

# **BioMODEL ENGINEERING**

## **- A PETRI NET PERSPECTIVE -**

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on sabbatical leave from  
**Brandenburg University of Technology Cottbus**  
**Dept. of CS**

## □ BACKGROUND

- > *Petri Nets - a crash course*
- > *some case studies*

## □ BIOMODEL ENGINEERING

- > *a unifying framework: QPN - SPN - CPN*
- > *relation SPN - CPN ?*

## □ OUR TOOL BOX

- > *Snoopy*
- > *Charlie*
- > *Marcie*

## □ LATEST NEWS

- > *colored framework*
- > *Generalized Hybrid Petri Nets*

## BACKGROUND

- > Petri Nets - a crash course
- > some case studies

## BIOMODEL ENGINEERING

- > a unifying framework: QPN - SPN - CPN
- > relation SPN - CPN ?

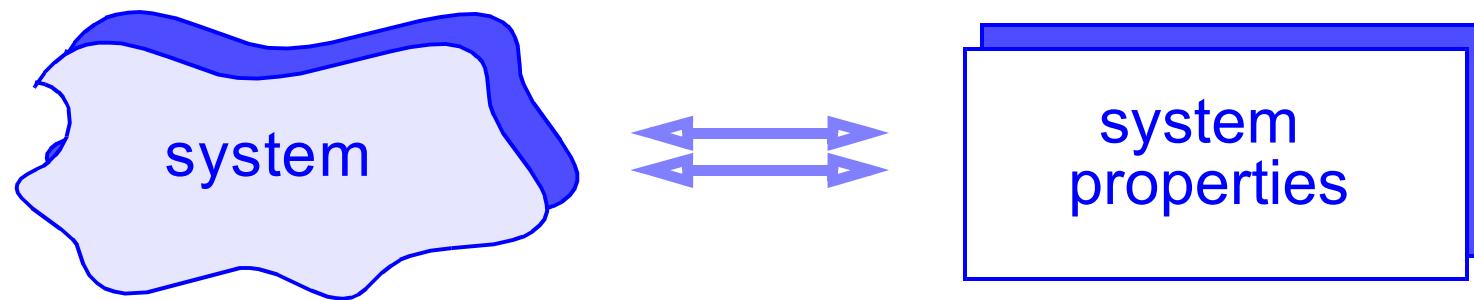
## OUR TOOL BOX

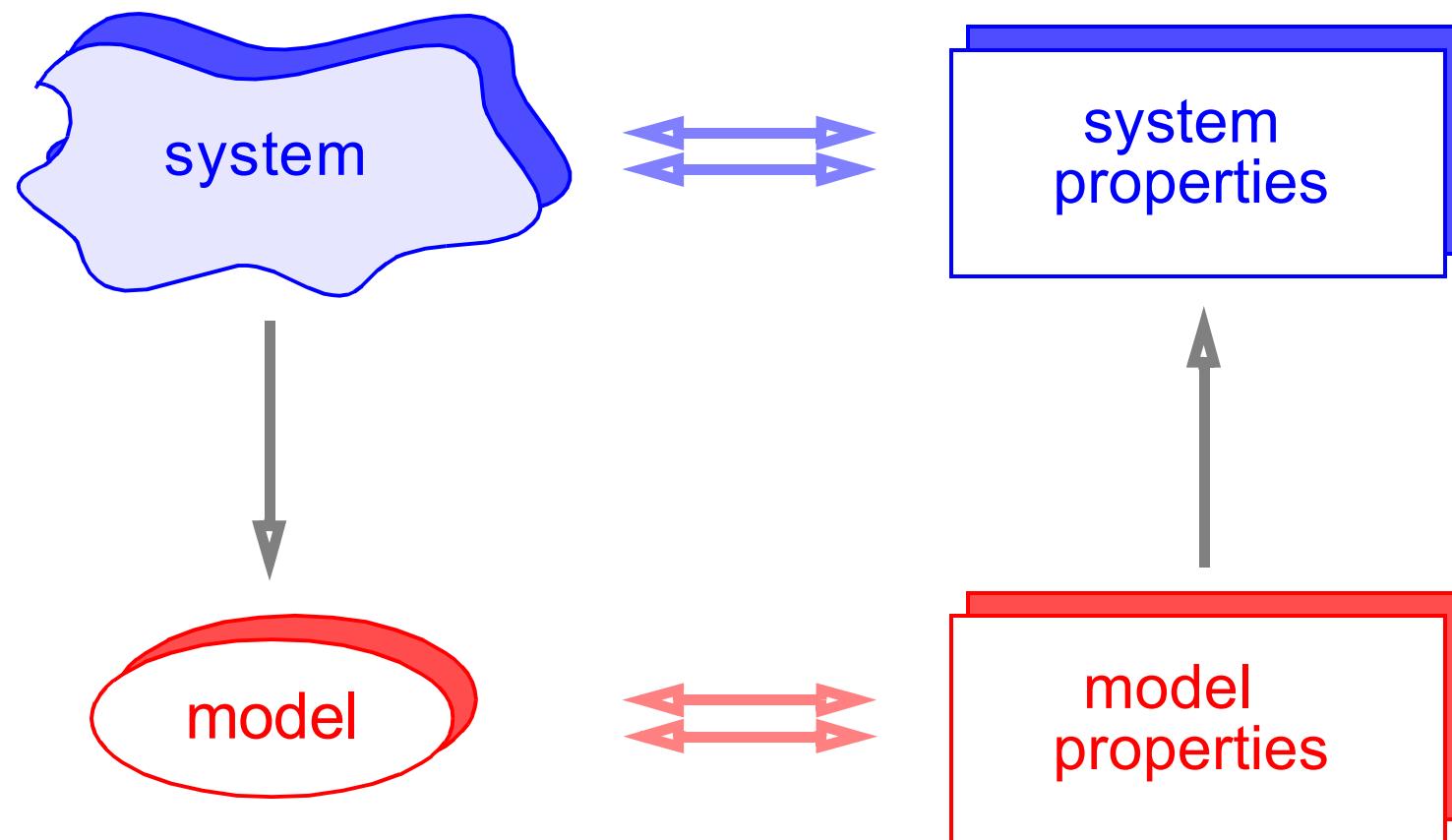
- > Snoopy
- > Charlie
- > Marcie

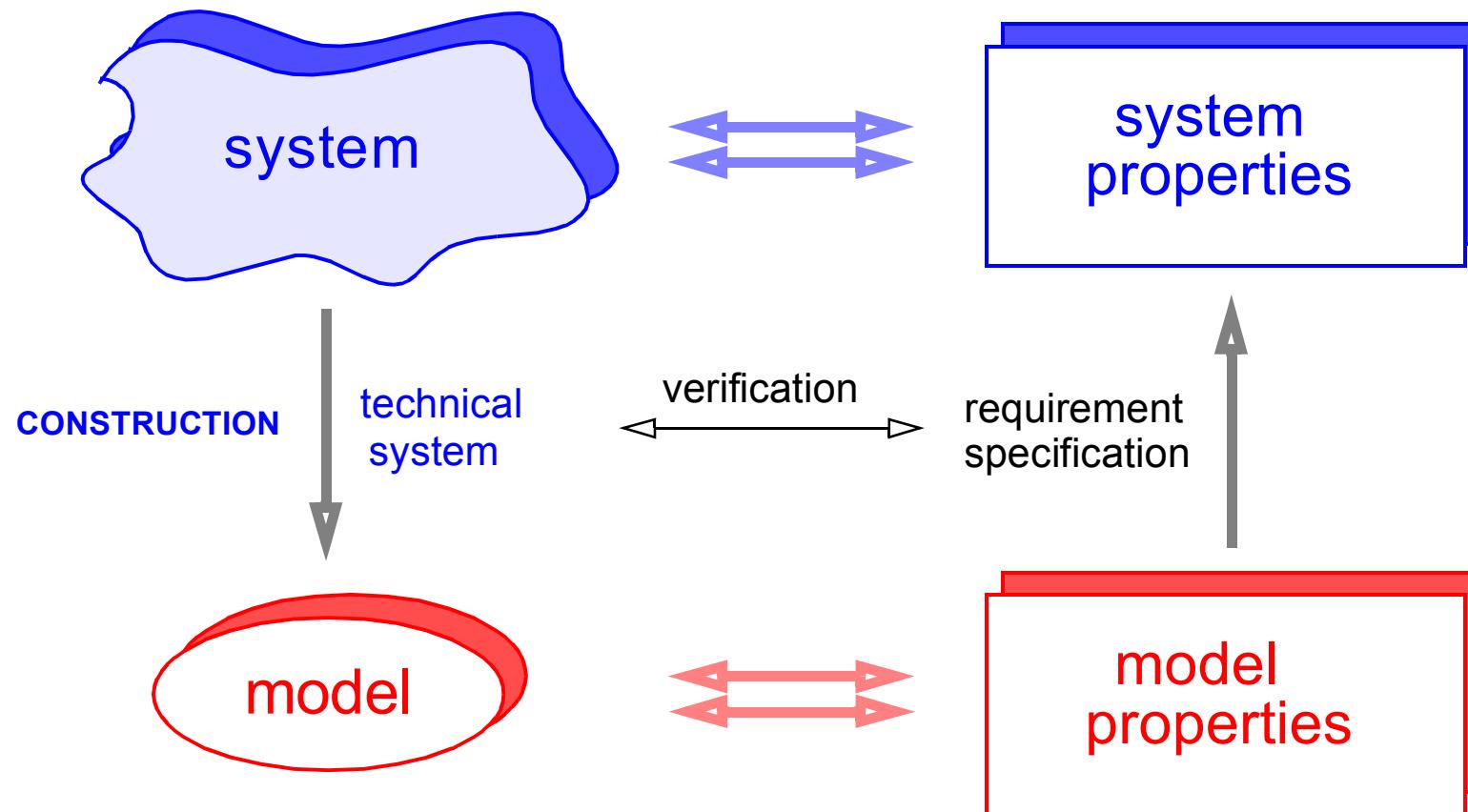
## LATEST NEWS

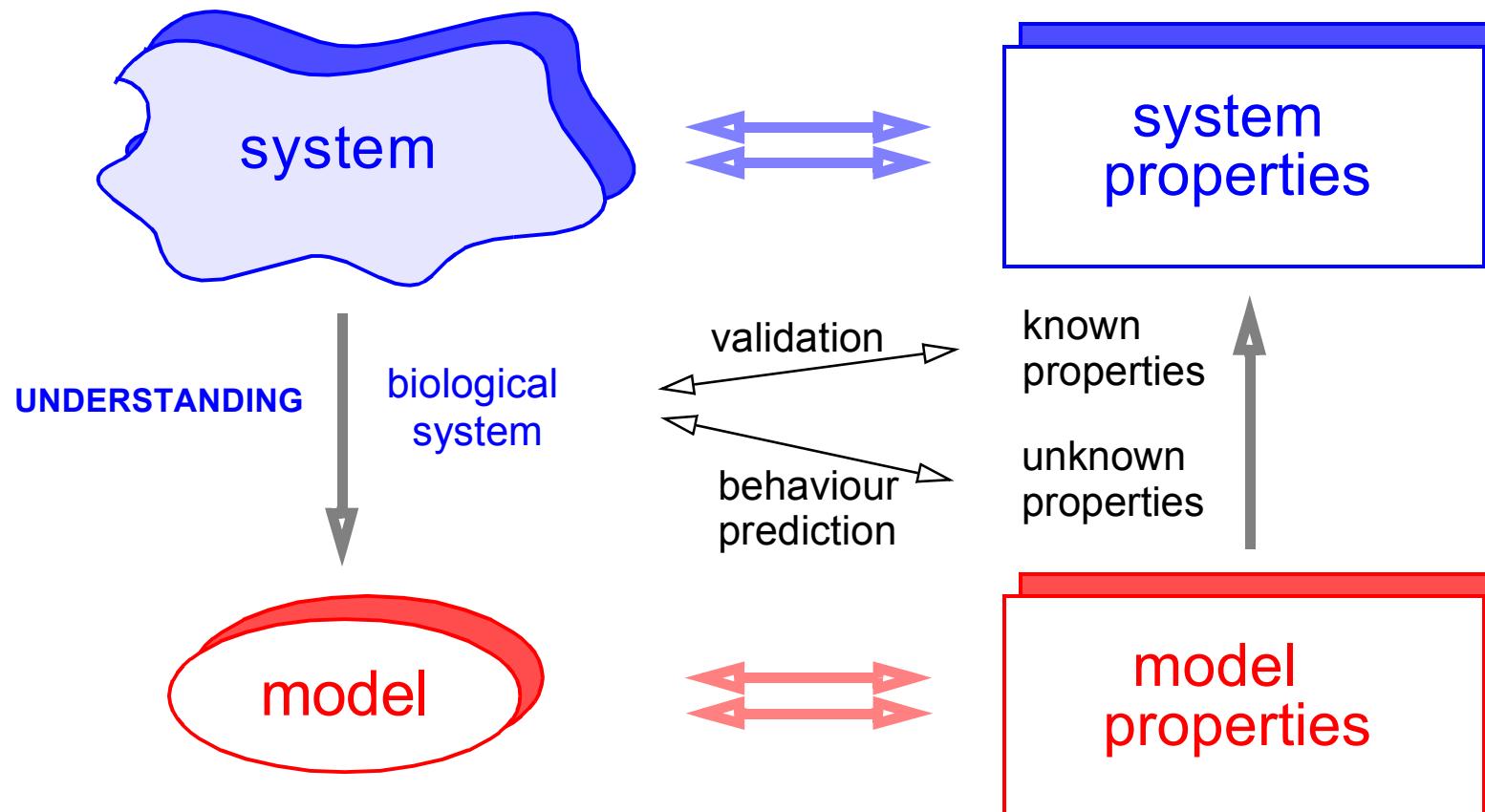
- > colored framework
- > Generalized Hybrid Petri Nets

# **BACKGROUND**

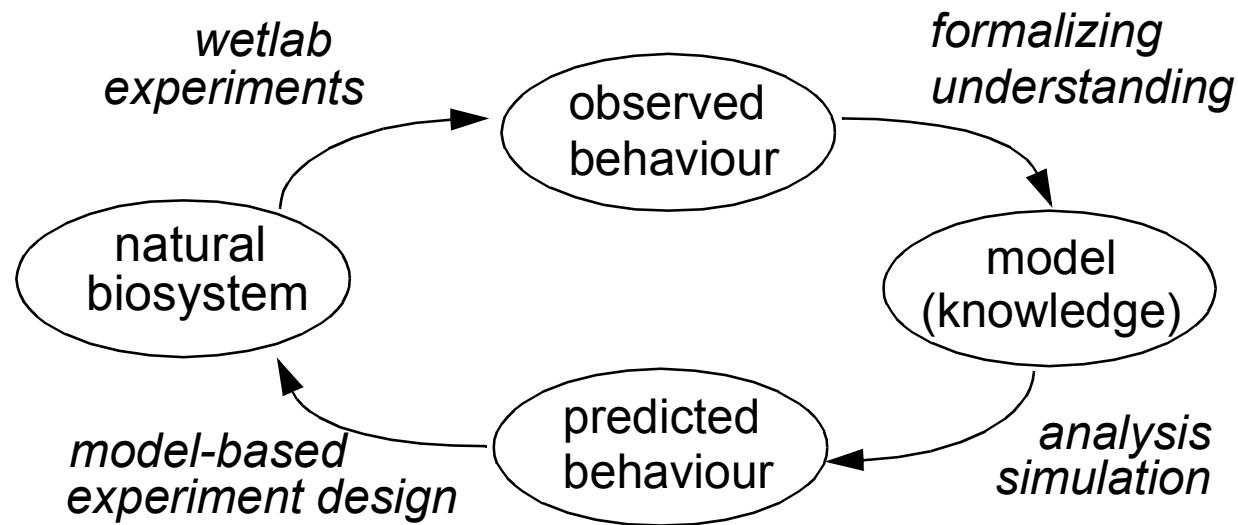




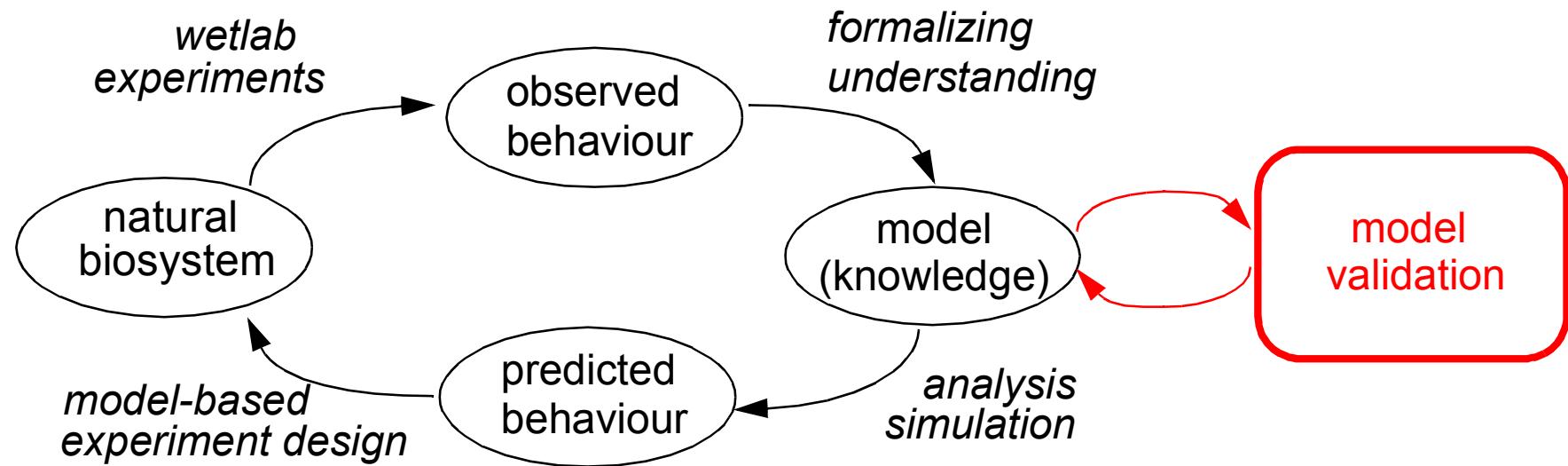




## MODELLING = FORMAL KNOWLEDGE REPRESENTATION

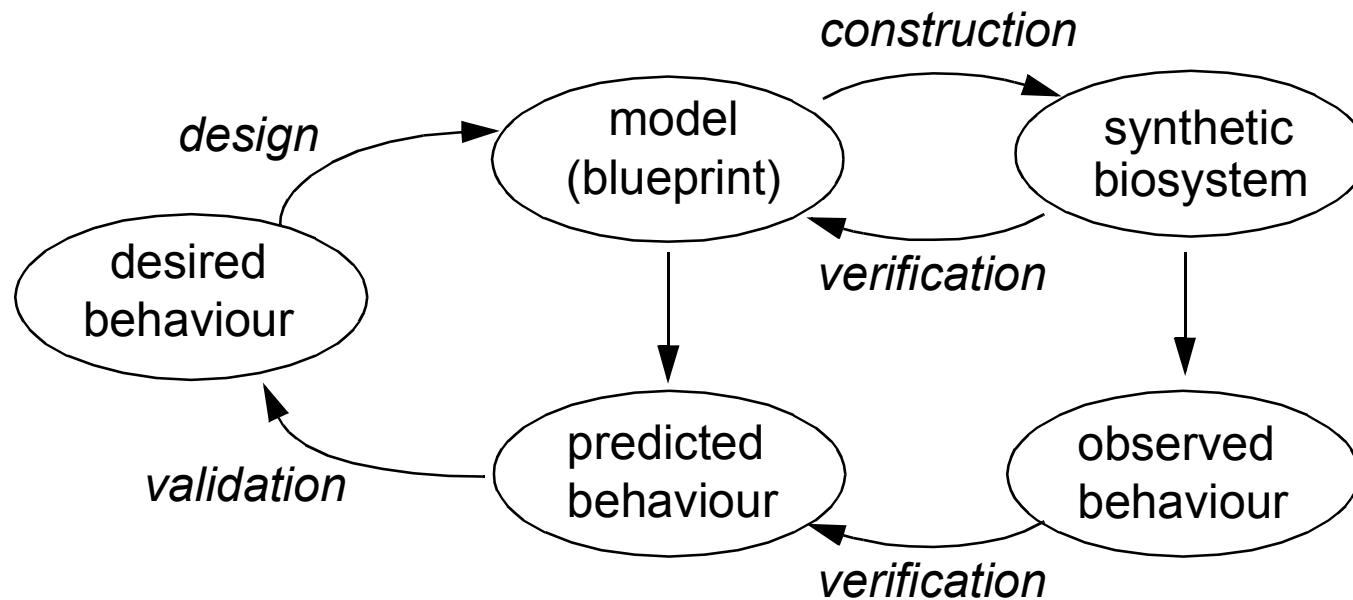


**MODELLING = FORMAL KNOWLEDGE REPRESENTATION**



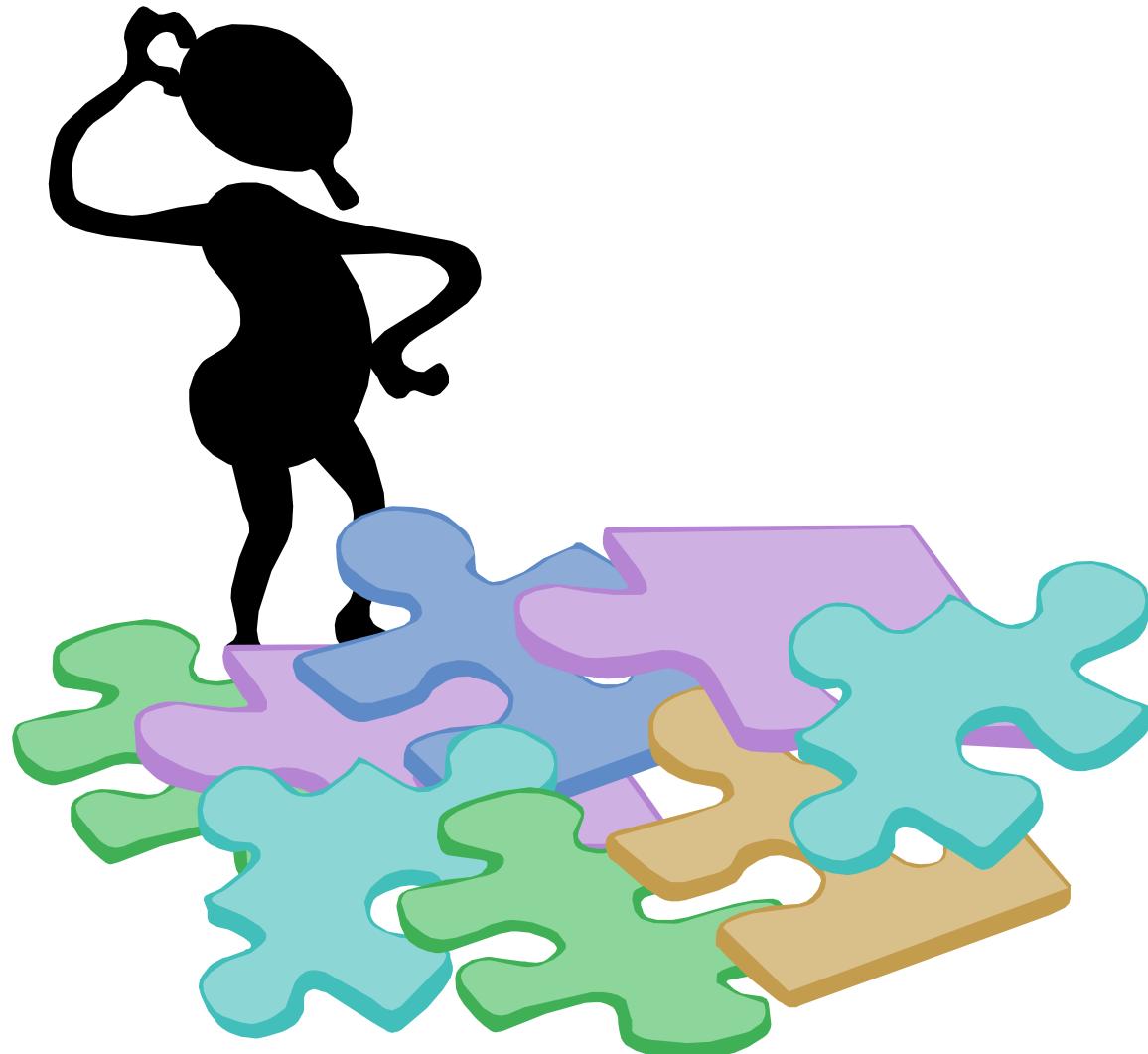
**MODEL VALIDATION = CONFIDENCE INCREASE**

## MODELLING = BLUEPRINT FOR SYSTEM CONSTRUCTION



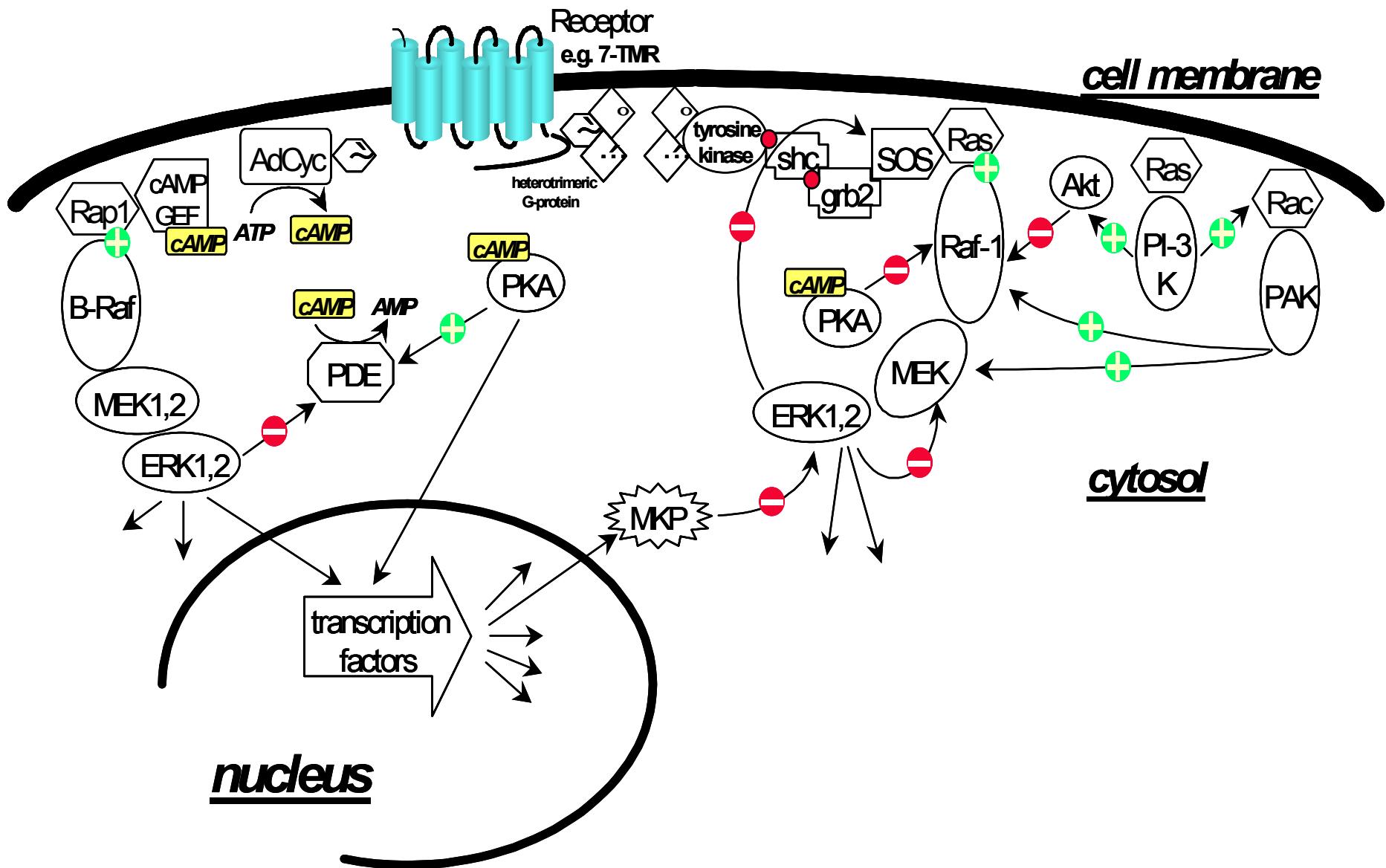
**RELIABLE AND ROBUST ENGINEERING REQUIRES VERIFIED MODELS**

