

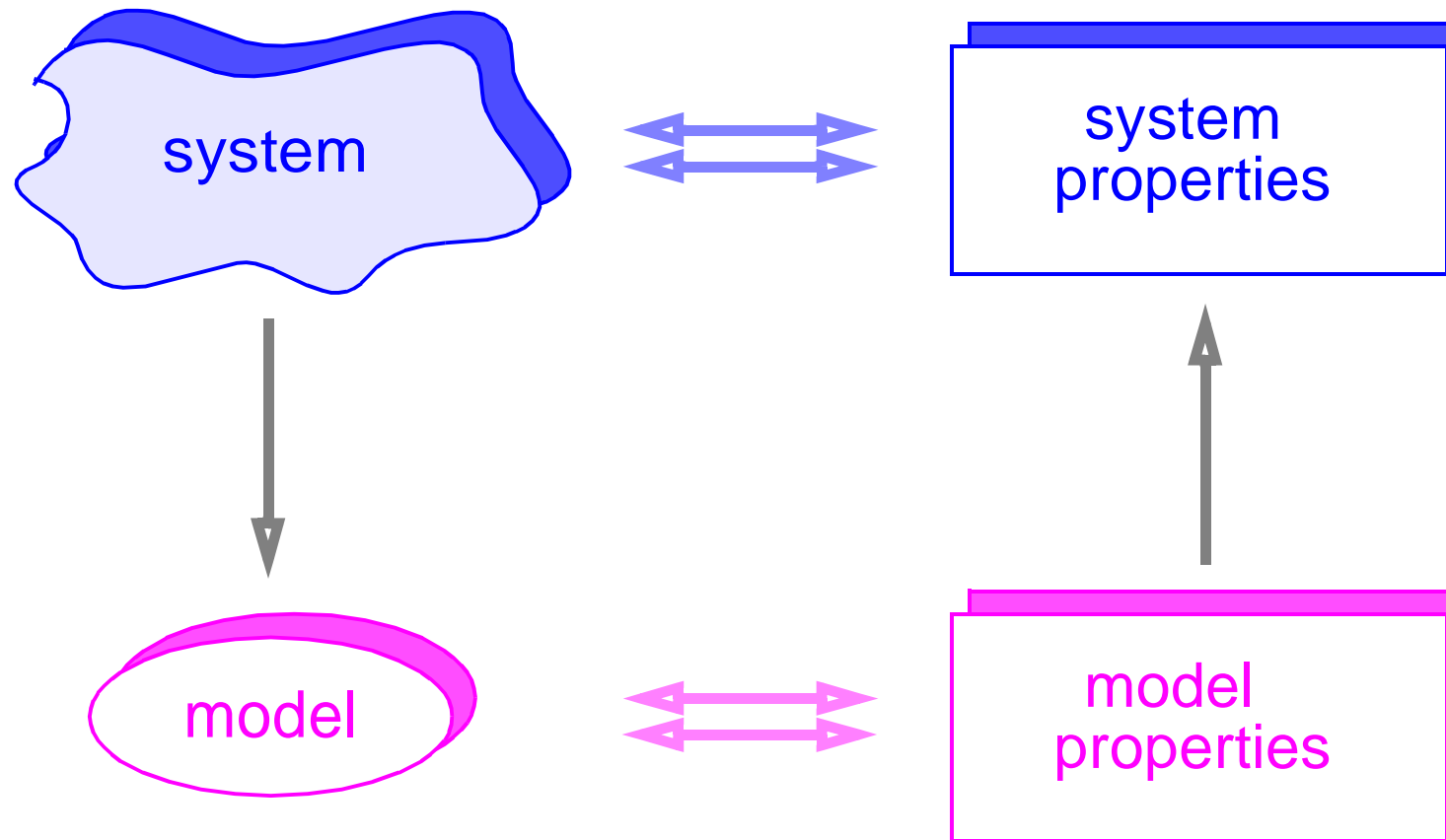
FROM PETRI NETS TO DIFFERENTIAL EQUATIONS

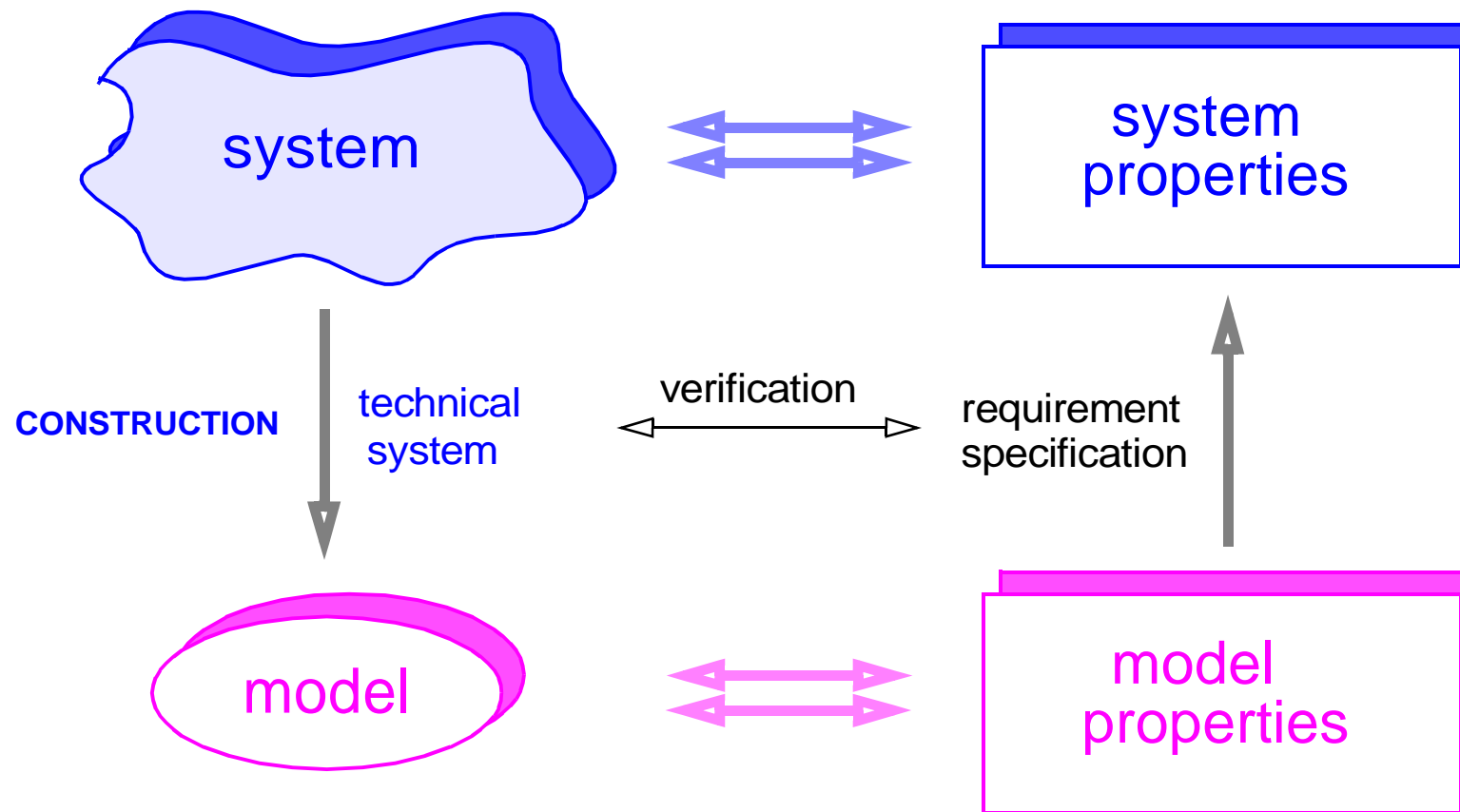
AN INTEGRATIVE APPROACH
FOR BIOCHEMICAL NETWORK ANALYSIS

