

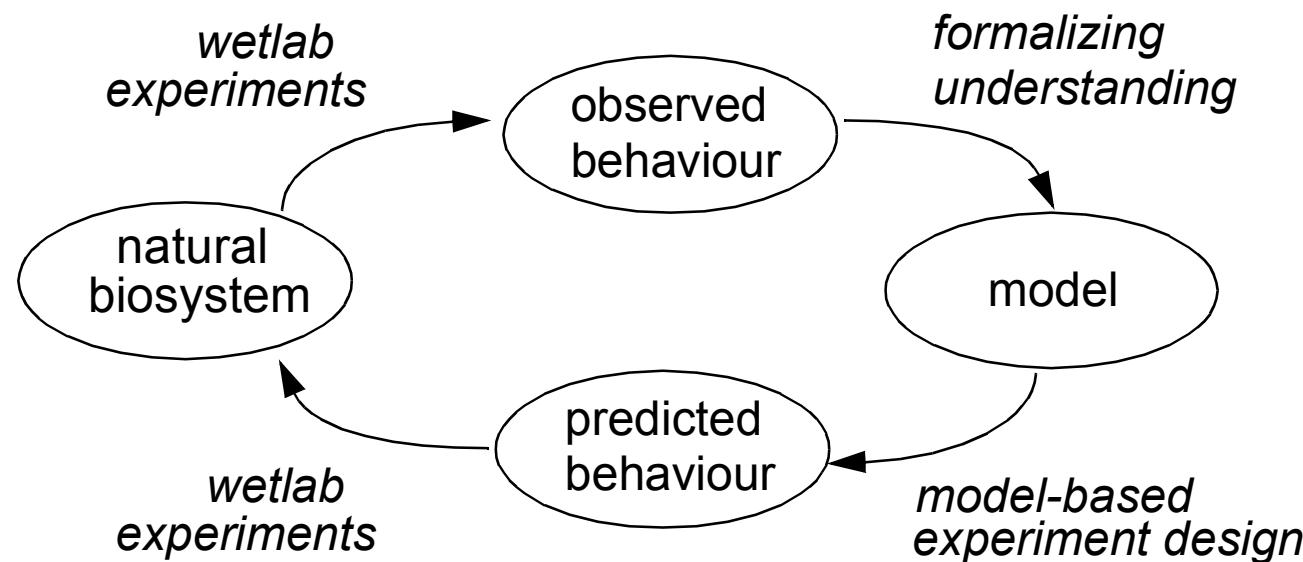
## **PART I - TALK 2**

# **PN-BASED ANALYSIS OF BIOCHEMICAL NETWORKS**

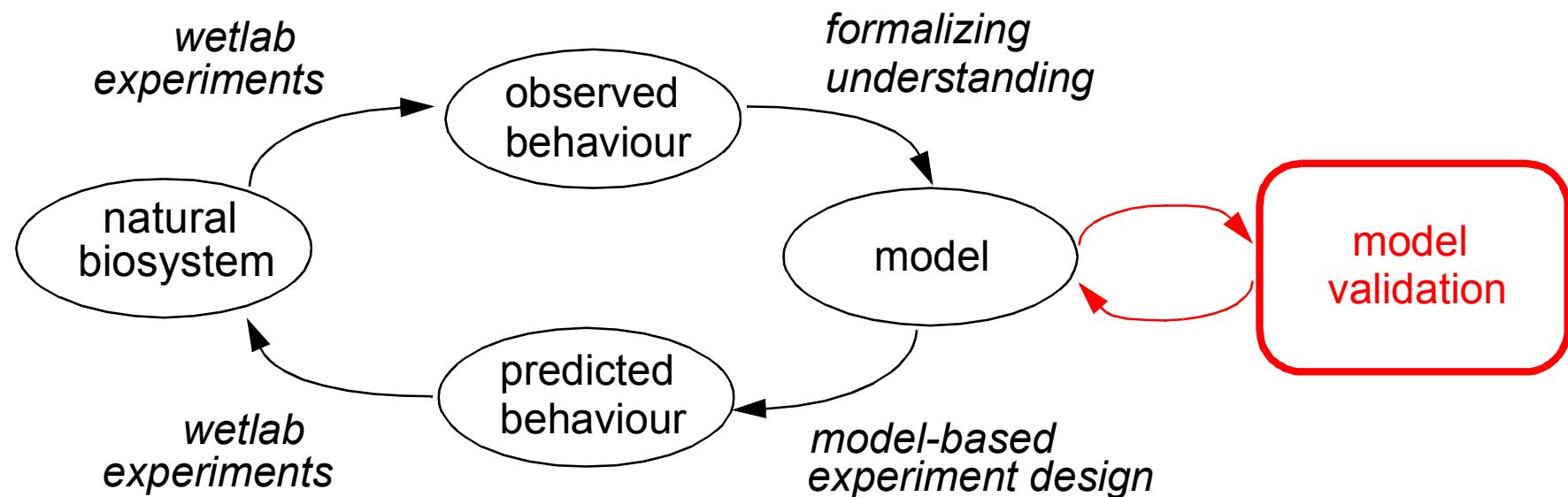
**Monika Heiner**

**Brunel University, on sabbatical