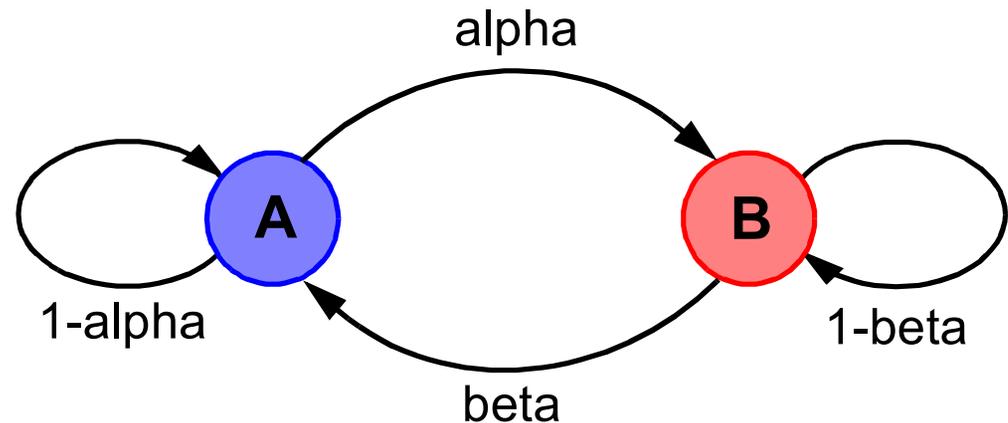
❑ **two cell types: phenotype A and B**

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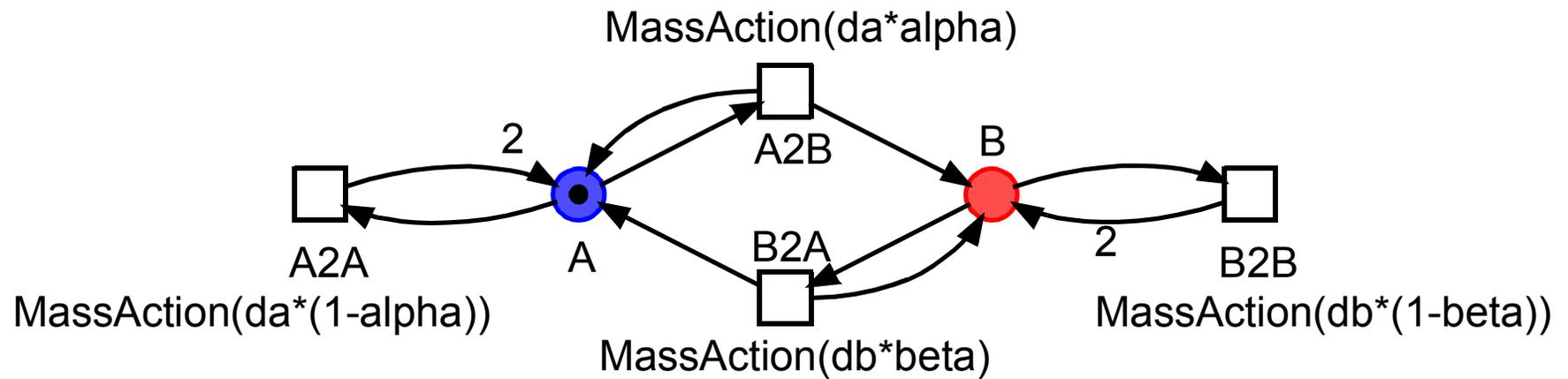
❑ **model parameters**

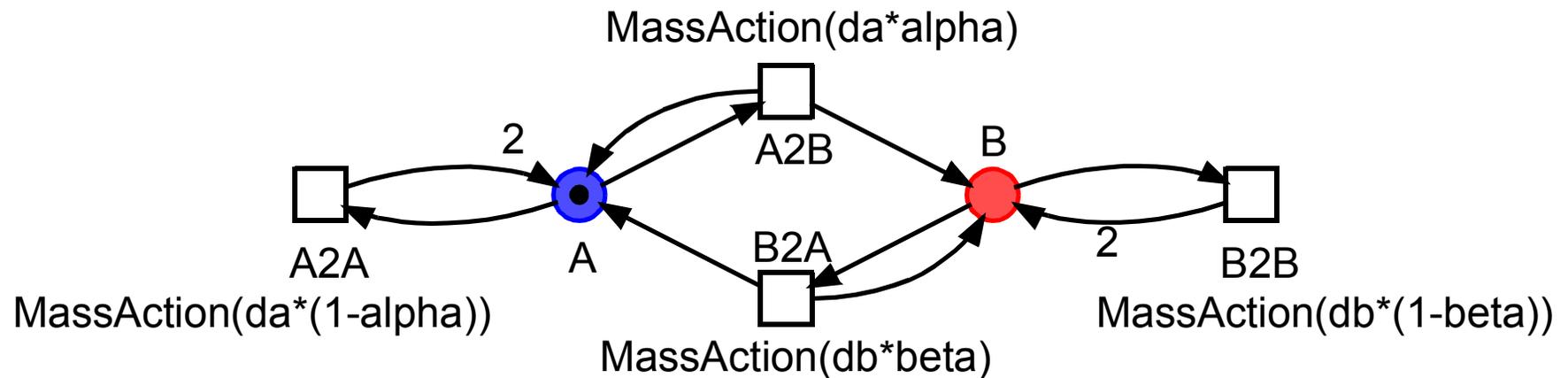
- > *alpha = beta - mutation rates*
- > *da, db - fitness of A, B*
- > *da/db - relative fitness*



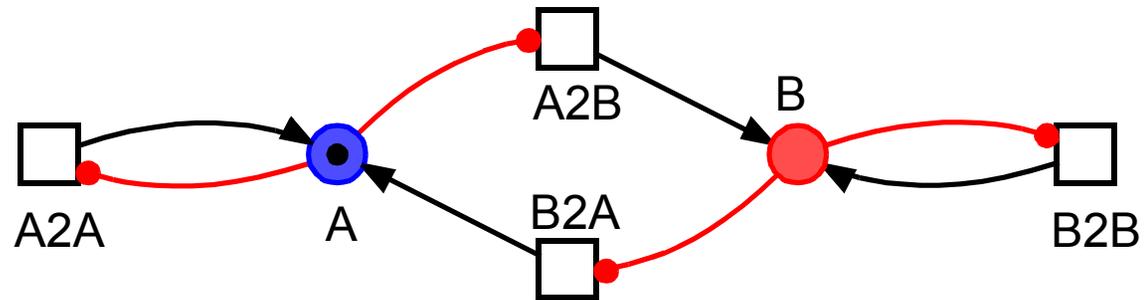
❑ **output**

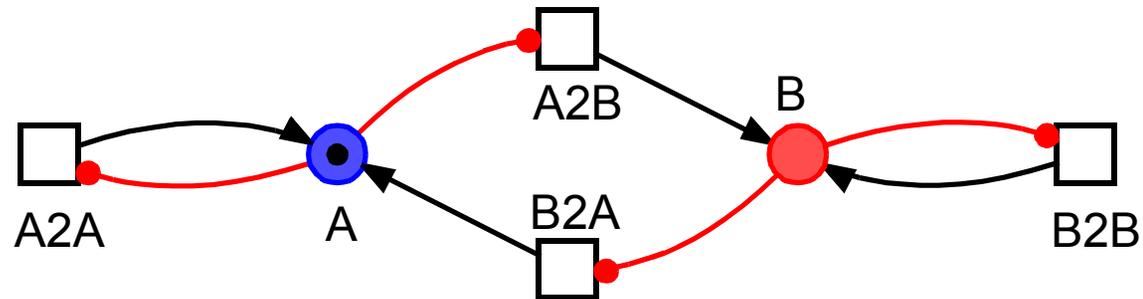
- > *total number of bacteria*
- > *proportion of A =  $A / (A + B)$*
- > *proportion of B =  $B / (A + B)$*