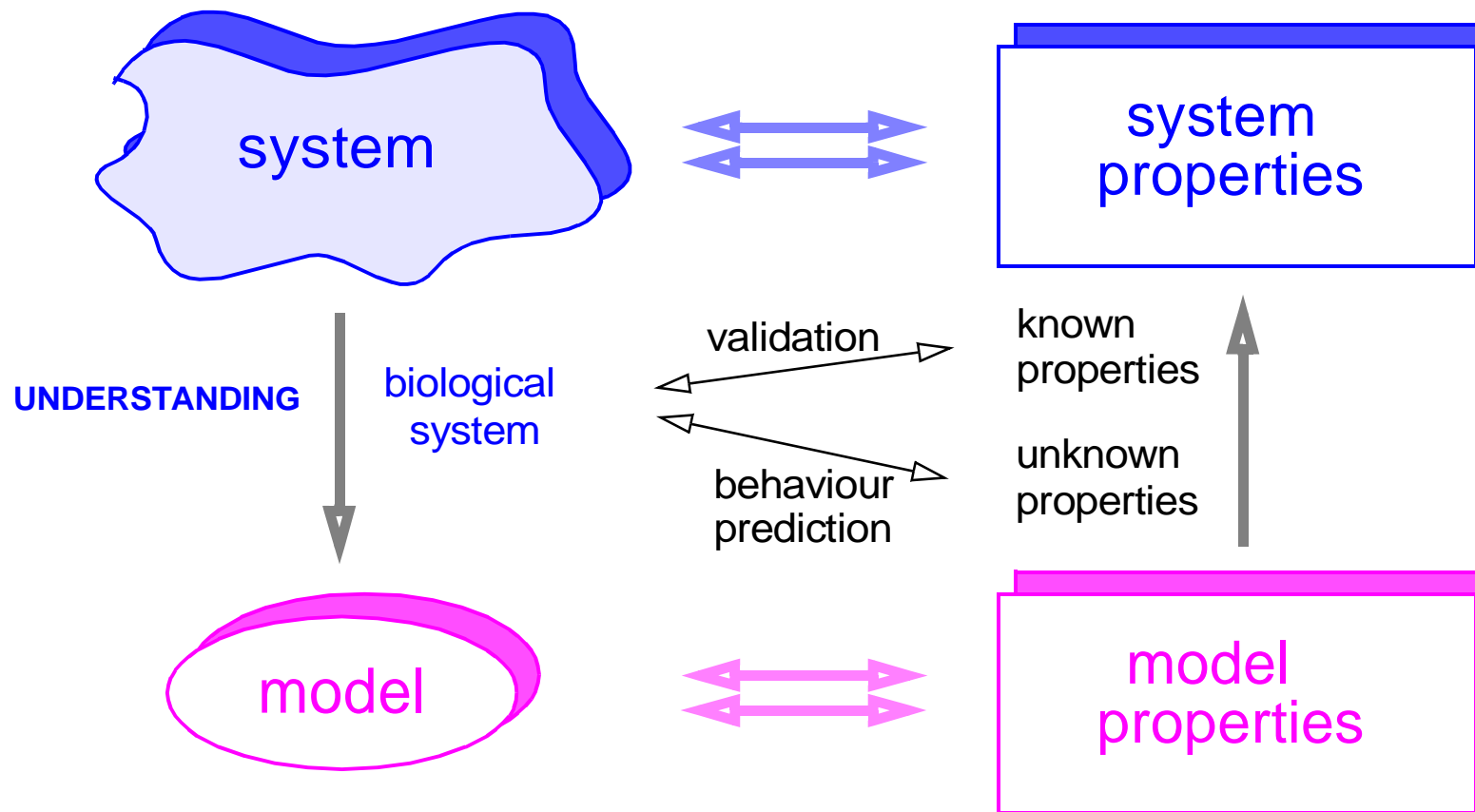
Monika Heiner

Brandenburg University of Technology Cottbus

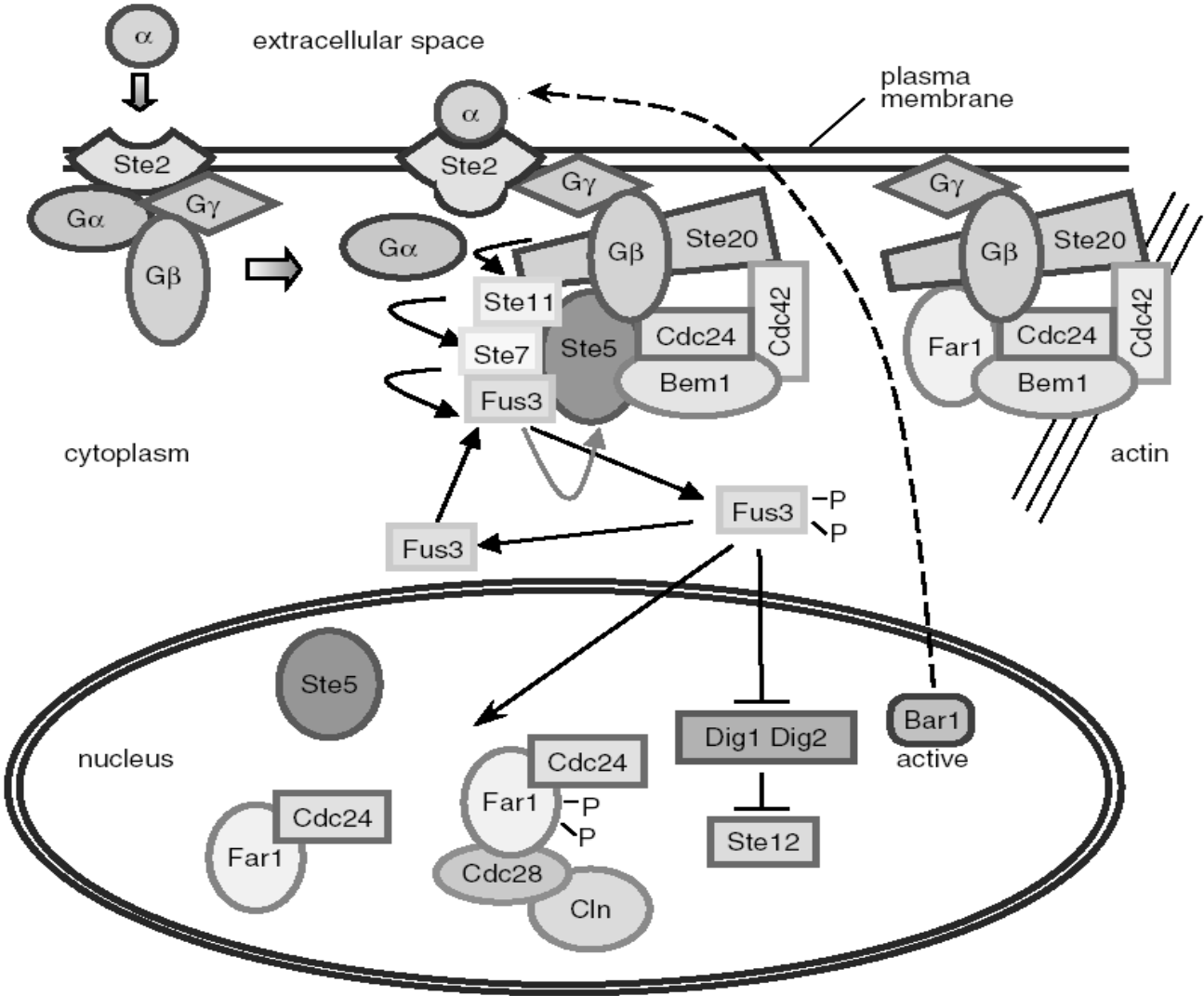
Dept. of CS

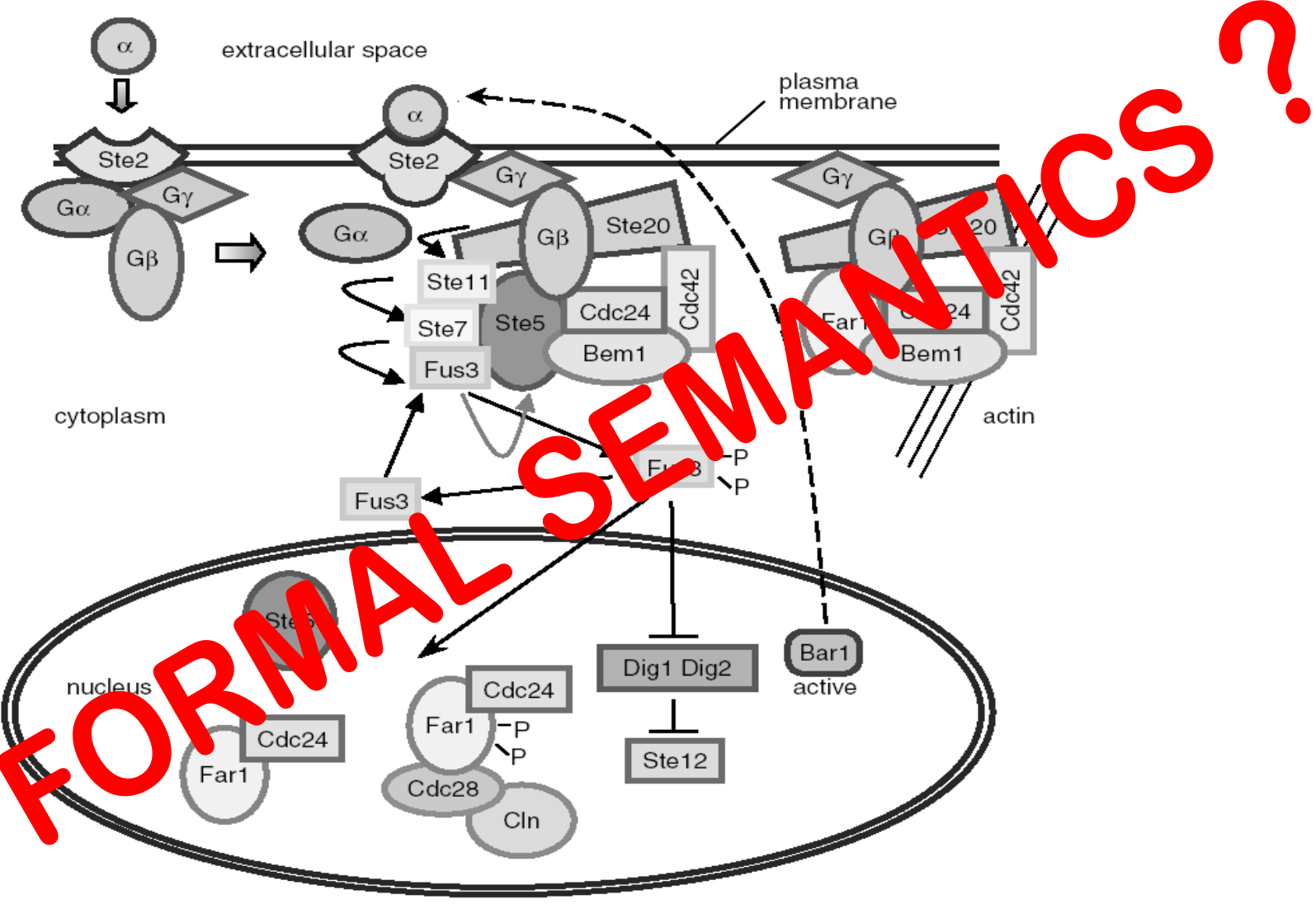






WHAT KIND OF MODEL SHOULD BE USED?





$$\begin{aligned} \frac{d\alpha}{dt} &= -v_1 \\ \frac{dSte2}{dt} &= -v_2 + v_3 - v_5 \\ \frac{dSte2_{active}}{dt} &= v_2 - v_3 - v_4 \\ \frac{dSst2_{active}}{dt} &= v_{46} - v_{47} \\ \frac{dG\alpha\beta\gamma}{dt} &= -v_6 + v_9 \\ \frac{dG\alpha GTP}{dt} &= v_6 - v_7 - v_8 \\ \frac{dG\alpha GDP}{dt} &= v_7 + v_8 - v_9 \\ \frac{dG\beta\gamma}{dt} &= v_6 - v_9 - v_{10} + v_{11} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\ &\quad - v_{42} + v_{43} \\ \frac{dSte5}{dt} &= -v_{12} + v_{13} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\ \frac{dSte11}{dt} &= -v_{12} + v_{13} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\ \frac{dSte7}{dt} &= -v_{14} + v_{15} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \\ \frac{dFus3}{dt} &= -v_{14} + v_{15} + v_{17} + v_{21} + v_{23} + v_{25} + v_{27} - v_{29} \\ &\quad + v_{30} + v_{33} \\ \frac{dSte20}{dt} &= -v_{18} + v_{19} + v_{21} + v_{23} + v_{25} + v_{27} + v_{32} \end{aligned}$$

$$\begin{aligned} v_1 &= \alpha[t] \cdot Bar|_{active}[t] \cdot k_1 \\ v_2 &= Ste2[t] \cdot \alpha[t] \cdot k_2 \\ v_3 &= Ste2_{active}[t] \cdot k_3 \\ v_4 &= Ste2_{active}[t] \cdot k_4 \\ v_5 &= Ste2[t] \cdot k_5 \\ v_6 &= Ste2_{active}[t] \cdot G\alpha\beta\gamma[t] \cdot k_6 \\ v_7 &= G\alpha GTP[t] \cdot k_7 \\ v_8 &= G\alpha GTP[t] \cdot Sst2_{active}[t] \cdot k_8 \\ v_9 &= G\alpha GDP[t] \cdot G\beta\gamma[t] \cdot k_9 \\ v_{10} &= G\beta\gamma[t] \cdot C[t] \cdot k_{10} \\ v_{11} &= D[t] \cdot k_{11} \\ v_{12} &= Ste5[t] \cdot Ste11[t] \cdot k_{12} \\ v_{13} &= A[t] \cdot k_{13} \\ v_{14} &= Ste7[t] \cdot Fus3[t] \cdot k_{14} \\ v_{15} &= B[t] \cdot k_{15} \\ v_{16} &= A[t] \cdot B[t] \cdot k_{16} \\ v_{17} &= C[t] \cdot k_{17} \\ v_{18} &= D[t] \cdot Ste20[t] \cdot k_{18} \end{aligned}$$

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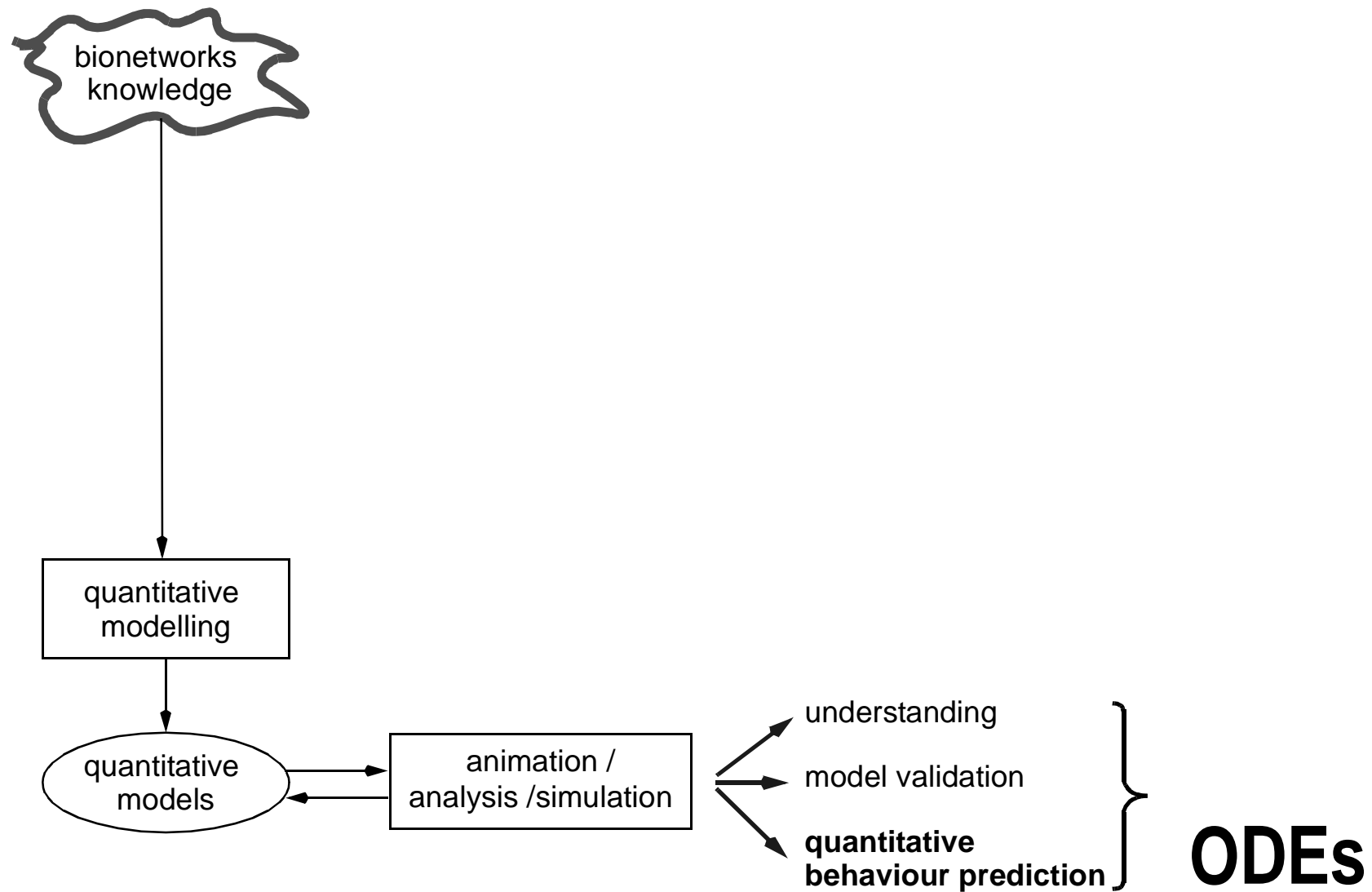
→ READABILITY ?