# **WHAT KIND OF MODEL SHOULD BE USED?**



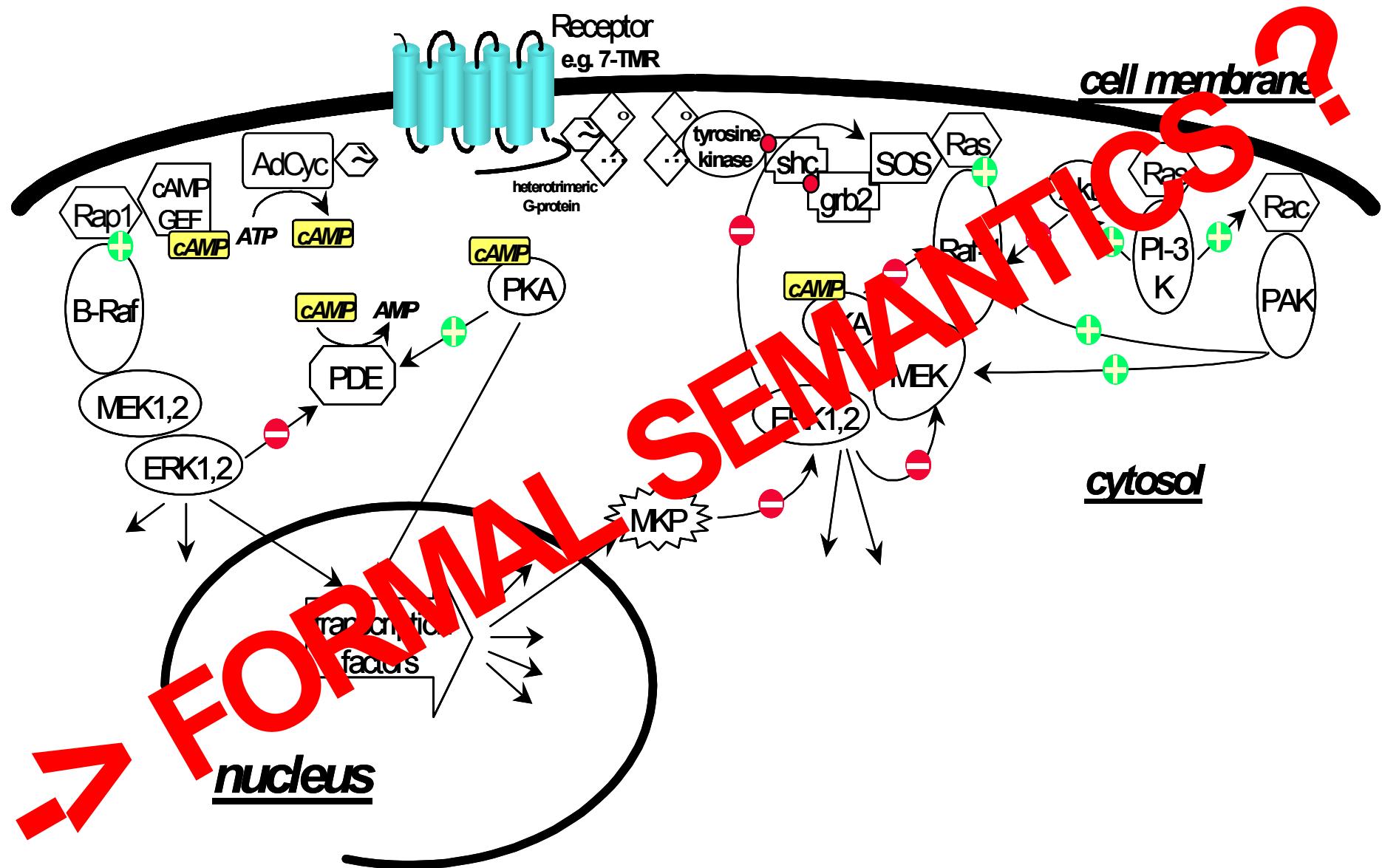
# NETWORK REPRESENTATIONS, Ex1

PN & BioModel Engineering



# NETWORK REPRESENTATIONS, Ex1

PN & BioModel Engineering



## NETWORK REPRESENTATIONS, Ex2

PN & BioModel Engineering

$$\begin{aligned}
 \frac{d\alpha}{dt} &= -v_1 \\
 \frac{d\text{Ste2}}{dt} &= -v_2 + v_3 - v_5 \\
 \frac{d\text{Ste2}_{\text{active}}}{dt} &= v_2 - v_3 - v_4 \\
 \frac{d\text{Sst2}_{\text{active}}}{dt} &= v_{46} - v_{47} \\
 \frac{dG\alpha\beta\gamma}{dt} &= -v_6 + v_9 \\
 \frac{dG\alpha\text{GTP}}{dt} &= v_6 - v_7 - v_8 \\
 \frac{dG\alpha\text{GDP}}{dt} &= v_7 + v_8 - v_9 \\
 \frac{dG\beta\gamma}{dt} &= v_6 - v_9 - v_{10} + v_{11} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\
 &\quad - v_{42} + v_{43} \\
 \frac{d\text{Ste5}}{dt} &= -v_{12} + v_{13} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\
 \frac{d\text{Ste11}}{dt} &= -v_{12} + v_{13} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\
 \frac{d\text{Ste7}}{dt} &= -v_{14} + v_{15} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\
 \frac{d\text{Fus3}}{dt} &= -v_{14} + v_{15} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} - v_{29} \\
 &\quad + v_{30} + v_{33} \\
 \frac{d\text{Ste20}}{dt} &= -v_{18} + v_{19} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32}
 \end{aligned}$$

$$\begin{aligned}
 v_1 &= \alpha[t] \cdot \text{Bar1}_{\text{active}}[t] \cdot k_1 \\
 v_2 &= \text{Ste2}[t] \cdot \alpha[t] \cdot k_2 \\
 v_3 &= \text{Ste2}_{\text{active}}[t] \cdot k_3 \\
 v_4 &= \text{Ste2}_{\text{active}}[t] \cdot k_4 \\
 v_5 &= \text{Ste2}[t] \cdot k_5 \\
 v_6 &= \text{Ste2}_{\text{active}}[t] \cdot G\alpha\beta\gamma[t] \cdot k_6 \\
 v_7 &= G\alpha\text{GTP}[t] \cdot k_7 \\
 v_8 &= G\alpha\text{GTP}[t] \cdot \text{Sst2}_{\text{active}}[t] \cdot k_8 \\
 v_9 &= G\alpha\text{GDP}[t] \cdot G\beta\gamma[t] \cdot k_9 \\
 v_{10} &= G\beta\gamma[t] \cdot C[t] \cdot k_{10} \\
 v_{11} &= D[t] \cdot k_{11} \\
 v_{12} &= \text{Ste5}[t] \cdot \text{Ste11}[t] \cdot k_{12} \\
 v_{13} &= A[t] \cdot k_{13} \\
 v_{14} &= \text{Ste7}[t] \cdot \text{Fus3}[t] \cdot k_{14} \\
 v_{15} &= B[t] \cdot k_{15} \\
 v_{16} &= A[t] \cdot B[t] \cdot k_{16} \\
 v_{17} &= C[t] \cdot k_{17} \\
 v_{18} &= D[t] \cdot \text{Ste20}[t] \cdot k_{18}
 \end{aligned}$$

## NETWORK REPRESENTATIONS, Ex2

PN & BioModel Engineering

$$\begin{aligned}
 \frac{d\alpha}{dt} &= -v_1 \\
 \frac{d\text{Ste2}}{dt} &= -v_2 + v_3 - v_5 \\
 \frac{d\text{Ste2}_{\text{active}}}{dt} &= v_2 - v_3 - v_4 \\
 \frac{d\text{Sst2}_{\text{active}}}{dt} &= v_{46} - v_{47} \\
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 \end{aligned}$$

$$\begin{aligned}
 v_1 &= \alpha[t] \cdot \text{Bar1}_{\text{active}}[t] \\
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 v_3 &= \text{Ste2}_{\text{active}}[t] \cdot k_3 \\
 v_4 &= \text{Ste2}_{\text{inactive}}[t] \cdot k_4 \\
 v_5 &= \text{Ste2}[t] \cdot k_5 \\
 v_6 &= \text{Ste2}_{\text{active}}[t] \cdot G\alpha\beta\gamma[t] \cdot k_6 \\
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 v_9 &= G\alpha\text{GDP}[t] \cdot G\beta\gamma[t] \cdot k_9 \\
 v_{10} &= G\beta\gamma[t] \cdot C[t] \cdot k_{10} \\
 v_{11} &= D[t] \cdot k_{11} \\
 v_{12} &= \text{Ste5}[t] \cdot \text{Ste11}[t] \cdot k_{12} \\
 v_{13} &= A[t] \cdot k_{13} \\
 v_{14} &= \text{Ste7}[t] \cdot \text{Fus3}[t] \cdot k_{14} \\
 v_{15} &= B[t] \cdot k_{15} \\
 v_{16} &= A[t] \cdot B[t] \cdot k_{16} \\
 v_{17} &= C[t] \cdot k_{17} \\
 v_{18} &= D[t] \cdot \text{Ste20}[t] \cdot k_{18}
 \end{aligned}$$

**READABILITY?**

**knowledge**

-> **PROBLEM 1**

-> *uncertain*

-> *growing, changing*

-> *distributed over independent data bases, papers, journals, . . .*

**various, mostly ambiguous representations**

-> **PROBLEM 2**

-> *verbose descriptions*

-> *diverse graphical representations*

-> *contradictory and / or fuzzy statements*

**network structure**

-> **PROBLEM 3**

-> *tend to grow fast*

-> *dense, apparently unstructured*

-> *hard to read*

- knowledge -> **PROBLEM 1**
  - > *uncertain*
  - > *growing, changing*
  - > *distributed over independent data bases, papers, journals, . . .*
  
- various, mostly ambiguous representations -> **PROBLEM 2**
  - > *verbose descriptions*
  - > *diverse graphical representations*
  - > *contradictory and / or fuzzy statements*
  
- network structure -> **PROBLEM 3**
  - > *tend to grow fast*
  - > *dense, apparently unstructured*
  - > *hard to read*

-> MODELS ARE FULL OF ASSUMPTIONS <-

- **readable**

- > *fault avoidance*
- > *informal = cartoon-like representations ?*

- **analysable**

- > *formal = mathematical representations*

- **executable**

- > *to experience the model*

- **unifying power**

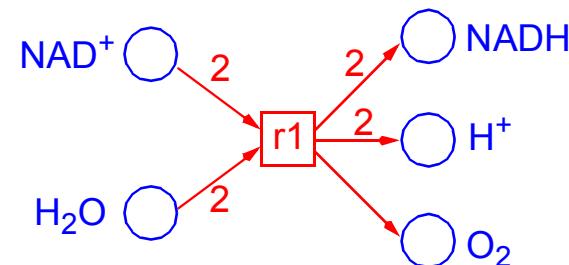
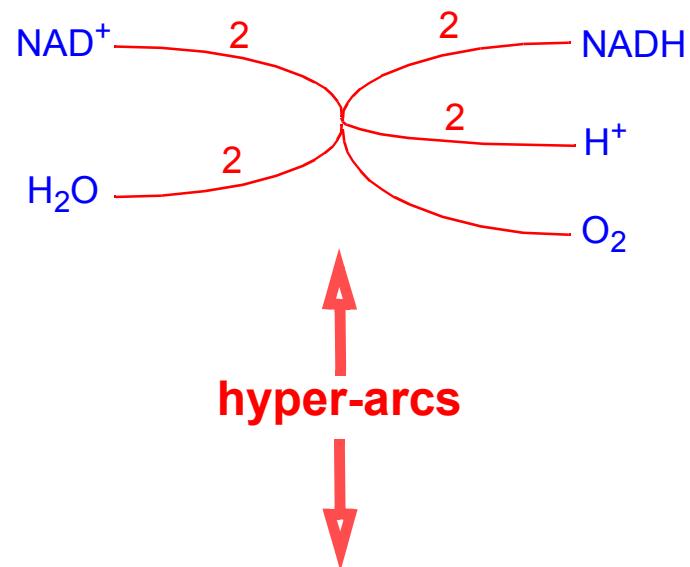
- > *high-level description for various analysis approaches*

... .

***ARE NETWORKS  
OF BIOCHEMICAL  
REACTIONS***

... .

***NATURALLY  
EXPRESSIBLE AS  
PETRI NETS***

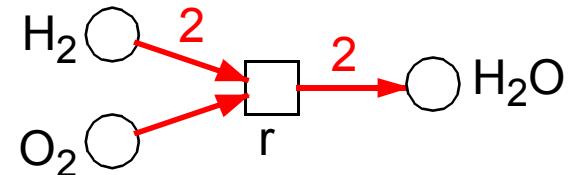
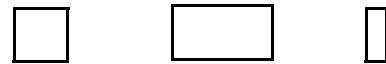


# **PETRI NETS**

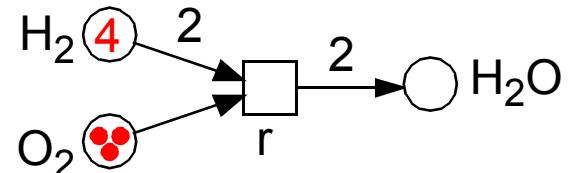
## **- A CRASH COURSE -**

- transitions**  
-> active system components   
  
- places**  
-> passive system components    
-> local conditions

- transitions  
-> active system components
  
- places  
-> passive system components  
-> local conditions
  
- weightes arcs  
-> causality

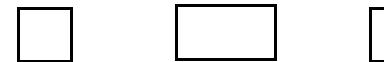


- transitions  
-> active system components
- places  
-> passive system components  
-> local conditions
- weightes arcs  
-> causality
- condition's state  
-> token(s) on its place
- system state  
-> marking



( $H_2=4$ ,  $O_2=3$ ,  $H_2O=0$ )

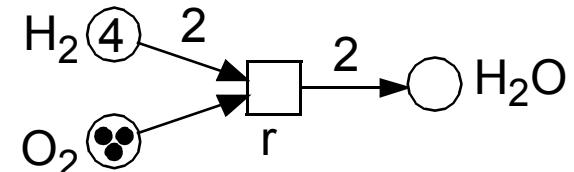
- transitions  
-> active system components



- places  
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- weightes arcs  
-> causality
- condition's state  
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- system state  
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( $H_2=4$ ,  $O_2=3$ ,  $H_2O=0$ )

$PN = (P, T, F, m_0)$ ,  $F: (P \times T) \cup (T \times P) \rightarrow N_0$ ,  $m_0: P \rightarrow N_0$

- Petri nets = bipartite directed multigraphs

C. A. PETRI

INTERPRETATIONS OF NET THEORY

GMD, INTERNAL REPORT 75-07, 2ND IMPROVED EDITION 1976

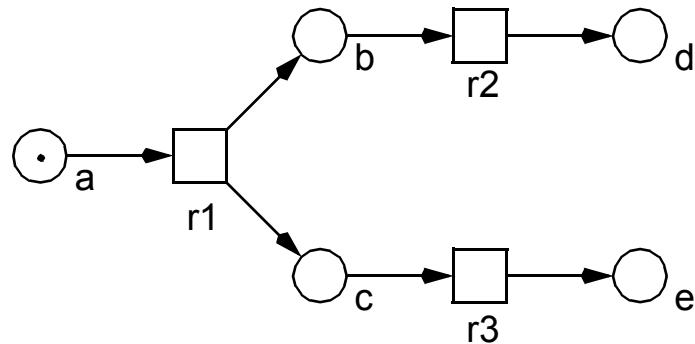
places	transitions
state elements	transitional elements
conditions	events/facts
statements	dependencies
model domains	specifications
chemical compounds	chemical reactions
open one-point sets	closed one-point sets
channels	offices
languages	translators
products	production activities

- **an action may happen, if** -> prerequisite
- > *all preconditions are fulfilled (corresponding to the arc weights);*
  
- **if an action happens, then** -> firing behaviour
- > *tokens are removed from all preconditions (corresponding to the arc weights), and*
- > *tokens are added to all postconditions (corresponding to the arc weights);*
  
- **action happens (firing of a transition)** -> model assumptions
- > *atomic*
- > *time-less*



# PARTIAL ORDER VERSUS INTERLEAVING SEMANTICS

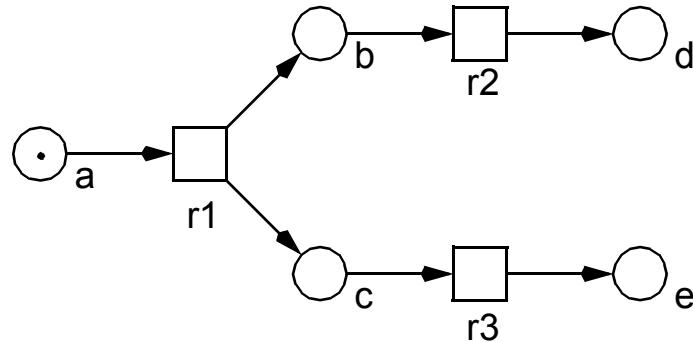
PN & BioModel Engineering



- ❑ **order between r1 - r2 and r1 - r3**
  - > *causality*                     $x < y [ x-y ]$
  - > *dependency*
  
- ❑ **no order between r2 , r3**
  - > *concurrency*                 $x \parallel y$
  - > *independency*

# PARTIAL ORDER VERSUS INTERLEAVING SEMANTICS

PN & BioModel Engineering



- ❑ order between  $r_1 - r_2$  and  $r_1 - r_3$ 
  - > causality  $x < y [x-y]$
  - > dependency
- ❑ no order between  $r_2 , r_3$ 
  - > concurrency  $x \parallel y$
  - > independency

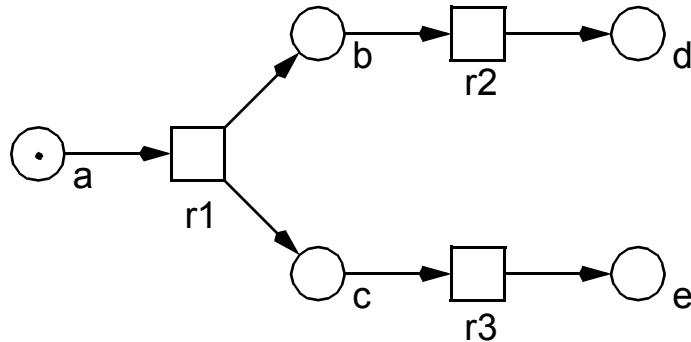
## ❑ possible interleaving runs

- >  $r_1 - r_2 - r_3$
- >  $r_1 - r_3 - r_2$
- > *totally ordered runs*

-> **INTERLEAVING SEMANTICS**  
all totally ordered runs

# PARTIAL ORDER VERSUS INTERLEAVING SEMANTICS

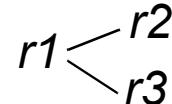
PN & BioModel Engineering



- possible interleaving runs
  - >  $r1 - r2 - r3$
  - >  $r1 - r3 - r2$
  - > totally ordered runs
- > INTERLEAVING SEMANTICS  
all totally ordered runs

- order between  $r1 - r2$  and  $r1 - r3$ 
  - > causality  $x < y [ x-y ]$
  - > dependency
- no order between  $r2 , r3$ 
  - > concurrency  $x \parallel y$
  - > independency

- one partial order run



“true concurrency semantics”

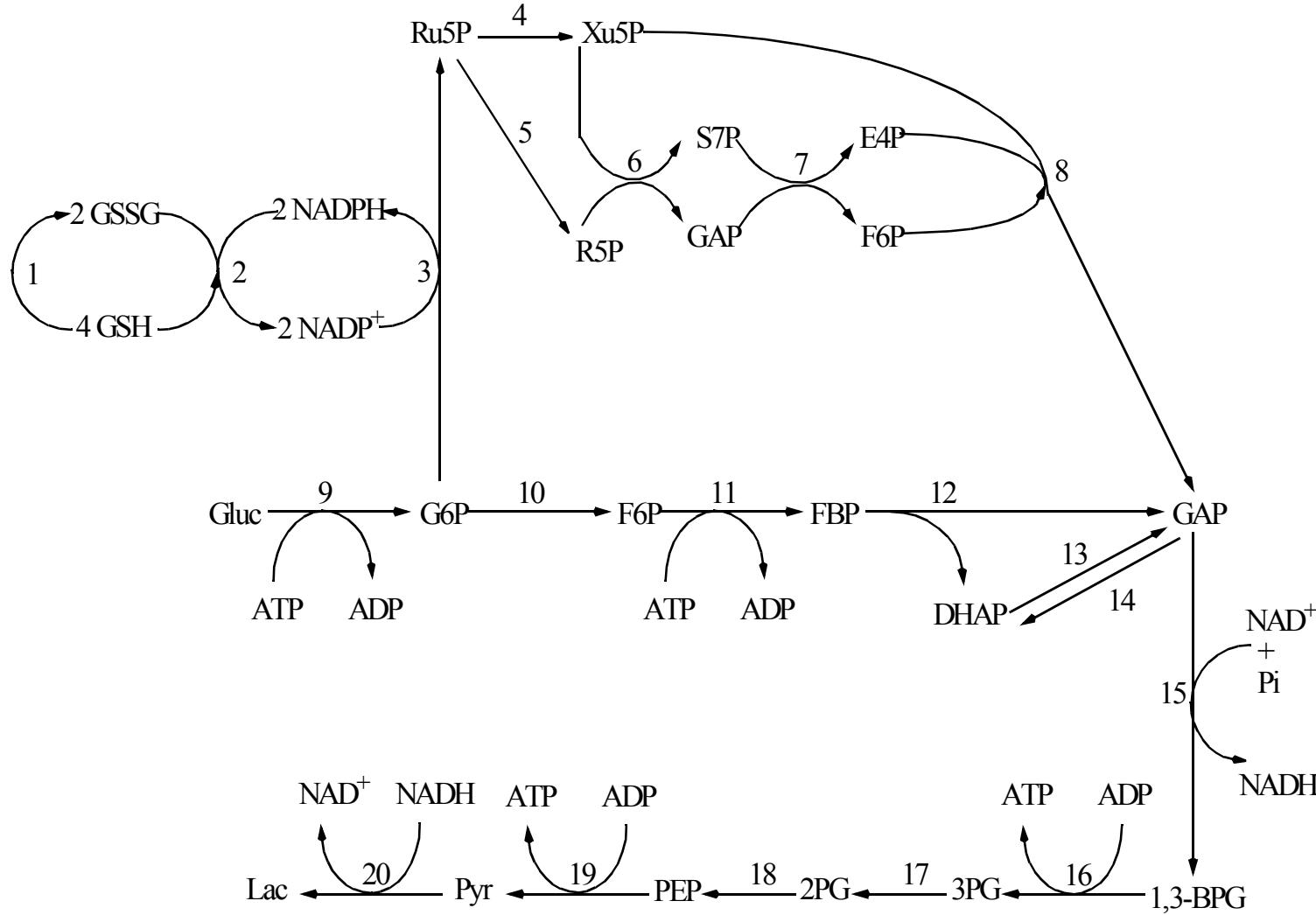
- > PARTIAL ORDER SEMANTICS  
all partially ordered runs

# **BIO PETRI NETS - SOME EXAMPLES**

# Ex1 - Glycolysis and Pentose Phosphate Pathway

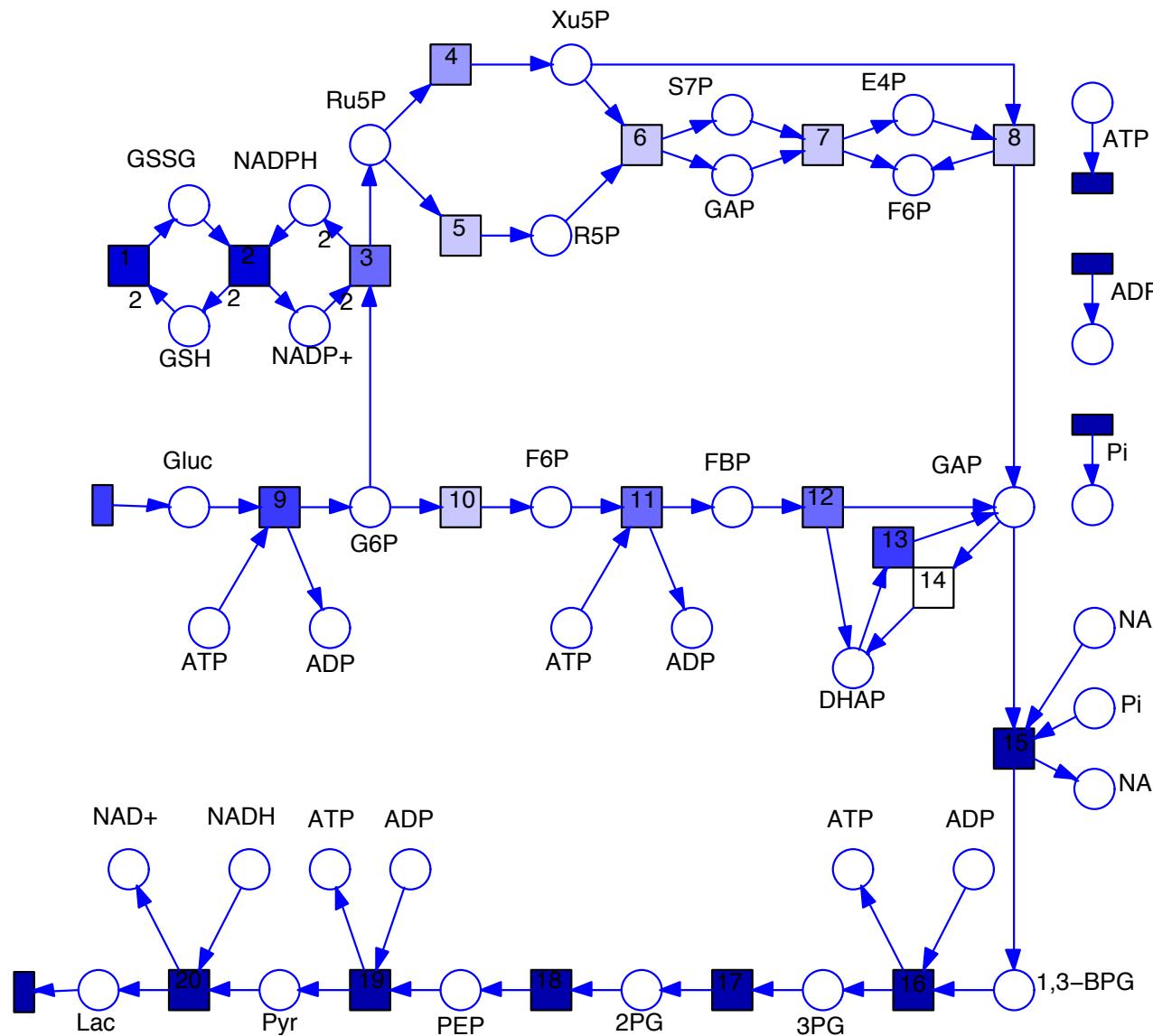
PN & BioModel Engineering

[Reddy 1993]



# Ex1 - Glycolysis and Pentose Phosphate Pathway

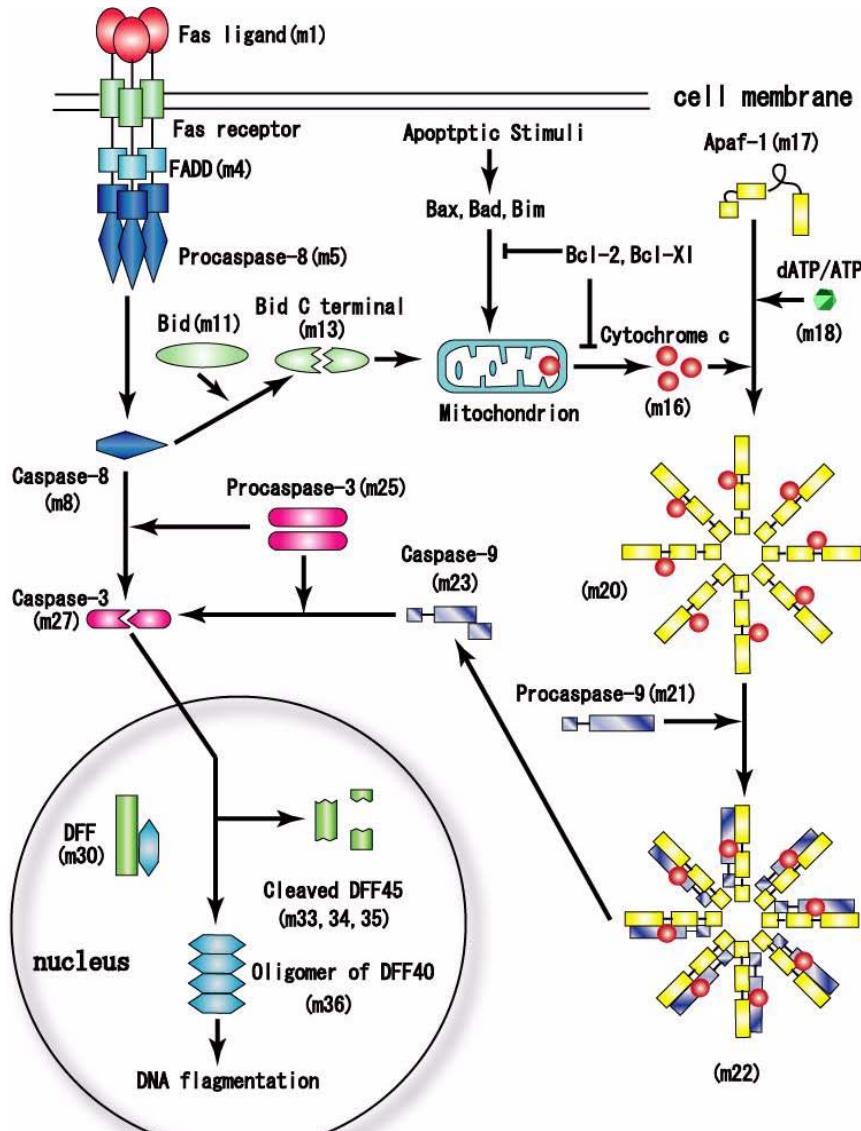
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[Reddy 1993]  
[Heiner 1998]