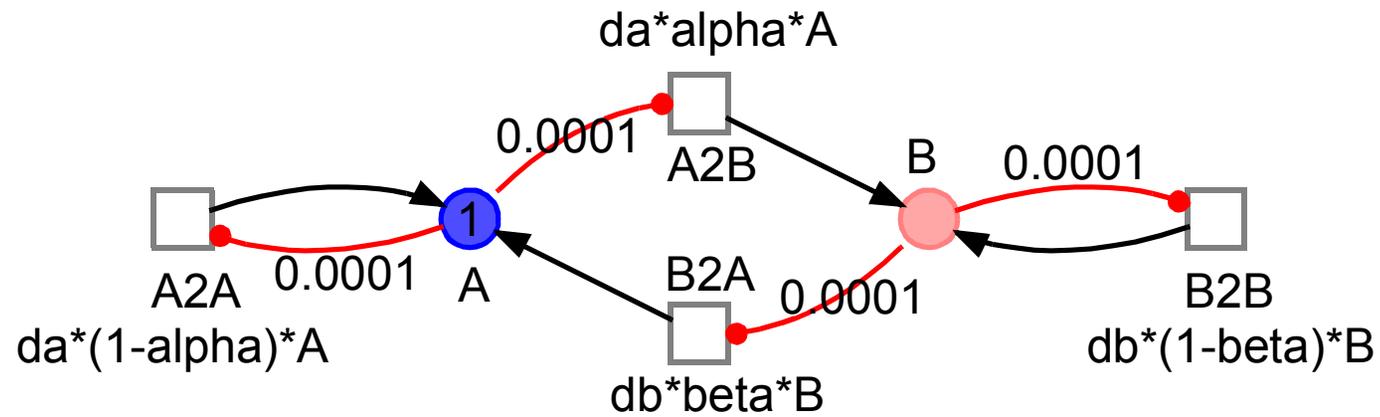
$$\begin{aligned} \frac{dA}{dt} &= da \cdot (1 - \alpha) \cdot A + db \cdot \beta \cdot B \\ \frac{dB}{dt} &= db \cdot (1 - \beta) \cdot B + da \cdot \alpha \cdot A \end{aligned}$$

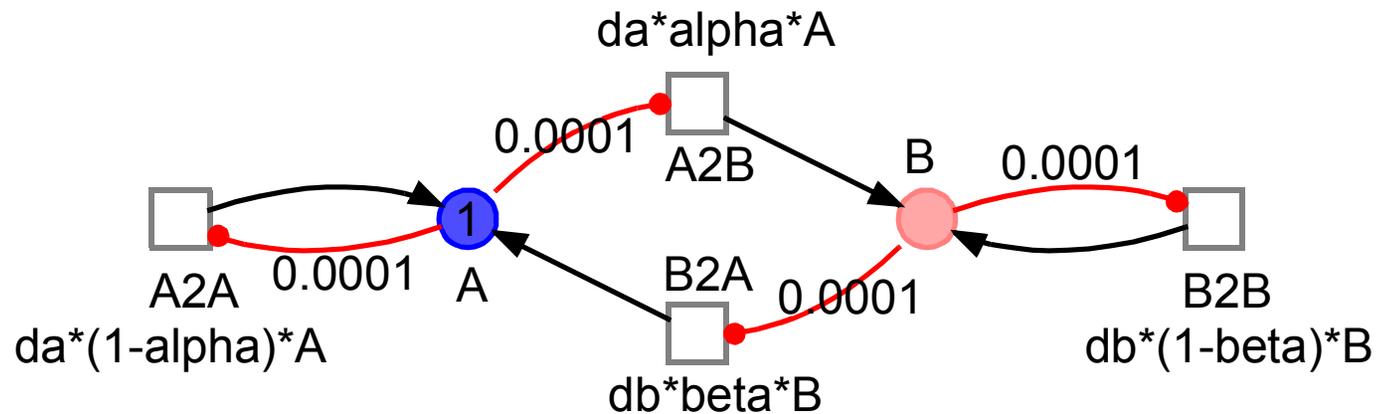




$$\begin{aligned} dA/dt = & da * (1 - \alpha) * A * \text{read}(1,A) \\ & + db * \beta * B * \text{read}(1,B) \end{aligned}$$

$$\begin{aligned} dB/dt = & db * (1 - \beta) * B * \text{read}(1,B) \\ & + da * \alpha * A * \text{read}(1,A) \end{aligned}$$