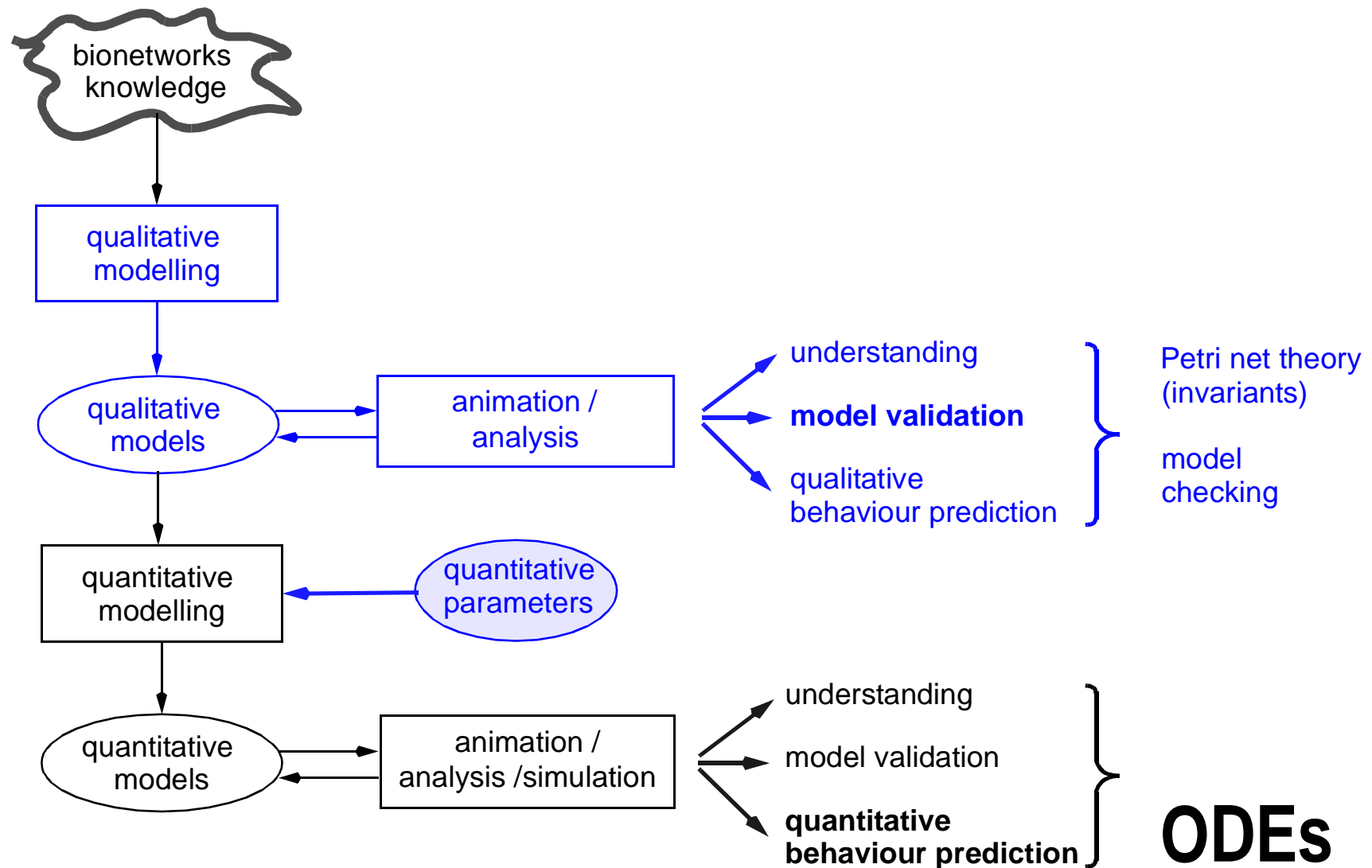
- ❑ **various, mostly ambiguous representations** **-> PROBLEM 1**
 - > *verbose descriptions*
 - > *diverse graphical representations*
 - > *contradictory and / or fuzzy statements*

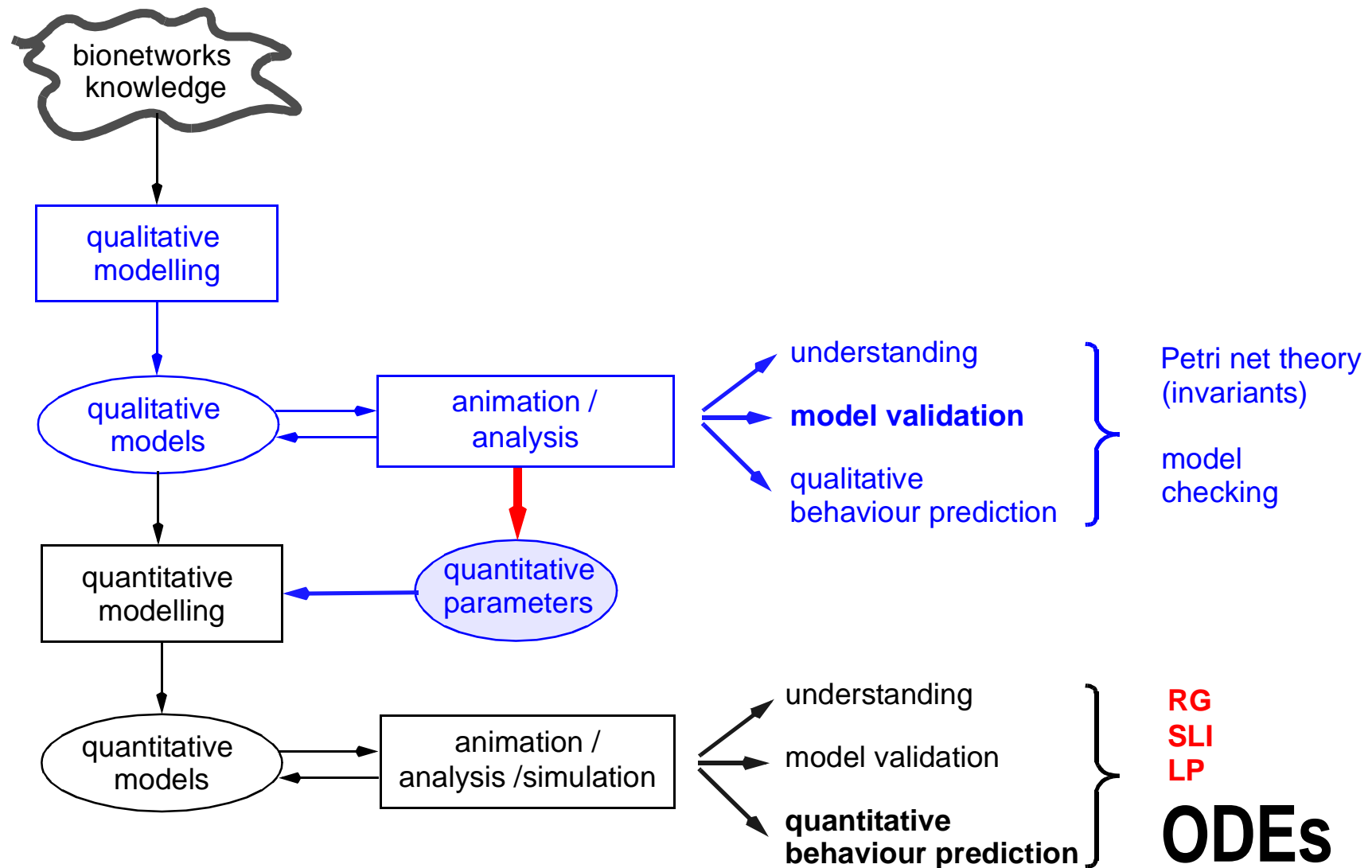
- ❑ **knowledge** **-> PROBLEM 2**
 - > *uncertain*
 - > *growing, changing*
 - > *distributed over various data bases, papers, journals, . . .*

- ❑ **network structures** **-> PROBLEM 3**
 - > *tend to grow fast*
 - > *dense, apparently unstructured*
 - > *hard to read*

-->> models are full of ASSUMPTIONS <<--

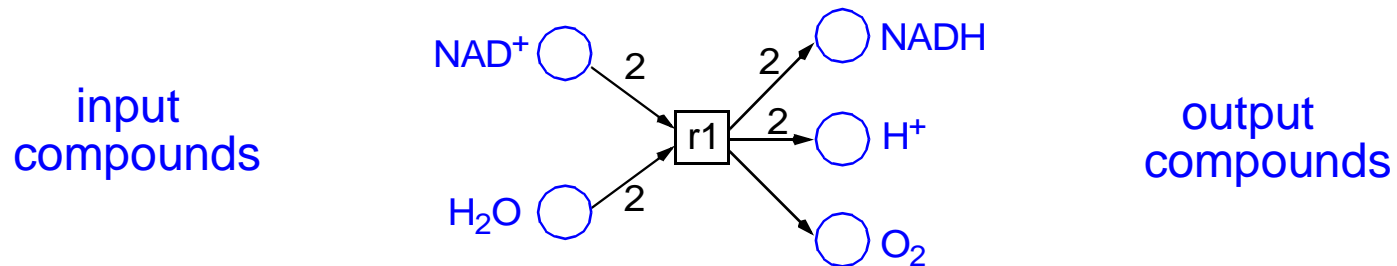
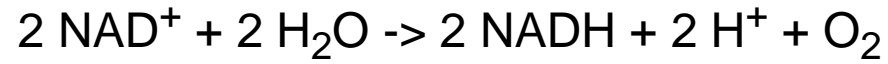