## Ex2: APOPTOSIS IN MAMMALIAN CELLS

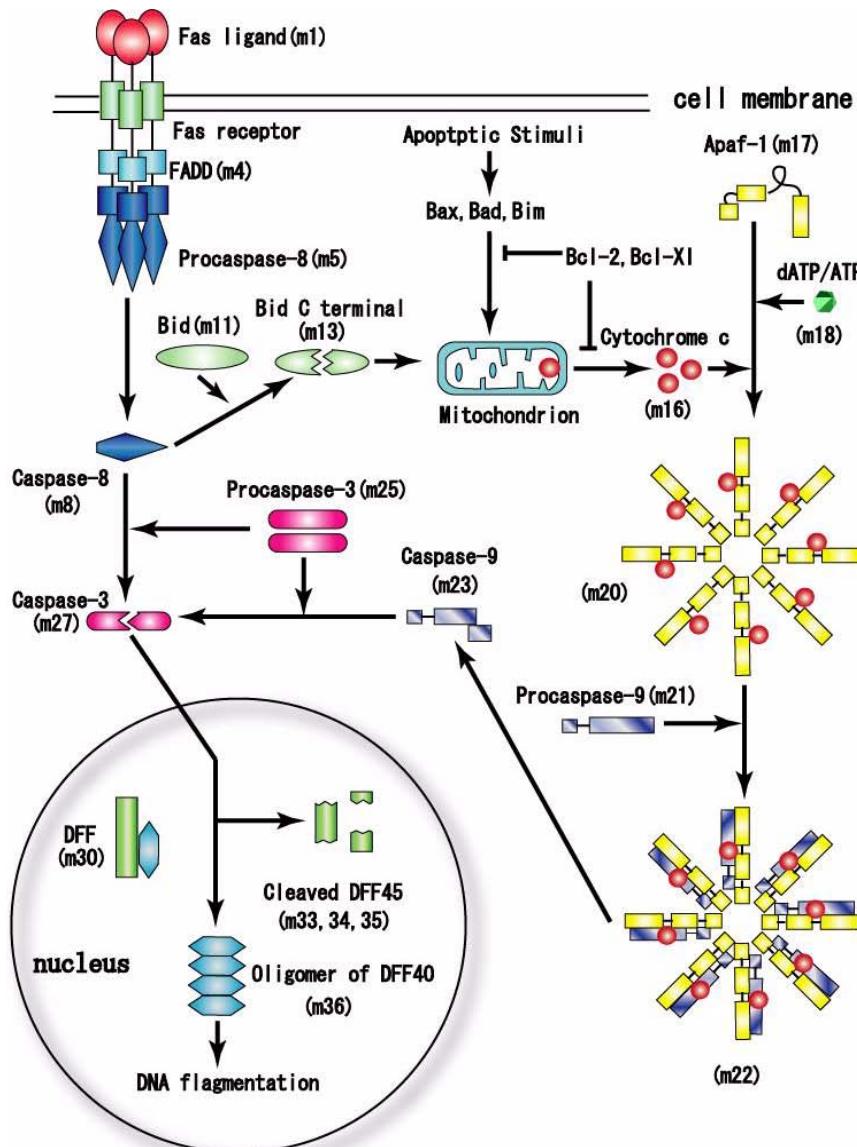
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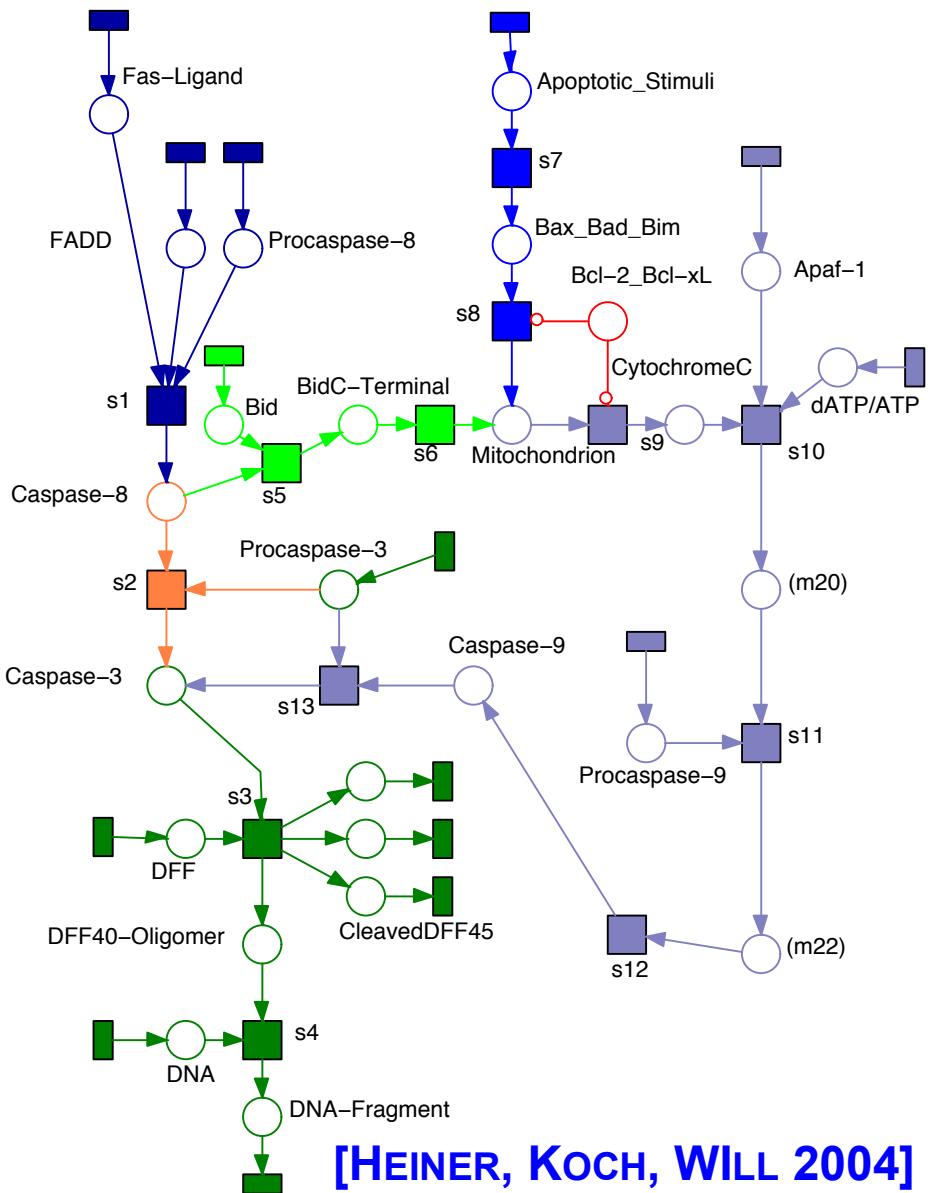
[GON 2003]

## Ex2: APOPTOSIS IN MAMMALIAN CELLS

PN & BioModel Engineering



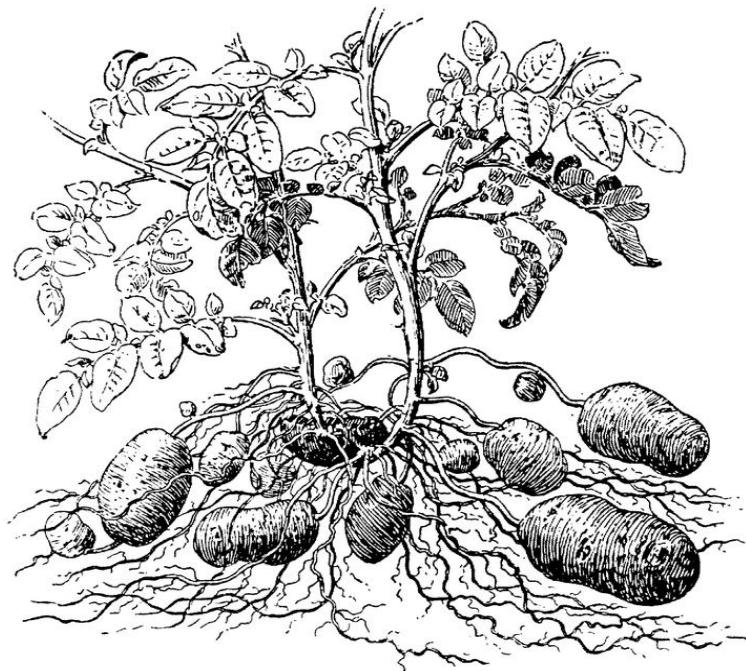
[GON 2003]



[HEINER, KOCH, WILL 2004]

# Ex3 - Carbon Metabolism in Potato Tuber

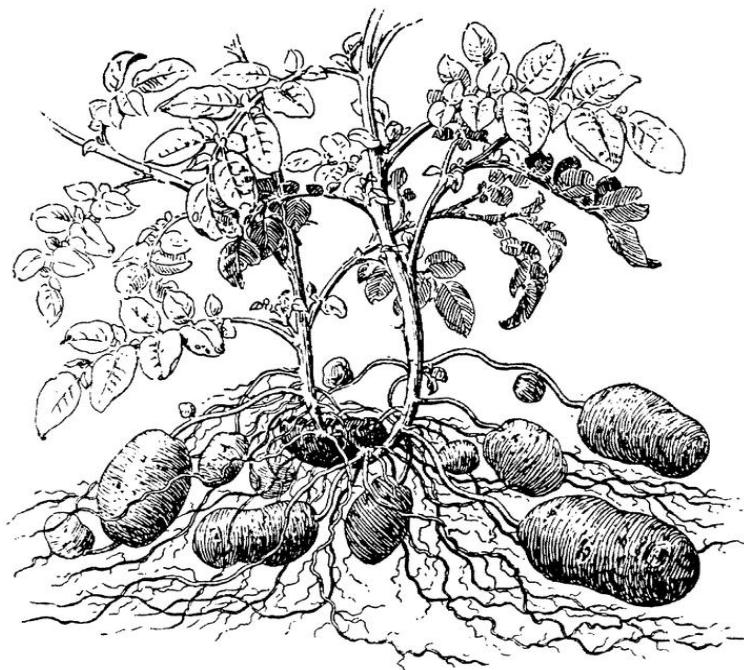
PN & BioModel Engineering



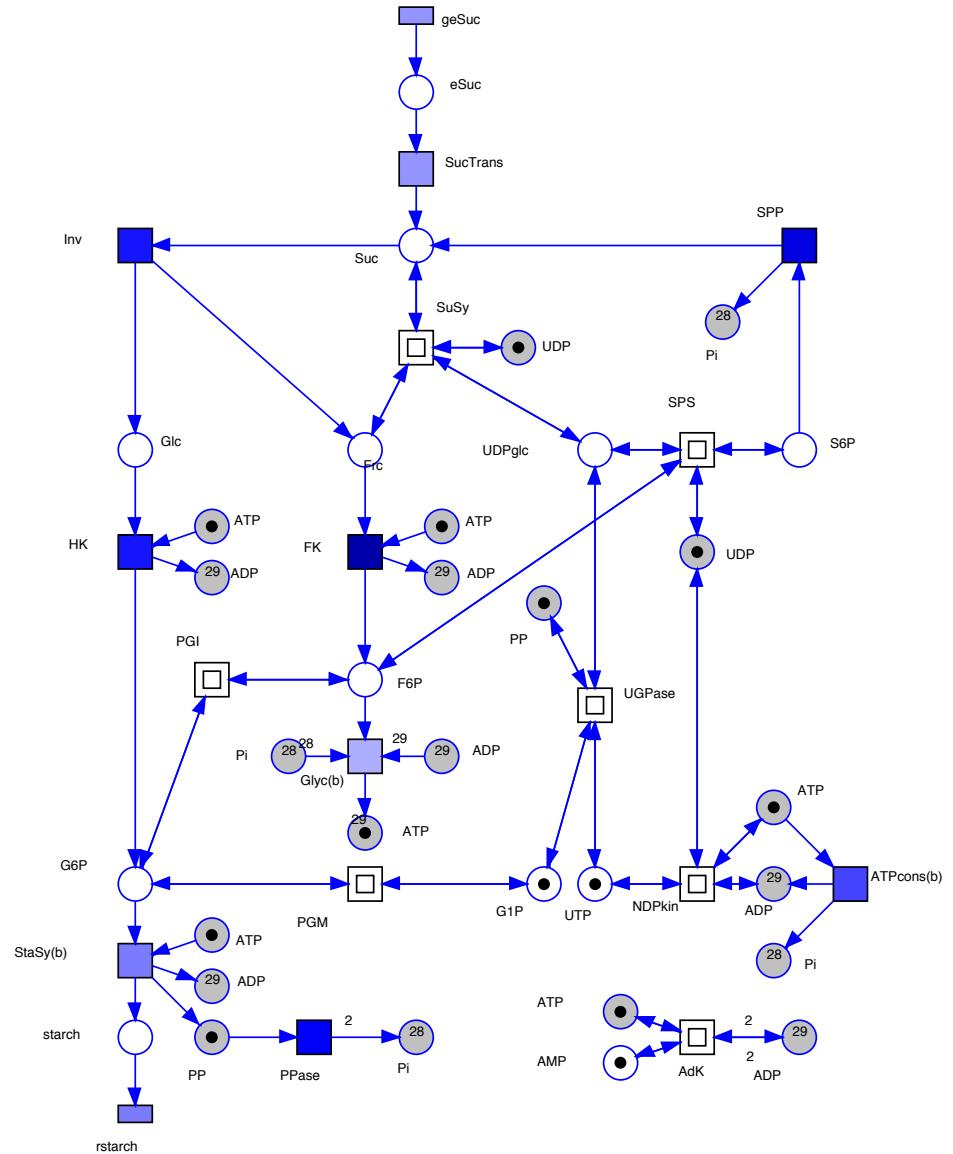
[Koch, JUNKER, HEINER 2005]

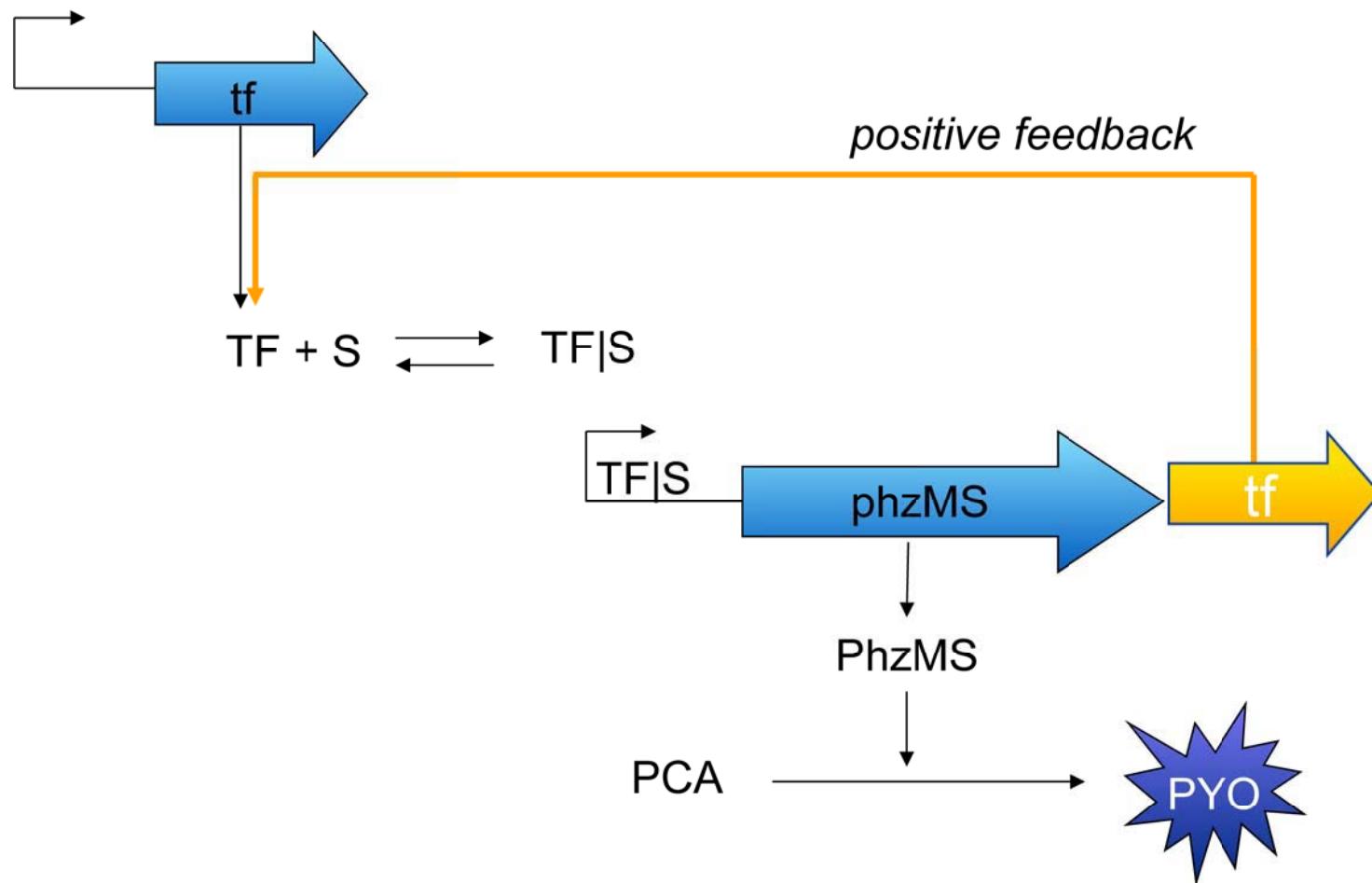
# Ex3 - Carbon Metabolism in Potato Tuber

PN & BioModel Engineering



[Koch, JUNKER, HEINER 2005]

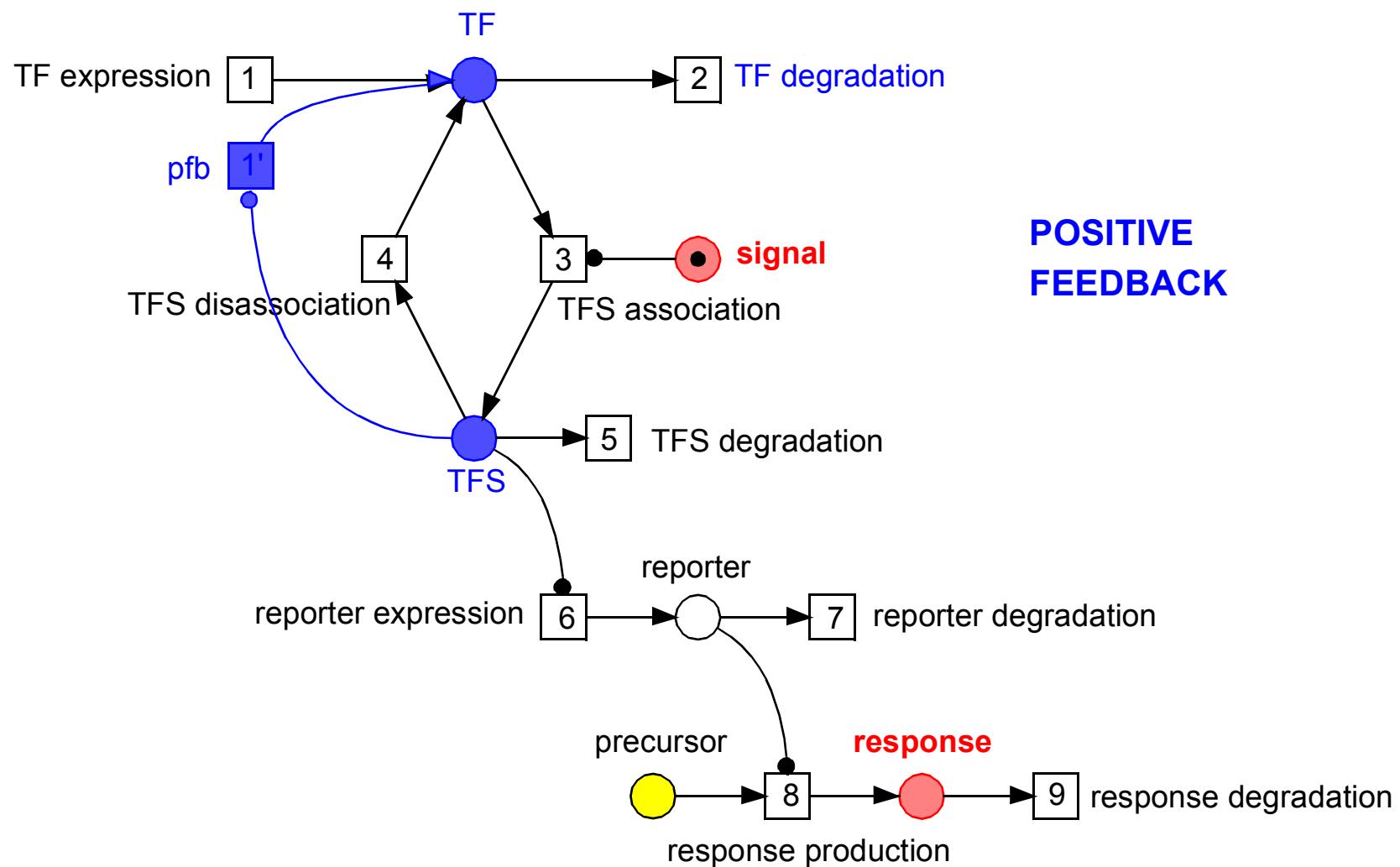




[GILBERT, HEINER, ROSSER, FULTON, GU, TRYBILLO 2008]

## Ex4 - BIOSENSOR

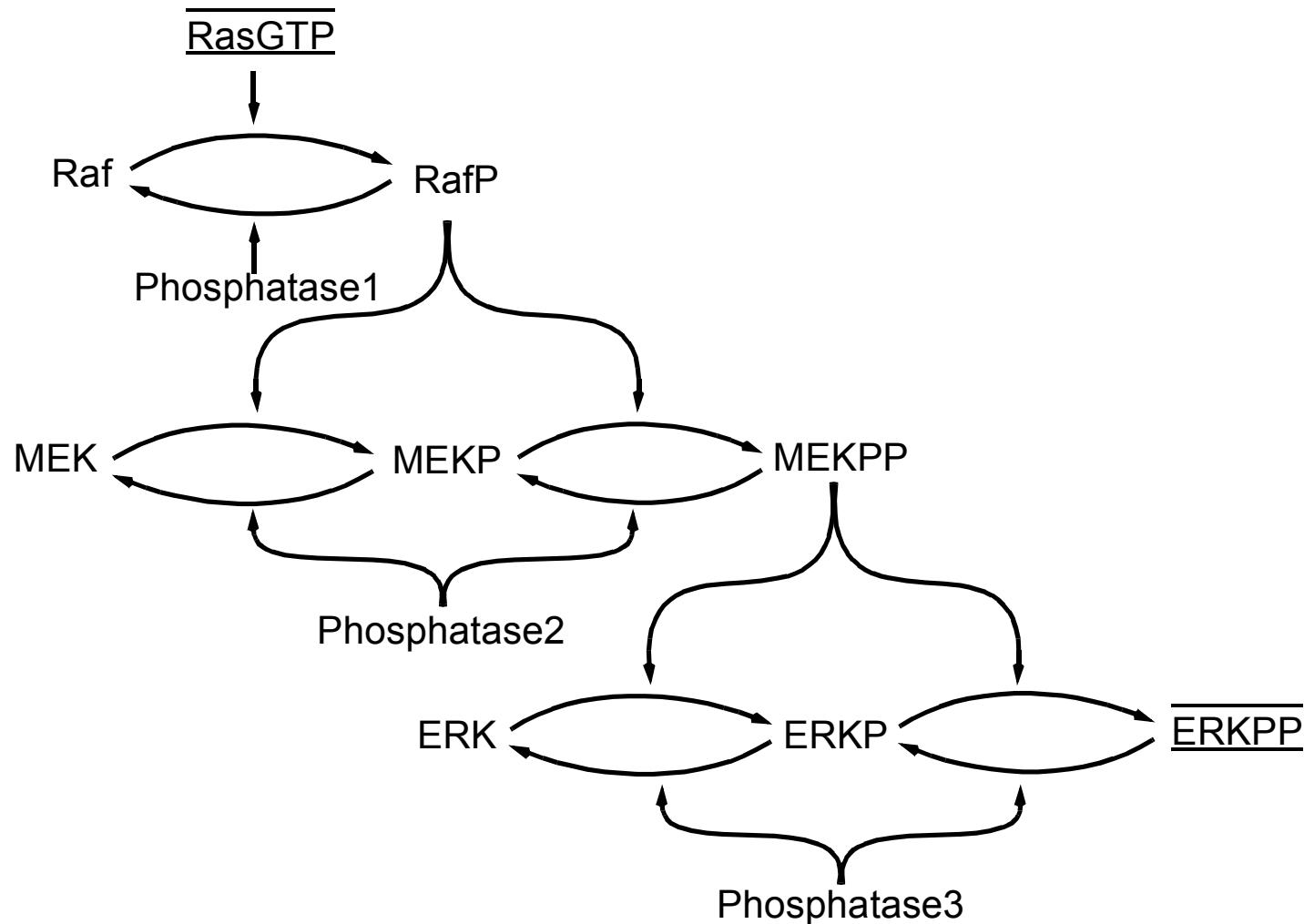
PN & BioModel Engineering



[GILBERT, HEINER, ROSSER, FULTON, GU, TRYBILLO 2008]

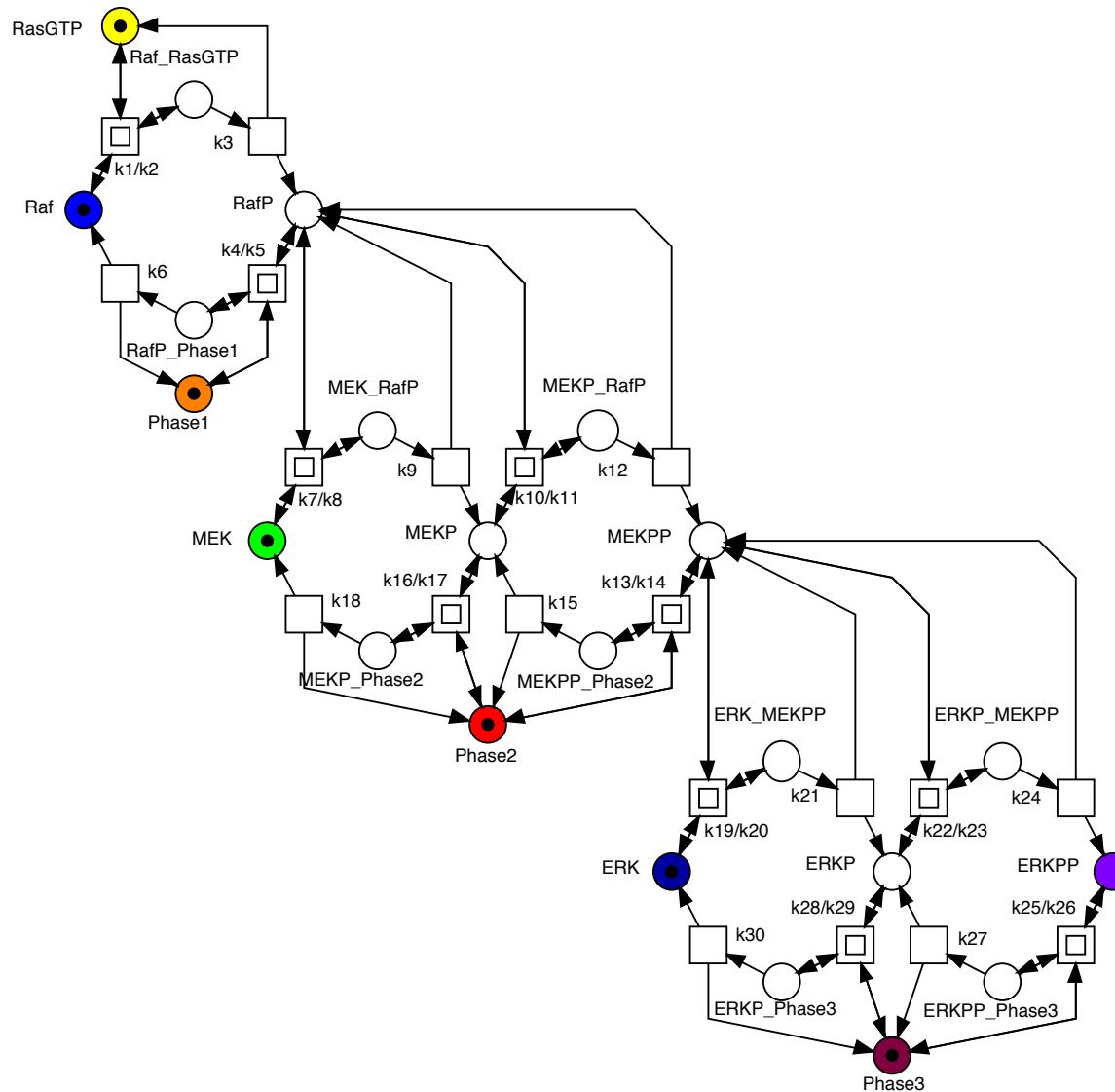
## Ex5 - SIGNALLING CASCADE

PN & BioModel Engineering



## Ex5 - SIGNALLING CASCADE

PN & BioModel Engineering

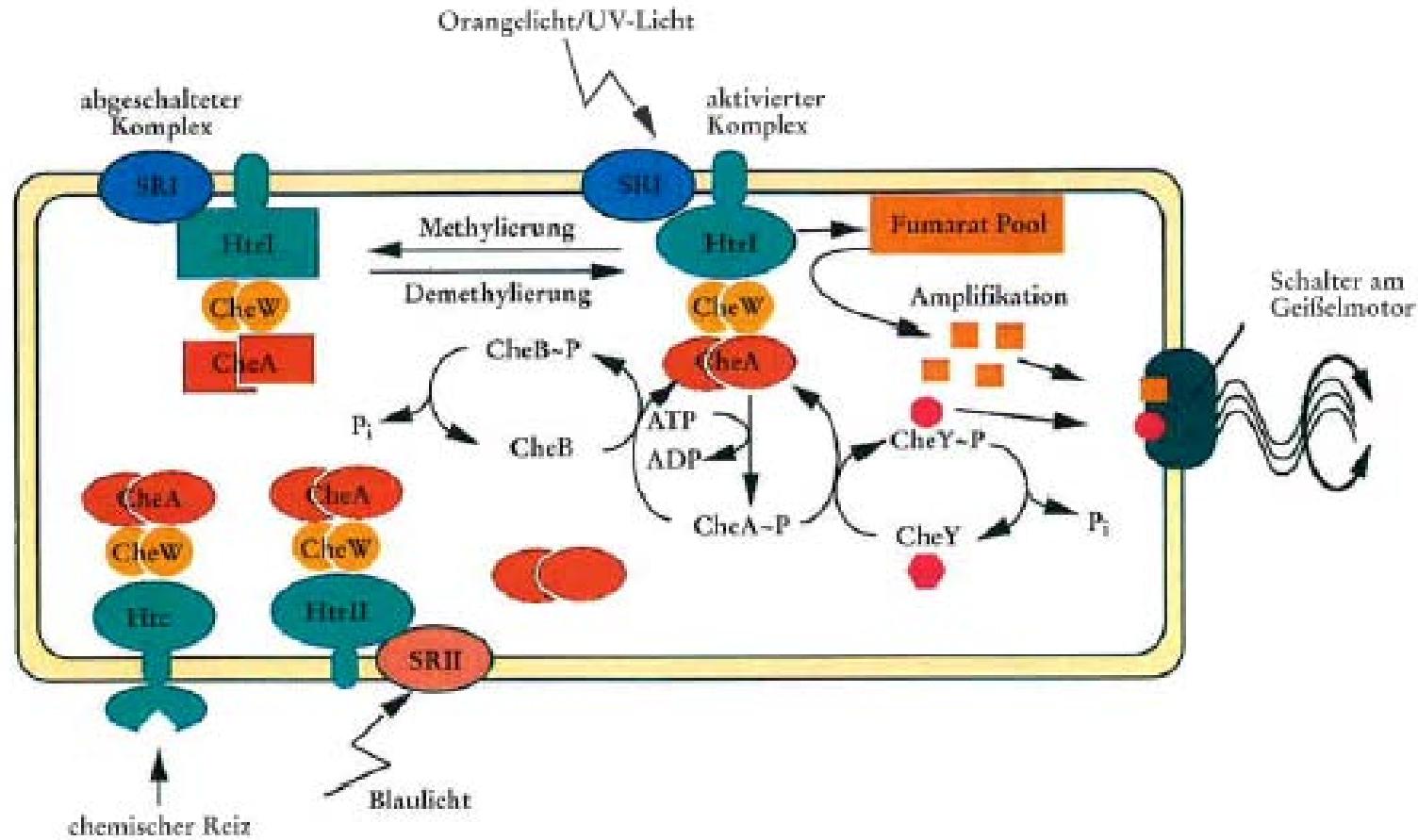


[GILBERT,  
HEINER,  
LEHRACK 2007]

[HEINER,  
GILBERT,  
DONALDSON 2008]