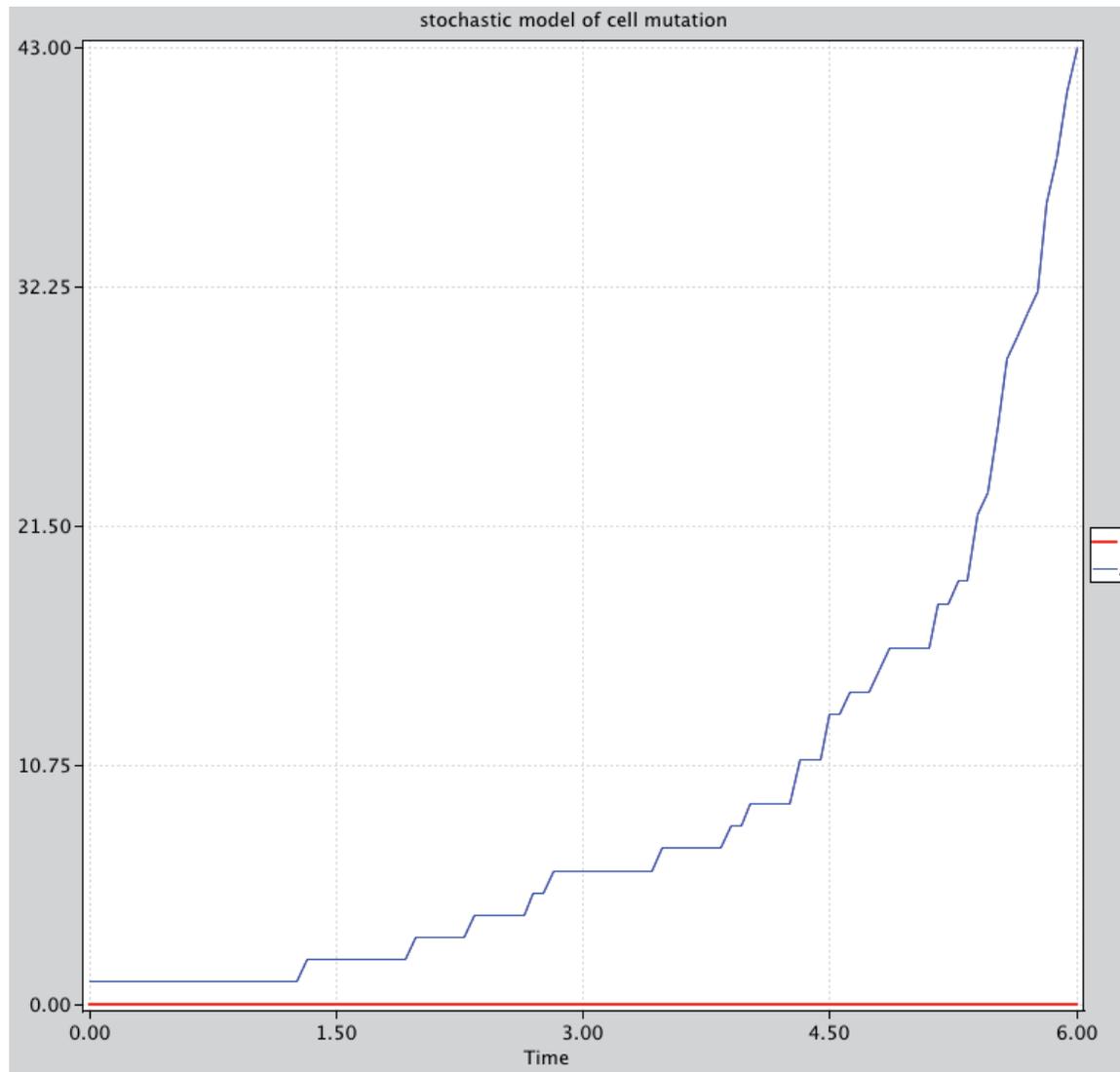


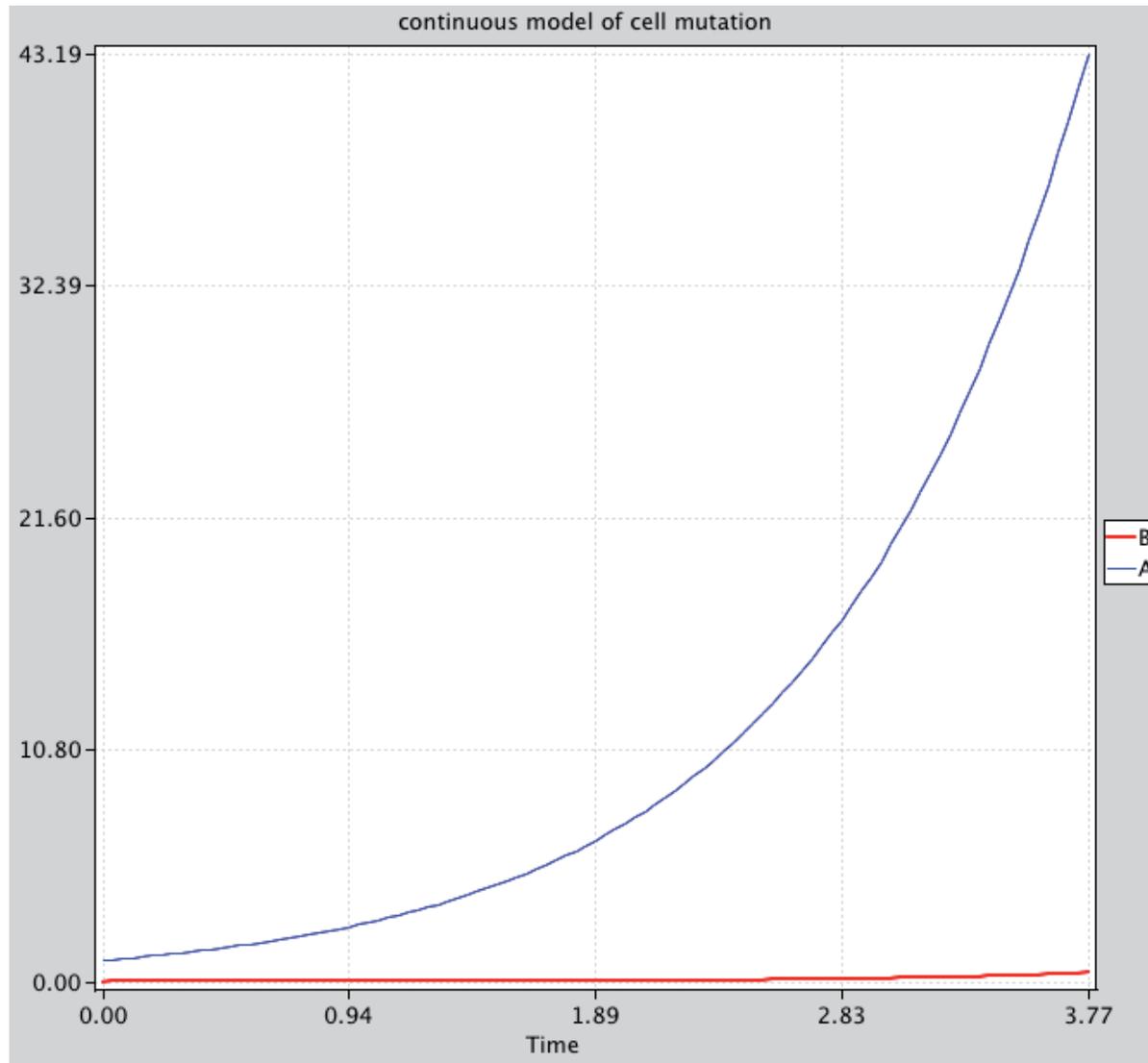


$$\begin{aligned} \frac{dA}{dt} = & da * (1 - \alpha) * A * \text{read}(0.0001, A) \\ & + db * \beta * B * \text{read}(0.0001, B) \end{aligned}$$

$$\begin{aligned} \frac{dB}{dt} = & db * (1 - \beta) * B * \text{read}(0.0001, B) \\ & + da * \alpha * A * \text{read}(0.0001, A) \end{aligned}$$

# MULTICELL COLONY, STOCHASTIC PLOT





- ❑ **cellDivision0a.spcontped**  
-> *take a look on the unreduced ODEs and simulate them*
  
- ❑ **cellDivision0a.spcontped**  
-> *take a look on the reduced ODEs and simulate them*
  
- ❑ **cellDivision0b.spcontped**  
-> *take a look on the ODEs and simulate them*
  
- ❑ **cellDivision0c.spcontped**  
-> *take a look on the ODEs and simulate them*  
-> *repeat with varied parameters and different read arc weights;*

**MAIN LESSON LEARNT:**

**NEVER BLINDLY TRUST NUMERICAL SIMULATION RESULTS.**

# SUMMARY

## □ representation of bionetworks by Petri nets

-> *partial order representation*

-> *formal semantics*

-> *unifying view*

-> *better comprehension*

-> *sound analysis techniques*

## ❑ representation of bionetworks by Petri nets

- > *partial order representation*
- > *formal semantics*
- > *unifying view*

- > *better comprehension*
- > *sound analysis techniques*

## ❑ purposes

- > *animation*
- > *model validation against consistency criteria*
- > *qualitative / quantitative behaviour prediction*

- > *to experience the model*
- > *to increase confidence*
- > *experiment design,  
new insights*