leave from  
Brandenburg University of Technology Cottbus**

## MODELLING = FORMAL KNOWLEDGE REPRESENTATION



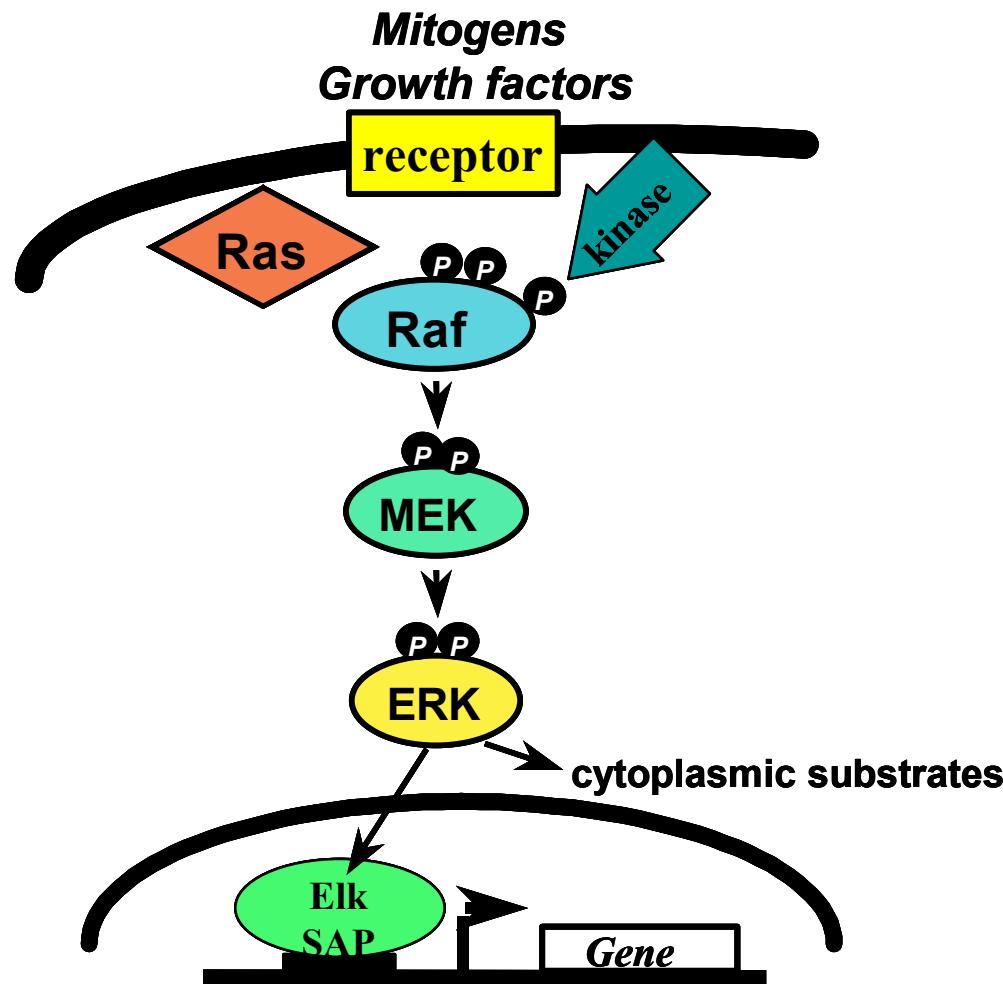
**MODELLING = FORMAL KNOWLEDGE REPRESENTATION**