# Ex6 - SWITCH CYCLE HALOBACTERIUM SALINARUM

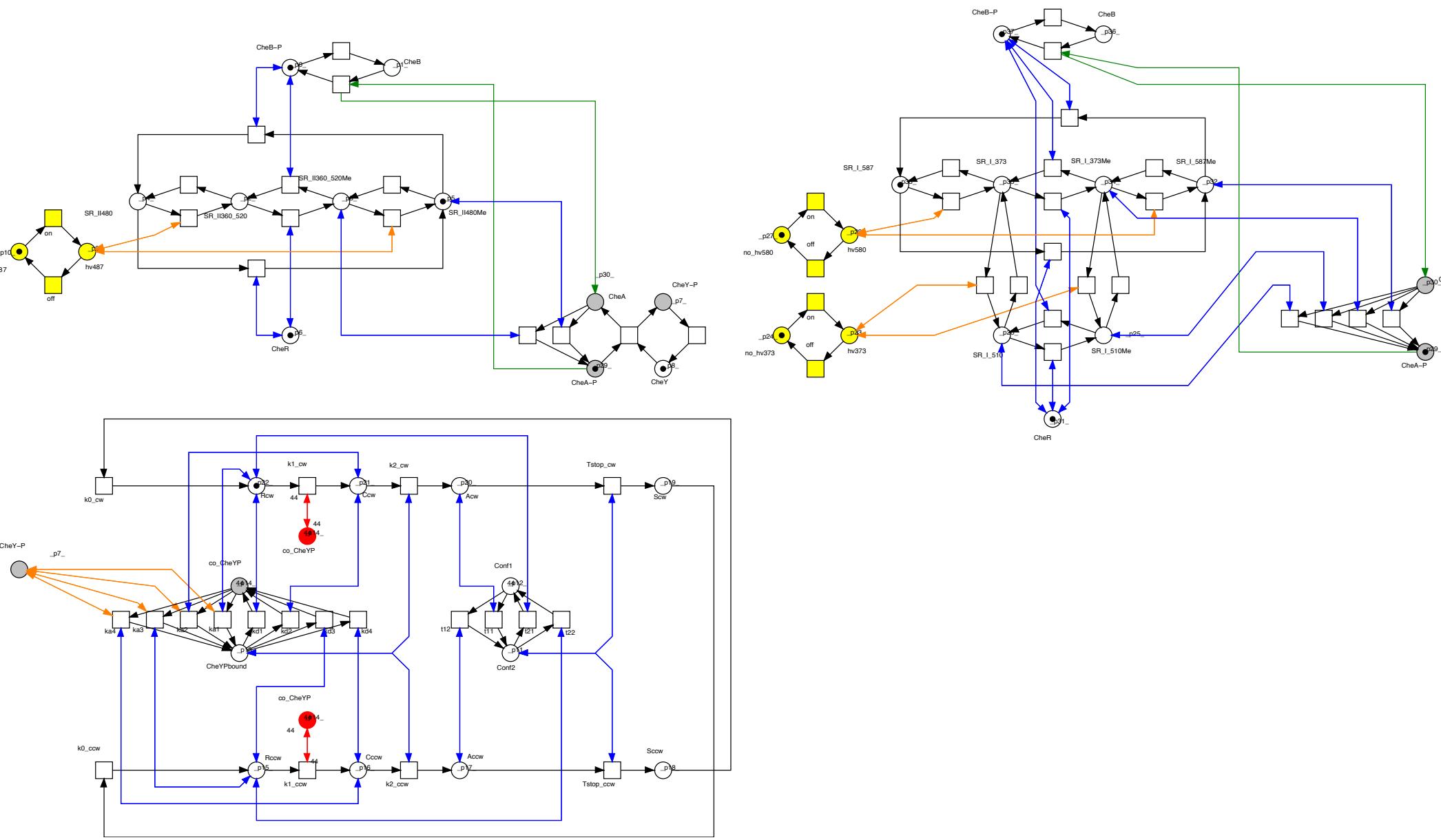
PN & BioModel Engineering



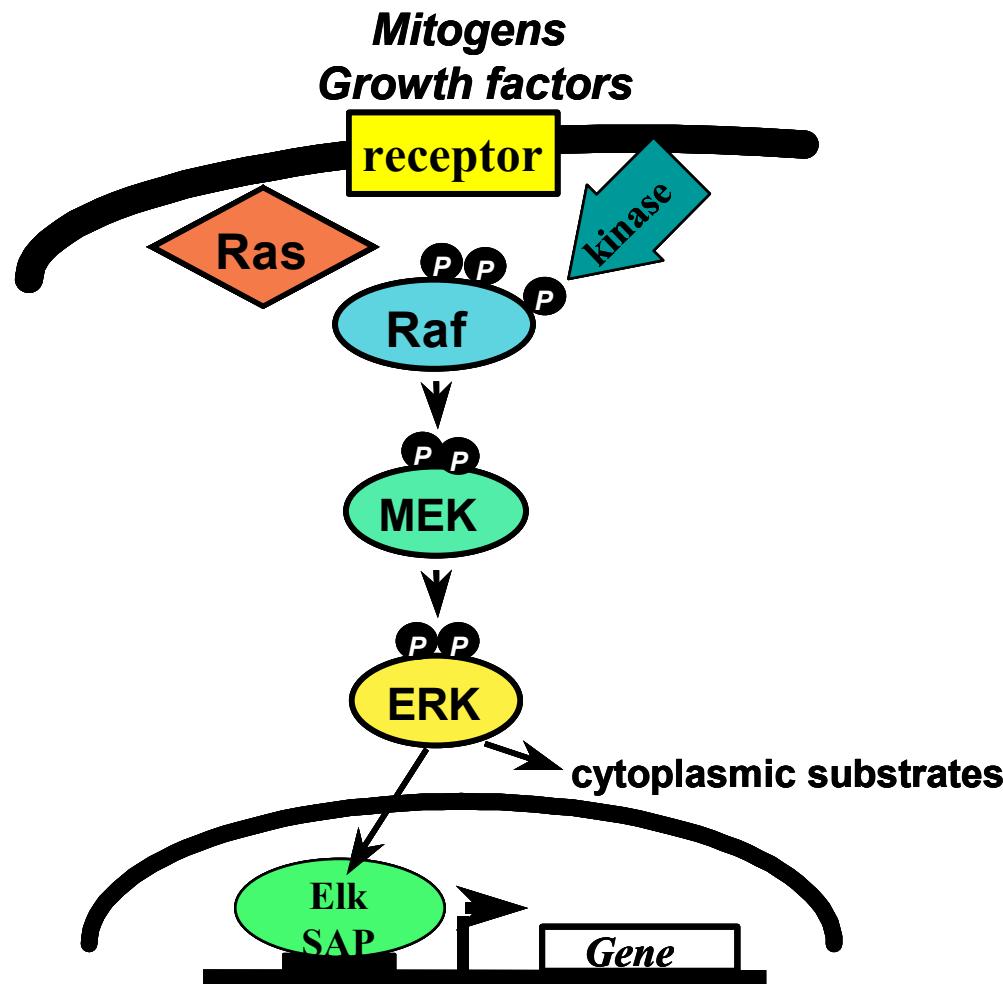
[MARWAN, OESTERHELT 1999]

# Ex6 - SWITCH CYCLE HALOBACTERIUM SALINARUM

PN & BioModel Engineering

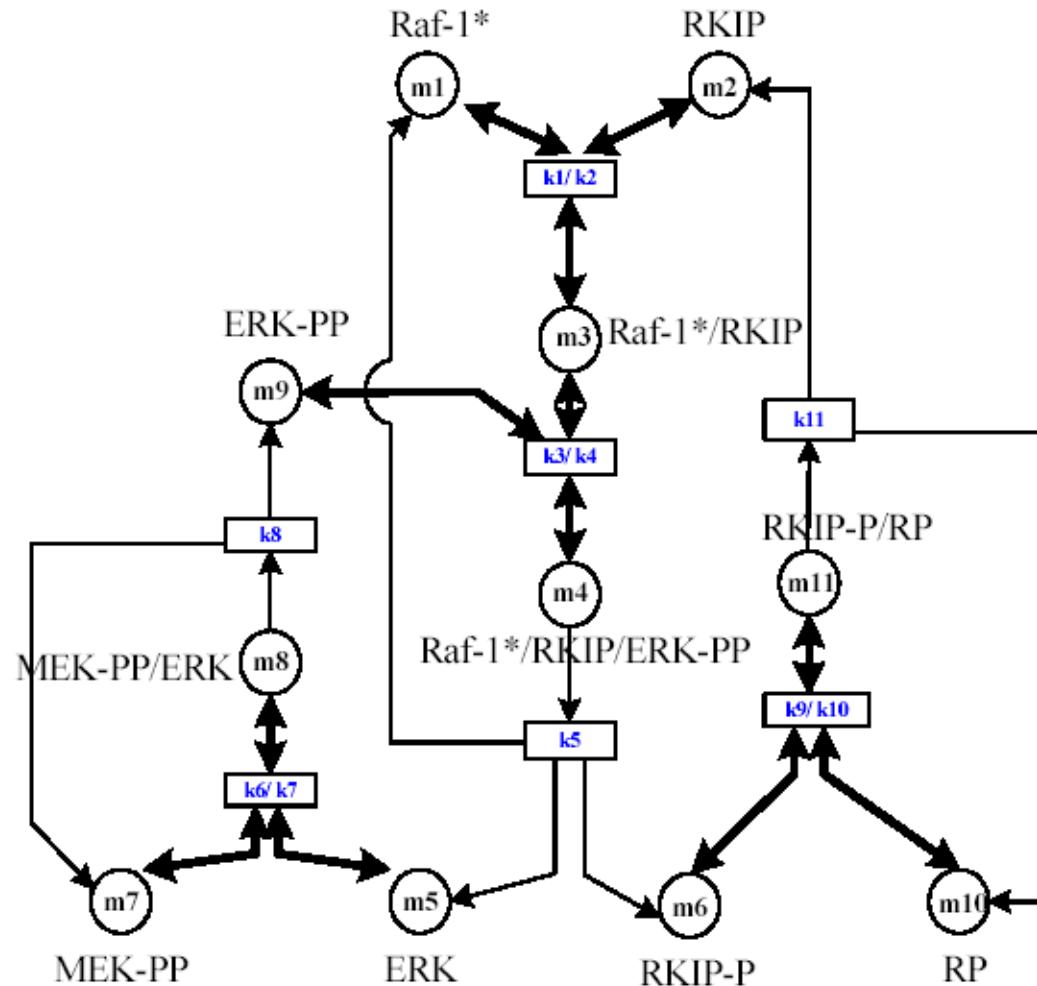


...one pathway...



## Ex7 - RKIP SIGNALLING PATHWAY

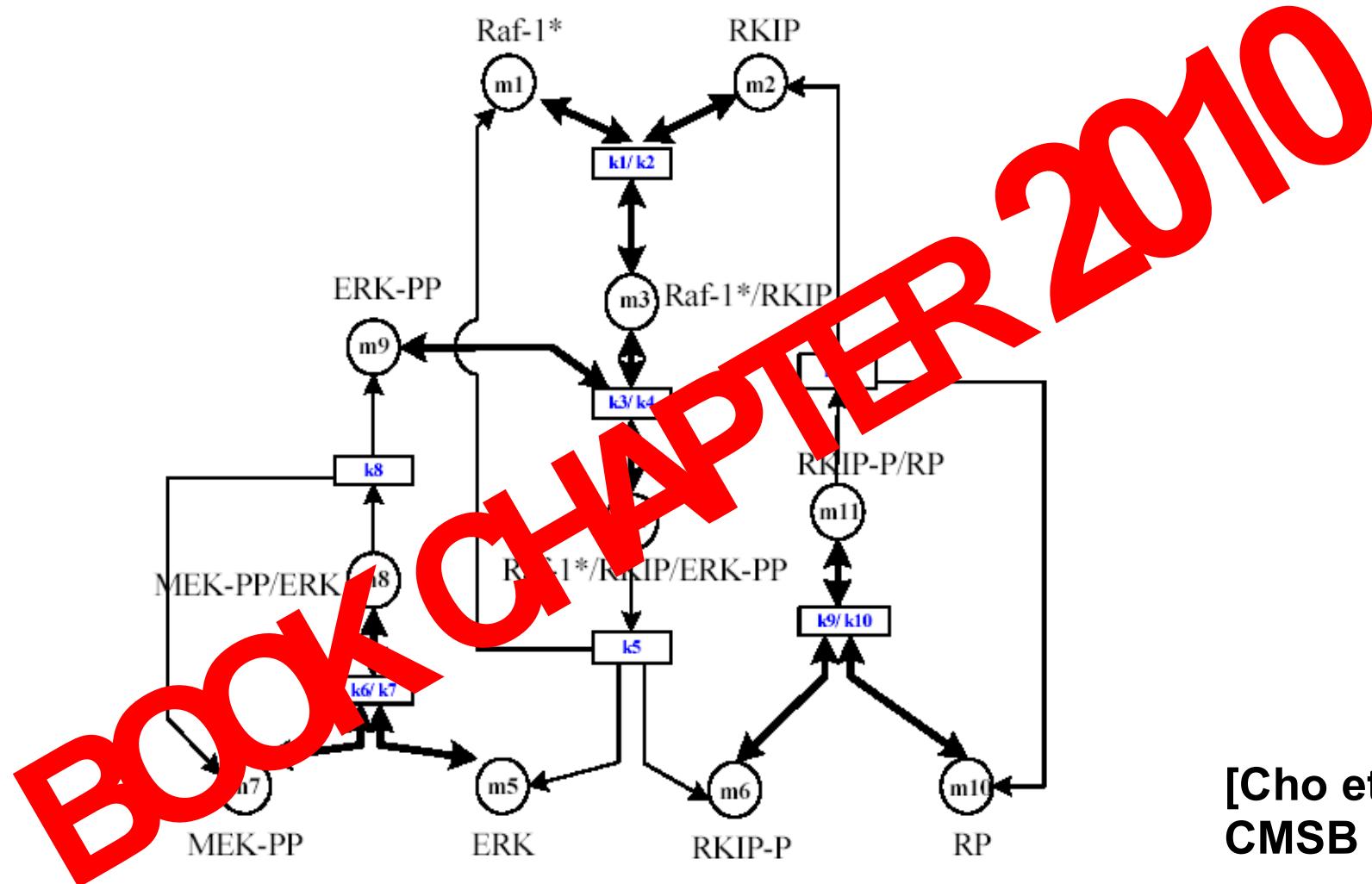
PN & BioModel Engineering



[Cho et al.,  
CMSB 2003]

## Ex7 - RKIP SIGNALLING PATHWAY

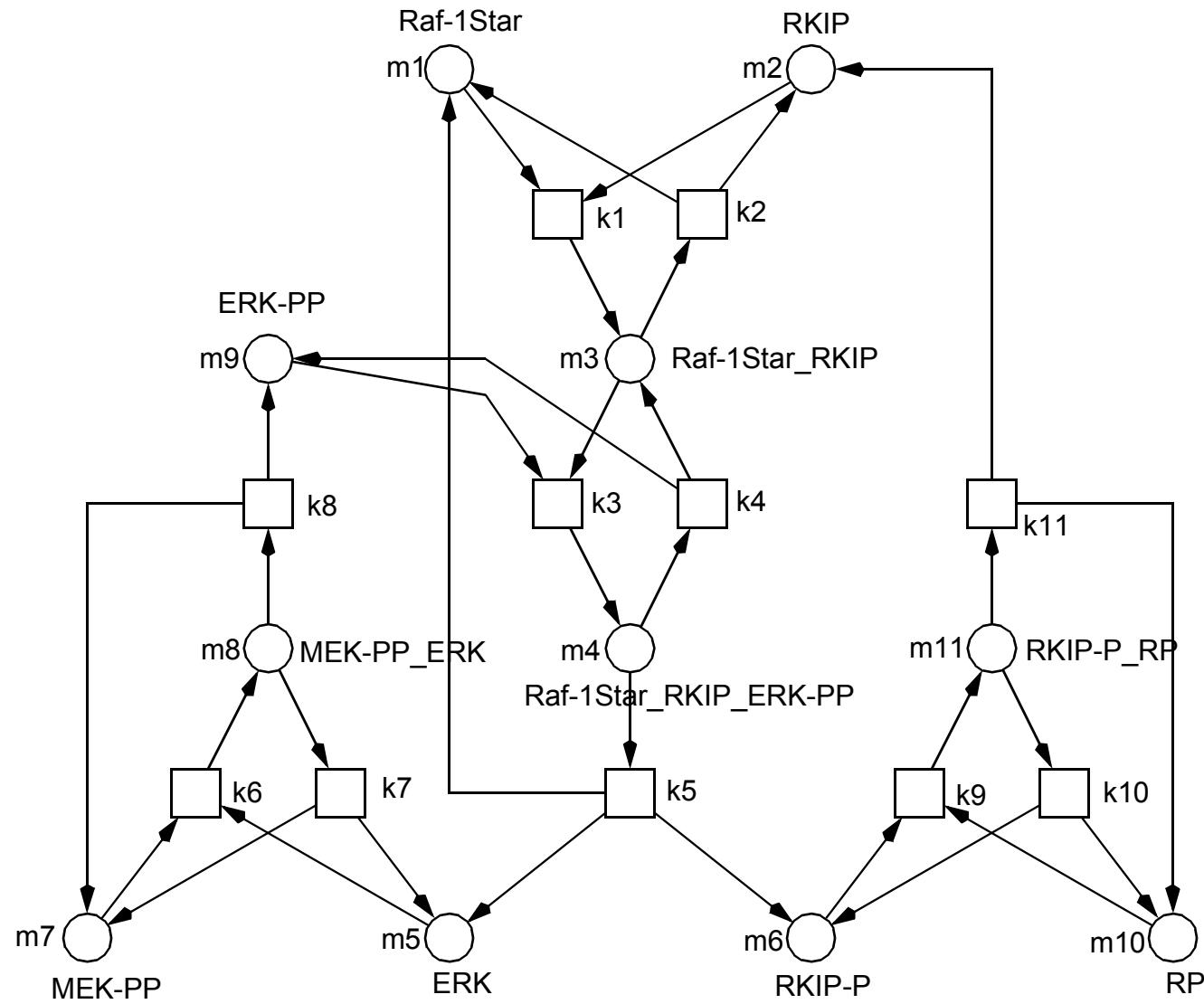
PN & BioModel Engineering



[Cho et al.,  
CMSB 2003]

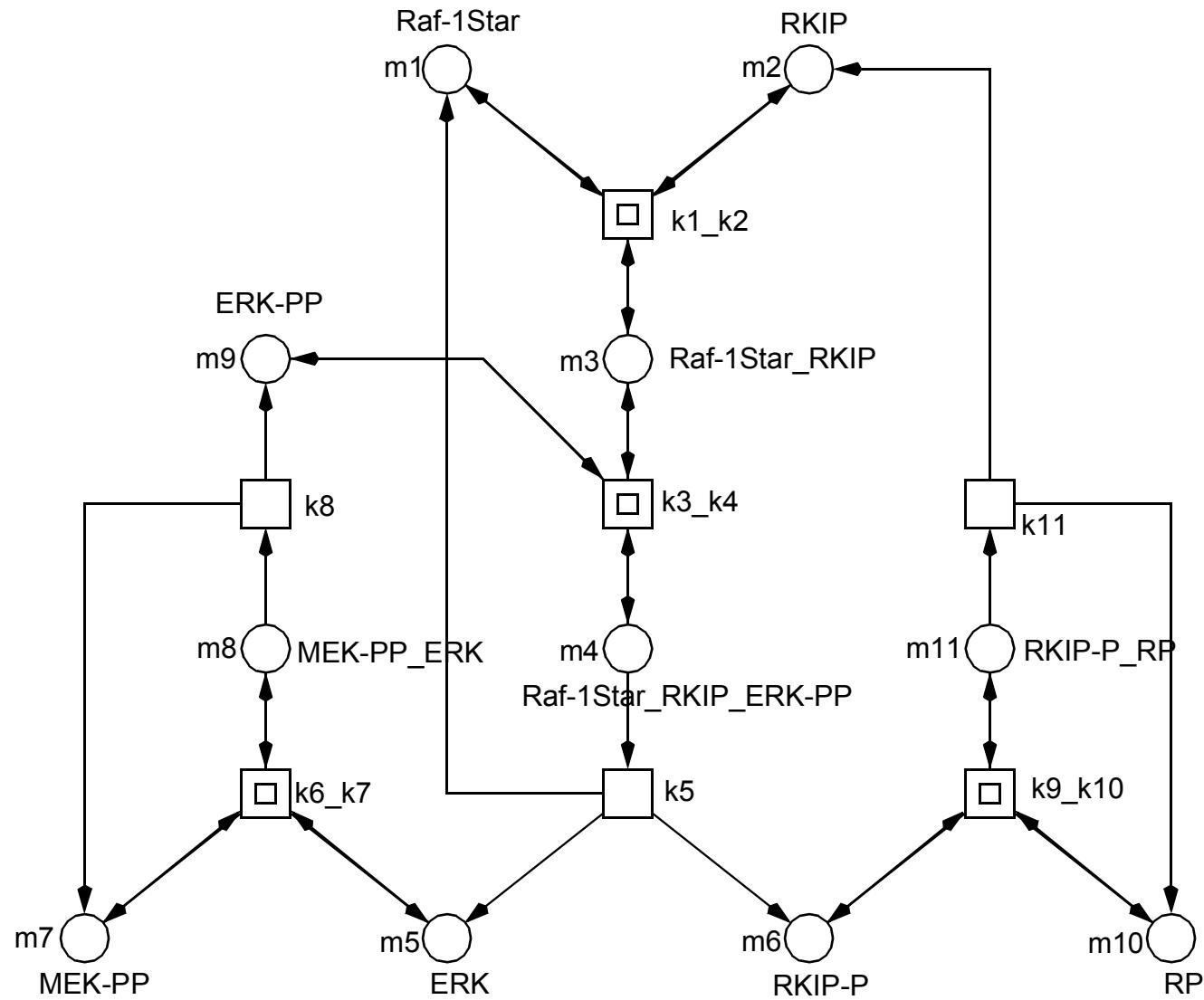
# Ex7 - RKIP SIGNALLING PATHWAY, PETRI NET