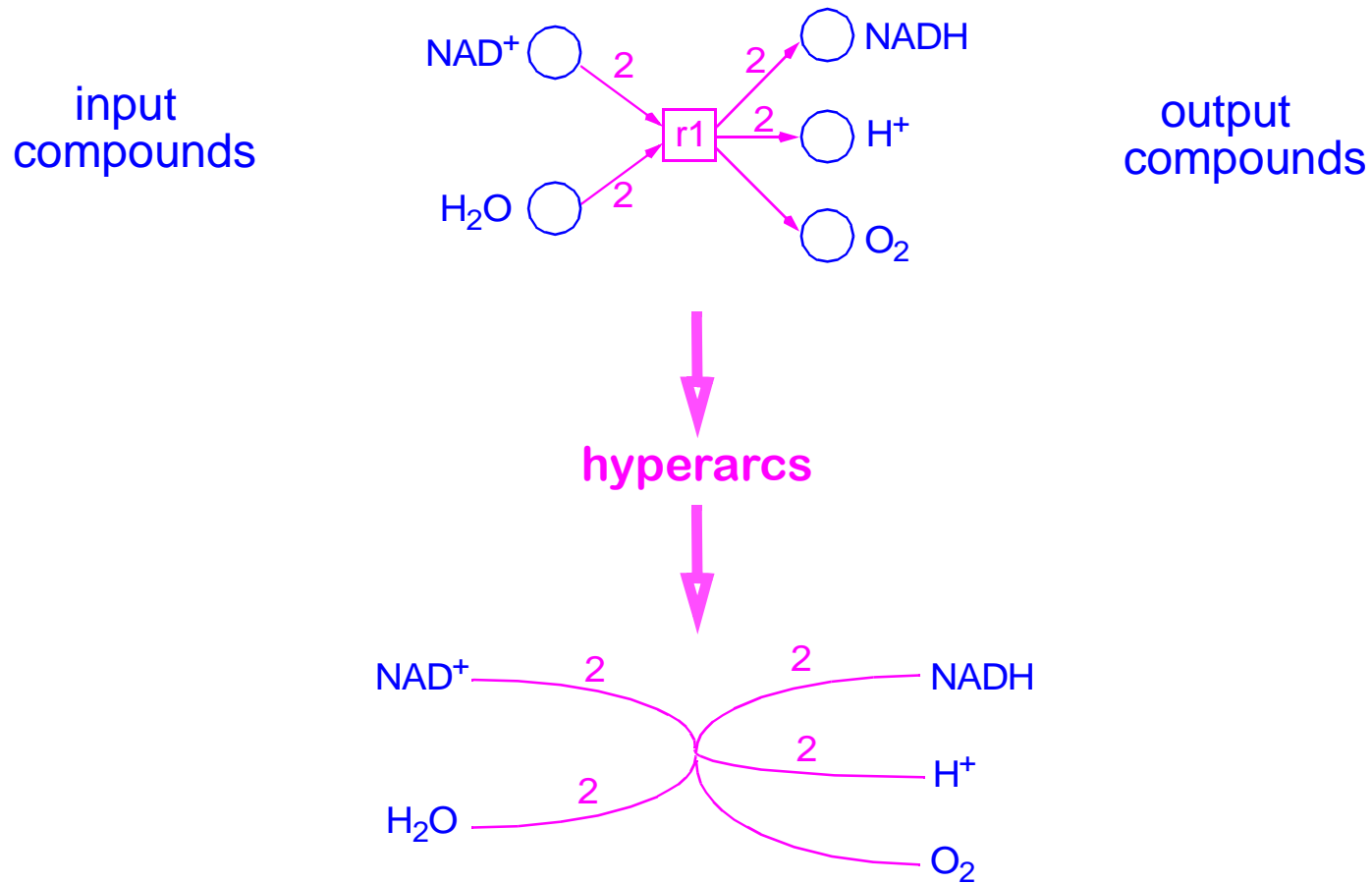
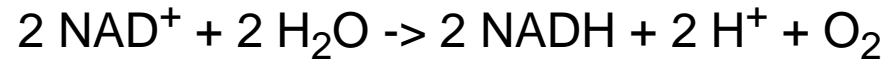


PETRI NETS - AN INFORMAL CRASH COURSE

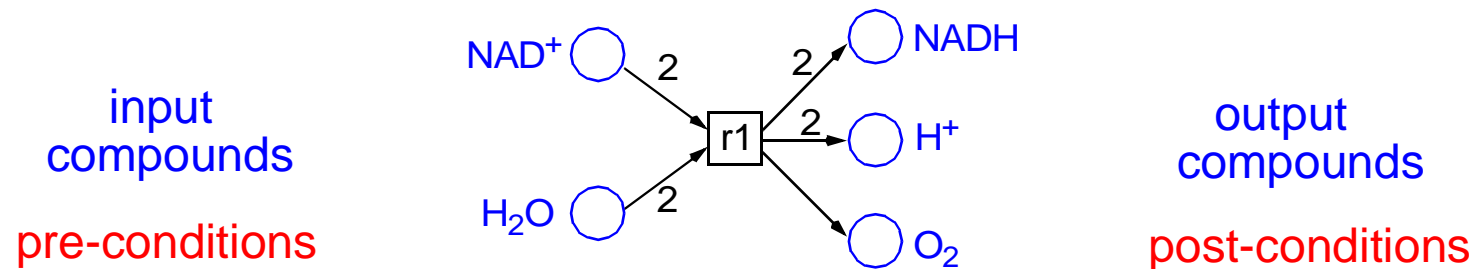
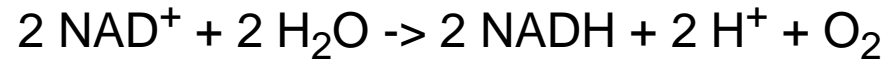
□ atomic actions → Petri net transitions → chemical reactions



□ atomic actions → Petri net transitions → chemical reactions

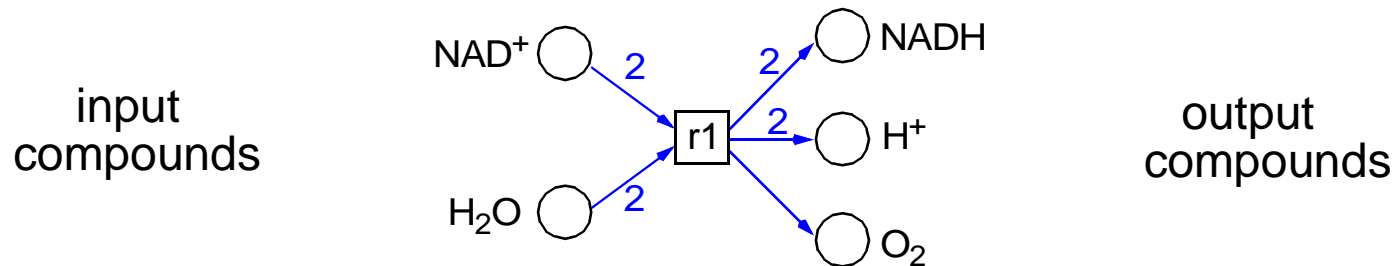
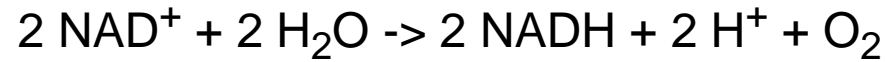


□ atomic actions → Petri net transitions → chemical reactions



□ local conditions → Petri net places → chemical compounds

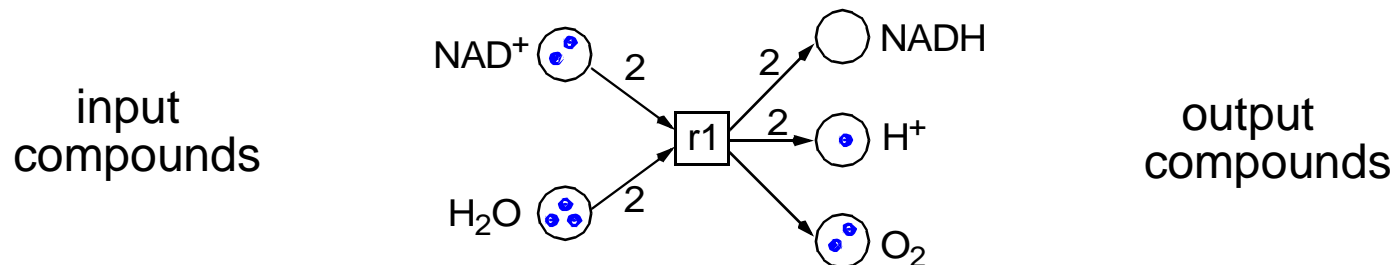
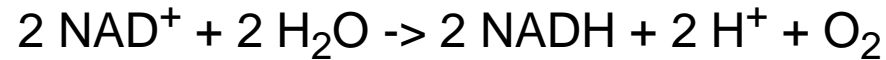
□ atomic actions → Petri net transitions → chemical reactions



□ local conditions → Petri net places → chemical compounds

□ multiplicities → Petri net arc weights → stoichiometric relations

□ atomic actions -> Petri net transitions -> chemical reactions



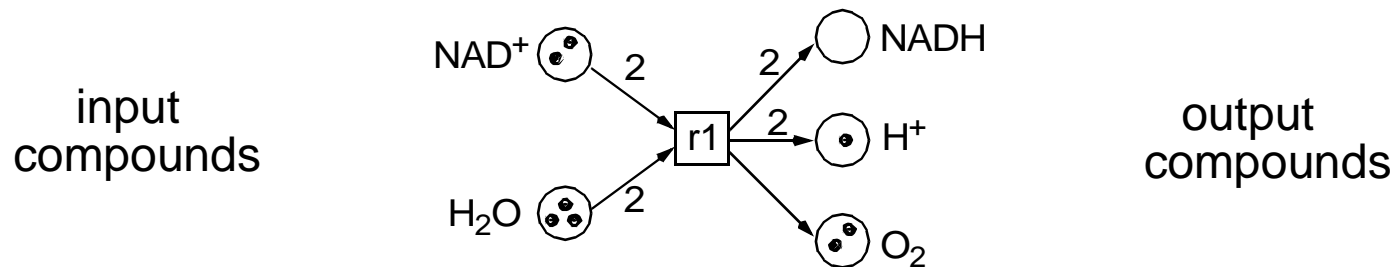
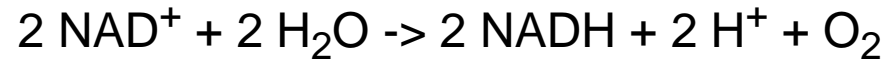
□ local conditions -> Petri net places -> chemical compounds

□ multiplicities -> Petri net arc weights -> stoichiometric relations

□ condition's state -> token(s) in its place -> available amount (e.g. mol)

□ system state -> marking -> compounds distribution

□ atomic actions → Petri net transitions → chemical reactions



□ local conditions → Petri net places → chemical compounds

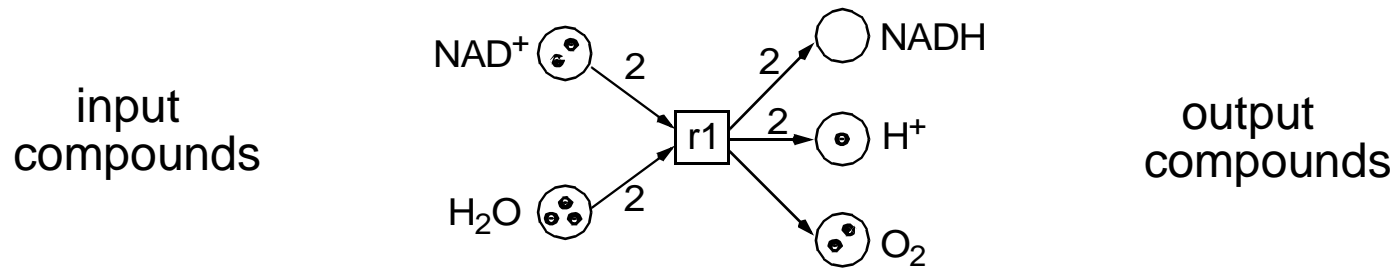
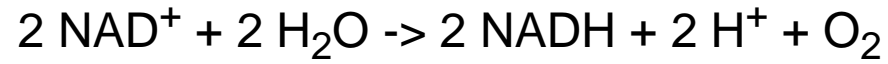
□ multiplicities → Petri net arc weights → stoichiometric relations

□ condition's state → token(s) in its place → available amount (e.g. mol)