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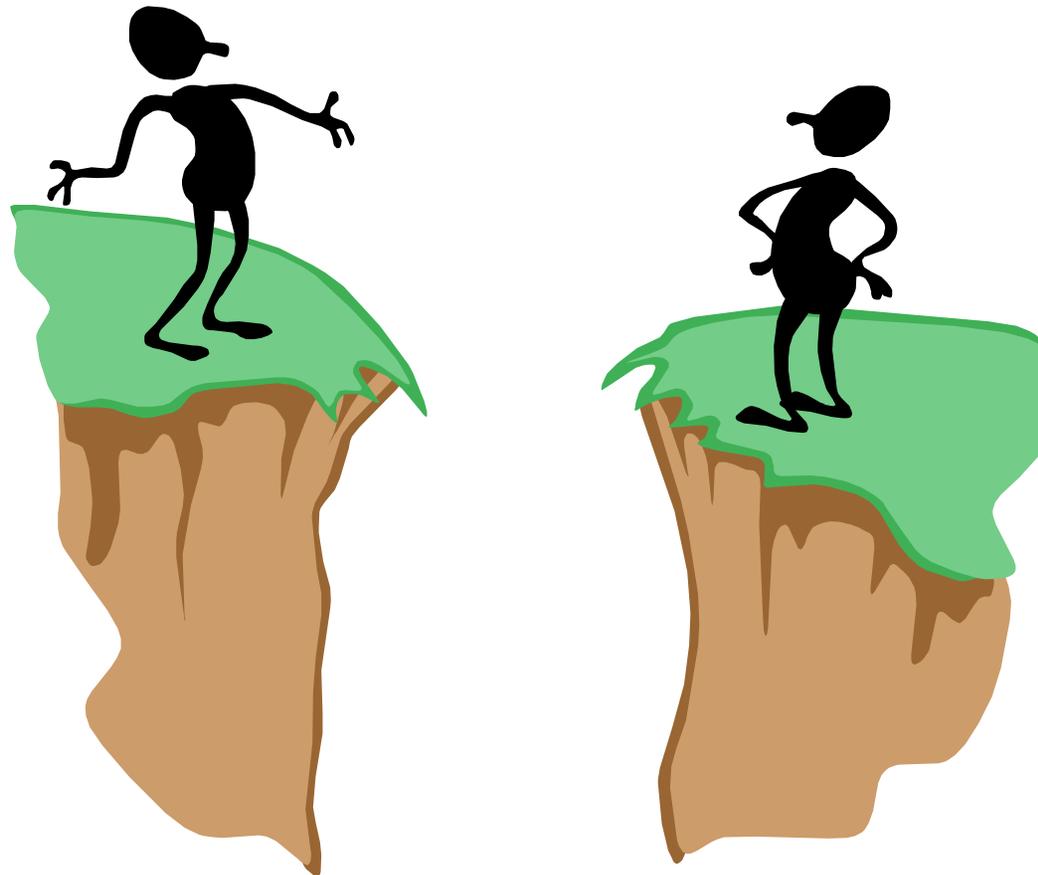
- > *animation*
- > *model validation against consistency criteria*
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- > *to experience the model*
- > *to increase confidence*
- > *experiment design, new insights*

## ❑ step-wise model development

- > *qualitative model*
- > *discrete quantitative model*
- > *continuous quantitative model*

- > *discrete Petri nets*
- > *stochastic Petri nets*
- > *continuous Petri nets = ODEs*



**THANKS !**

**[HHTTP://WWW-DSSZ.INFORMATIK.TU-COTTBUS.DE](http://www-dssz.informatik.tu-cottbus.de)**

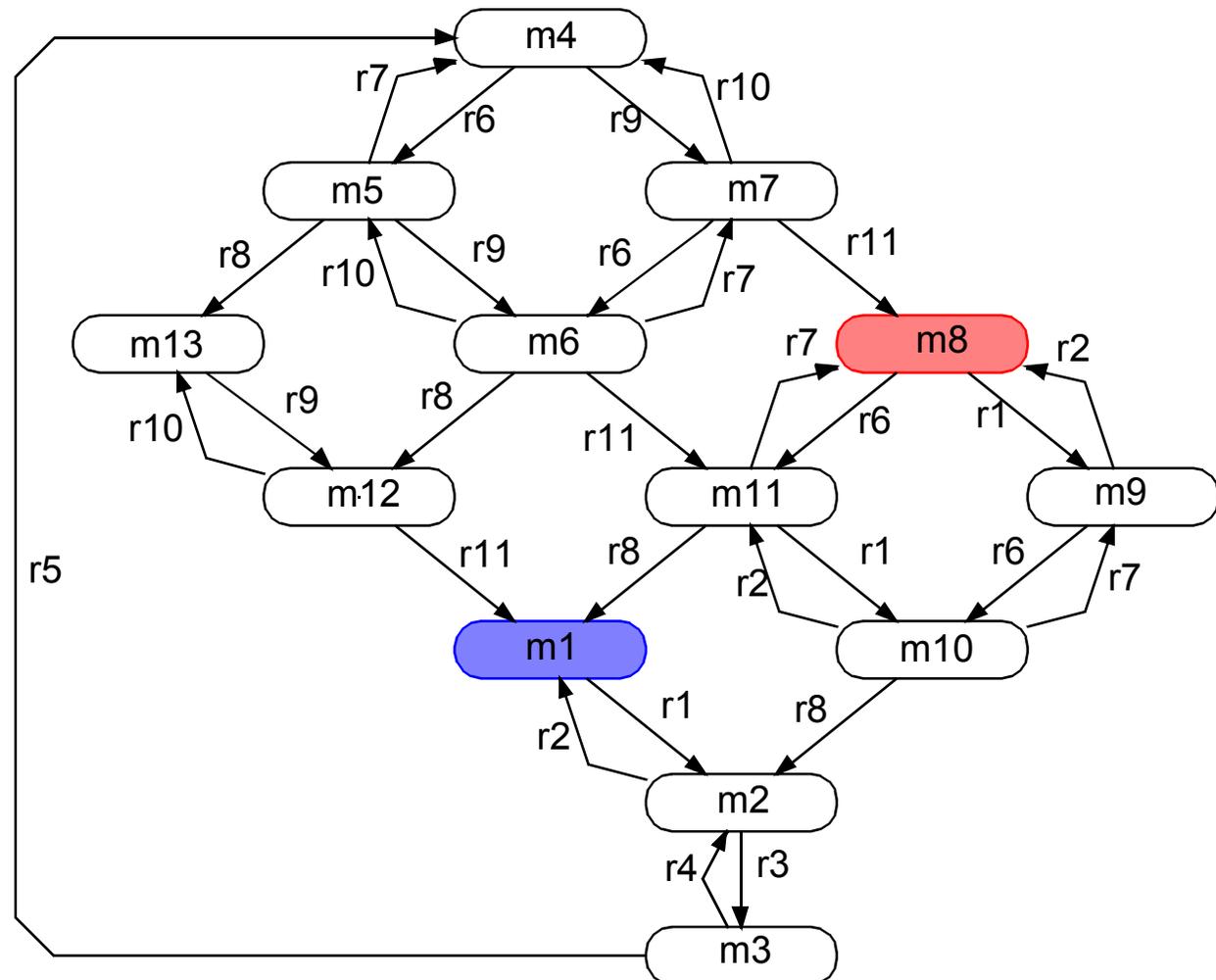
**SUPPLEMENT:**

**ABOUT THE RELATION**

**SPN - CPN**

# EX1 - RKIP, REACHABILITY GRAPH (STS)

- ❑ simple algorithm
- ❑ nodes : system states
- ❑ arcs : the (single) firing transition
- ❑ single step firing rule