PN & BioModel Engineering



# Ex7 - RKIP SIGNALLING PATHWAY, HIERARCHICAL PETRI NET

PN & BioModel Engineering

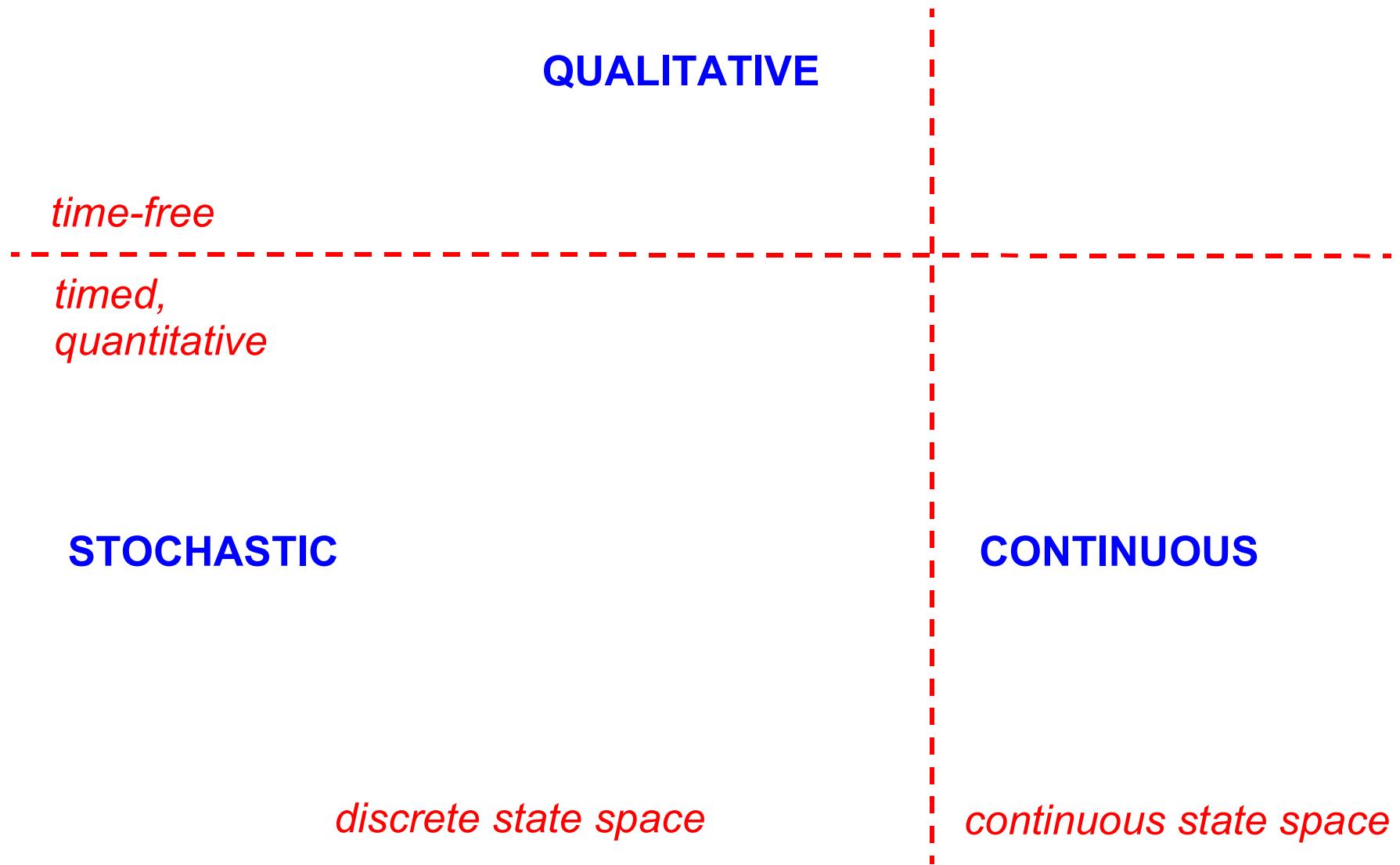


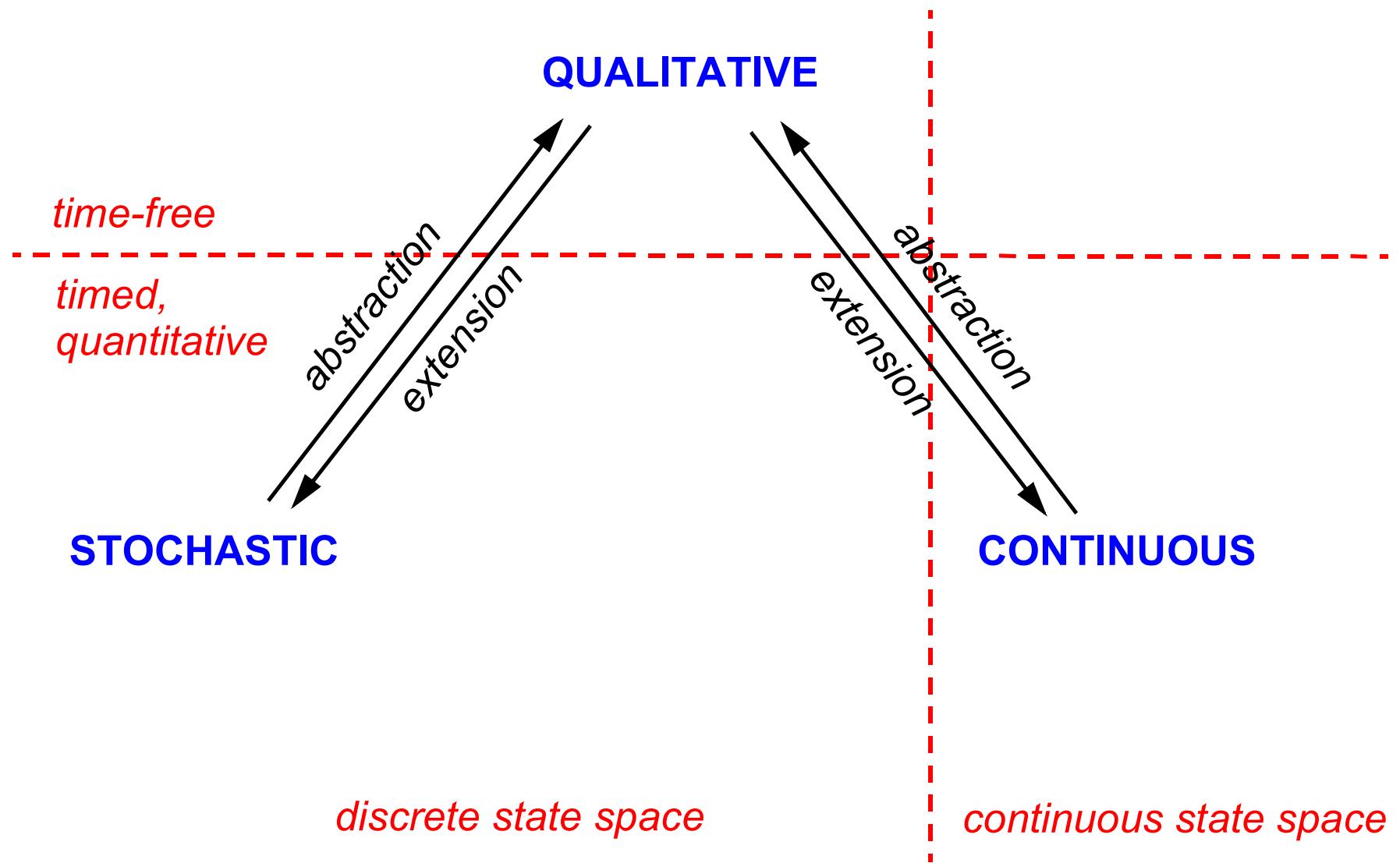
# THE FRAMEWORK

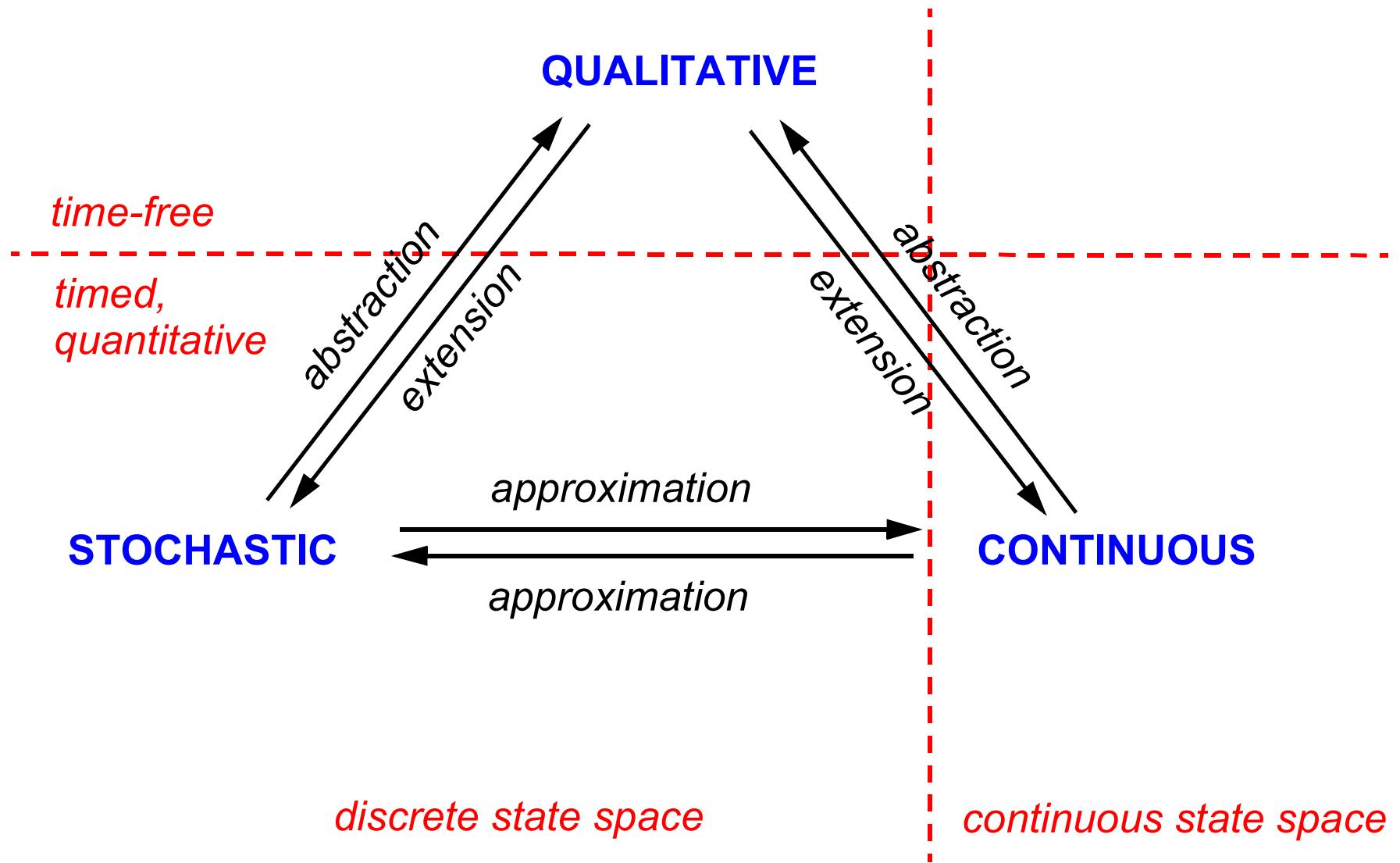
**QUALITATIVE**

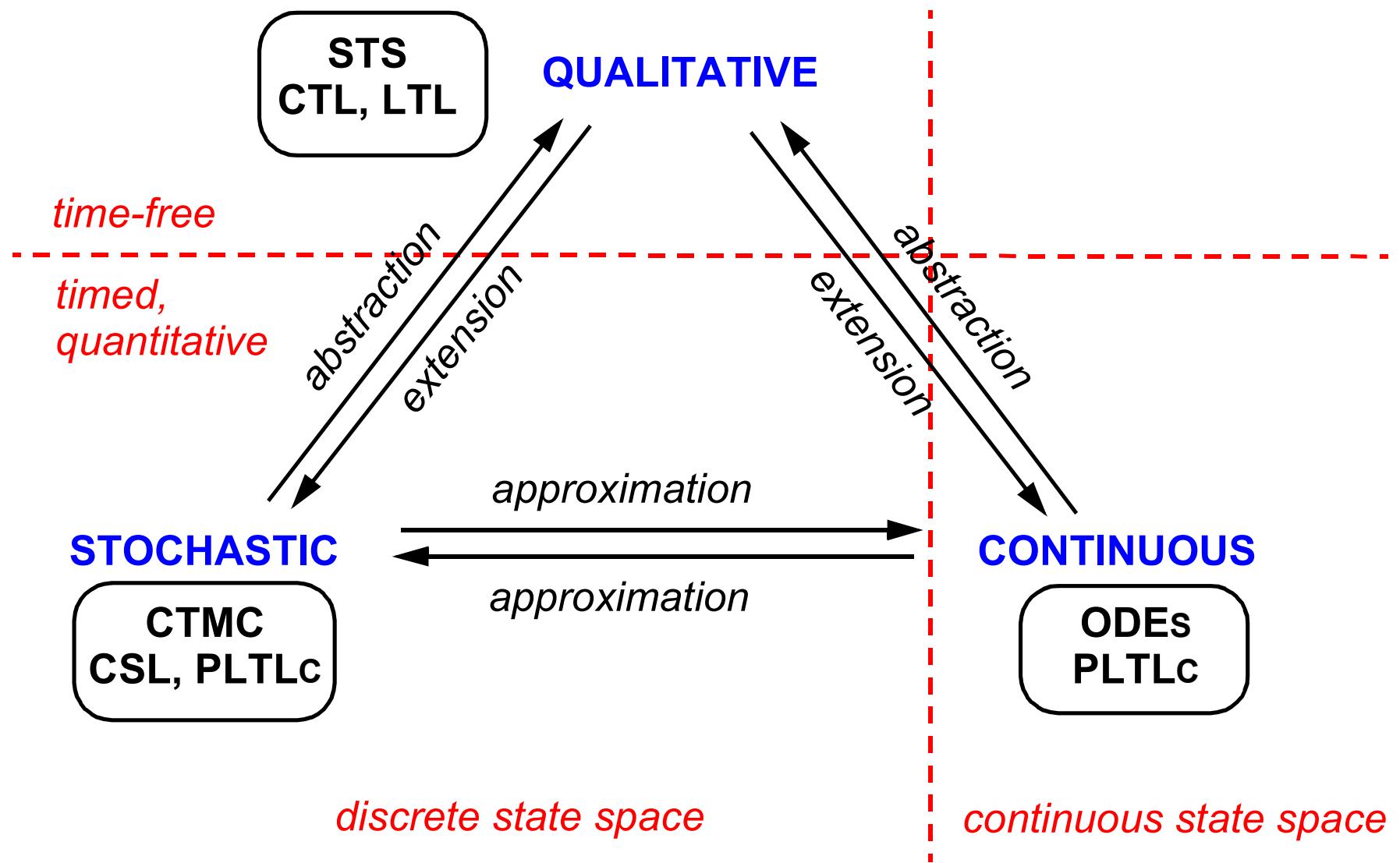
**STOCHASTIC**

**CONTINUOUS**

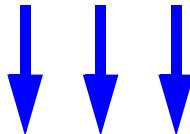








## THREE MODELS SHARING STRUCTURE



**QUANTITATIVE MODEL = QUALITATIVE MODEL**

**+**

**QUANTITATIVE PARAMETERS  
(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*
- **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*

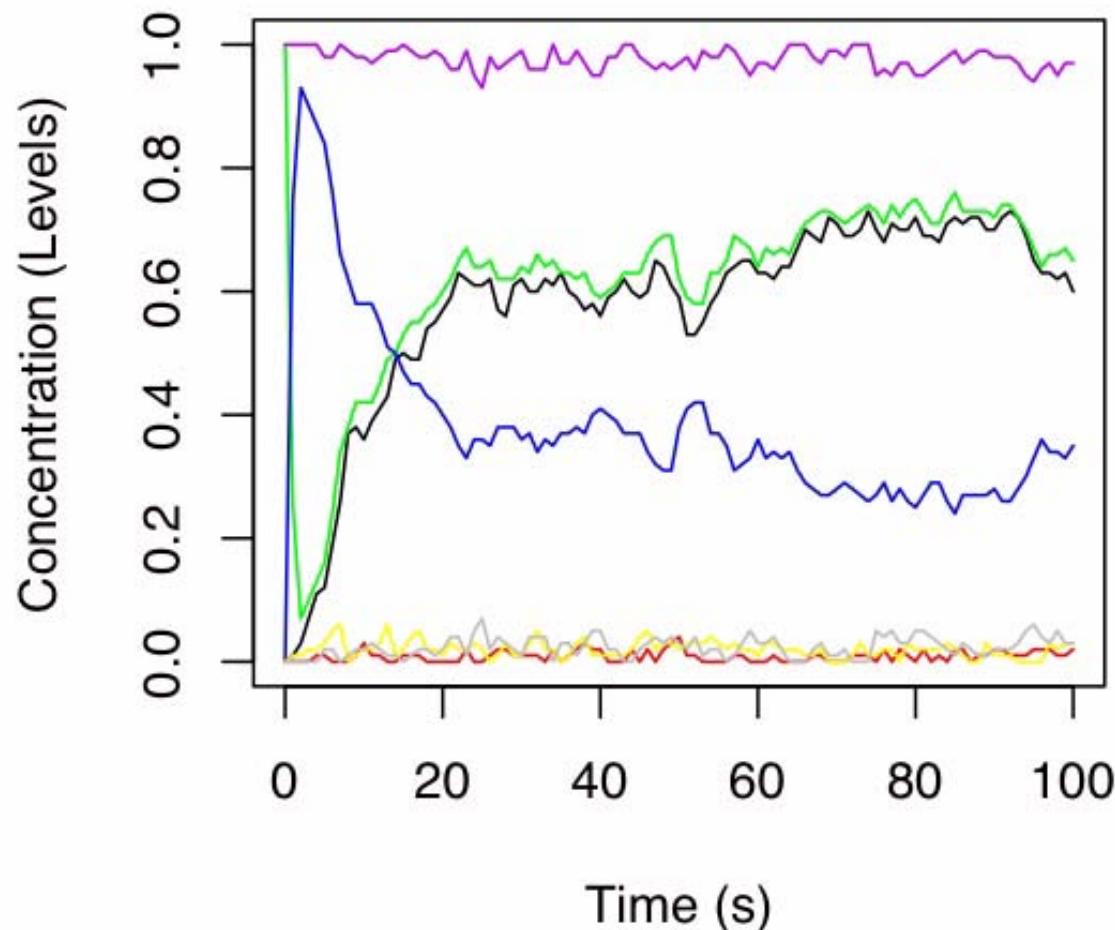
- *molecules semantics*

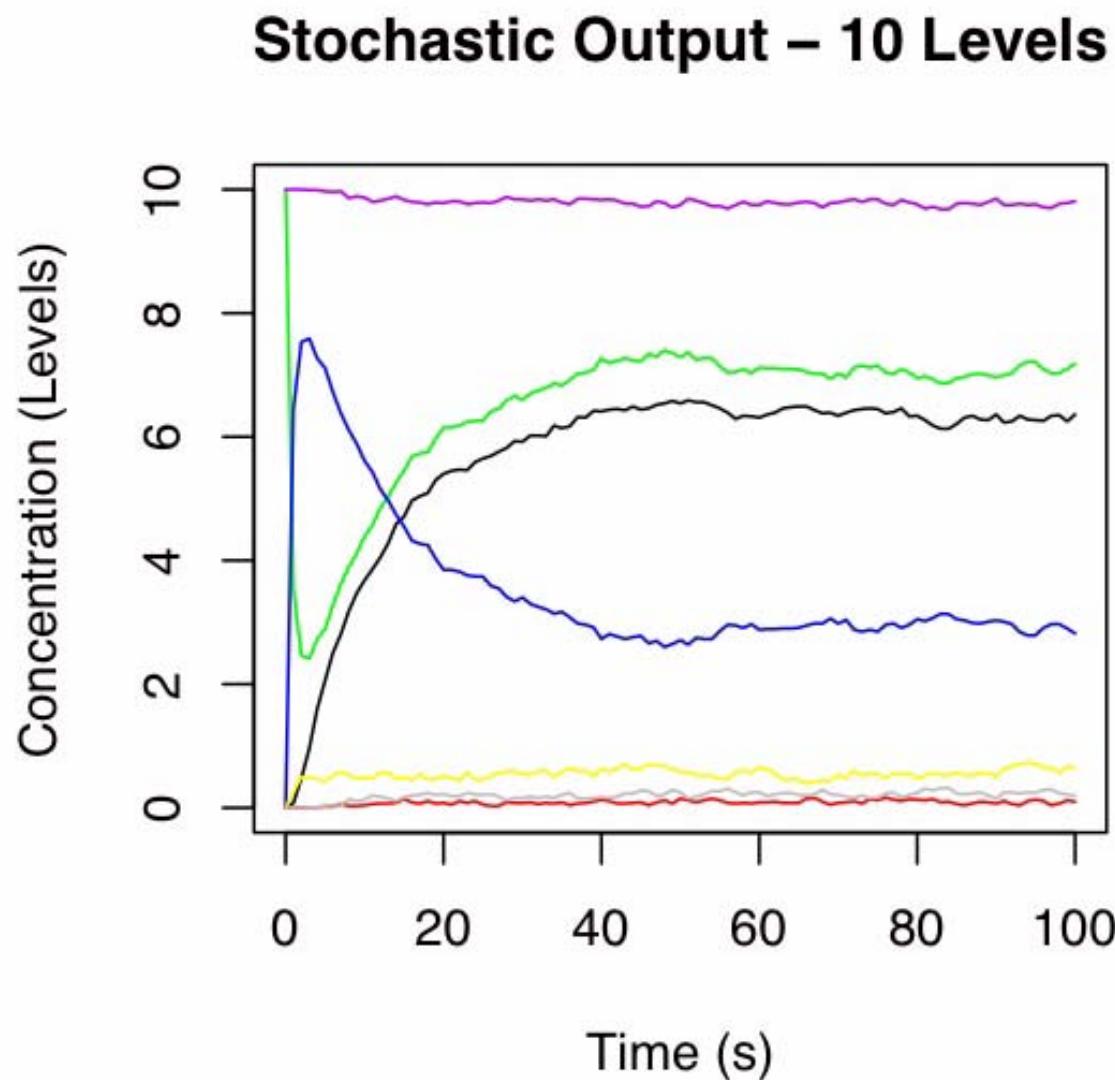
$$h_t := \textcolor{red}{c_t} \cdot \prod_{p \in \bullet t} \binom{m(p)}{f(p, t)}$$

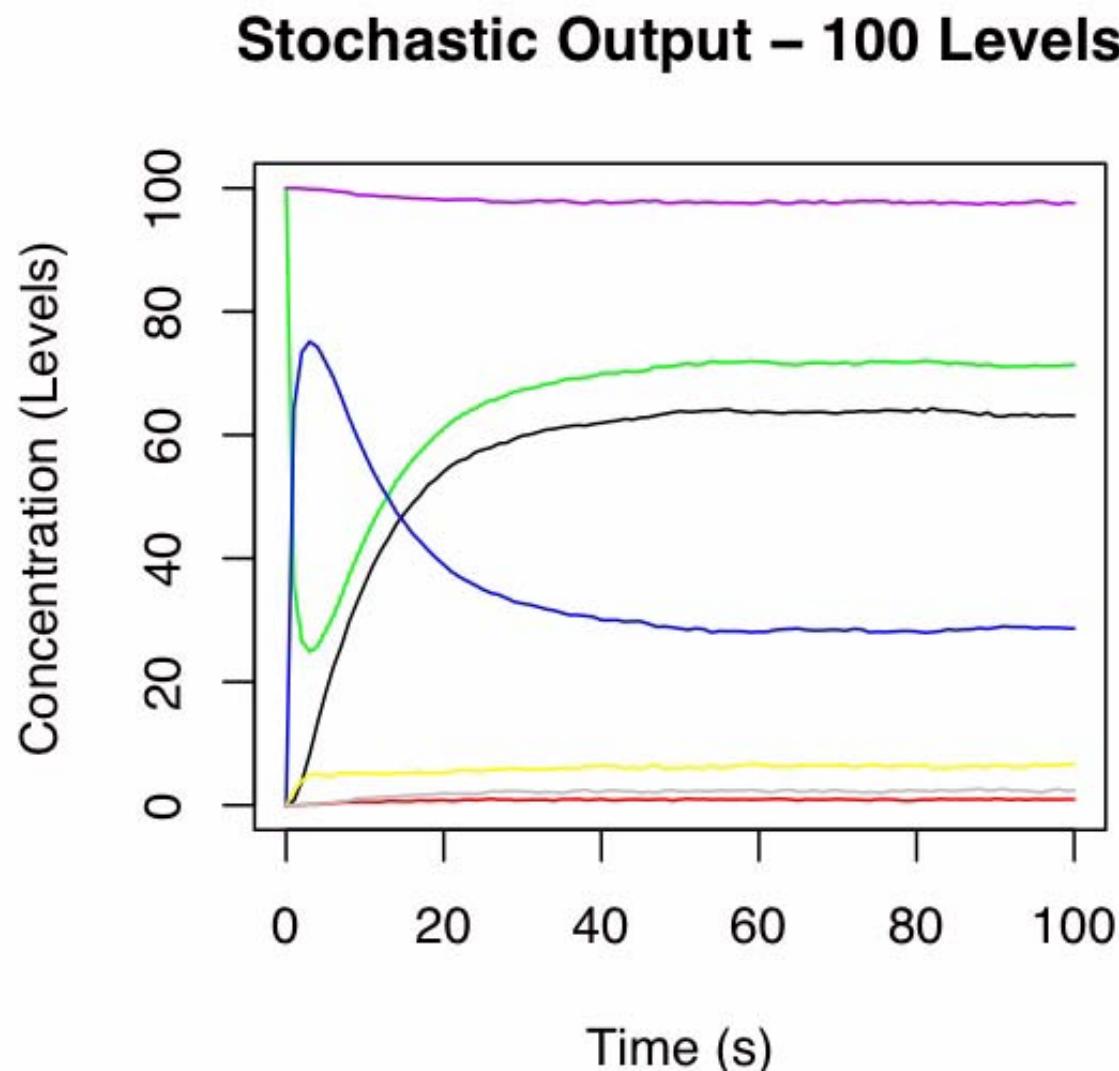
- *concentration levels semantics*

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

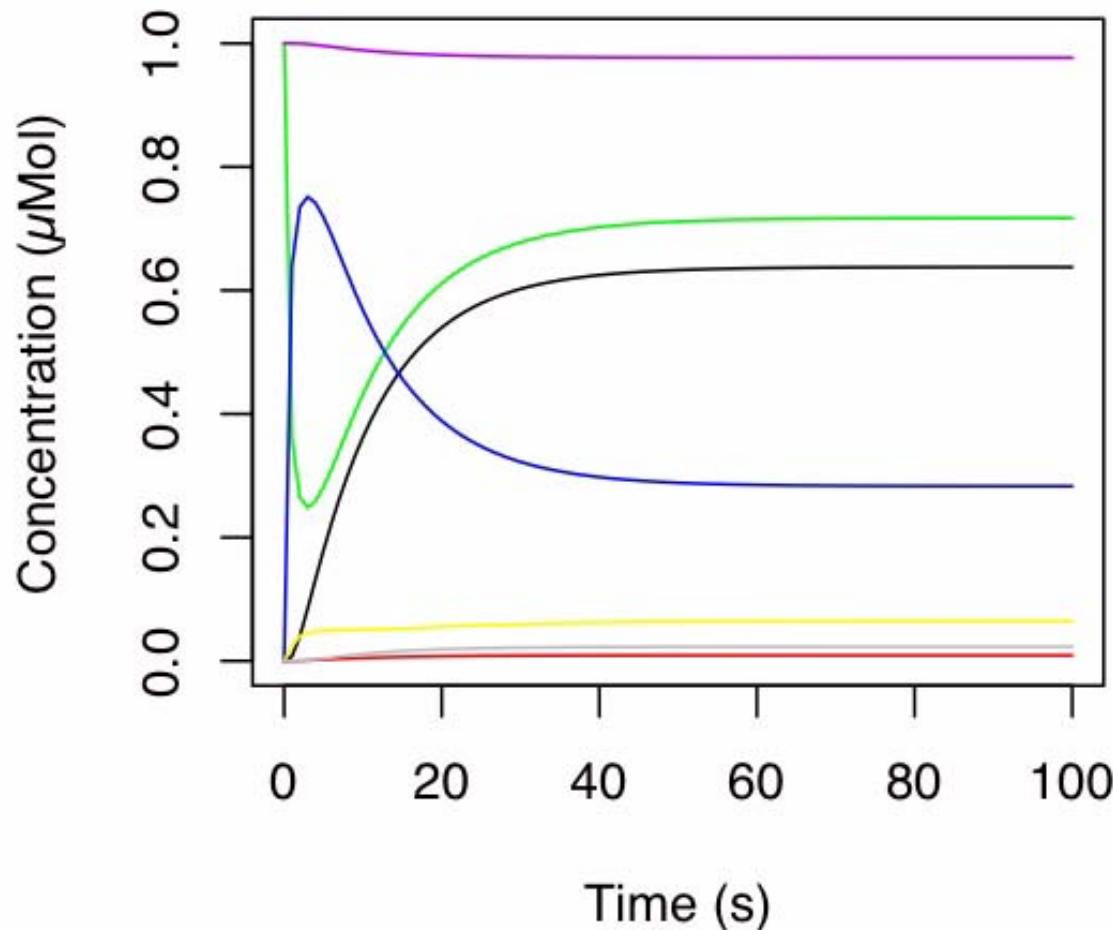
## Stochastic Output – 1 Level







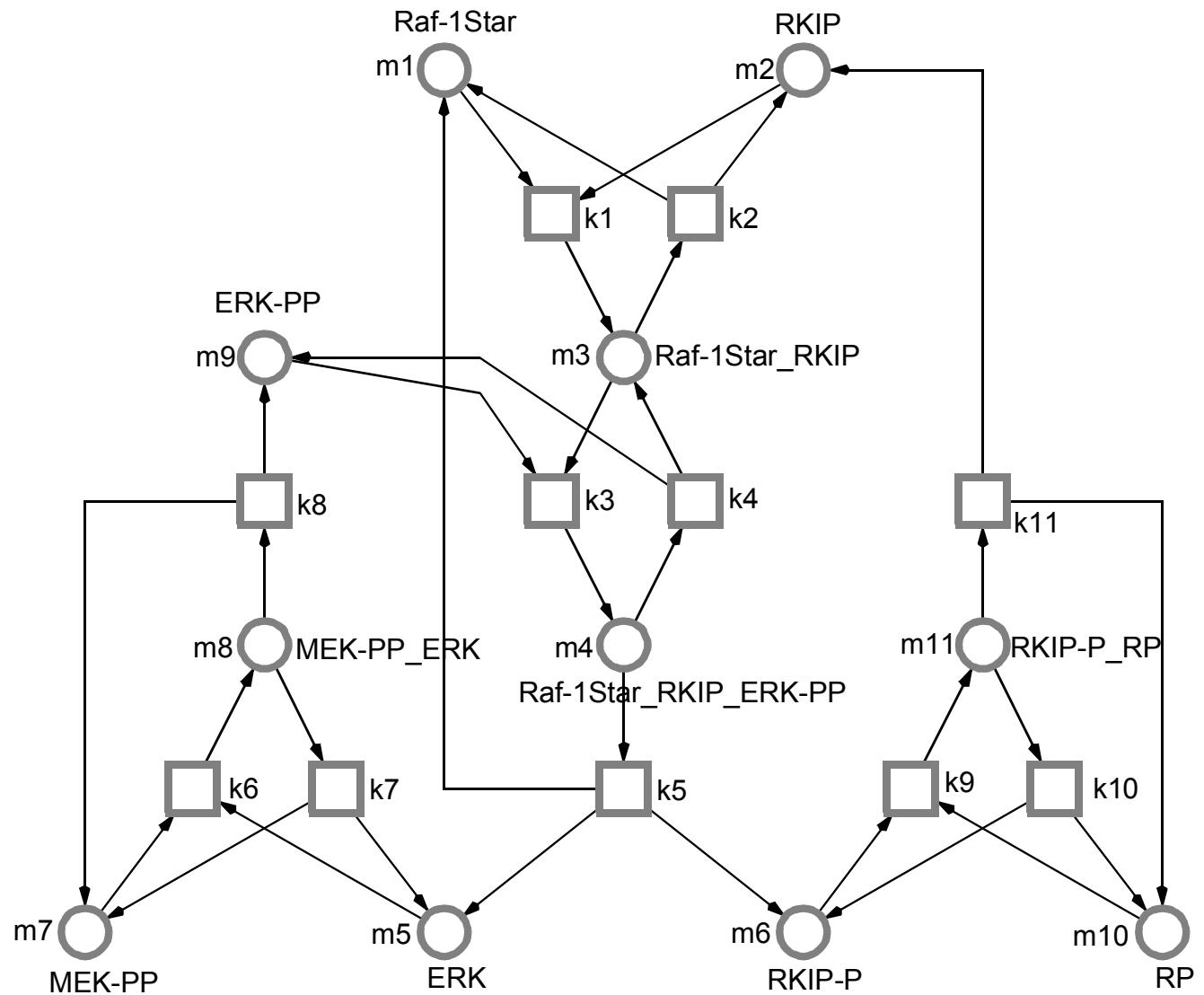
### Deterministic Output



- **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*
- **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