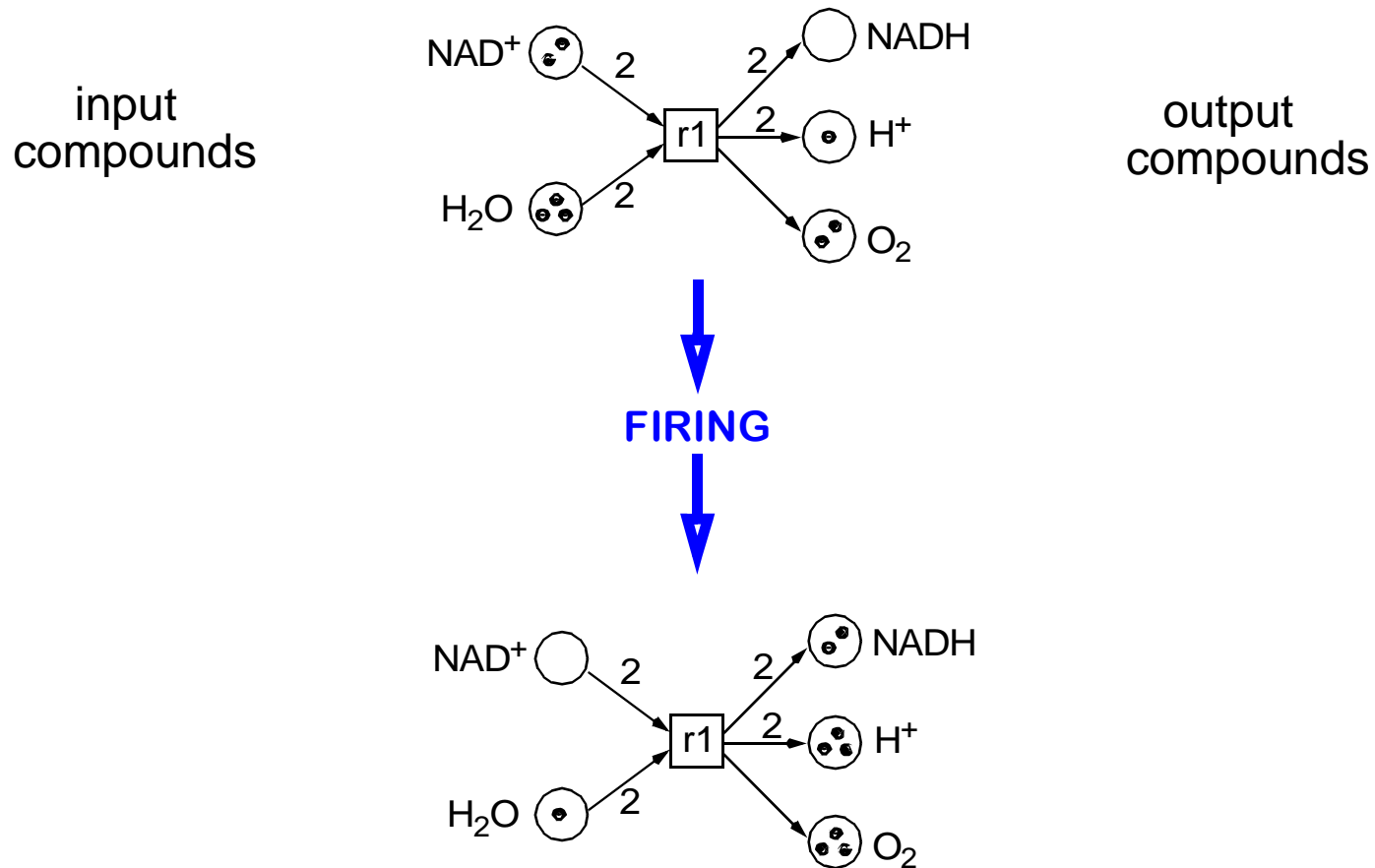
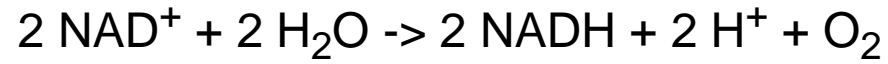
□ system state → marking → compounds distribution

□ $\text{PN} = (\text{P}, \text{T}, \text{F}, \text{m}_0)$, $\text{F}: (\text{P} \times \text{T}) \cup (\text{T} \times \text{P}) \rightarrow \mathbb{N}_0$, $\text{m}_0: \text{P} \rightarrow \mathbb{N}_0$

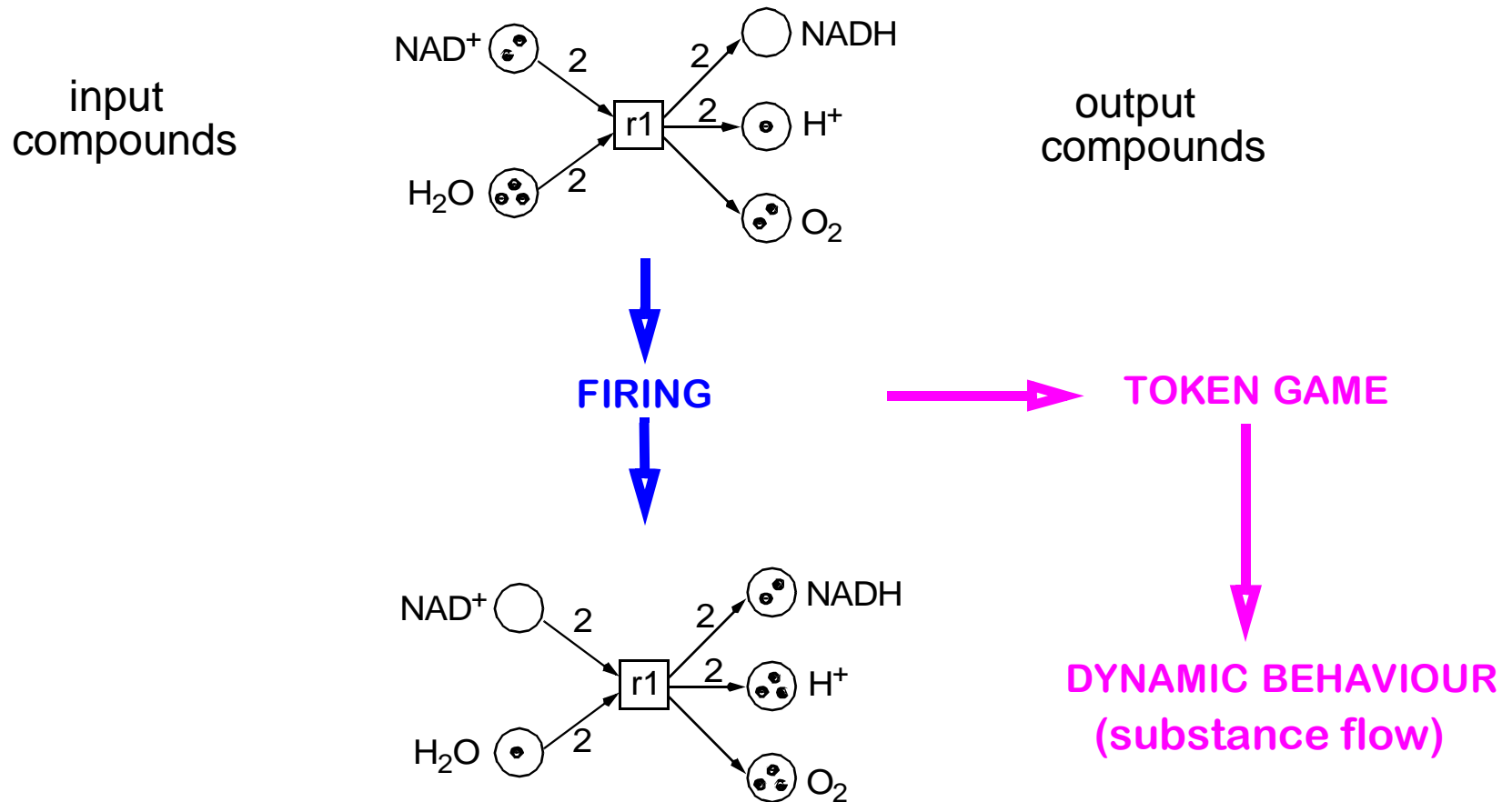
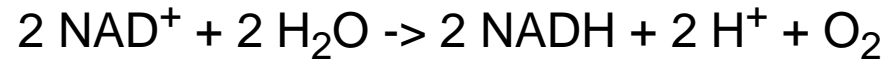
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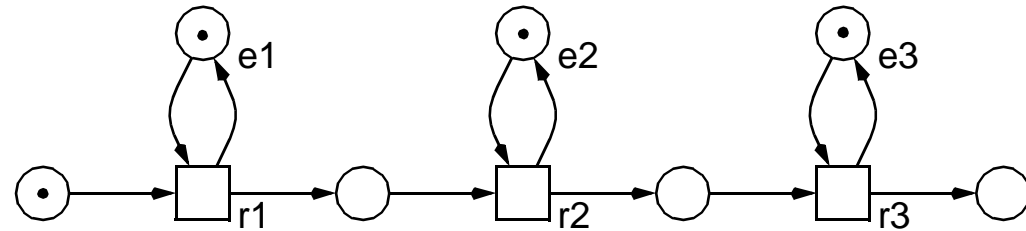