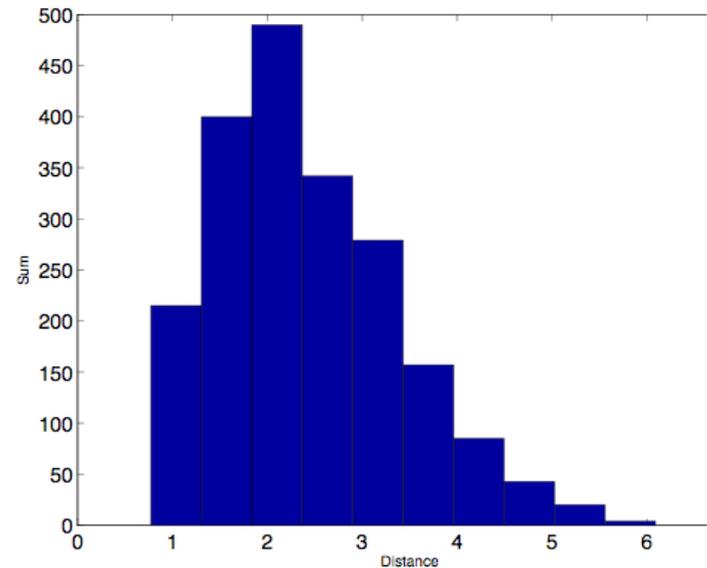
# EX1 - RKIP, QUANTITATIVE ANALYSIS

Species	S1	S2	S3	S4	S5	S6	S7	S8	S9	S10	S11	S12	S13
Raf-1*	1	0	0	1	1	1	1	1	0	0	1	1	1
RKIP	1	0	0	0	0	0	0	1	0	0	1	0	0
Raf-1*_RKIP	0	1	0	0	0	0	0	0	1	1	0	0	0
Raf-1*_RKIP_ERK-PP	0	0	1	0	0	0	0	0	0	0	0	0	0
ERK	0	0	0	1	0	0	1	1	1	0	0	0	0
RKIP-P	0	0	0	1	1	0	0	0	0	0	0	0	1
MEK-PP	1	1	1	1	0	0	1	1	1	0	0	1	1
MEK-PP_ERK	0	0	0	0	1	1	0	0	0	1	1	0	0
ERK-PP	1	1	0	0	0	0	0	0	0	0	0	1	1
RP	1	1	1	1	1	0	0	1	1	1	1	0	1
RKIP-P_RP	0	0	0	0	0	1	1	0	0	0	0	1	0