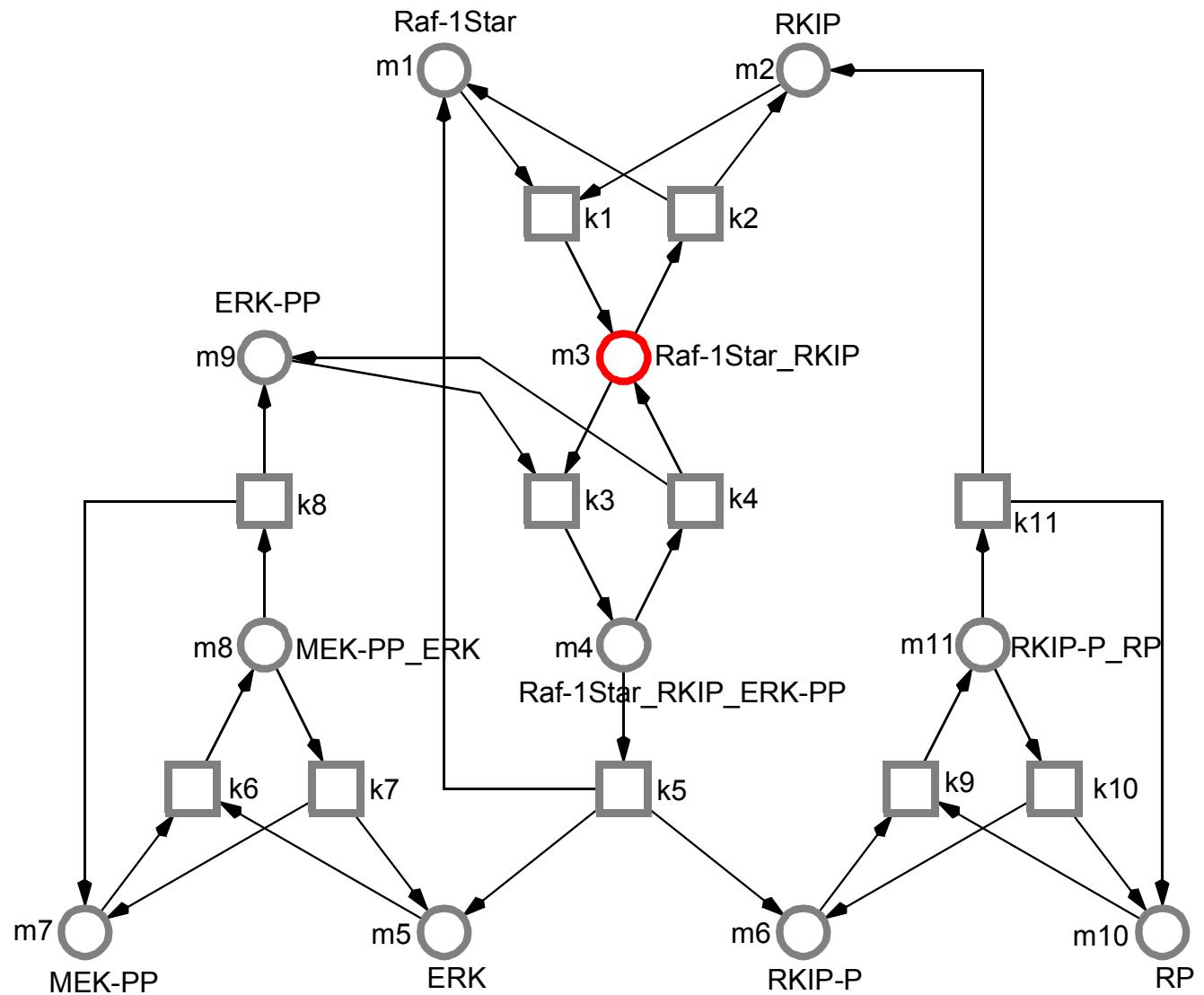
PN & BioModel Engineering



# CONTINUOUS PETRI NET DEFINES ODES

PN & BioModel Engineering

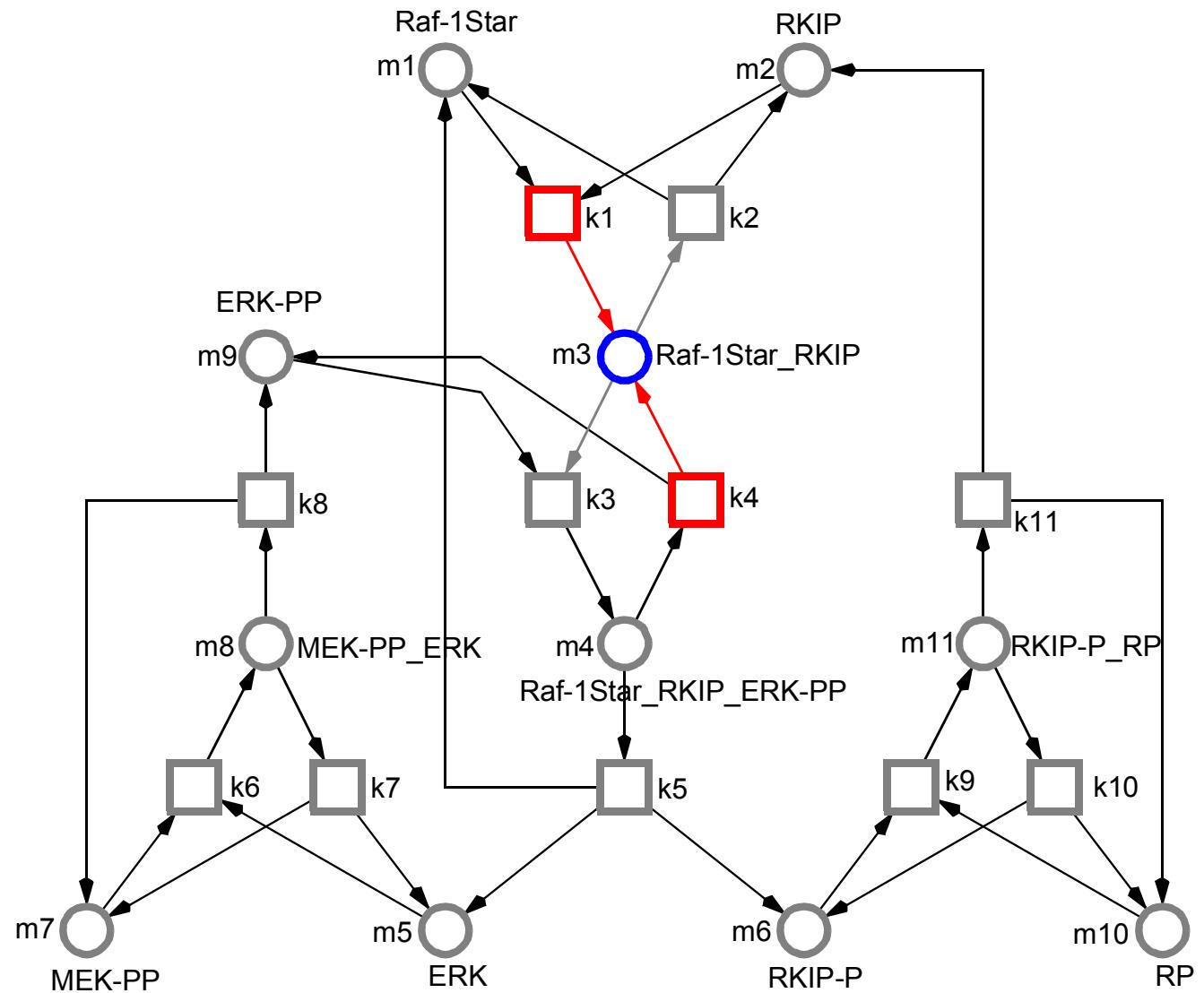
$$\frac{dm_3}{dt} =$$



# CONTINUOUS PETRI NET DEFINES ODES

PN & BioModel Engineering

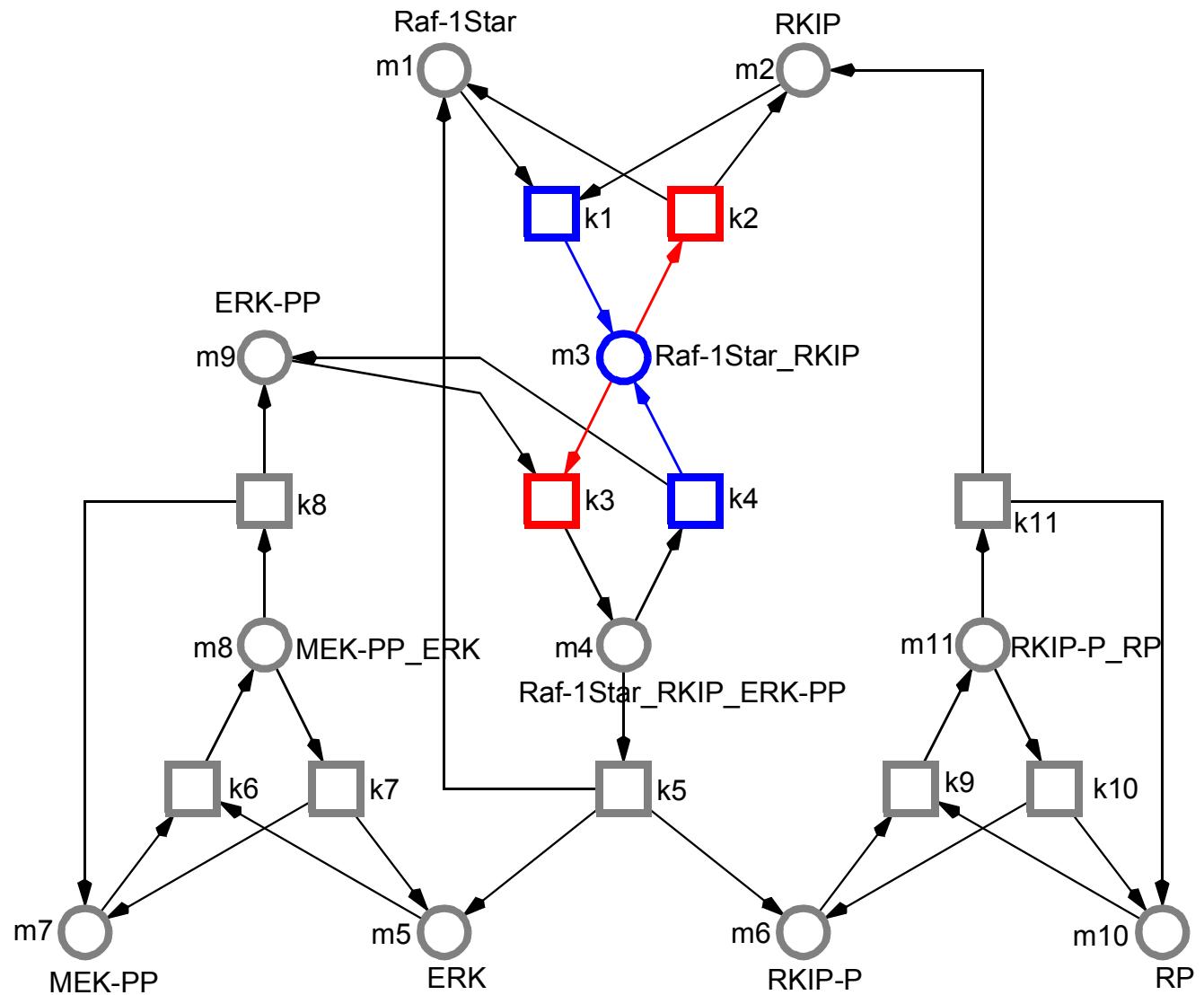
$$\frac{dm_3}{dt} = + r_1 \\ + r_4$$



# CONTINUOUS PETRI NET DEFINES ODES

PN & BioModel Engineering

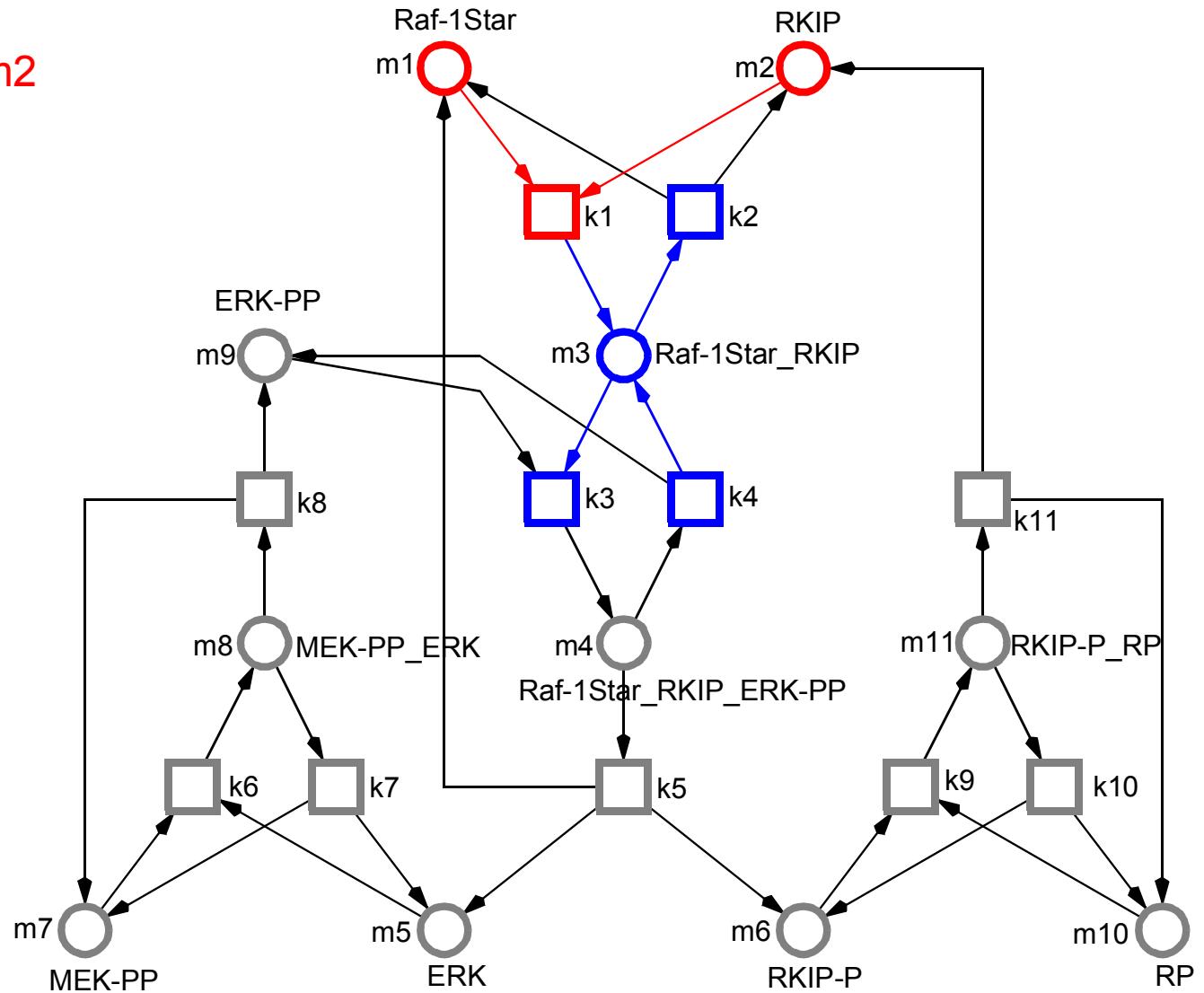
$$\frac{dm_3}{dt} = + r_1 \\ + r_4 \\ - r_2 \\ - r_3$$



# CONTINUOUS PETRI NET DEFINES ODES

PN & BioModel Engineering

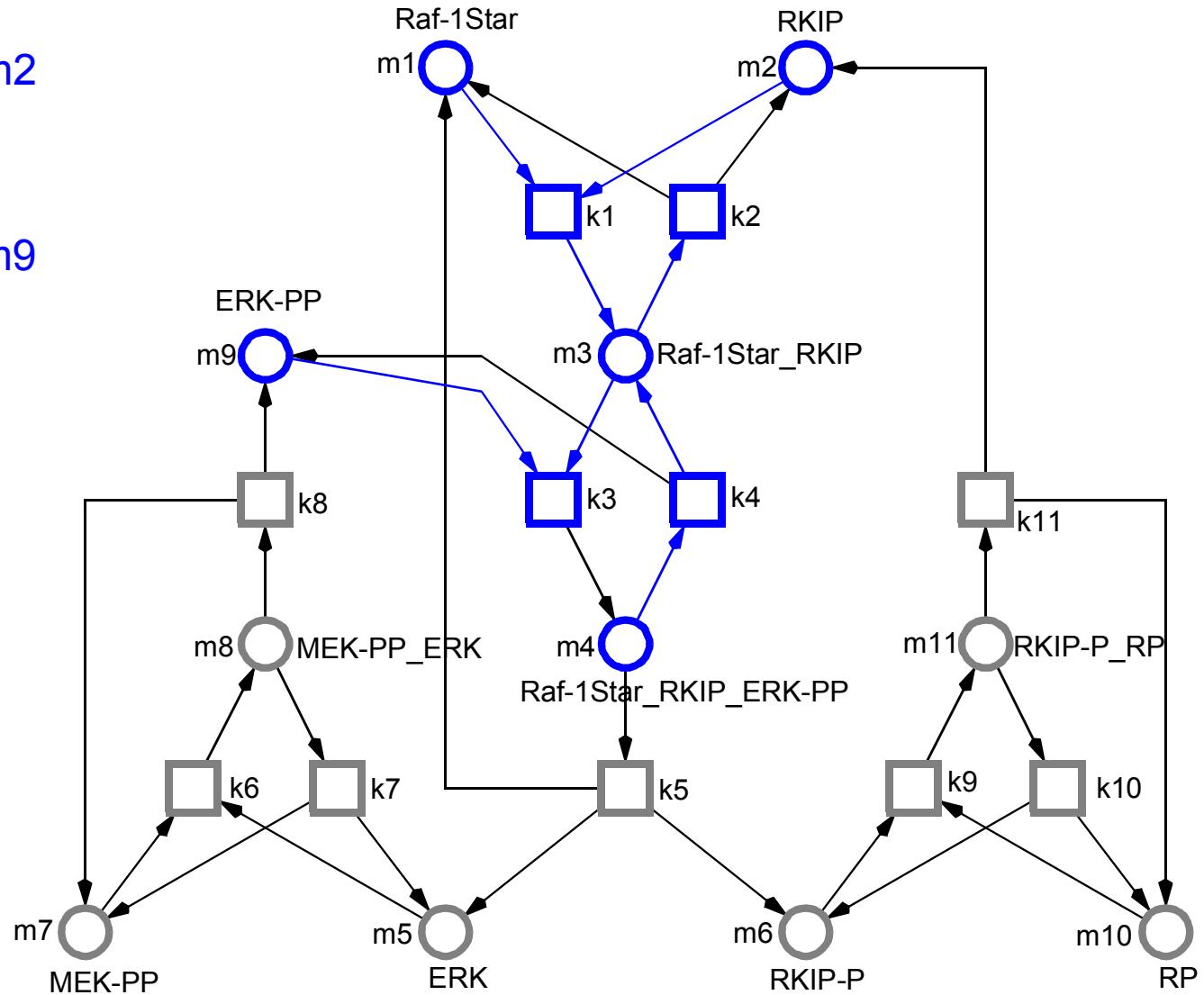
$$\frac{dm_3}{dt} = + k_1 * m_1 * m_2 \\ + r_4 \\ - r_2 \\ - r_3$$



# CONTINUOUS PETRI NET DEFINES ODES

PN & BioModel Engineering

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

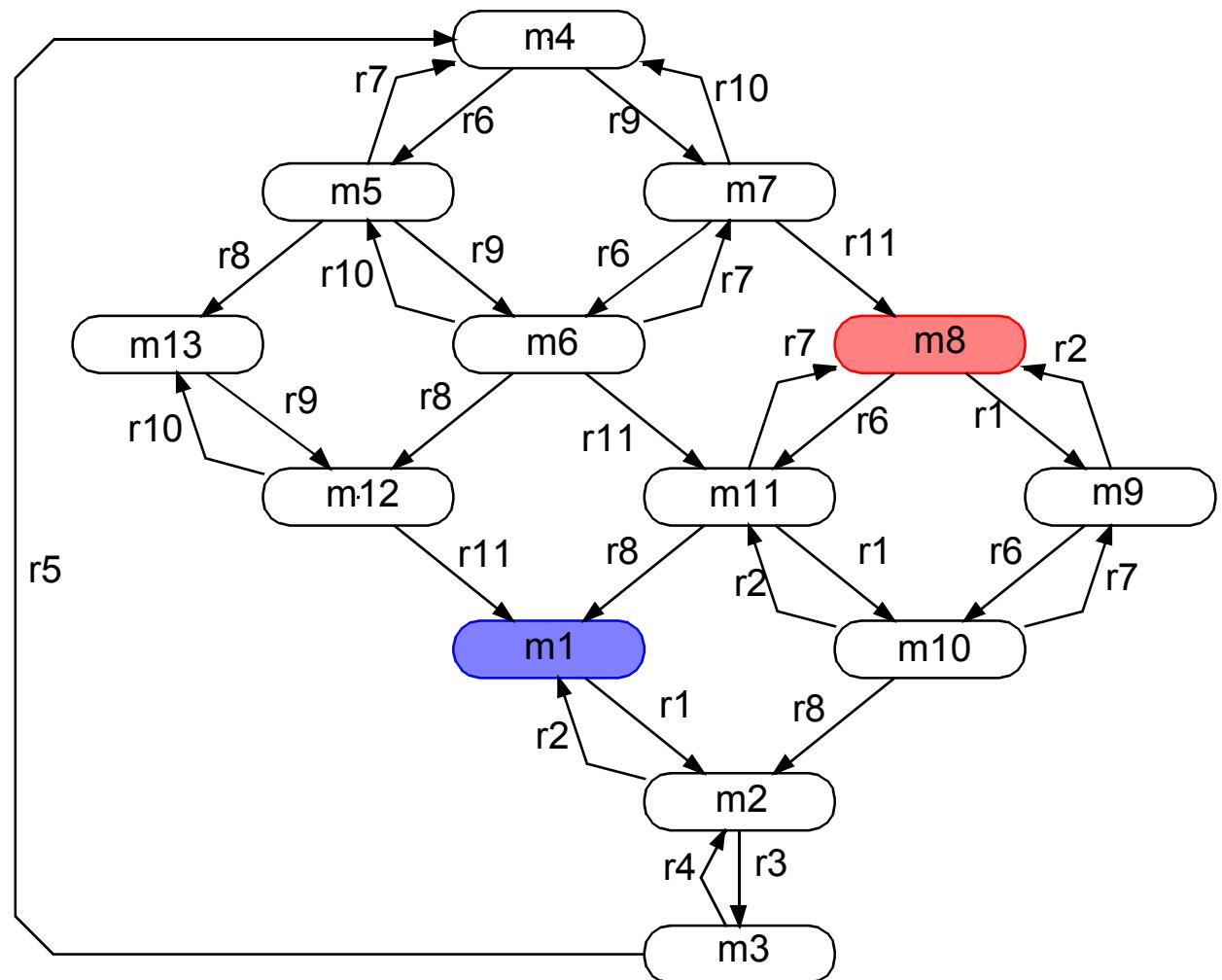


# **ABOUT THE RELATION SPN - CPN**

# Ex1 - RKİP, REACHABILITY GRAPH (STS)

PN & BioModel Engineering

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



# Ex1 - RKIP, QUANTITATIVE ANALYSIS

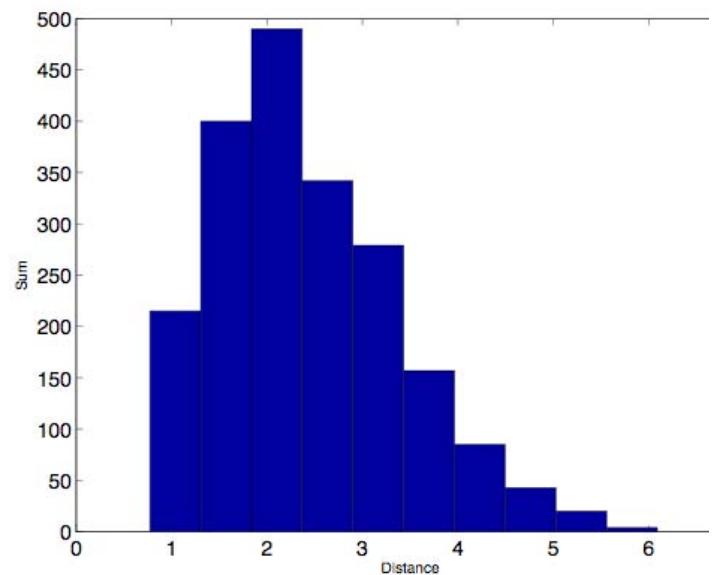
PN & BioModel Engineering

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

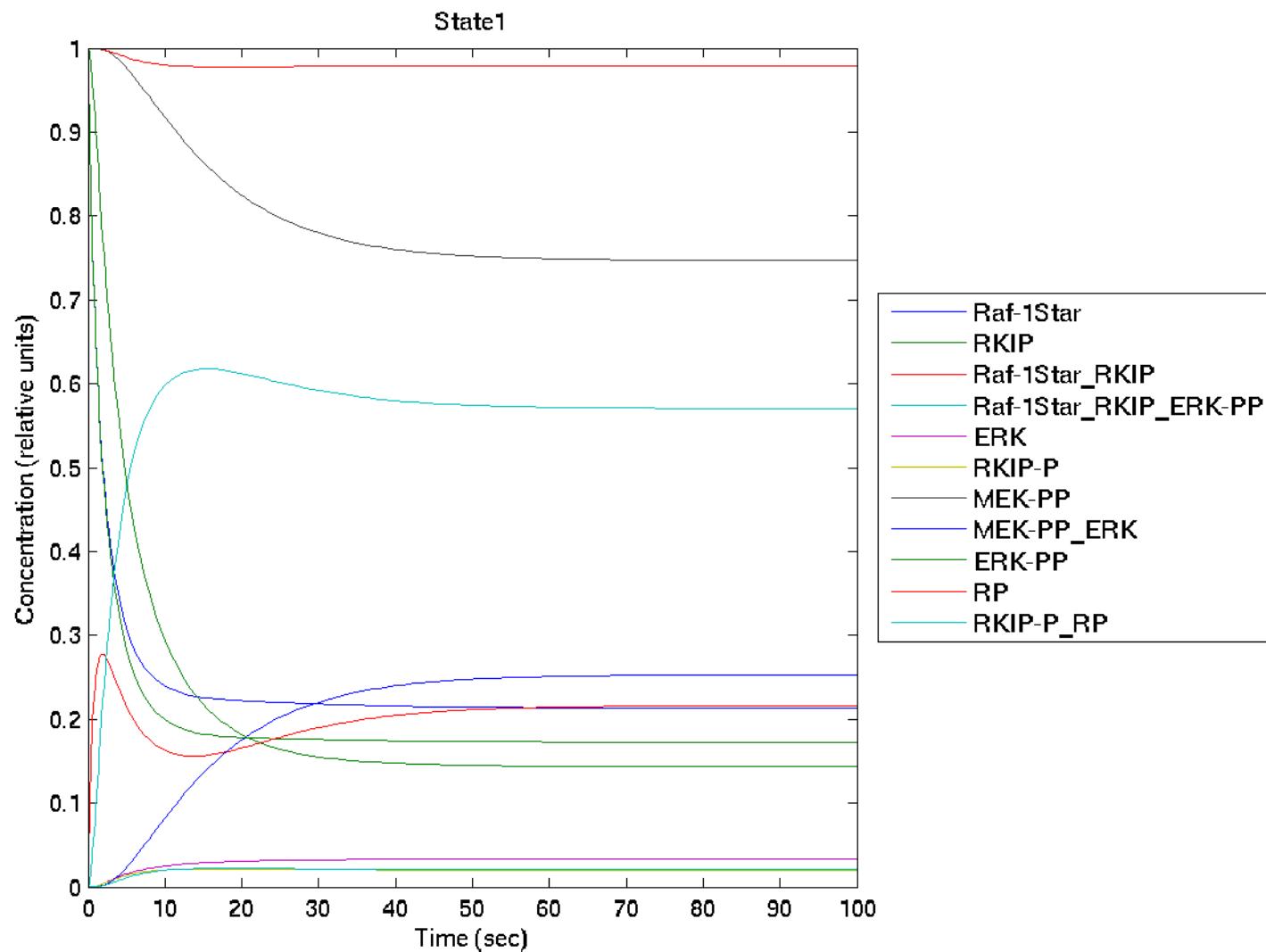


Distribution of ‘bad’ steady states as Euclidean distances from the ‘good’ final steady state