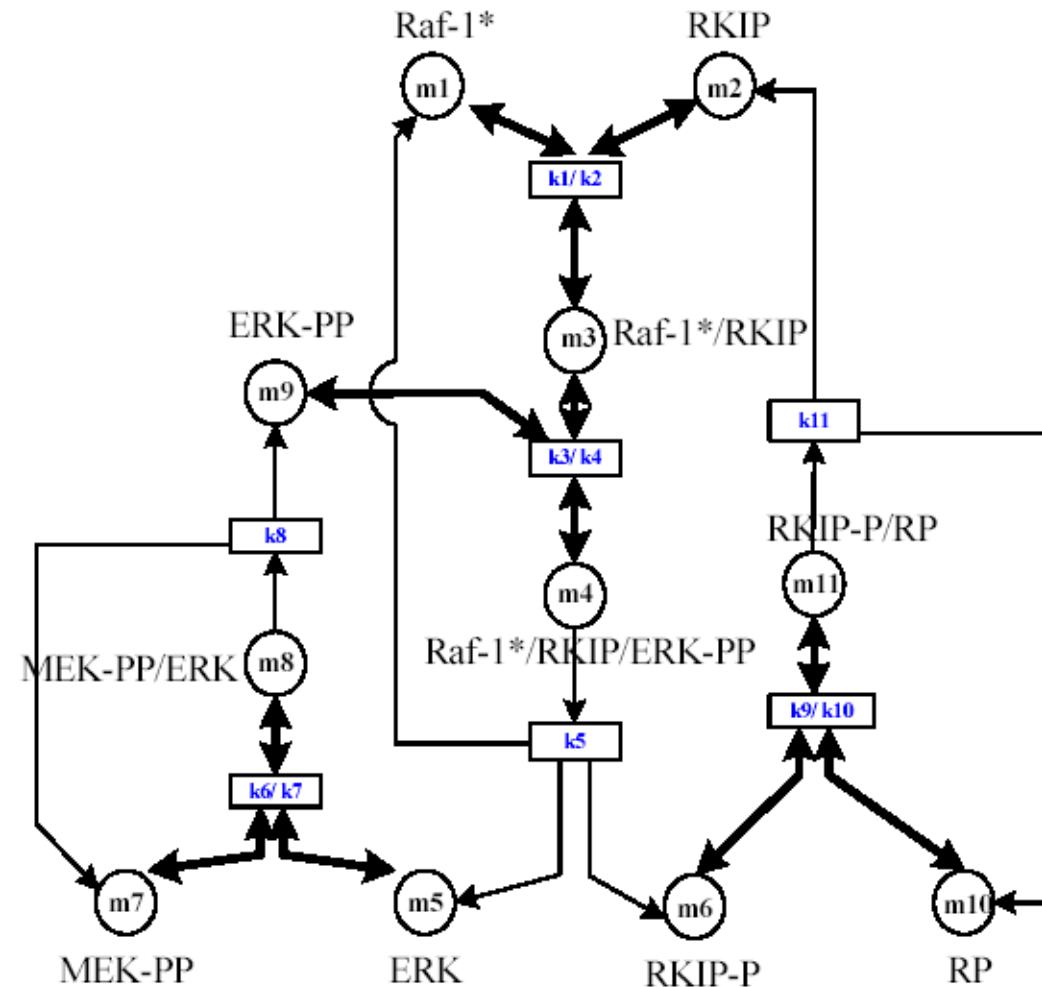


**MODEL VALIDATION = CONFIDENCE INCREASE**