Cho et al

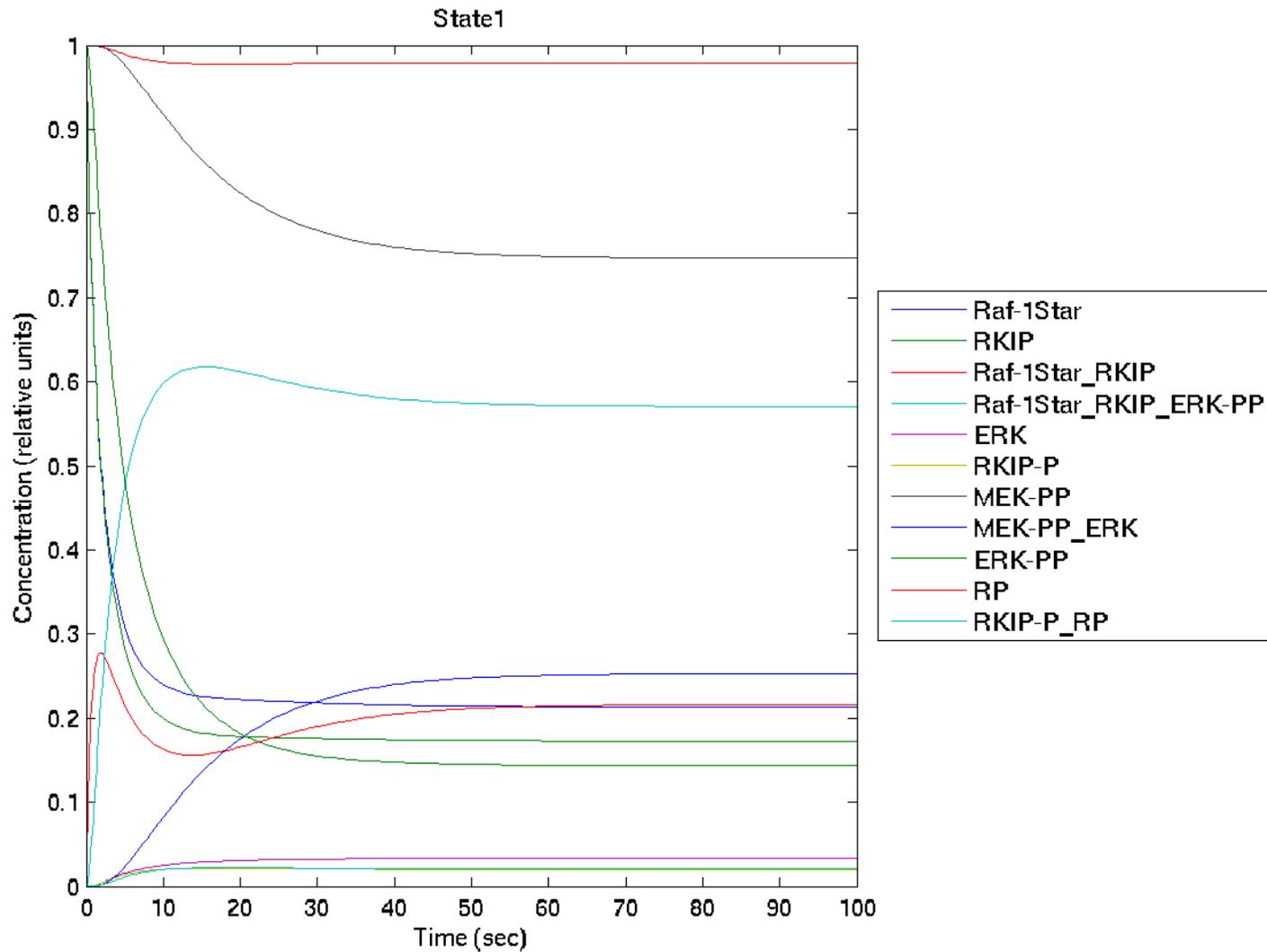
Biochemist

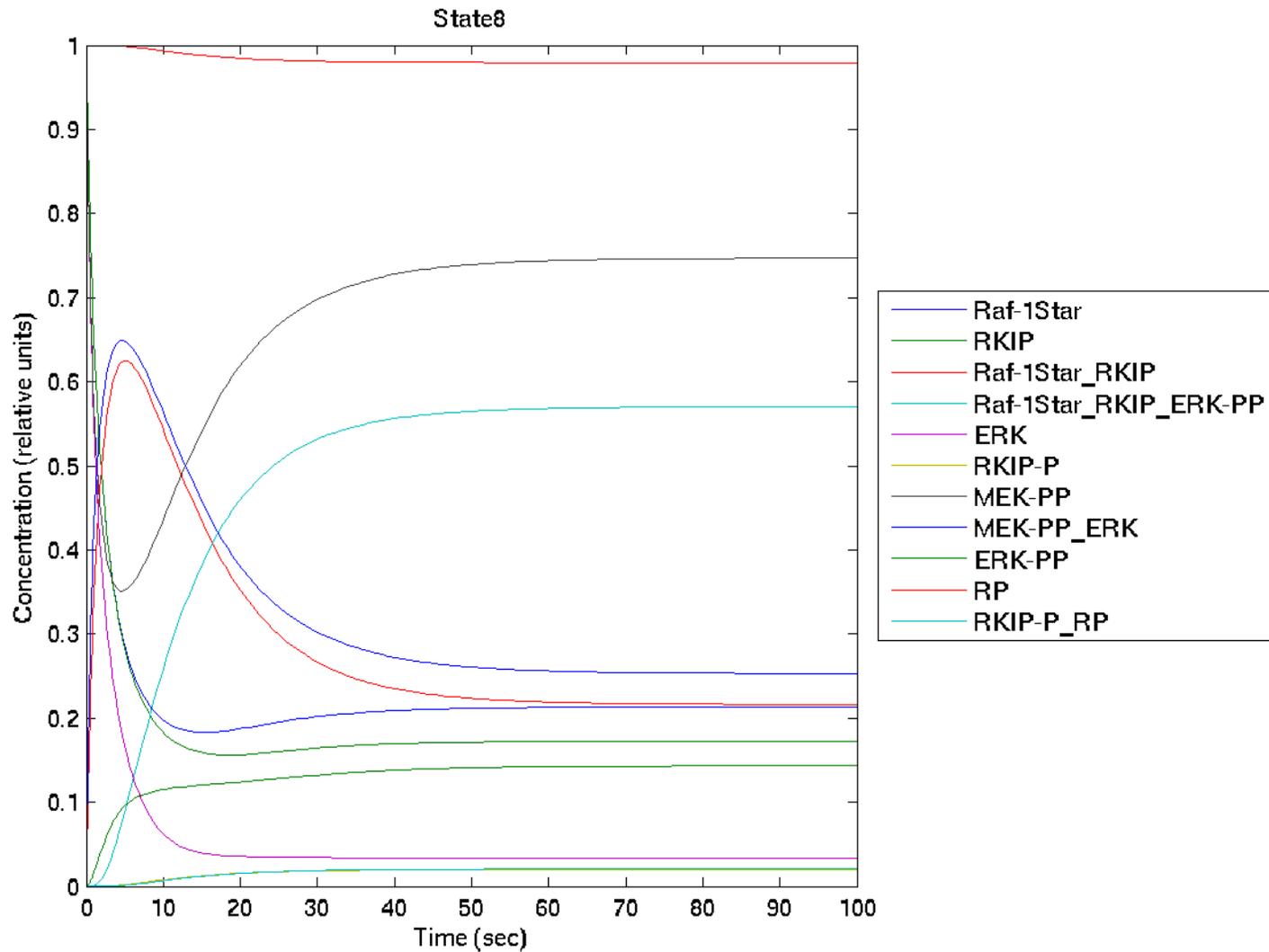
13 "good" state configurations

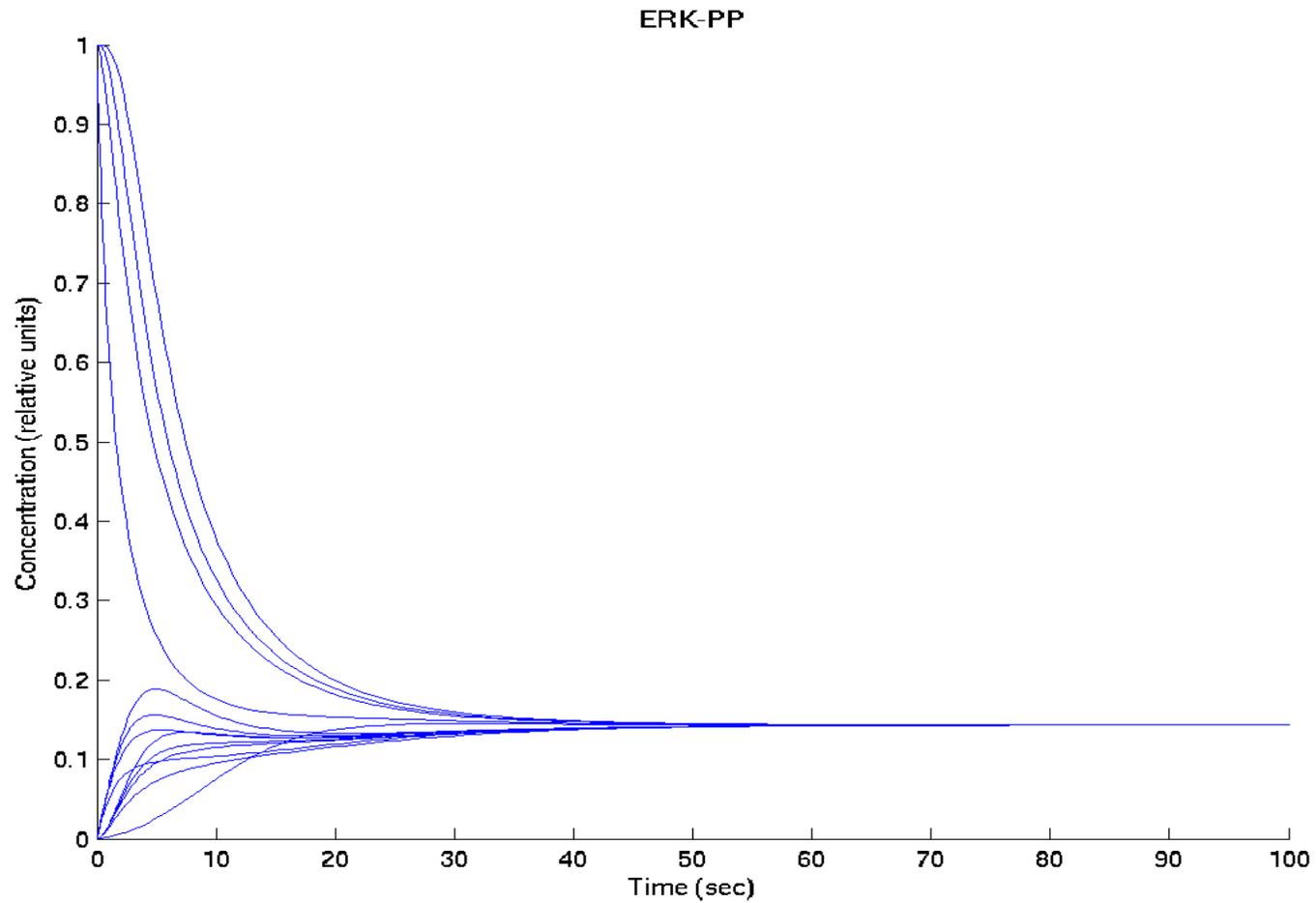


Distribution of 'bad' steady states as euclidean distances from the 'good' final steady state