the “bad” ones

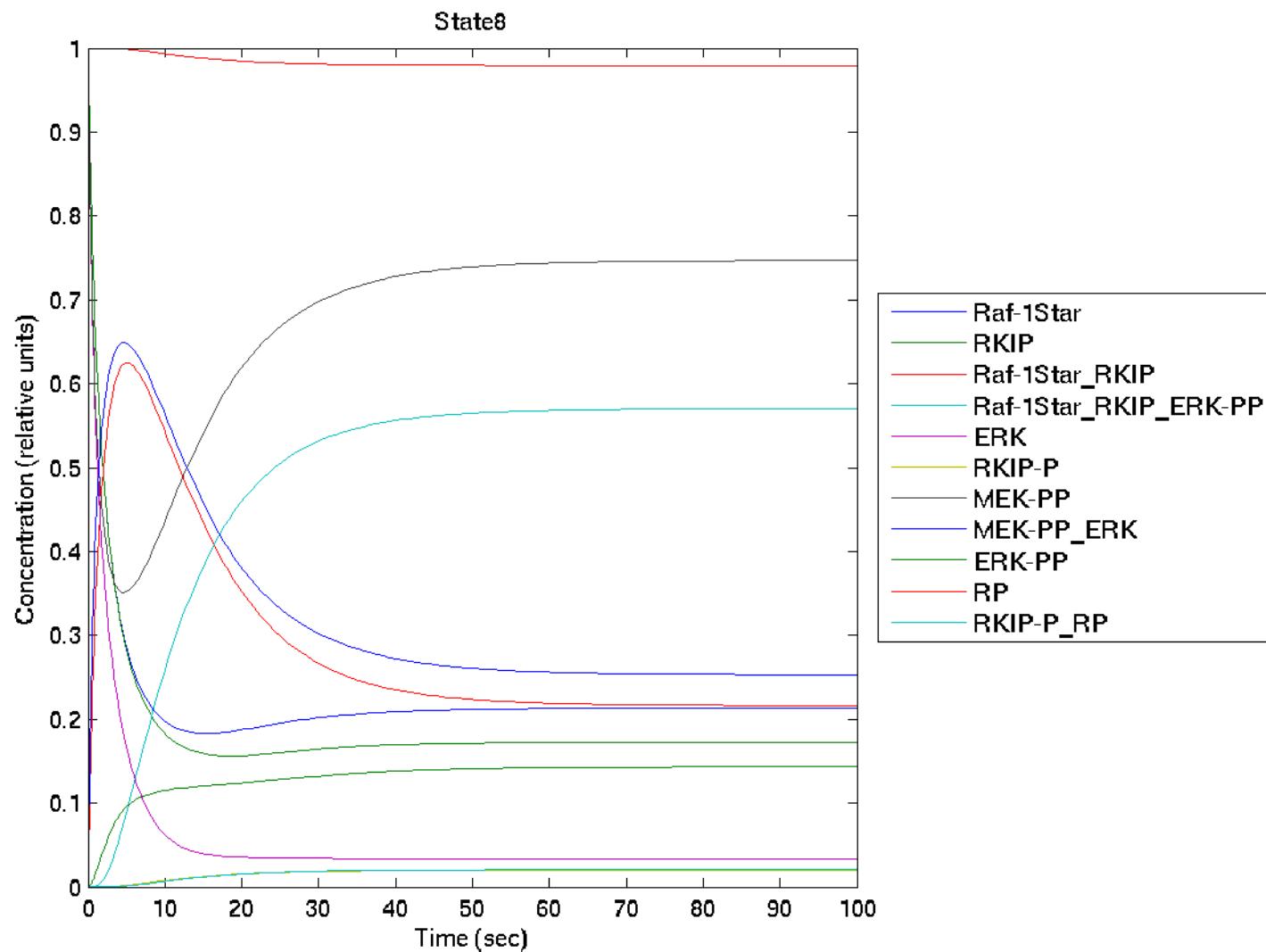
# Ex1 - RKIP, QUANTITATIVE ANALYSIS

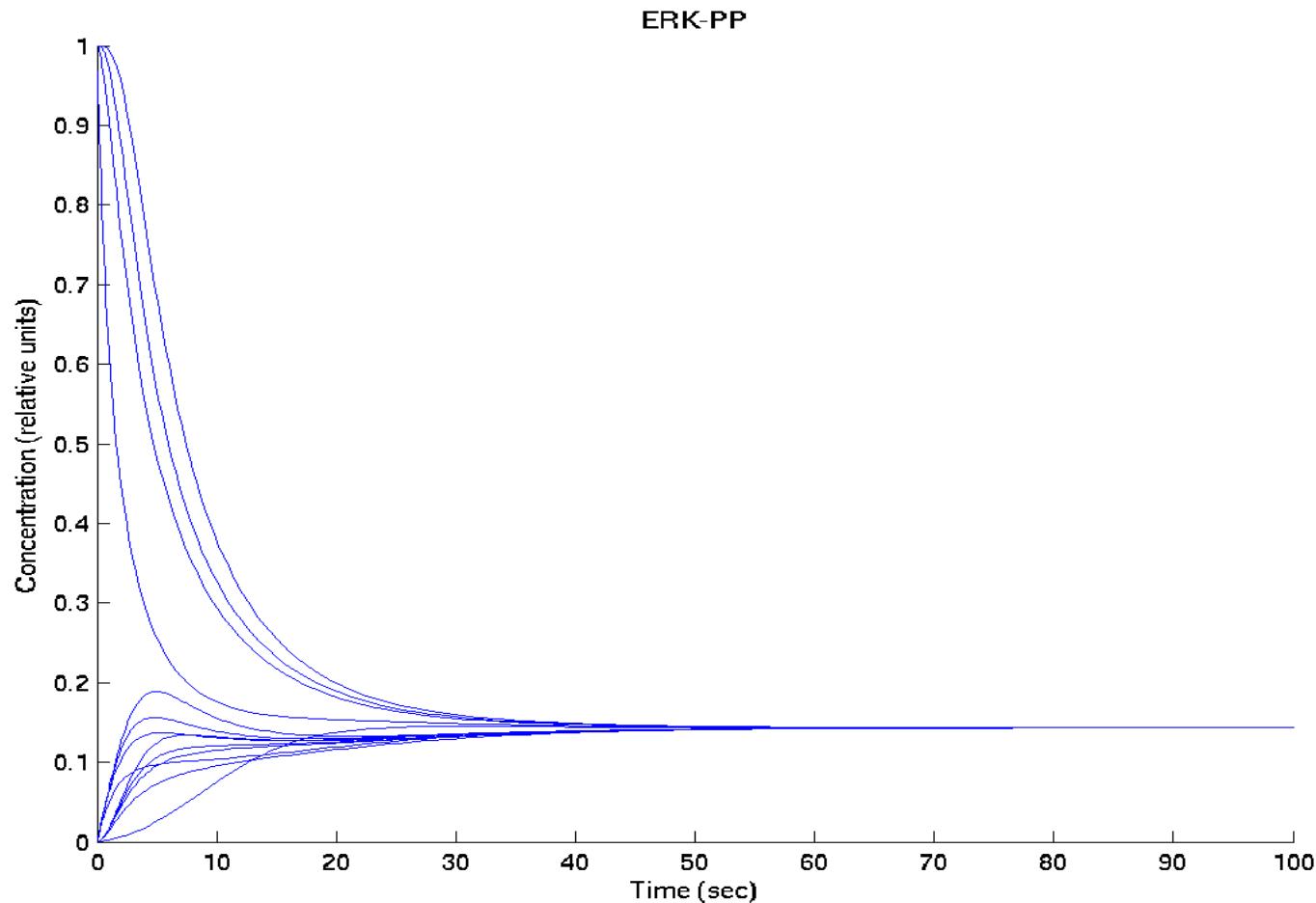
PN & BioModel Engineering



# Ex1 - RKIP, QUANTITATIVE ANALYSIS

PN & BioModel Engineering

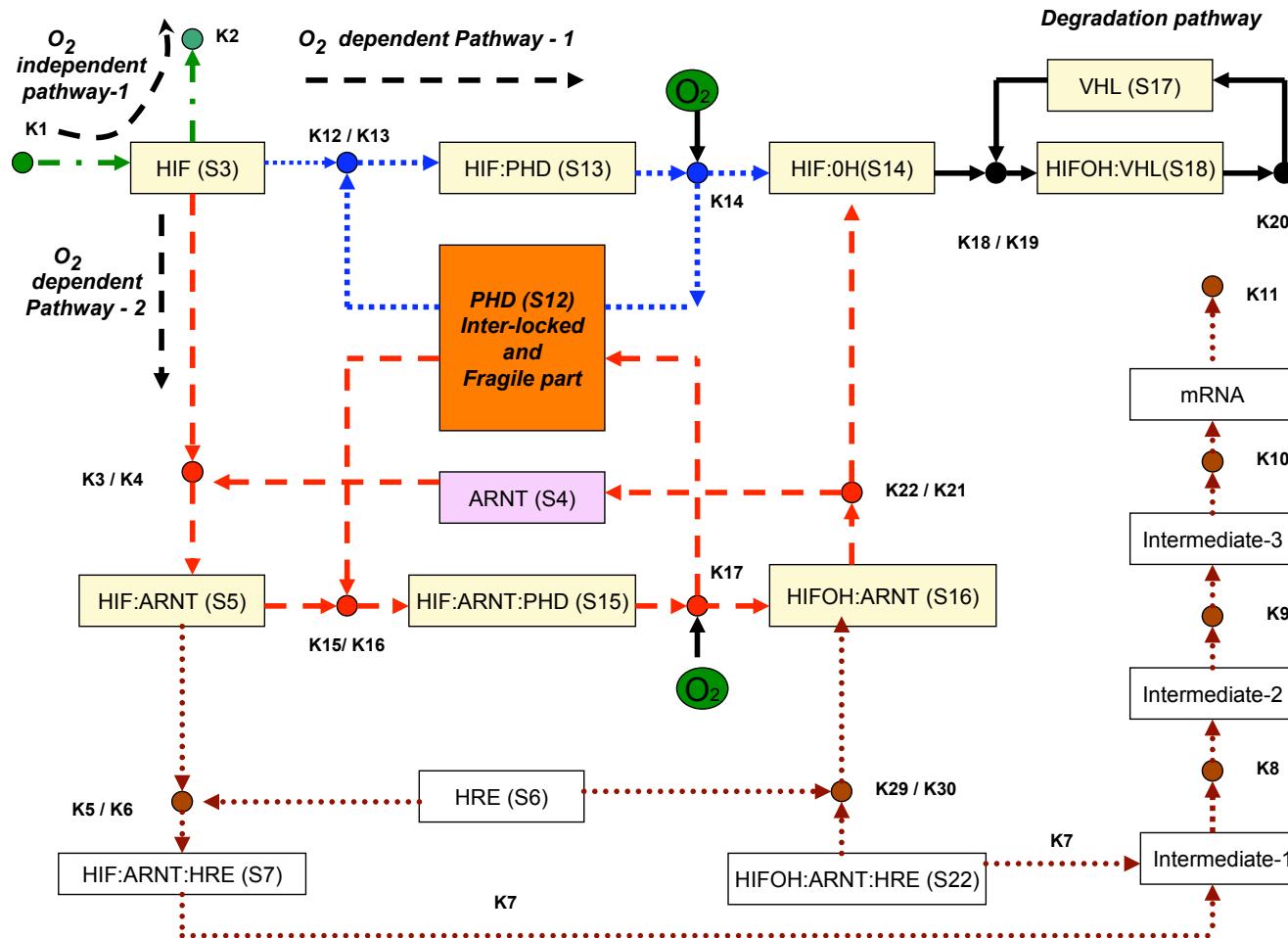




## Ex2 - HYPOXIA

PN & BioModel Engineering

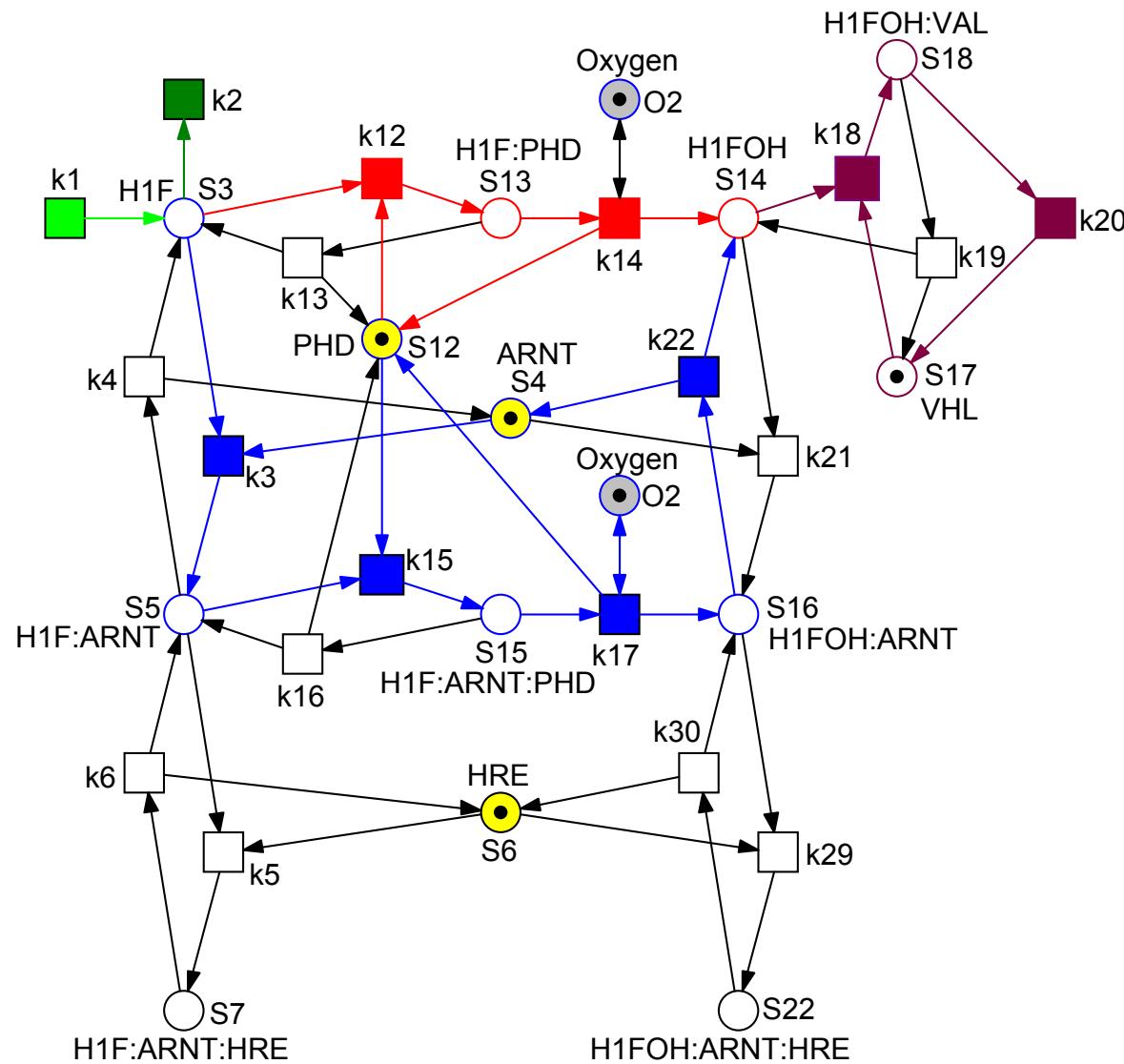
[YU ET AL. 2007]



## Ex2 - HYPOXIA

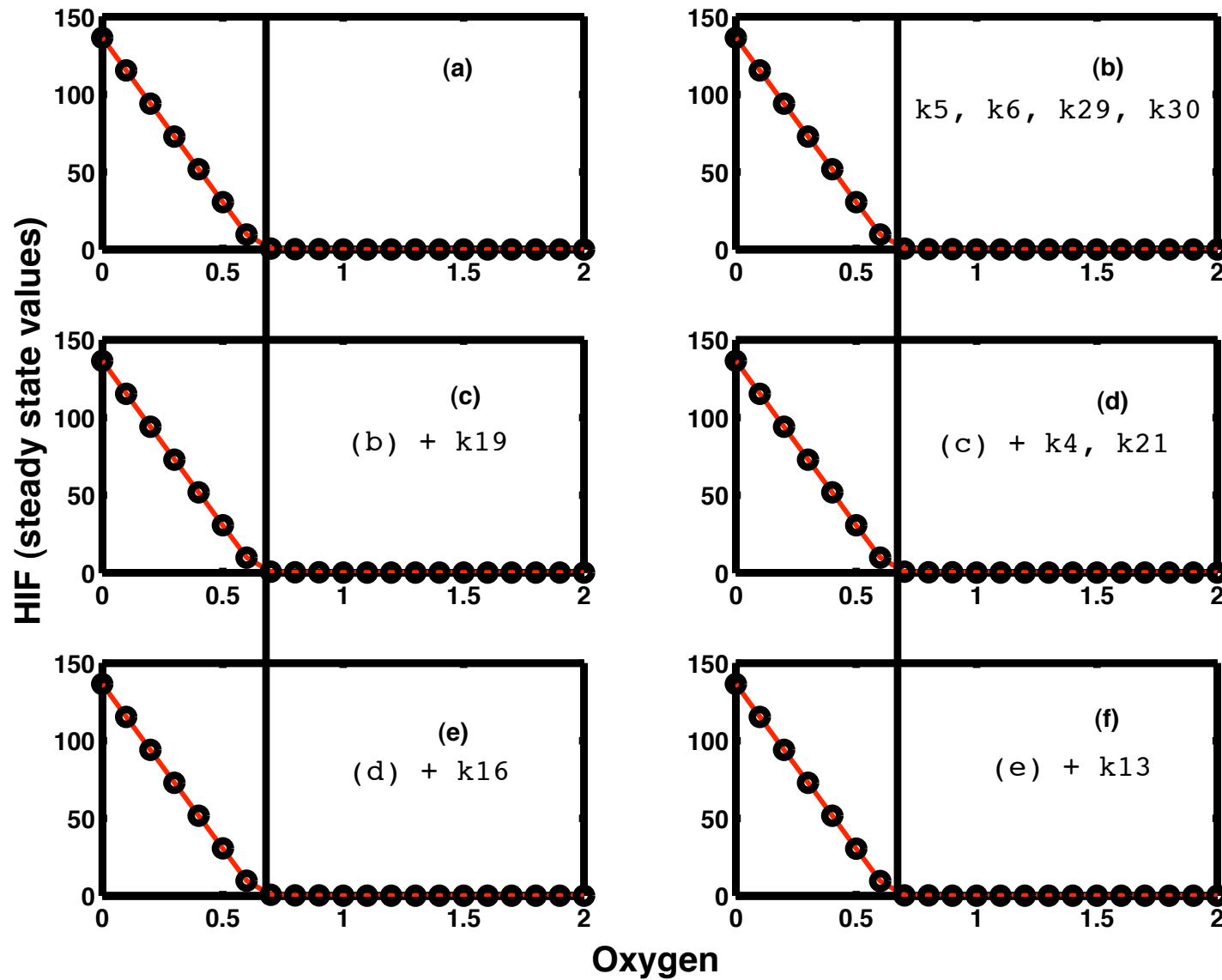
PN & BioModel Engineering

[HEINER; SRIRAM 2010]

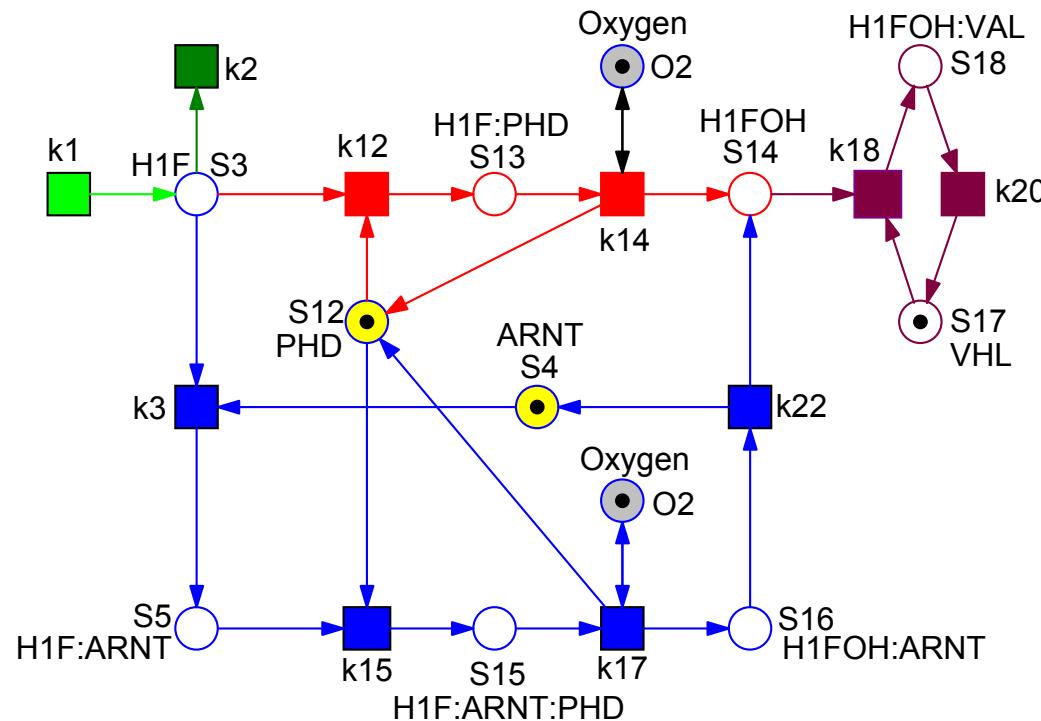


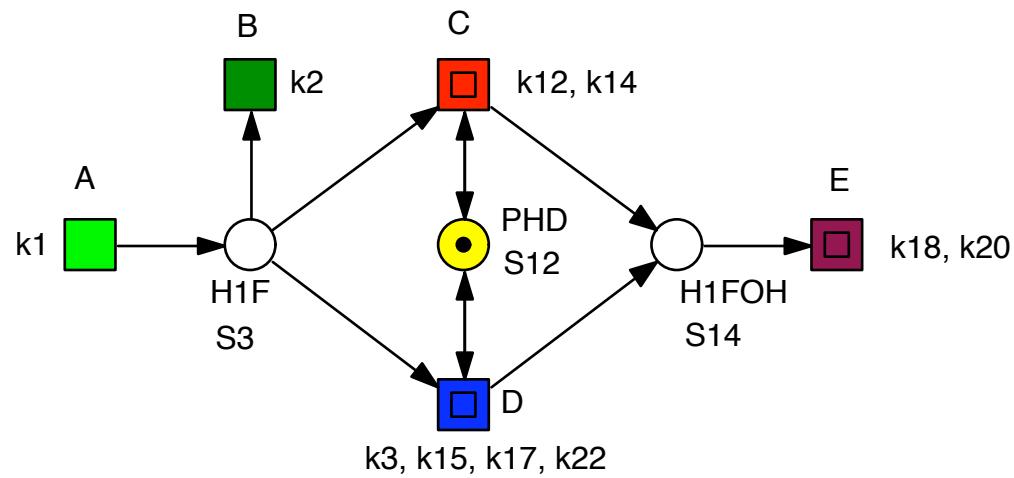
## Ex2 - HYPOXIA

PN & BioModel Engineering

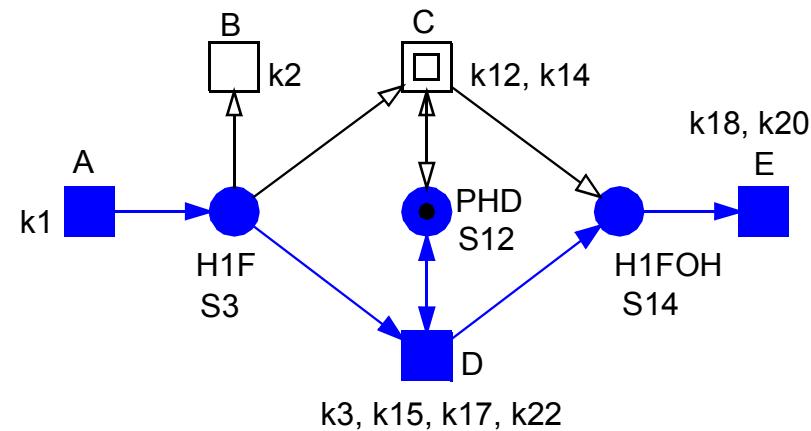
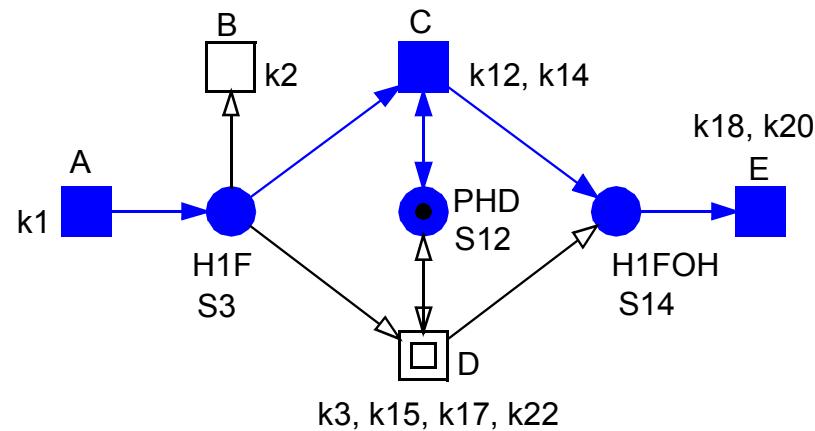
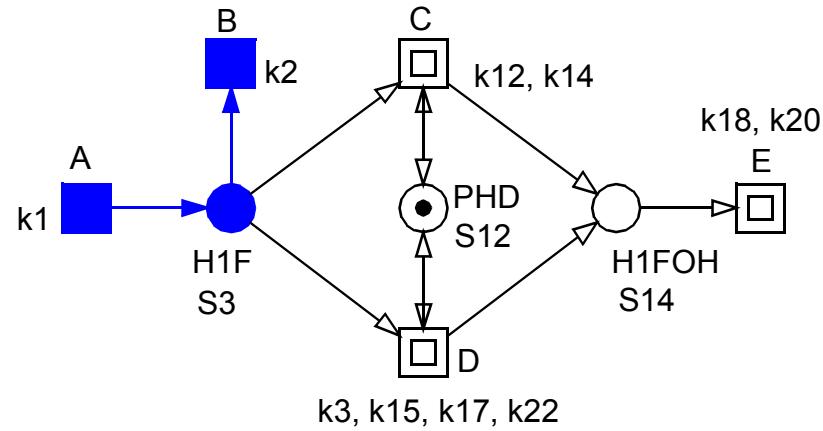


## Ex2 - HYPOXIA





## Ex2 - HYPOXIA



**BUT,**

**THERE ARE MANY EXAMPLES  
WHERE THE TRANSITION SPN -> CPN  
COMES WITH COUNTERINTUITIVE EFFECTS.**

# **OUR TOOL Box**

## □ **SNOOPY**

- > *modelling and animation/simulation of hierarchical graphs,*  
*e.g. (extended) fault trees,*  
*various Petri net classes, e.g. QPN, XQPN, SPN, XSPN, CPN, TPN,*  
*...,*  
*free style graphs*

## □ **CHARLIE**

- > *QPN, XQPN, Time/Timed Petri nets (TPN)*
- > *mostly standard analysis techniques of Petri net theory*

## □ **MARCIE**

- > *XQPN, SPN, XSPN*
- > *symbolic and simulative model checking*

-> *modelling and animation/simulation of hierarchical graphs*

**history**

- > *predecessor: Petri Net Editor PED, 1992 - 2004*
- > *initial implementation concepts 1997*
- > *core implementation 2004*
- > *many Master Theses*

**platform-independent**

- > *implementation in C++*
- > *wxWidgets (<http://www.wxwidgets.org>)*

**supported operating systems**

- > *MAC OS X*
- > *Windows*
- > *Linux (selected distributions)*

**now about 120,000 lines of code (without running extensions)**

- > standard analysis techniques of Petri net theory -> QPN
- > inspired by INA

- structural analysis**

- > net classes: SM, SG, FC, EFS, ES
  - > siphon/trap property, rank theorem

- analysis based on incidence matrix**

- > structural boundedness test
  - > place/transition invariants
  - > Abstract Dependent Transition sets (ADT sets)

- reachability/coverability graph**

- > explicite model checking -> RG of XQPN

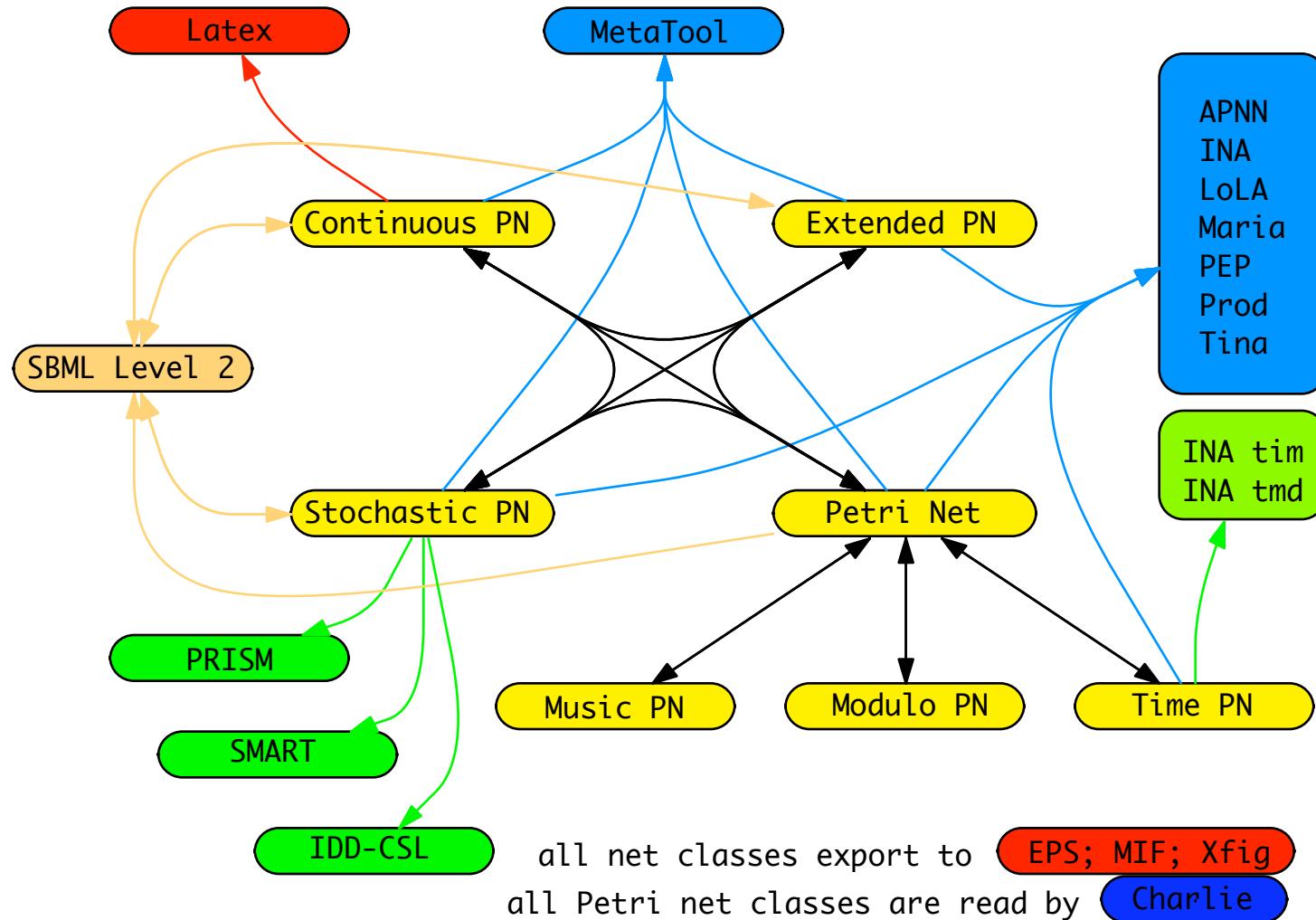
- structural reduction**

- Java thread programming with GUI**

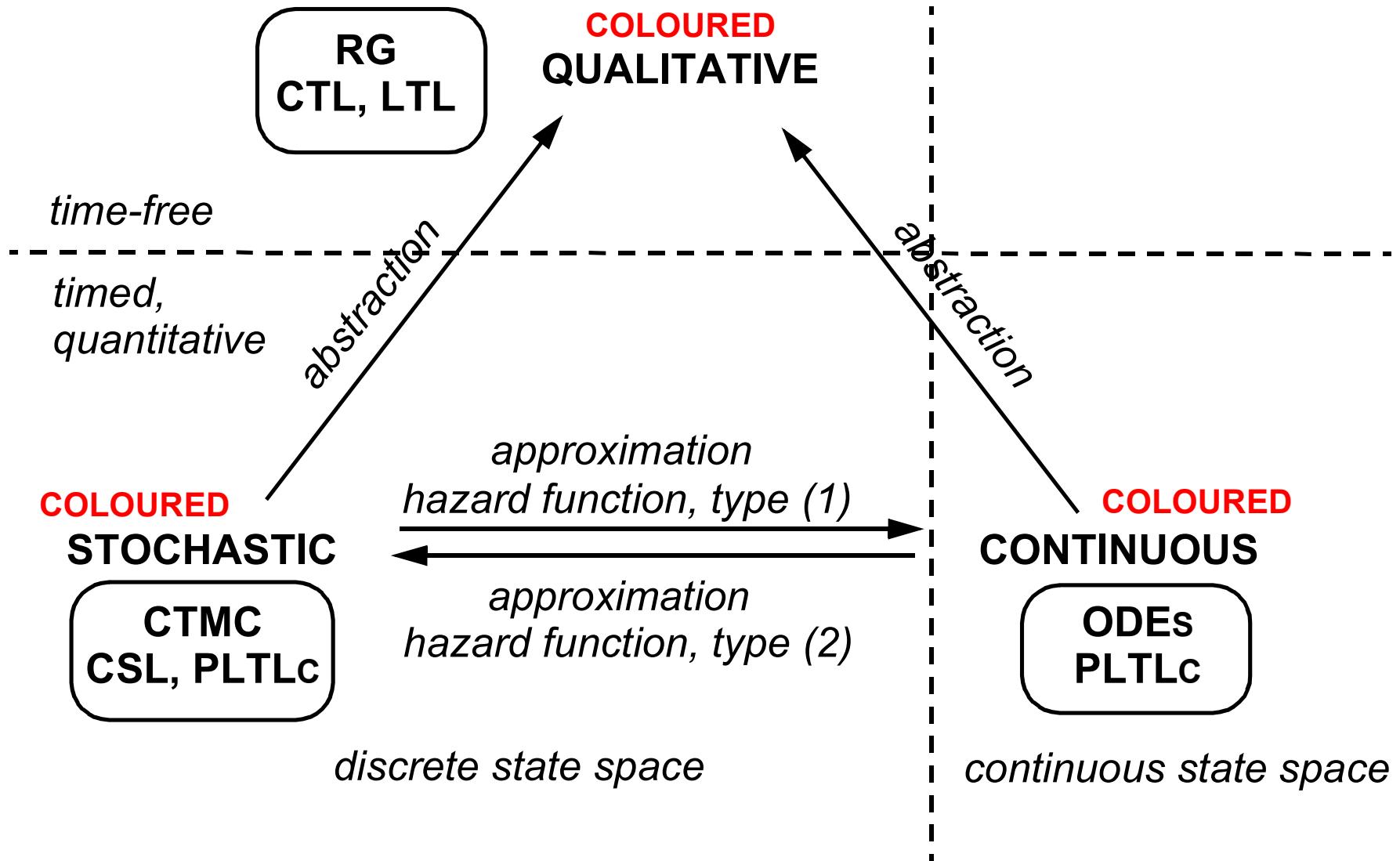
- **Model checking And Reachability analysis done effiCIEntly**
  - > *symbolic state space analysis (strongly connected components)*
    - > *Interval Decision Diagrams*
  - > *model checking of XQPN, SPN, XSPN*
- **XQPN: symbolic CTL model checking**
- **SPN: symbolic CSL model checking**
  - > “matrix free” transient and steady state analysis
  - > parallelized
  - > full CSL model checking + rewards
- **XSPN: simulative PLTL model checking**
  - > distributed
  - > in-line/ off-line traces
- **command line tool, written in C++**

# SNOOPY'S EXPORT FEATURES

PN & BioModel Engineering

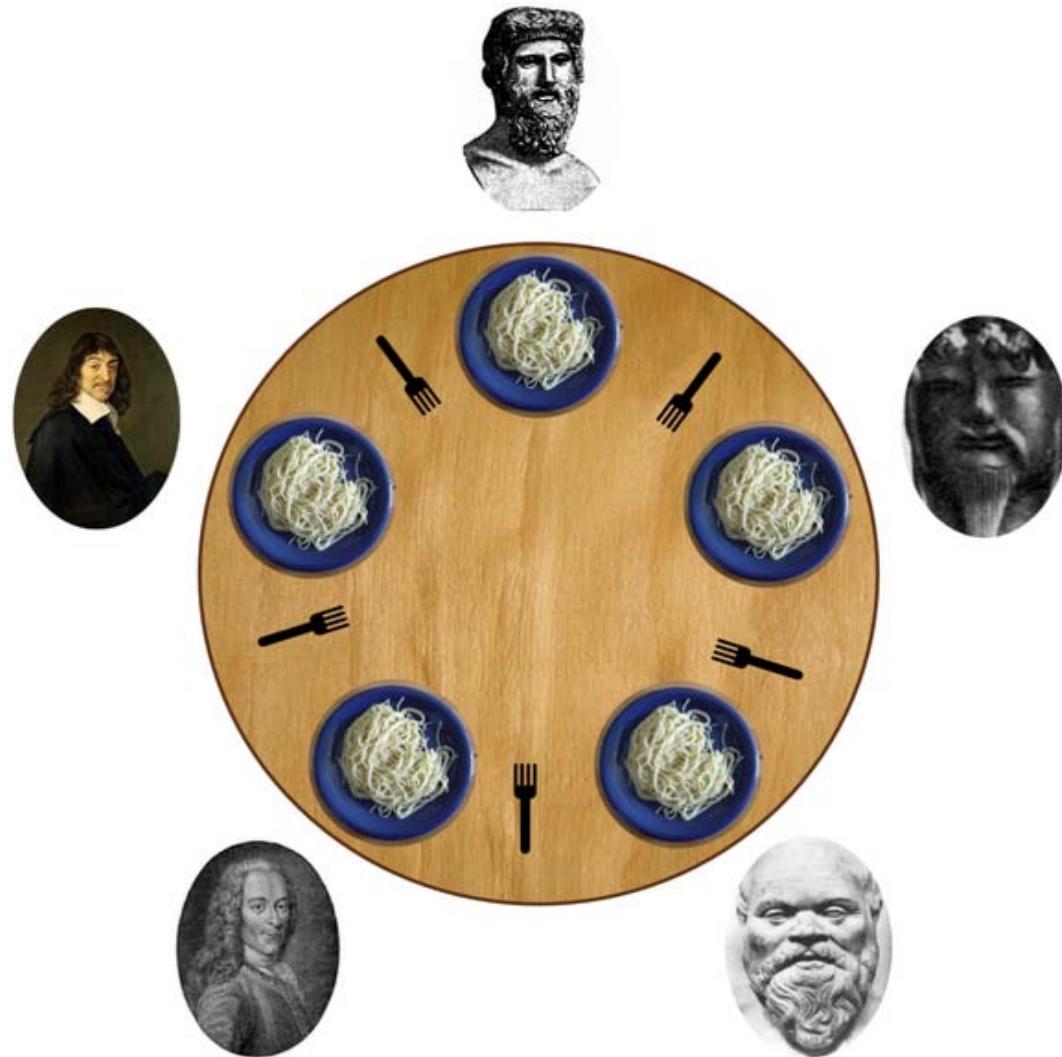


# LATEST NEWS



# COLOURED PETRI NETS, EXAMPLE

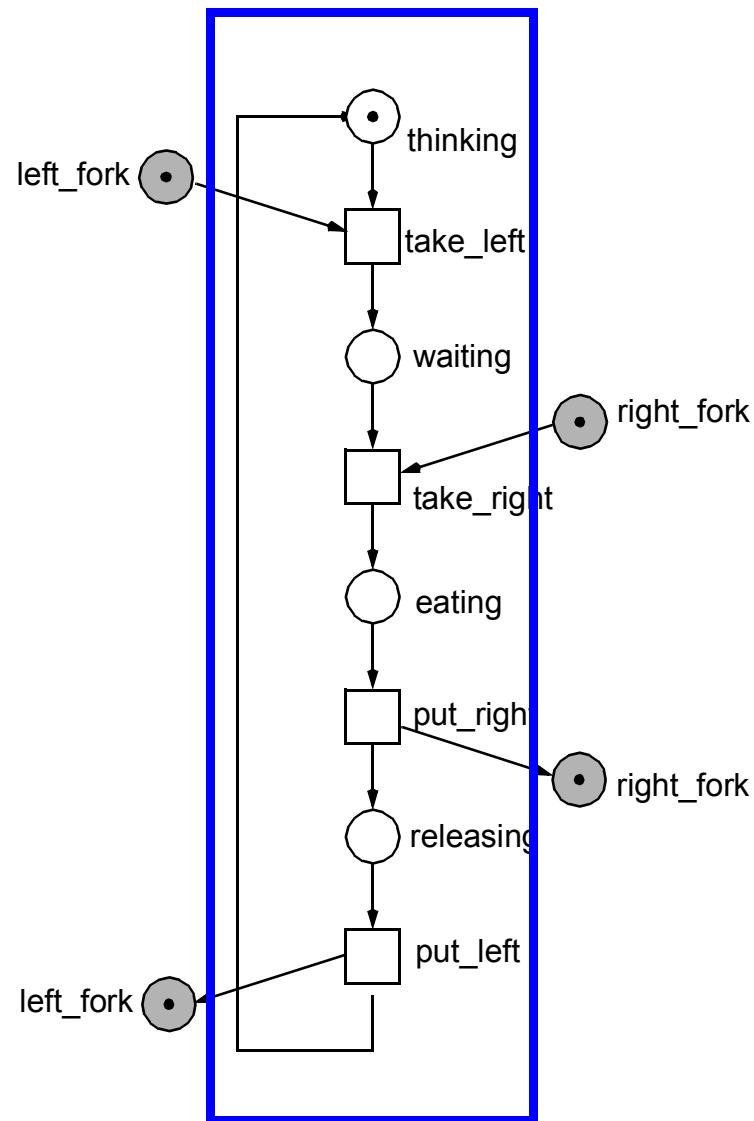
PN & BioModel Engineering



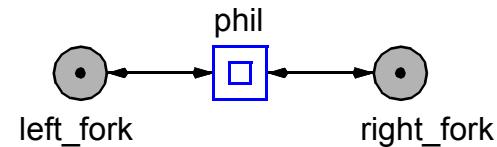
[http://en.wikipedia.org/wiki/Dining\\_philosophers\\_problem](http://en.wikipedia.org/wiki/Dining_philosophers_problem)