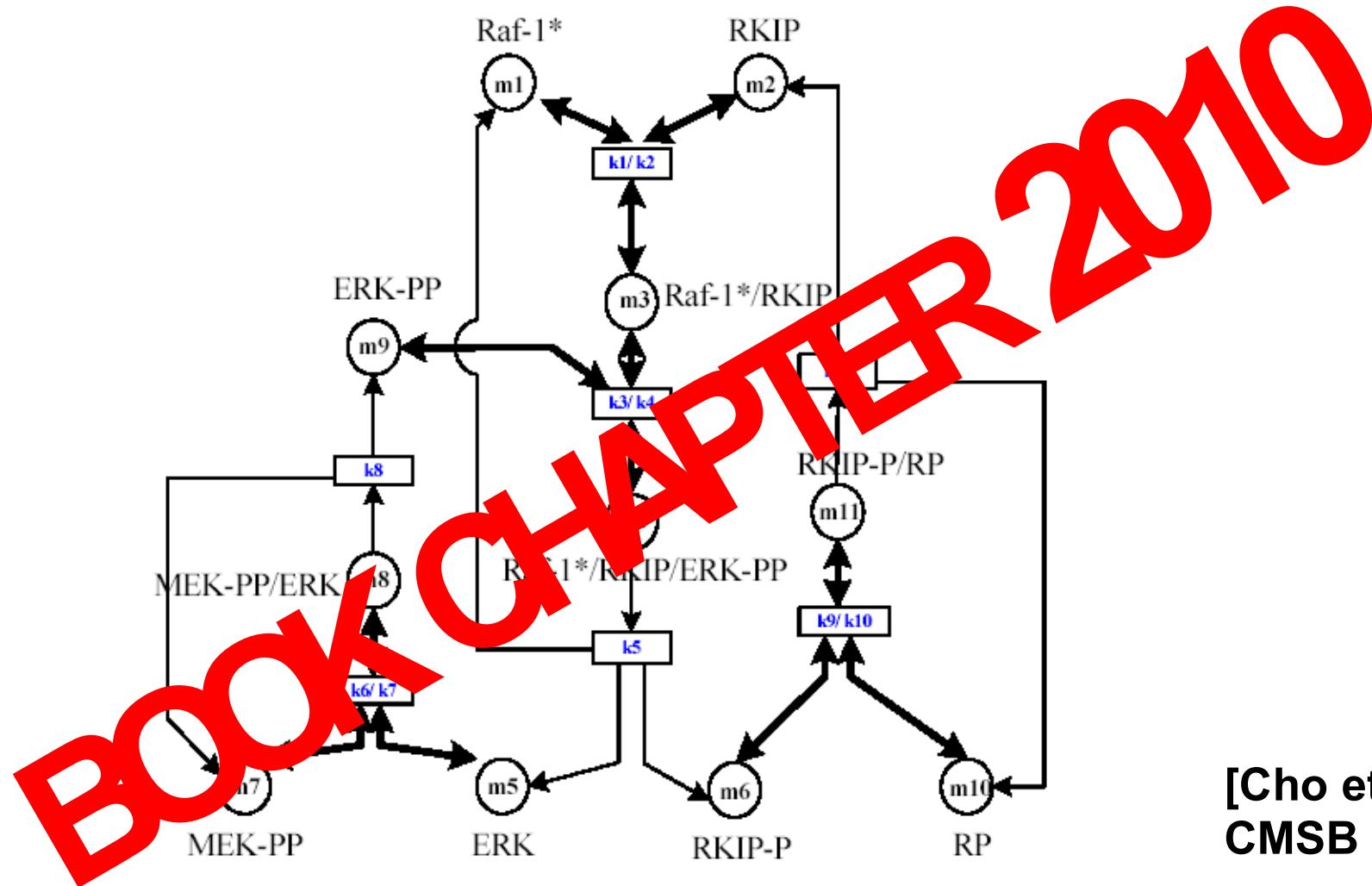
# A CASE STUDY

...one pathway...