the "bad" ones

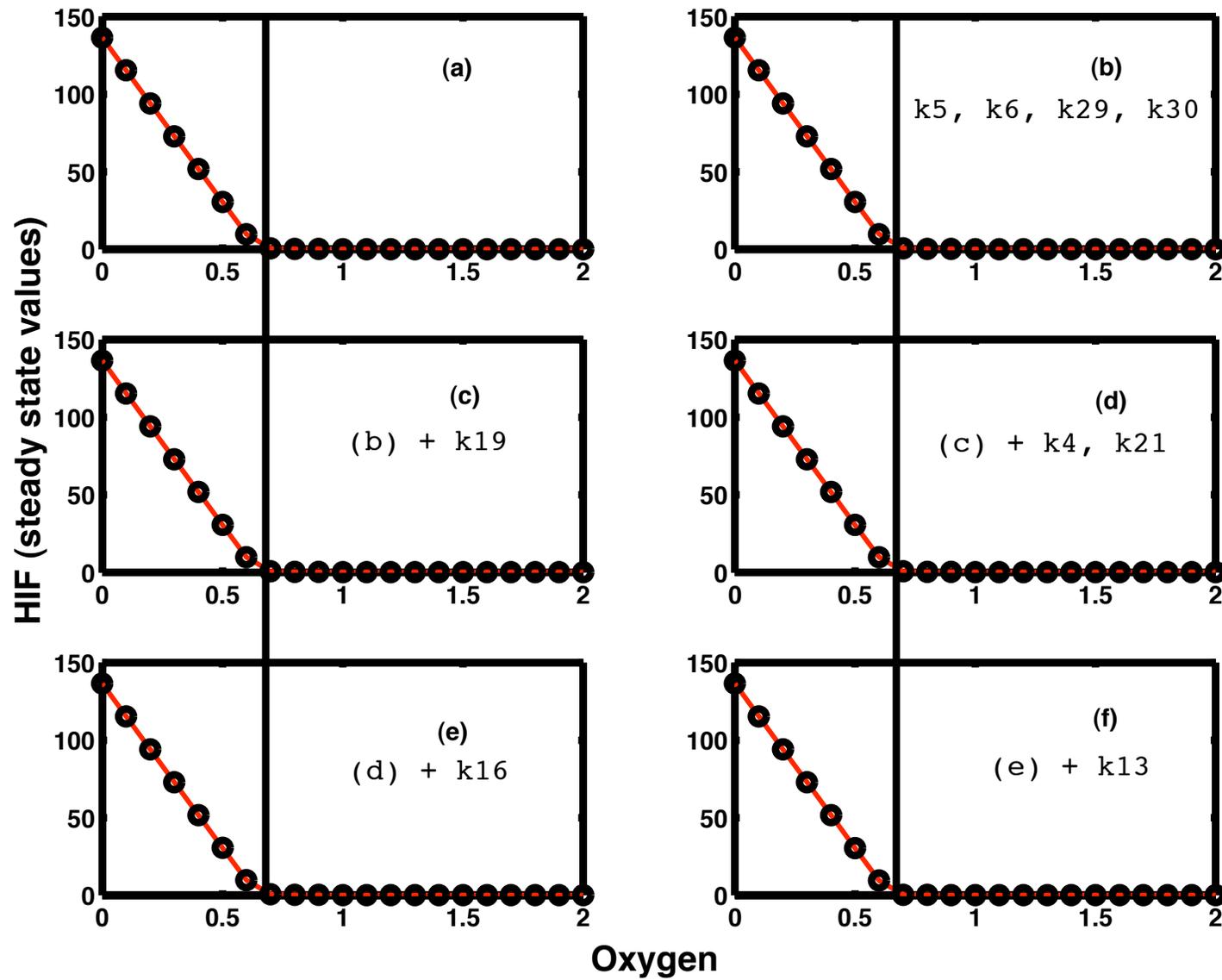




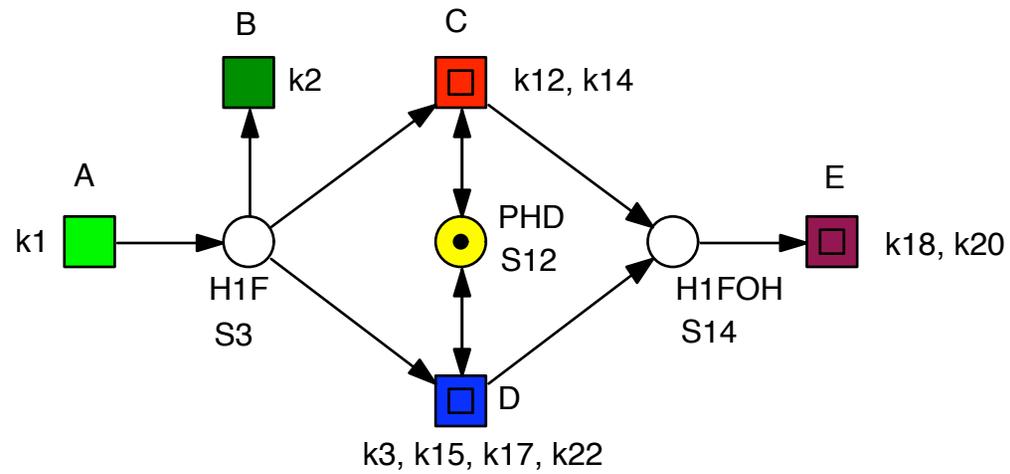




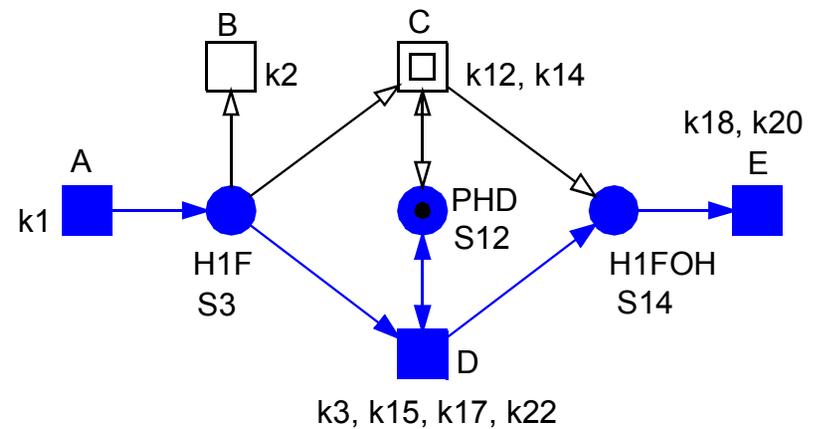
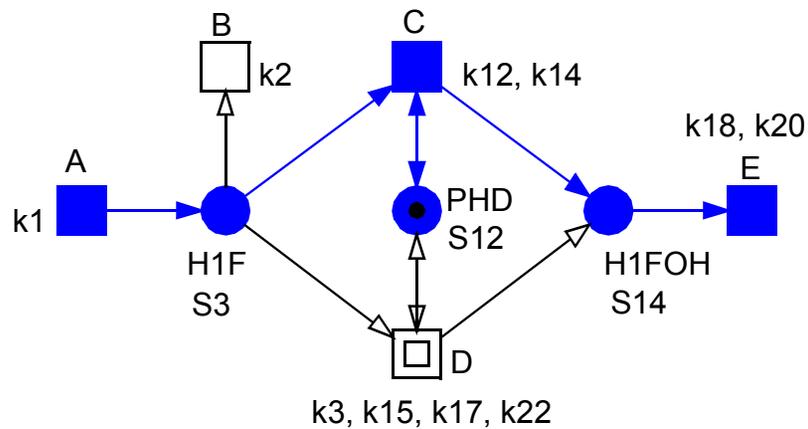
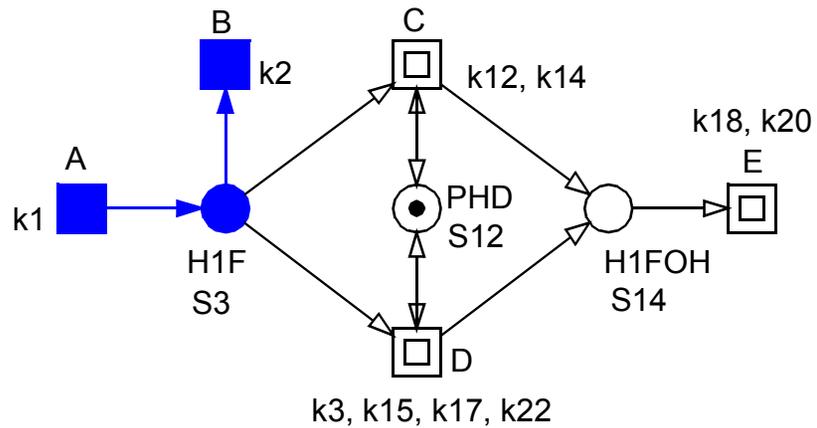