# COLOURED PETRI NETS, EXAMPLE

PN & BioModel Engineering



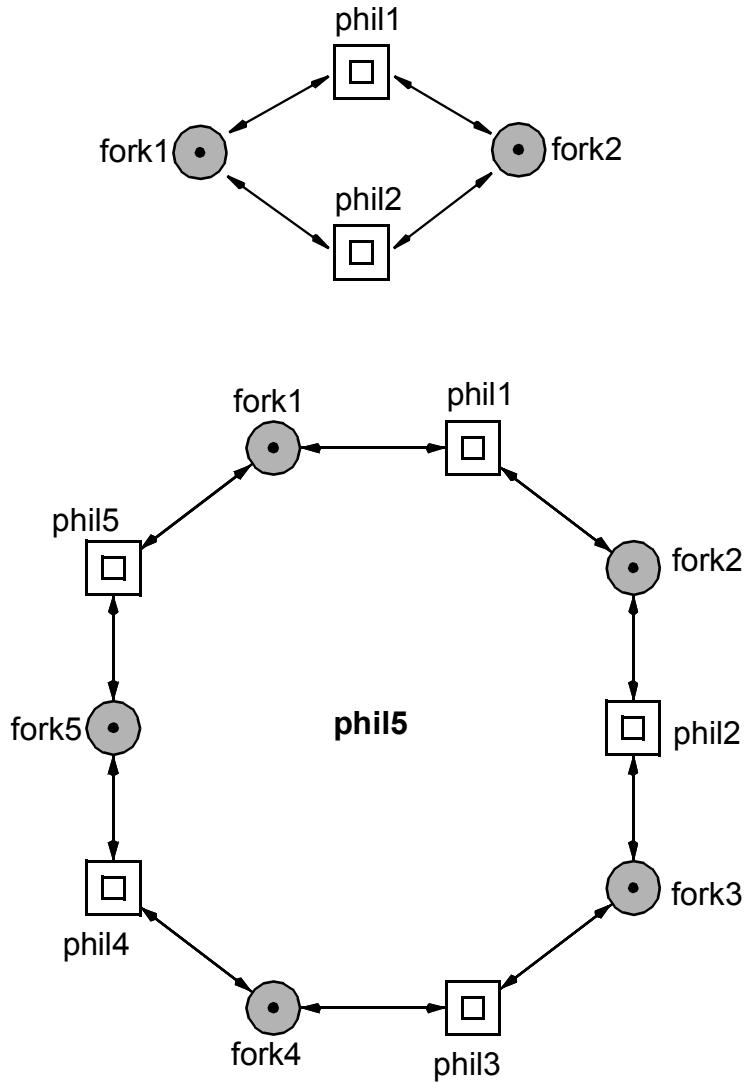
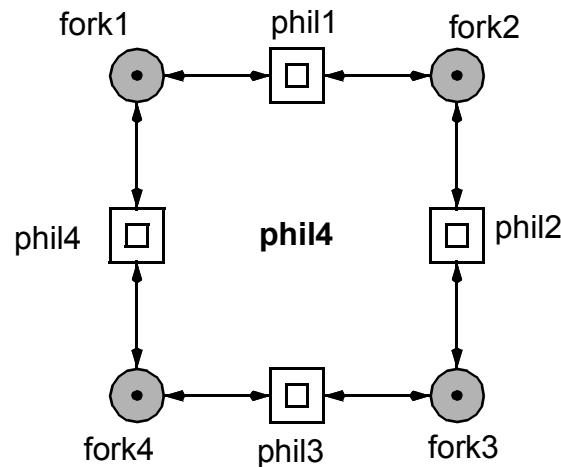
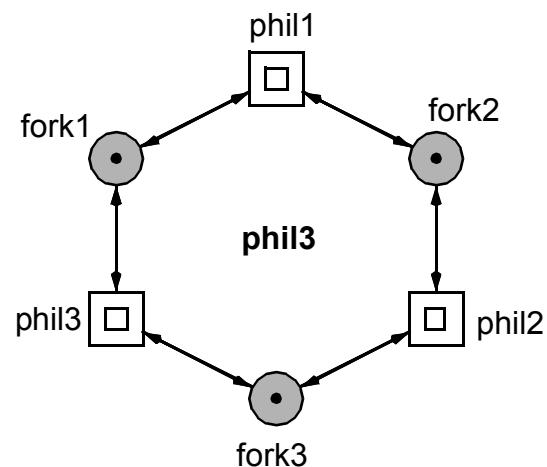
PHILOSOPHER PATTERN



# COLOURED PETRI NETS, EXAMPLE

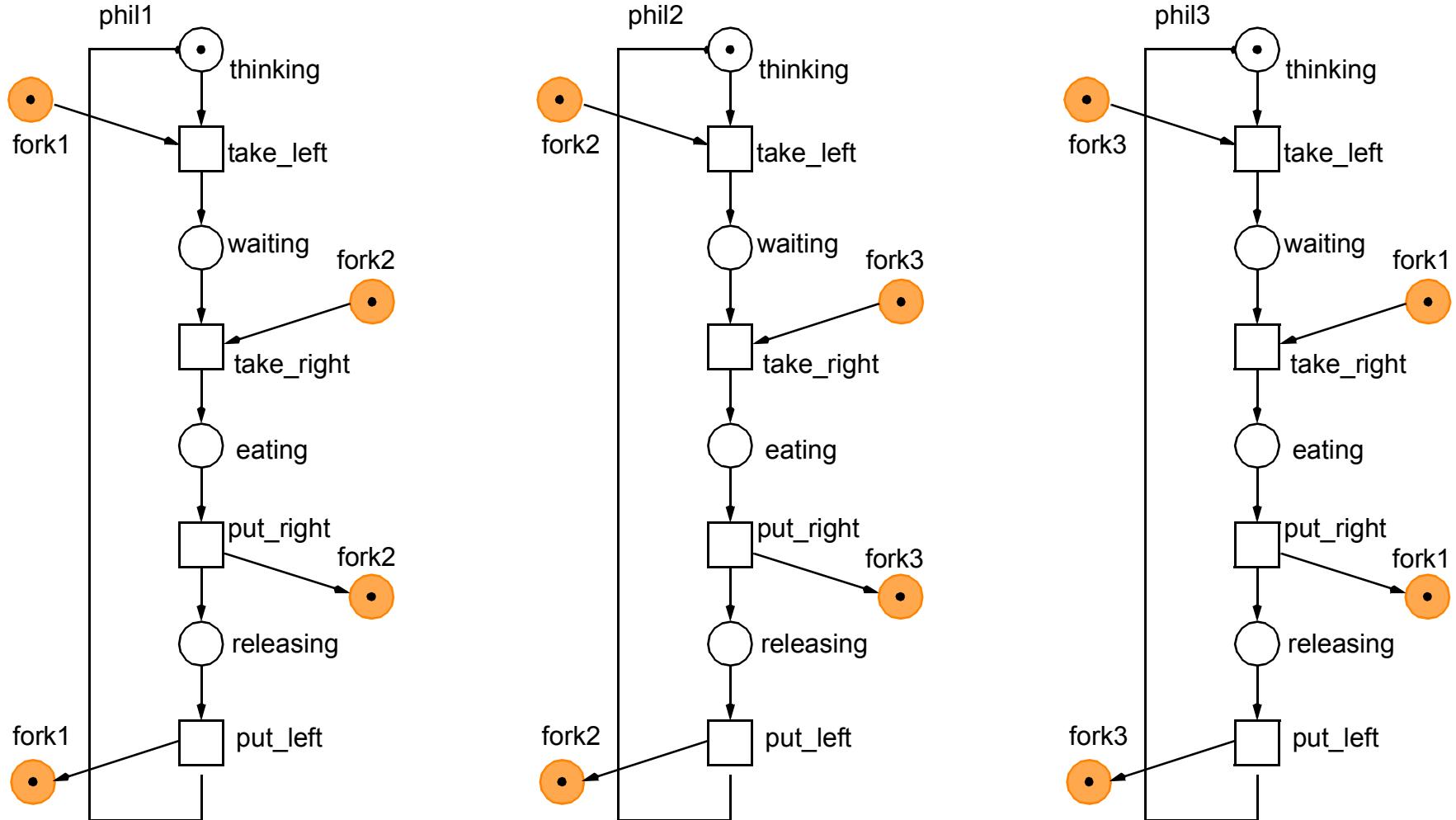
PN & BioModel Engineering

## SYSTEMS OF PHILOSOPHERS



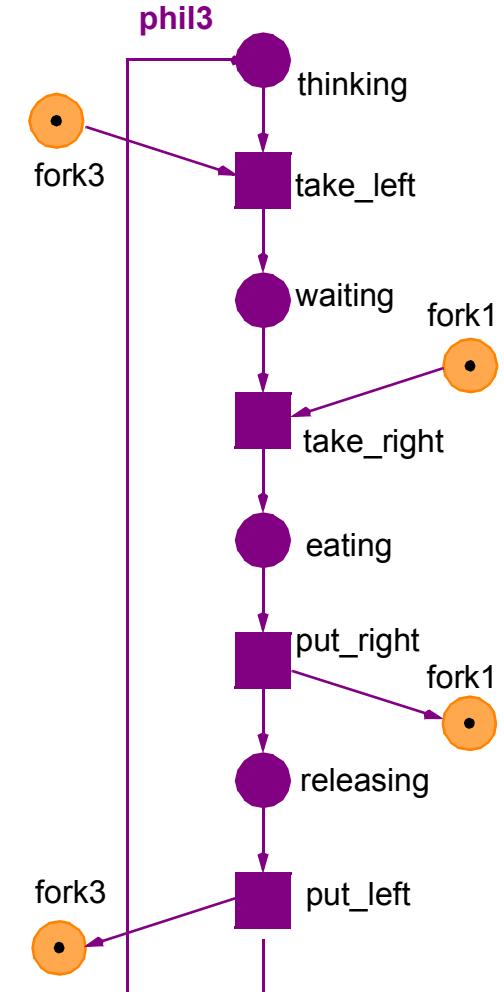
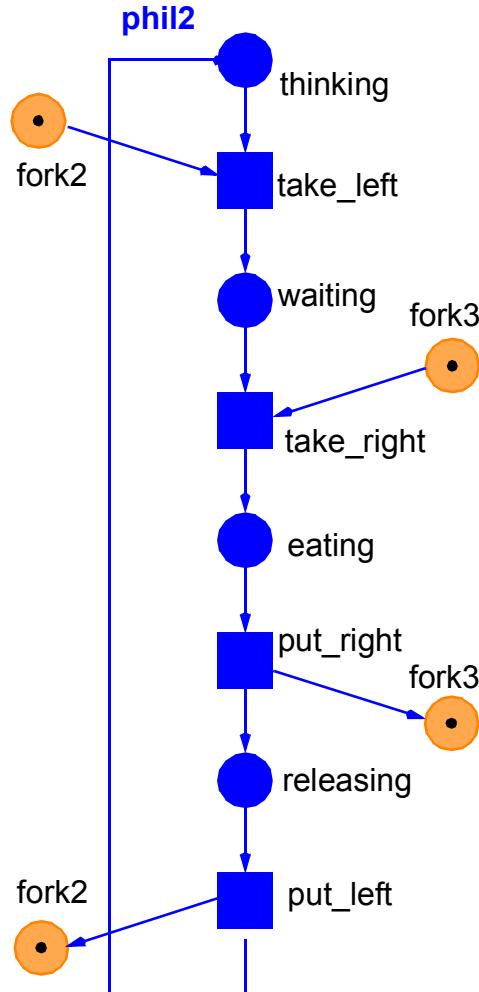
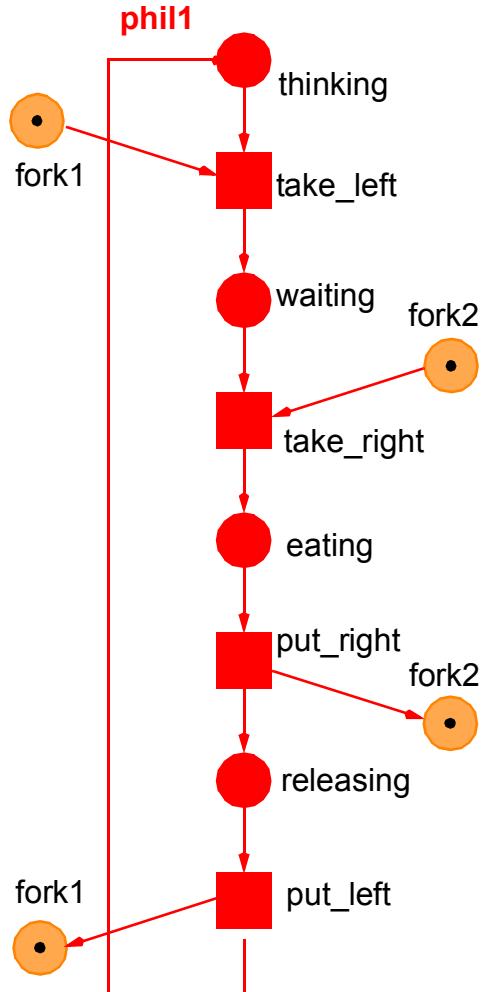
# COLOURED PETRI NETS, EXAMPLE

PN & BioModel Engineering



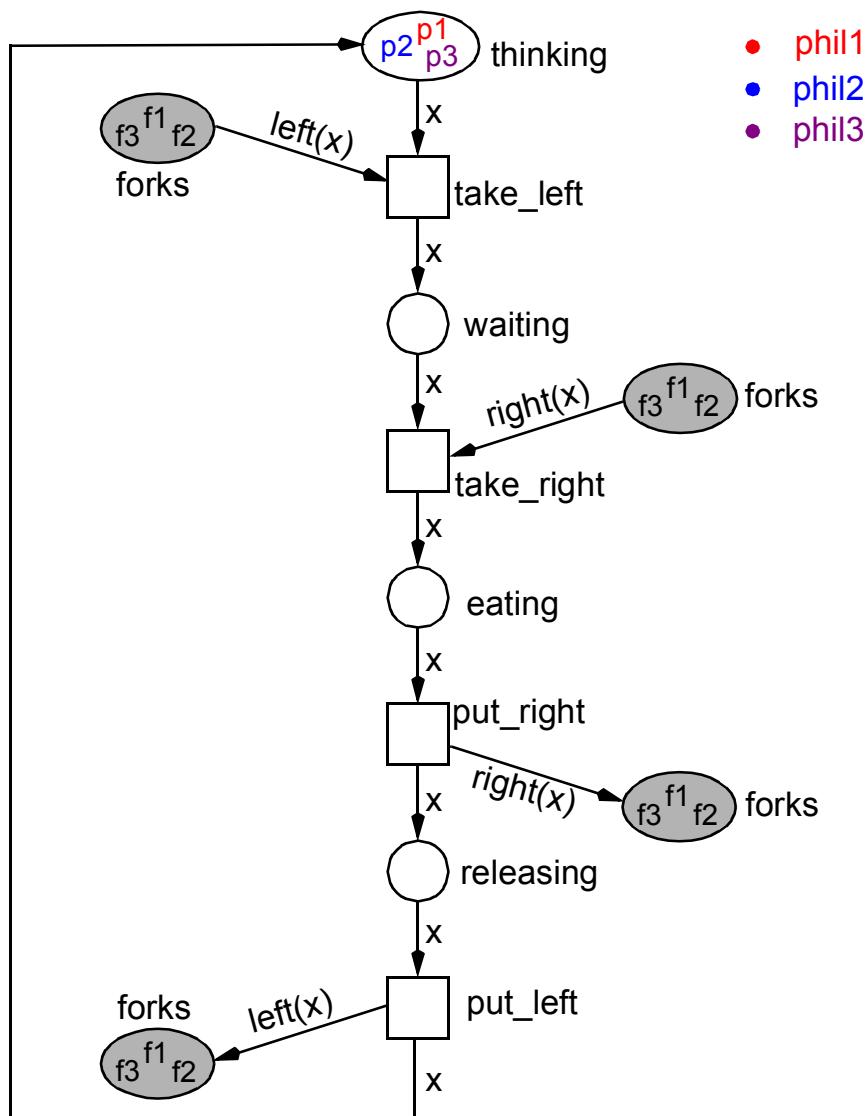
# COLOURED PETRI NETS, EXAMPLE

PN & BioModel Engineering



# COLOURED PETRI NETS, EXAMPLE

PN & BioModel Engineering



- phil1
- phil2
- phil3

const  $N = 3$  // number of phils

colour\_set Phils =  $p\{1 \dots N\}$   
 colour\_set Forks =  $f\{1 \dots N\}$

colour\_set(thinking) = Phils

...

colour\_set(releasing) = Phils

colour\_set(forks) = Forks

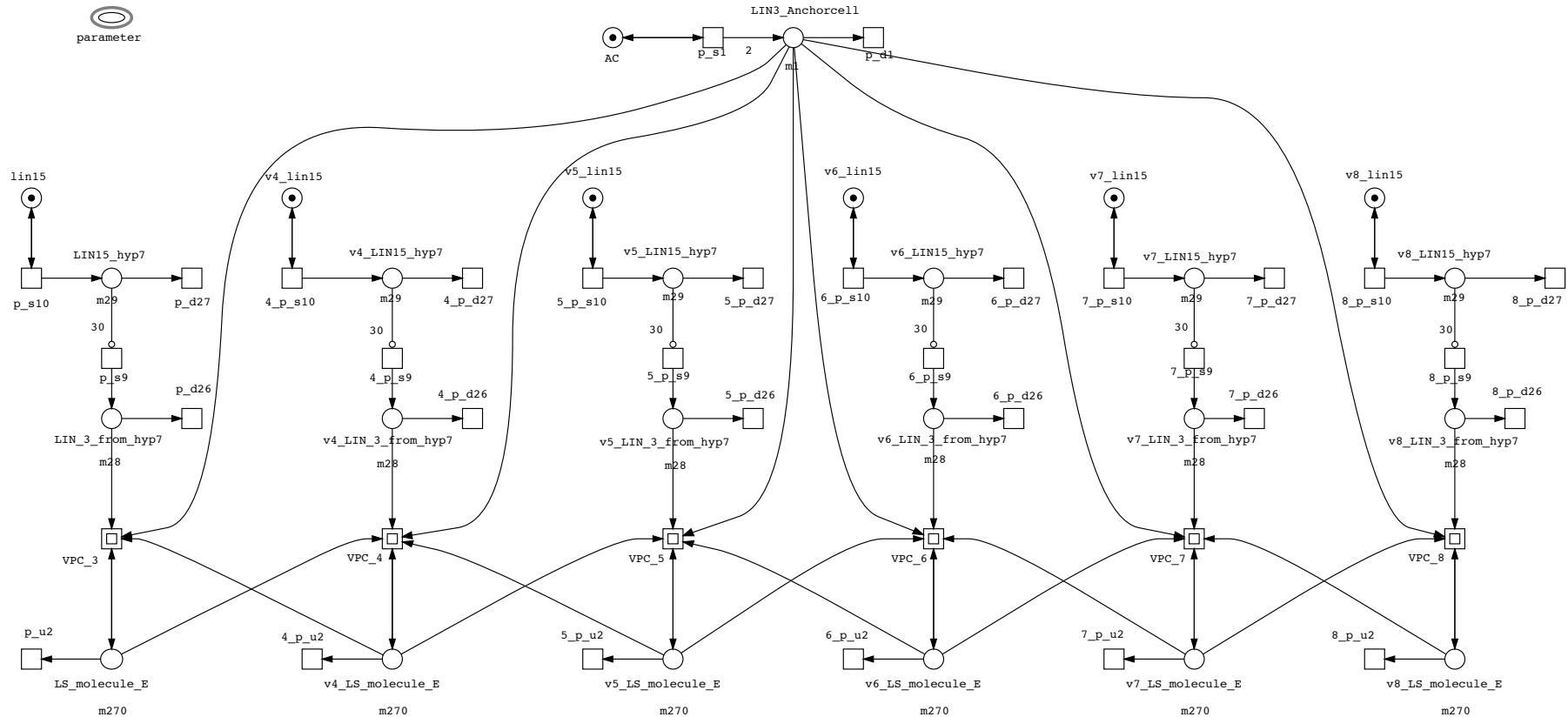
var  $x : \text{Philis}$

$\text{left}(x) = x$

$\text{right}(x) = (x \bmod n) + 1$

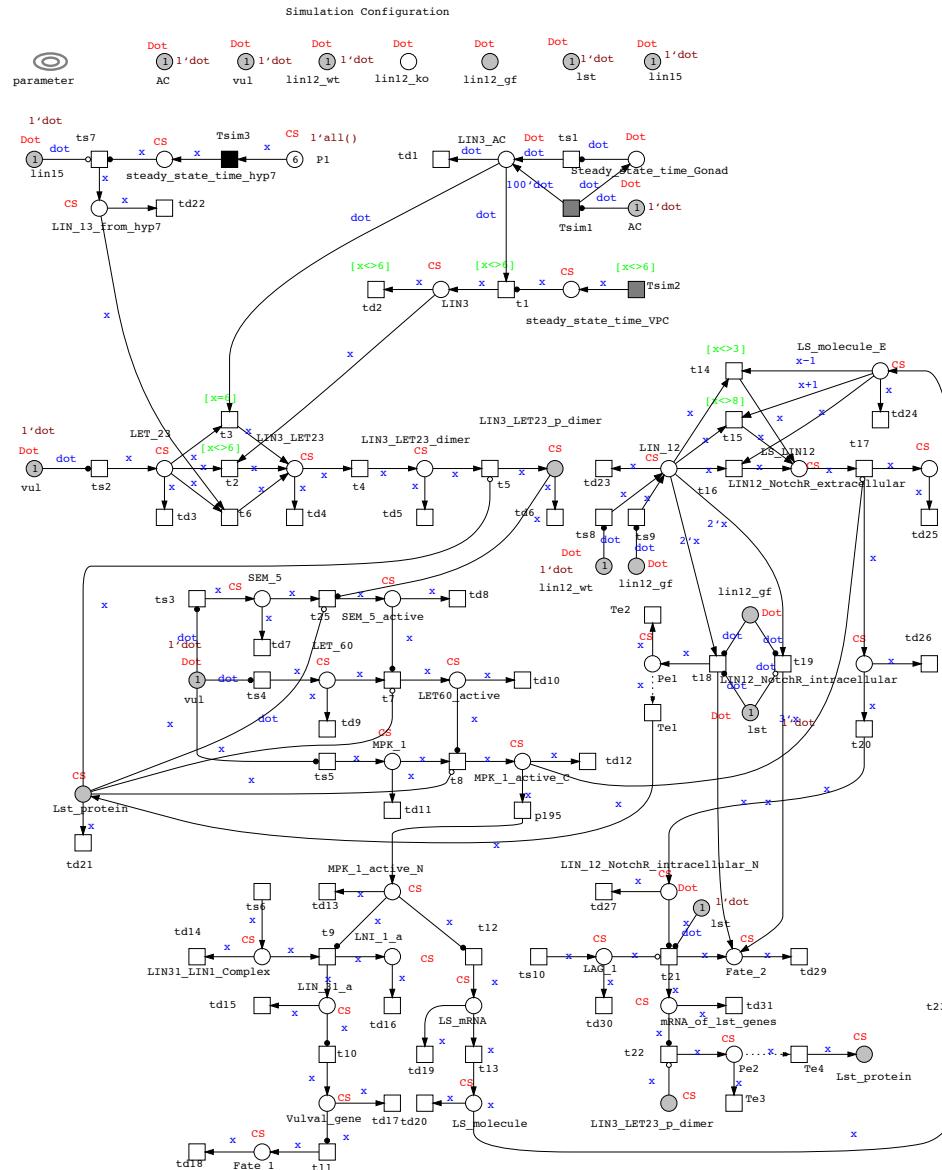
- **get multiple copies of patterns**
  - > *Halo model, new order of net sizes*
- **differentiate between submodels within a master net**
  - > *T-invariants*
  - > *generated models in conformance with wet-lab data*
  - > *mutants*
  - > *algorithmic folding*
- **encode locality**
  - > *Ca channel models*
  - > *cell tissue + communication between cells*
  - > *motility, gradients, . . .*
- **dynamic membrane systems**
- ...

[LI ET AL. 2009]  
 [BONZANNI ET AL. 2009]



# Ex - C. ELEGANCE

PN & BioModel Engineering



## □ Extended Generalized Stochastic Petri Nets (XSPN)

- > *discrete places*
- > *discrete transitions: stochastic, immediate, deterministically delayed, scheduled*
- > *special arcs: read, inhibitor, equal, reset*

## □ Continuous Petri Nets (CPN)

- > *continuous places*
- > *continuous transitions*
- > *special arcs: read, inhibitor*



**GHPN = XSPN + CPN**

## □ hybrid simulation engine

- > *static partitioning*
- > *dynamic partitioning*

# **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*

## □ representation of bionetworks by Petri nets

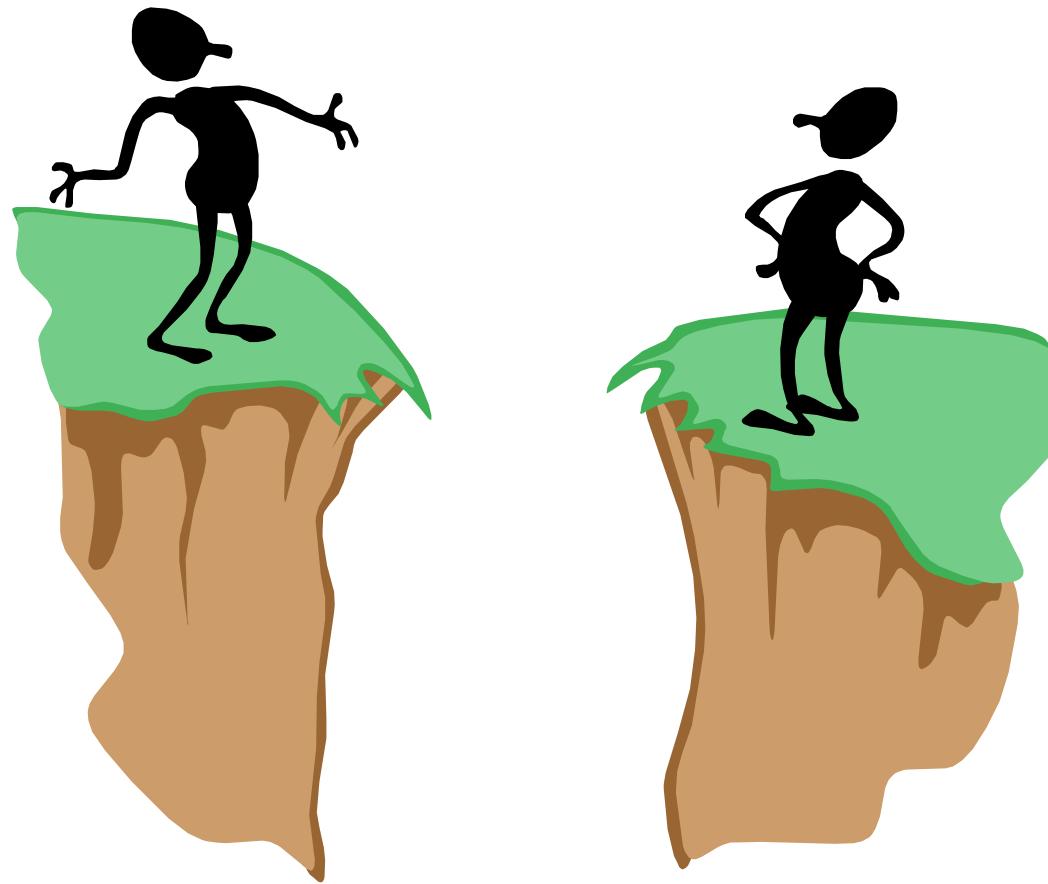
- > *partial order representation*
- > *formal semantics*
- > *unifying view*
- > *better comprehension*
- > *sound analysis techniques*

## □ purposes

- > *animation*
- > *model validation against consistency criteria*
- > *qualitative / quantitative behaviour prediction*
- > *to experience the model*
- > *to increase confidence*
- > *experiment design, new insights*

## □ step-wise model development

- > *qualitative model*
- > *discrete quantitative model*
- > *continuous quantitative model*
- > *discrete Petri nets*
- > *stochastic Petri nets*
- > *continuous Petri nets = ODEs*



**THANKS !**

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