□ atomic actions → Petri net transitions → chemical reactions



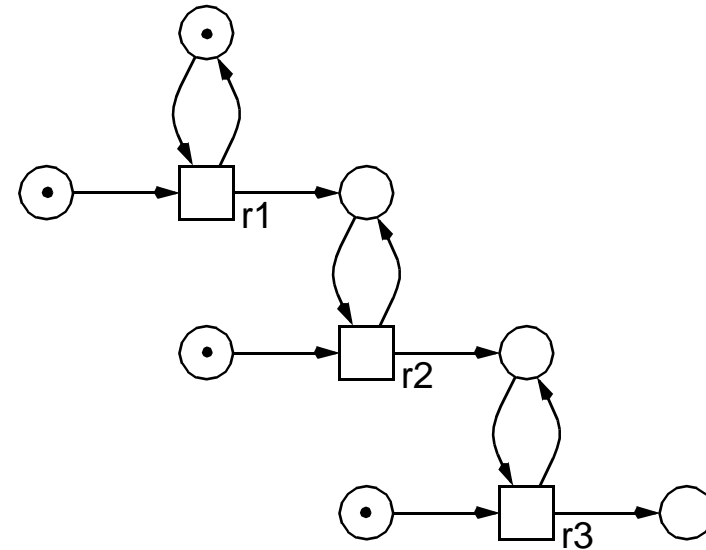
□ metabolic networks

-> *substance flows*

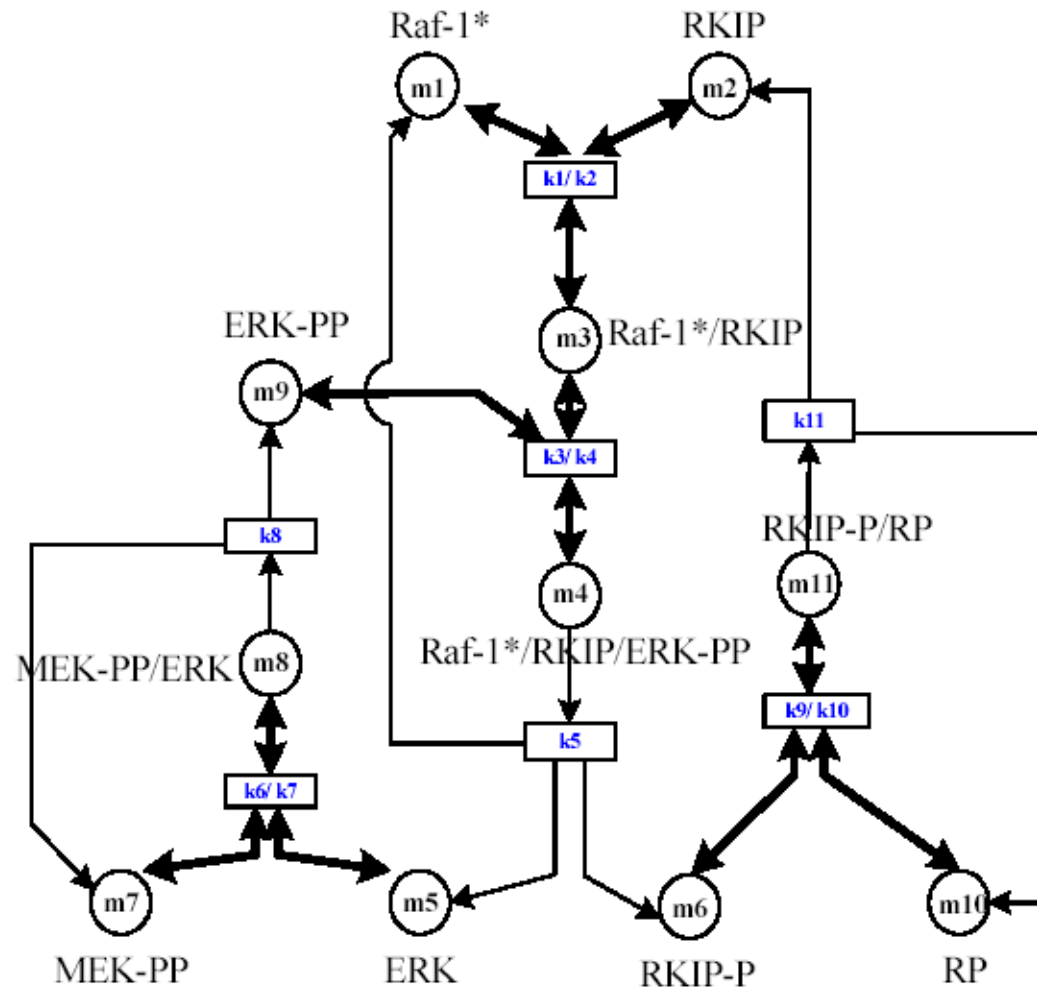


□ signal transduction networks

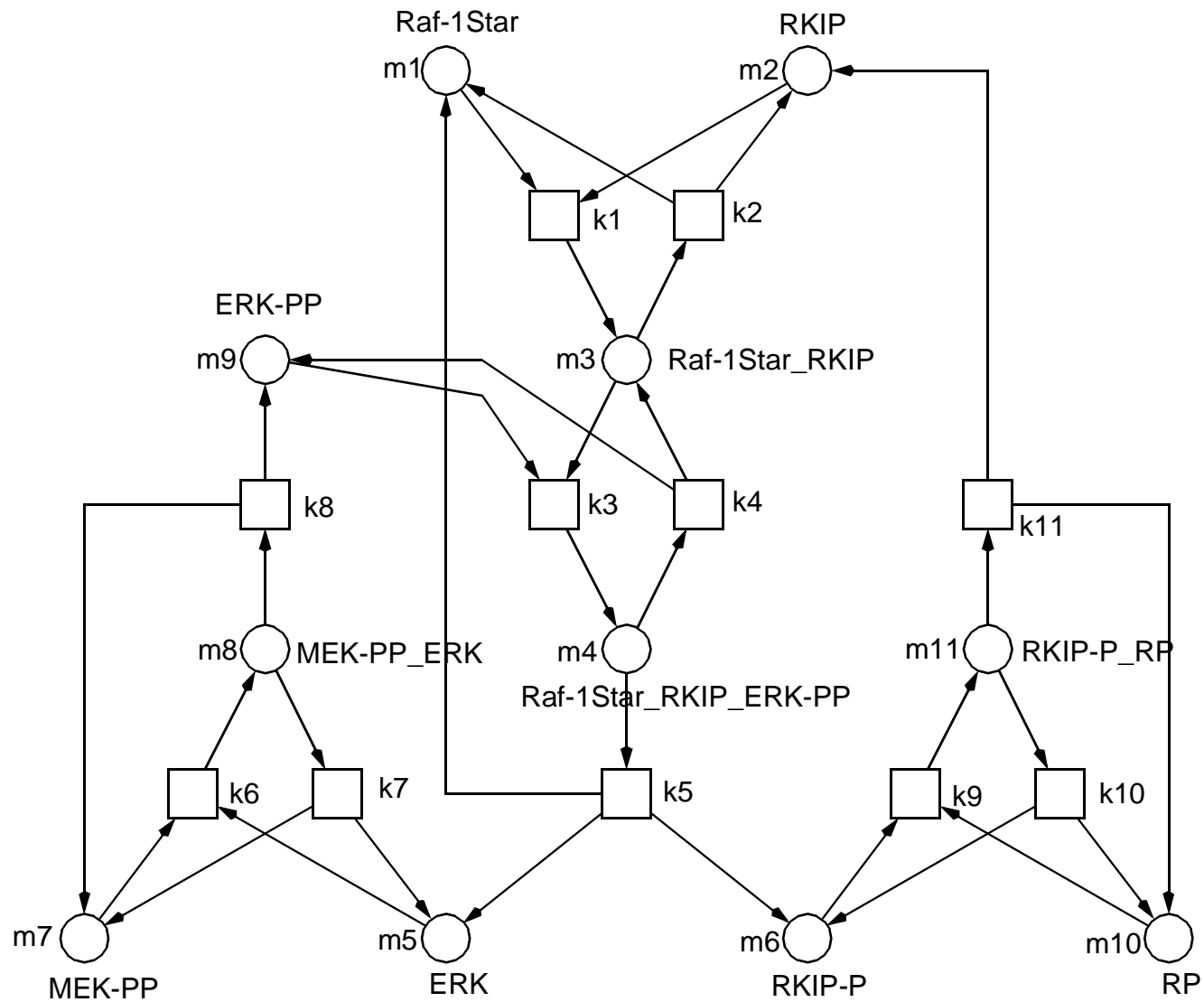
-> *signal flows*



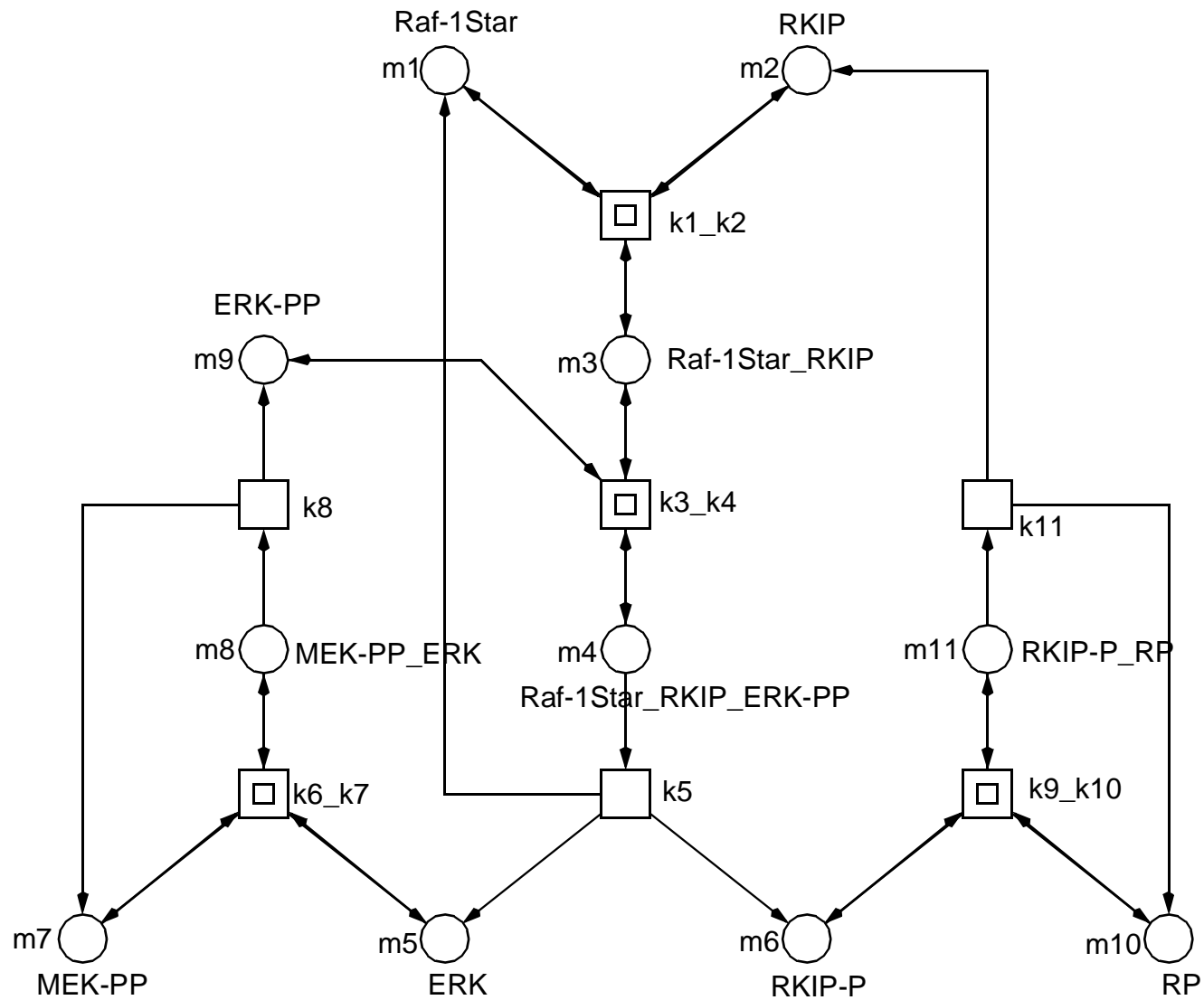
THE RUNNING EXAMPLE - THE RKIP PATHWAY



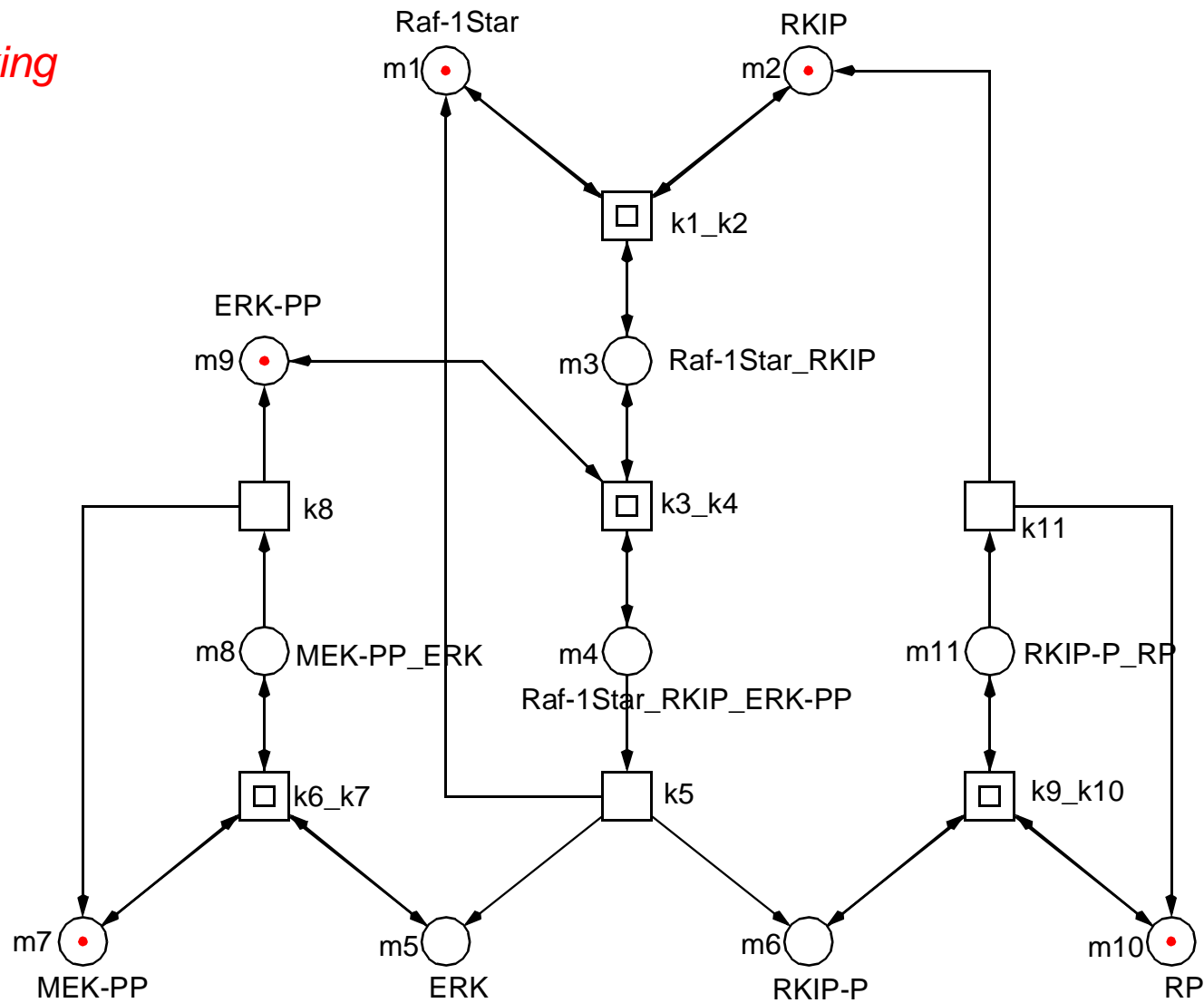
[Cho et al.,
CMSB 2003]



THE RKIP PATHWAY, HIERARCHICAL PETRI NET



initial marking



❑ biochemical networks

-> *networks of (abstract) chemical reactions*

❑ biochemically interpreted Petri net

-> *partial order sequences of chemical reactions (= elementary actions)
transforming input into output compounds / signals
[respecting the given stoichiometric relations, if any]*

-> *set of all pathways
from the input to the output compounds / signals
[respecting the stoichiometric relations, if any]*

❑ pathway

-> *self-contained partial order sequence of elementary (re-) actions*

❑ typical basic assumption

-> *steady state behaviour*

QUALITATIVE ANALYSES

- ❑ **static analyses** → **no state space construction**
 - > *structural properties (graph theory)*
 - > *P / T - invariants (linear algebra),*

- ❑ **dynamic analyses** → **total/partial state space construction (RG)**
 - > *analysis of **general** behavioural system properties,*
e.g. boundedness, liveness, reversibility, . . .