# Ex2 - HYPOXIA



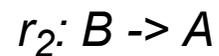
**BUT,**  
**THERE ARE MANY EXAMPLES**  
**WHERE THE TRANSITION SPN  $\rightarrow$  CPN**  
**COMES WITH COUNTERINTUITIVE EFFECTS.**

- **ACR: steady state value of variable (place) does not depend on total mass, only on kinetic constants**      *-> [SHINAR, FEINBERG 2010]*

- **simple example**      **mass-action kinetics**



$$v_1(r_1) = k_1AB$$



$$v_2(r_2) = k_2B$$

- ❑ **ACR: steady state value of variable (place) does not depend on total mass, only on kinetic constants** → [SHINAR, FEINBERG 2010]

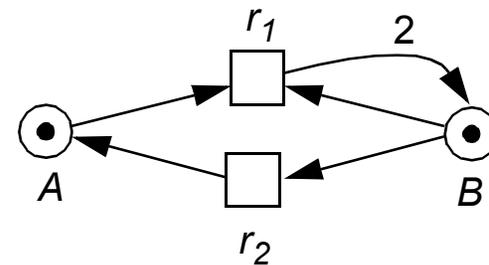
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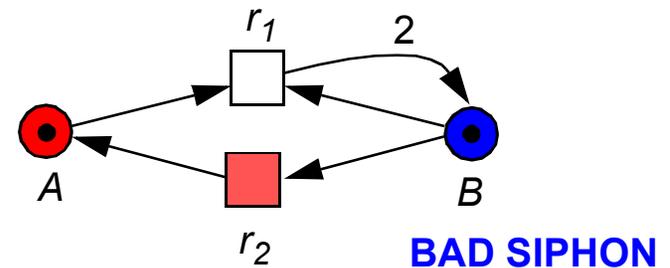
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$$CPI: m_0(A) + m_0(B) = total$$

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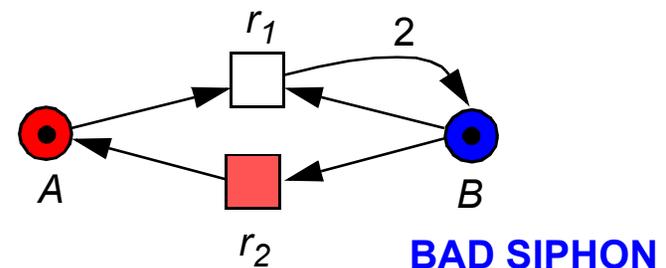


$$v_2(r_2) = k_2B$$

- **ODEs**

$$dA/dt = v_2 - v_1 = k_2B - k_1AB$$

$$dB/dt = v_1 - v_2 = k_1AB - k_2B$$



$$CPI: m_0(A) + m_0(B) = total$$

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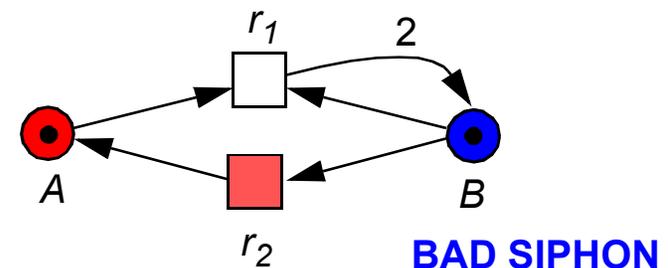
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- **steady state**

$$dA/dt = k_2B - k_1AB = 0$$

$$dB/dt = k_1AB - k_2B = 0$$



$$CPI: m_0(A) + m_0(B) = total$$

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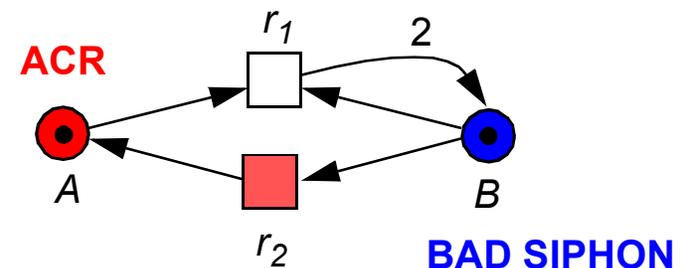
- **steady state**

$$dA/dt = k_2B - k_1AB = 0$$

$$dB/dt = k_1AB - k_2B = 0$$

$$\rightarrow \text{steady\_state}(A) = k_2/k_1$$

$$\text{steady\_state}(B) = \text{total} - k_2/k_1$$



$$CPI: m_0(A) + m_0(B) = \text{total}$$

# ABSOLUT CONCENTRATION ROBUSTNESS (ACR)

- **ACR: steady state value of variable (place) does not depend on total mass, only on kinetic constants** → [SHINAR, FEINBERG 2010]

- **simple example**      **mass-action kinetics**



$$v_1(r_1) = k_1AB$$



$$v_2(r_2) = k_2B$$

- **ODEs**

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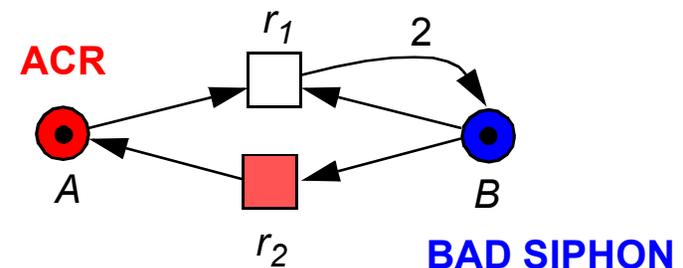
- **steady state**

$$dA/dt = k_2B - k_1AB = 0$$

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$$\rightarrow \text{steady\_state}(A) = k_2/k_1$$

$$\text{steady\_state}(B) = \text{total} - k_2/k_1$$



$$CPI: m_0(A) + m_0(B) = \text{total}$$