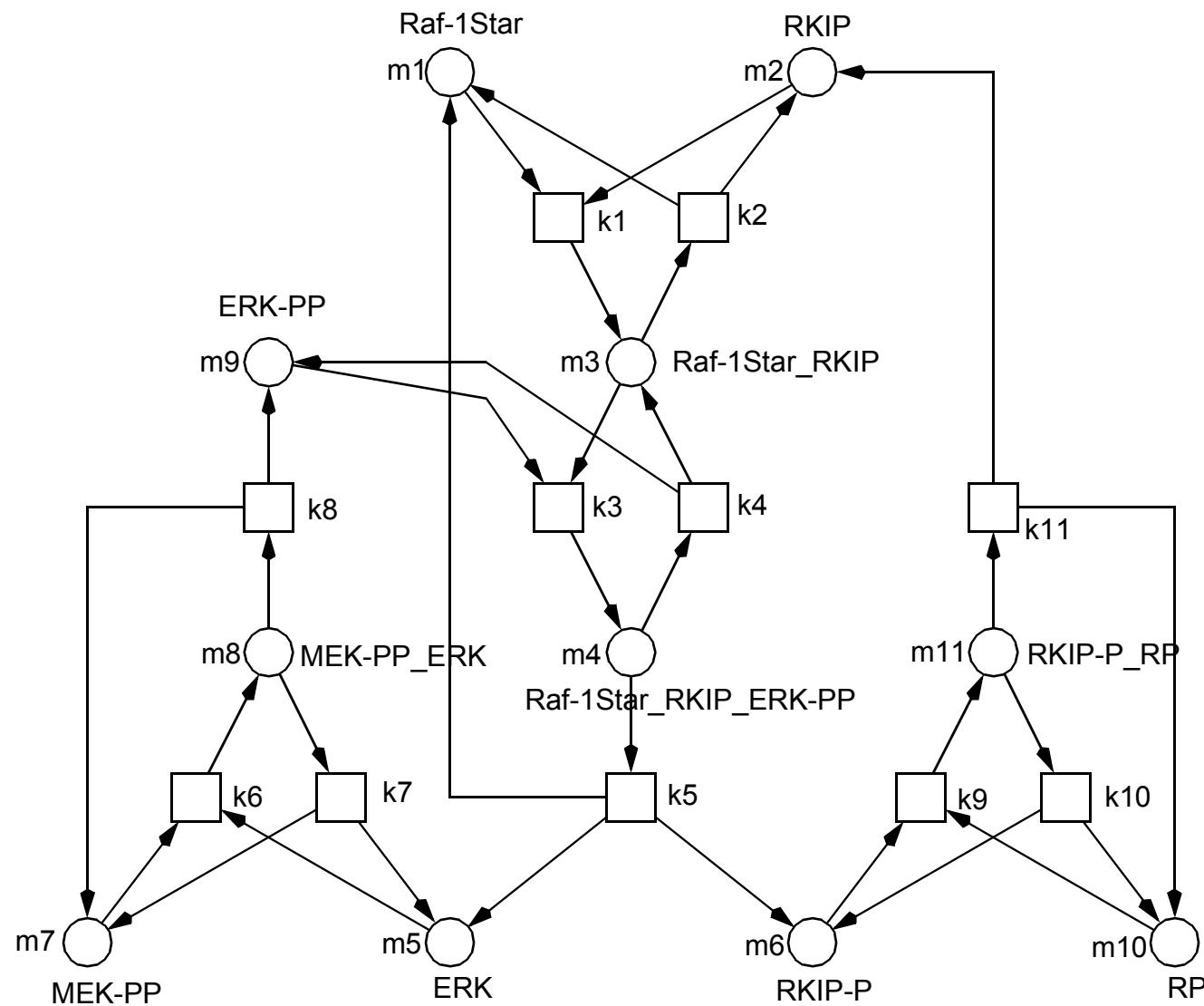
[Cho et al.,  
CMSB 2003]



[Cho et al.,  
CMSB 2003]