 - > *model checking of **special** behavioural system properties,*
e.g. reachability of a given (sub-) system state [with constraints],
reproducibility of a given (sub-) system state [with constraints]

 - expressed in temporal logics (CTL / LTL),*
very flexible, powerful query language

- a representation of the net structure

=> stoichiometric matrix

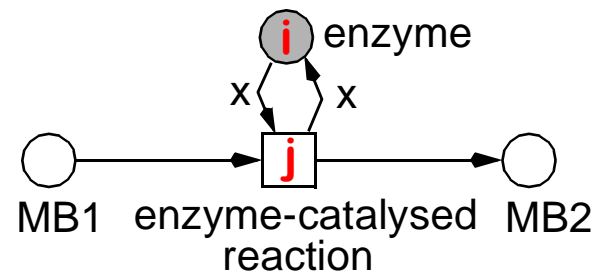
$$C =$$

P \ T	t1	...	tj	...	tm
p1					
pi			cij		
⋮			Δtj		
pn					

$$c_{ij} = (p_i, t_j) = F(t_j, p_i) - F(p_i, t_j) = \Delta t_j(p_i)$$

$$\Delta t_j = \Delta t_j(^*)$$

- matrix entry c_{ij} :
token change in place p_i by firing of transition t_j
- matrix column Δt_j :
vector describing the change of the whole marking by firing of t_j
- side-conditions are neglected



$$c_{ij} = 0$$

□ Lautenbach, 1973

□ T-invariants

-> integer solutions x of

$$Cx = 0, x \neq 0, x \geq 0$$

-> *multisets of transitions*

-> *Parikh vector*

□ minimal T-invariants

-> *there is no T-invariant with a smaller support*

-> *sets of transitions*

-> *gcd of all entries is 1*

□ any T-invariant is a non-negative linear combination of minimal ones

-> *multiplication with a positive integer*

-> *addition*

-> *Division by gcd*

$$kx = \sum_i a_i x_i$$

□ Covered by T-Invariants (CTI)

-> *each transition belongs to a T-invariant*

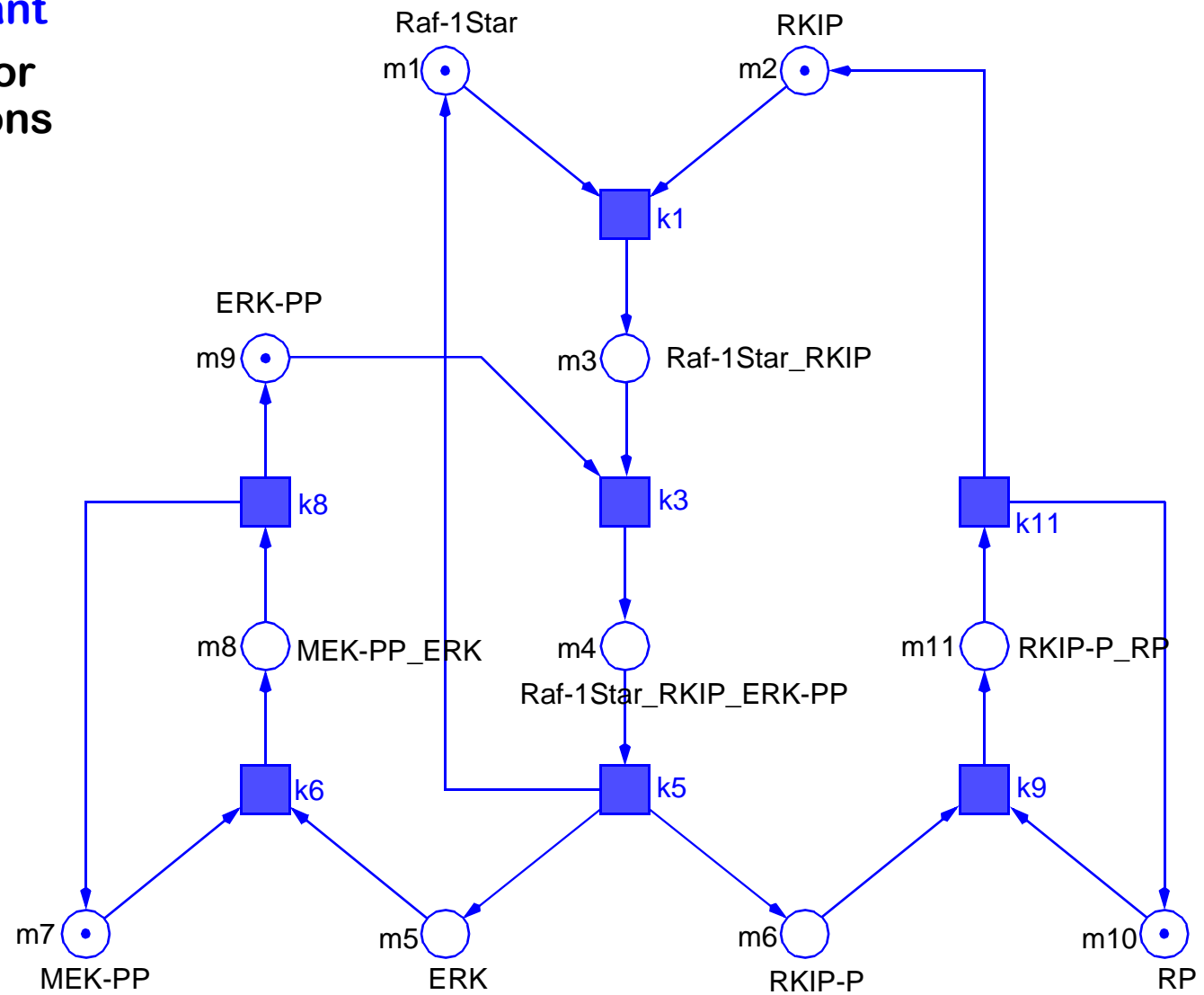
-> *BND & LIVE => CTI (necessary condition)*

- **T-invariants = (multi-) sets of transitions**
 - > *zero effect on marking*
 - > *reproducing a marking / system state*
 - > *steady state substance flows / reaction rates*
 - > *elementary modes [Schuster 1993]*

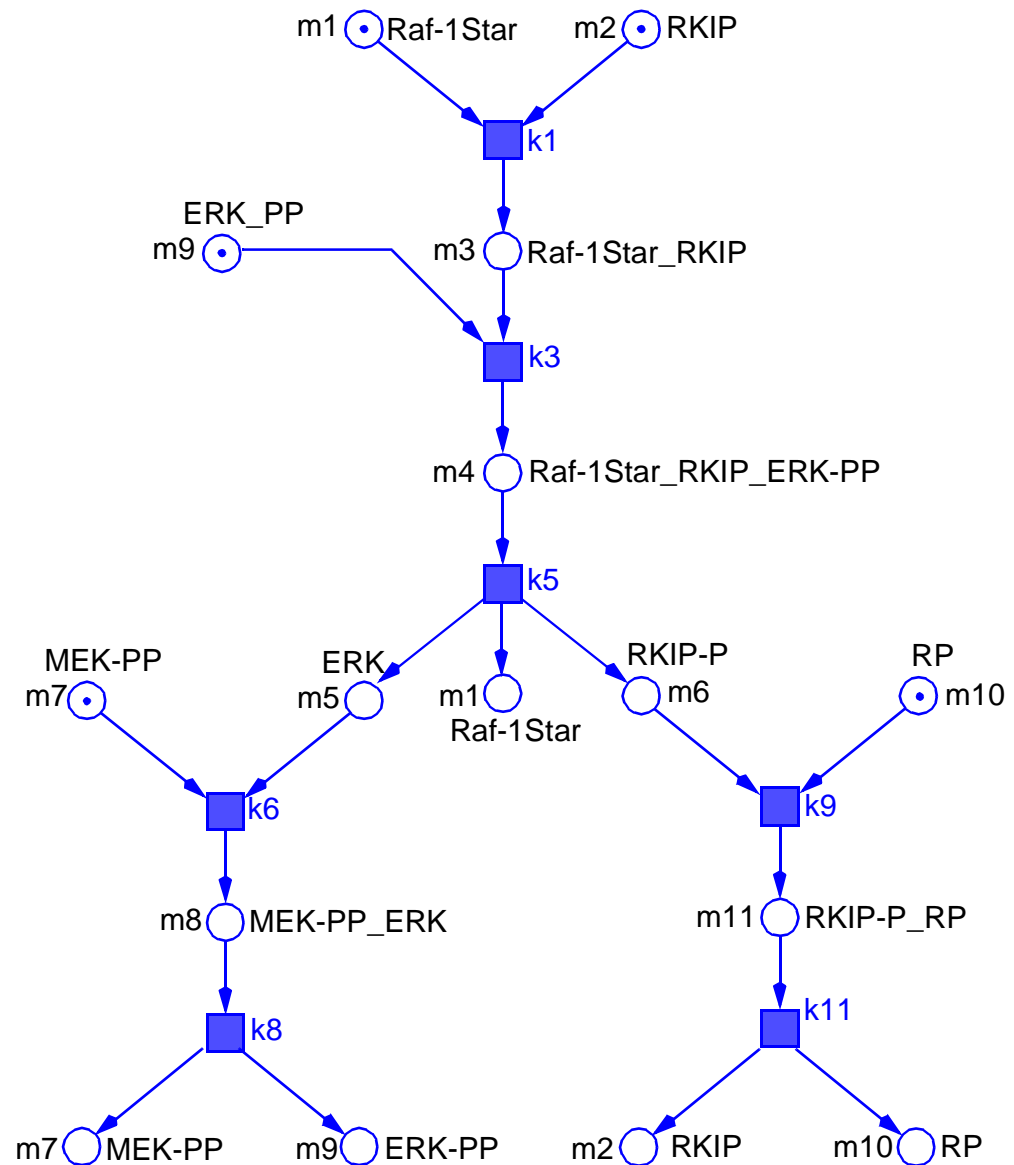
- **realizable T-invariants correspond to cycles in the RG**
 - > *RG: concurrent transitions -> all transitions' interleaving sequences*
 - > *if there are concurrent transitions in a realizable T-invariant, then there is a RG cycle for each interleaving sequence*
 - > *analogously for conflicts*

- **a T-invariant defines a subnet** **-> partial order structure**
 - > *the T-invariant's transitions (the support),*
 - + *all their pre- and post-places*
 - + *the arcs in between*
 - > *pre-sets of supports = post-sets of supports*

-> non-trivial T-invariant
 + four trivial ones for reversible reactions



- **partial order structure**
- **T-invariant's unfolding to describe its behaviour**
- **labelled condition / event net**
 - > *events*
 - *transition occurrences*
 - > *conditions*
 - *input / output compounds*
- **partial order semantics**
 - > *a net's all partial order runs*



□ Lautenbach, 1973

□ P-invariants

-> integer solutions y of

$$yC = 0, y \neq 0, y \geq 0$$

-> multisets of places

□ minimal P-invariants

-> there is no P-invariant with a smaller support

-> sets of places

-> gcd of all entries is 1

□ any P-invariant is a non-negative linear combination of minimal ones

-> multiplication with a positive integer

-> addition

-> Division by gcd

$$ky = \sum_i a_i y_i$$

□ Covered by P-Invariants (CPI)

-> each transition belongs to a P-invariant

-> CPI \Rightarrow BND (sufficient condition)

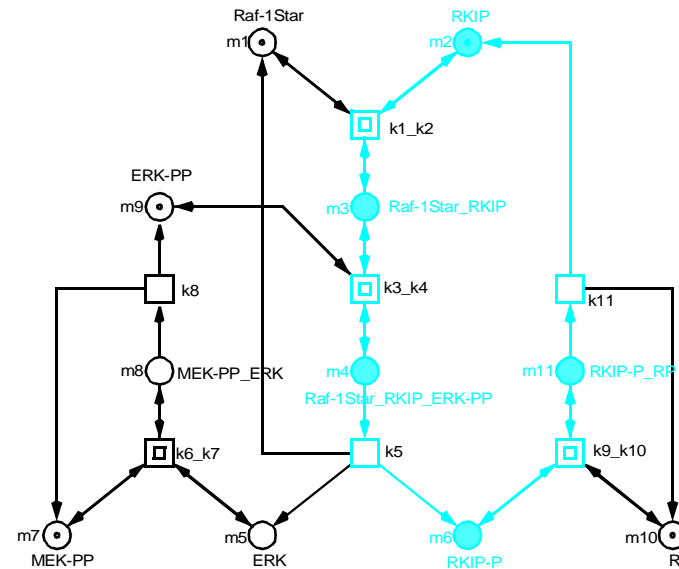
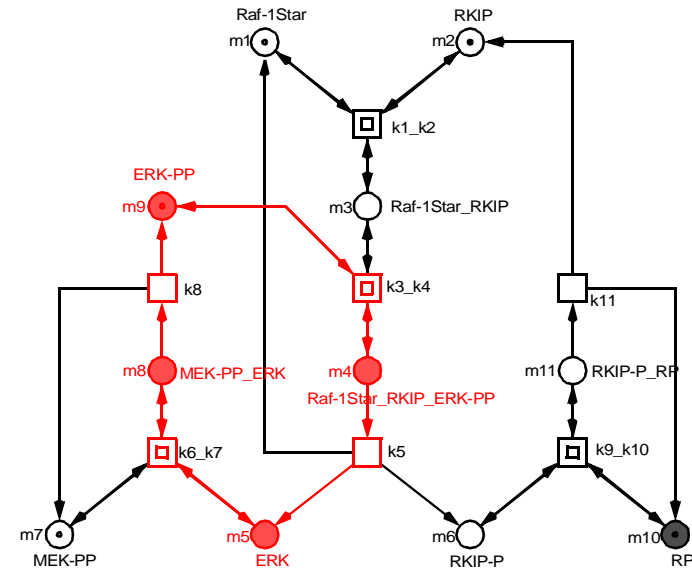
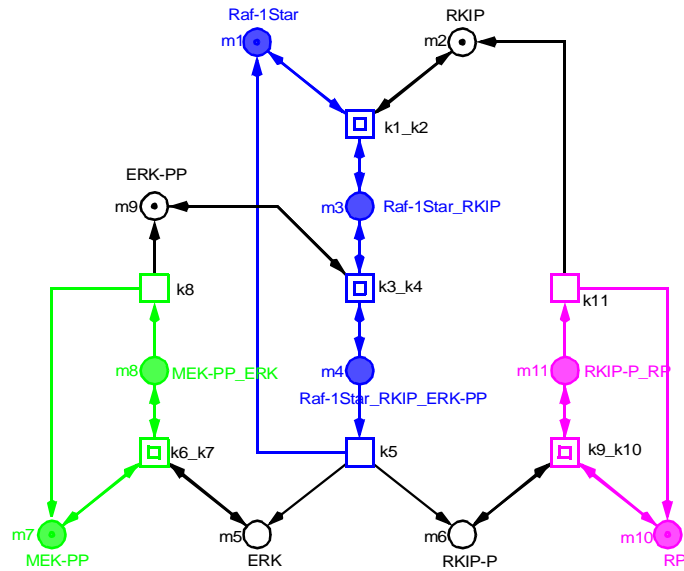
- the firing of any transition has no influence on the weighted sum of tokens on the P-invariant's places
 - > for all t : the effect of the arcs, removing tokens from a P-invariant's place is equal to the effect of the arcs, adding tokens to a P-invariant's place

- set of places with
 - > a constant weighted sum of tokens for all markings m reachable from m_0
$$ym = ym_0$$
 - > token / compound preservation

- a place belonging to a P-invariant is bounded
 - > CPI - sufficient condition for BND

- a P-invariant defines a subnet
 - > the P-invariant's places (the support),
+ all their pre- and post-transitions
+ the arcs in between
 - > pre-sets of supports = post-sets of supports

P-INVARIANTS, THE RKIP PATHWAY



P-INV1: MEK

P-INV2: RAF-1STAR

P-INV3: RP

P-INV4: ERK

P-INV5: RKIP

- ❑ each P-invariant gets at least one token
 - > *P-invariants are structural deadlocks and traps*

- ❑ all (non-trivial) T-invariants get realizable
 - > *to make the net live*

- ❑ minimal marking
 - > *minimization of the state space*

- ❑ assumption: top-to-bottom reading of the figure
 - > *but, all reachable markings are equivalent*
(= produce same state space)

-> UNIQUE INITIAL MARKING

□ structural properties

INA

ORD	HOM	NBM	PUR	CSV	SCF	CON	SC	Ft0	tF0	Fp0	pF0	MG	SM	FC	EFC	ES
Y	Y	Y	Y	N	N	Y	Y	N	N	N	N	N	N	N	N	Y
DTP	CPI	CTI	B	SB	REV	DSt	BSt	DTr	DCF	L	LV	L&S				
Y	Y	Y	Y	Y	Y	N	?	N	N	Y	Y	Y				

□ CPI

-> *structural bounded (SB)*

-> *each P-invariant represents a substance conservation subnet (cycle)*

□ CTI

-> *Live & BND -> CTI*

-> *4 trivial T-invariants for reversible reactions*

-> *1 non-trivial T-invariant describing the essential cyclic behaviour*

□ DTP & ES -> Live

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

- ❑ **unbounded Petri net -> infinite RG**
bounded Petri net -> finite RG

- ❑ **concurrency**
 - > *enumeration of all interleaving sequences* -> *interleaving semantics*

- ❑ **branching arcs in the RG**
 - > *conflict* **OR** *concurrency*

- ❑ **RG tend to be very large**
 - > *automatic evaluation necessary* -> *model checking*

- ❑ **worst case: over-exponential growth**
 - > *alternative analyses techniques ?*

❑ property 1

Is a given (sub-) marking (system state) reachable ?

$EF (ERK * RP);$

❑ property 2

Liveness of transition k8 ?

$AG EF (MEK-PP_ERK);$

❑ property 3

Is it possible to produce ERK-PP neither creating nor using MEK-PP ?

$E (! MEK-PP \ U ERK-PP);$

❑ property 4

Is there cyclic behaviour w.r.t. the presence / absence of RKIP ?

$EG ((RKIP \rightarrow EF (! RKIP)) * (! RKIP \rightarrow EF (RKIP)));$

❑ validation criterion 0

- > *all expected structural properties hold*
- > *all expected general behavioural properties hold*

❑ validation criterion 1

- > *CTI*
- > *no minimal T-invariant without biological interpretation*
- > *no known biological behaviour without corresponding T-invariant*

❑ validation criterion 2

- > *CPI*
- > *no minimal P-invariant without biological interpretation (?)*

❑ validation criterion 3

- > *all expected special behavioural properties hold*
- > *temporal-logic properties -> TRUE*

**NOW WE ARE READY
FOR SOPHISTICATED
QUANTITATIVE ANALYSES !**

- quantitative model = qualitative model + quantitative parameters

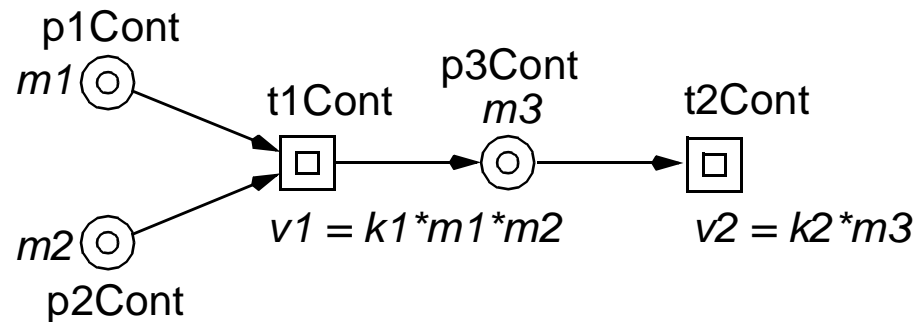
-> *BUT: quantitative parameters often unknown*

- typical quantitative parameters of bionetworks

-> *compound concentrations* -> *real numbers*

-> *reaction rates / fluxes* -> *concentration-dependent*

- continuous Petri nets

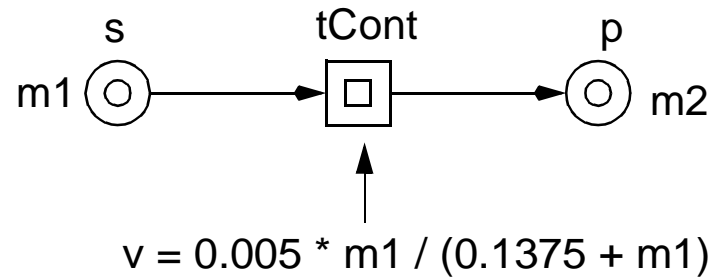


continuous nodes !

$$\left. \begin{aligned} d [p1Cont] / dt &= d [p2Cont] / dt = - v1 \\ d [p3Cont] / dt &= v1 - v2 \end{aligned} \right\}$$

ODEs

EXAMPLE - MICHAELIS-MENTEN REACTION



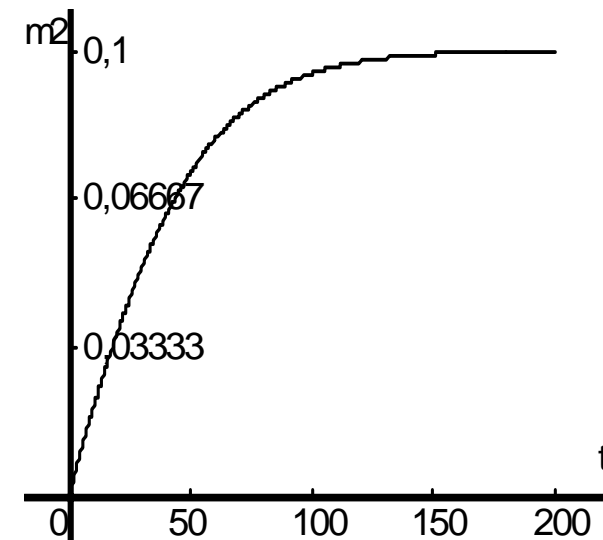
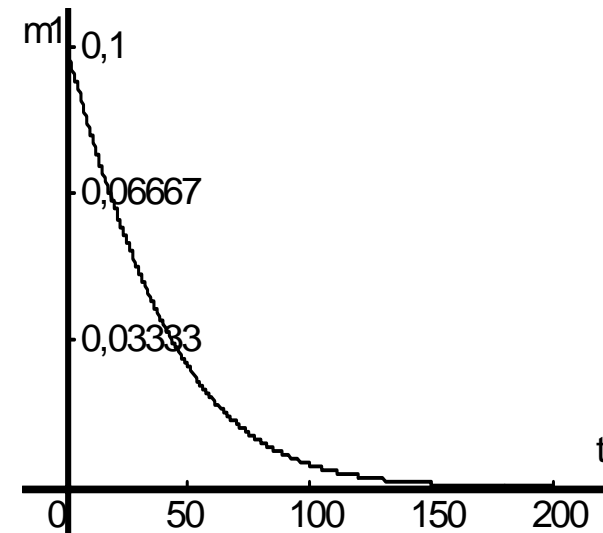
$V_{max} = 0.005$ (maximal reaction rate)

$K_m = 0.1375$ (Michaelis constant)

$$d[s]/dt = d[p]/dt = V_{max} * [s] / (K_m + [s])$$

$$dm1/dt = dm2/dt = V_{max} * m1 / (K_m + m1)$$

- > Visual Object Nets
- > GON / cell illustrator (?)



**THE QUALITATIVE MODEL
BECOMES
THE STRUCTURAL DESCRIPTION
OF THE QUANTITATIVE MODEL !**

❑ extensions

-> *read arcs*

-> *interleaving / partial order semantics*

-> *inhibitor arcs !?*

-> *Turing power !*

❑ efficient computation of minimal invariants

-> *exponential complexity*

-> *compositional / step-wise refinement approach (under development)*

❑ analysis of unbounded nets

-> *besides T-invariant analysis ?*

❑ model checking

-> *relevant properties ?*

❑ comparison: continuous / hybrid Petri nets \leftrightarrow ODEs

-> *Petri net simulation versus classical ODEs solver*

-> *is there a winner (for certain structures) ?*

❑ representation of bionetworks by Petri nets

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

-> *various sound analysis techniques*

❑ purposes

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

❑ two-step model development

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

❑ many challenging questions for analysis techniques

- > *qualitative as well as quantitative ones*

THANKS !