# RKIP PATHWAY, PETRI NET

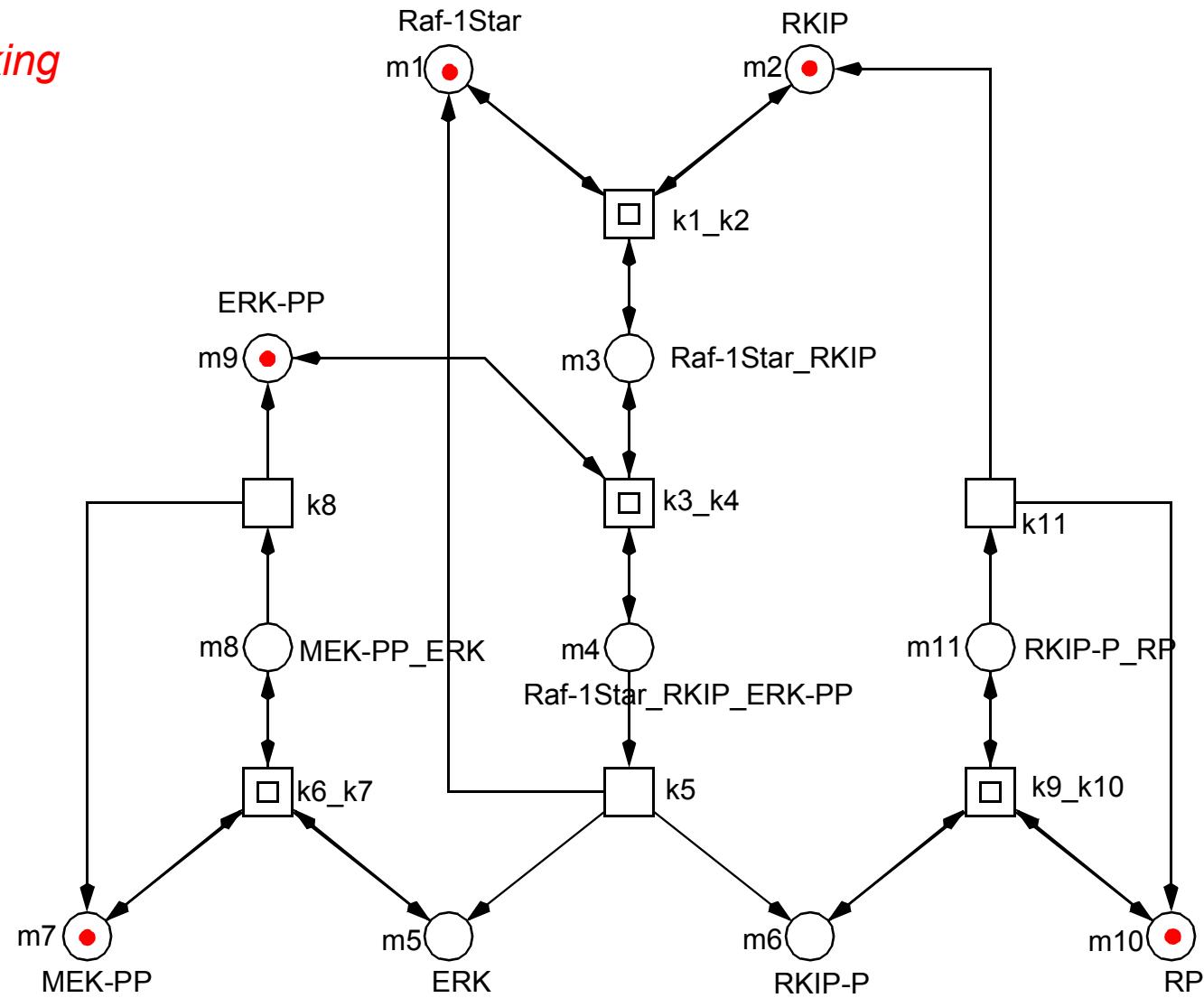
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# RKIP PATHWAY, HIERARCHICAL PETRI NET

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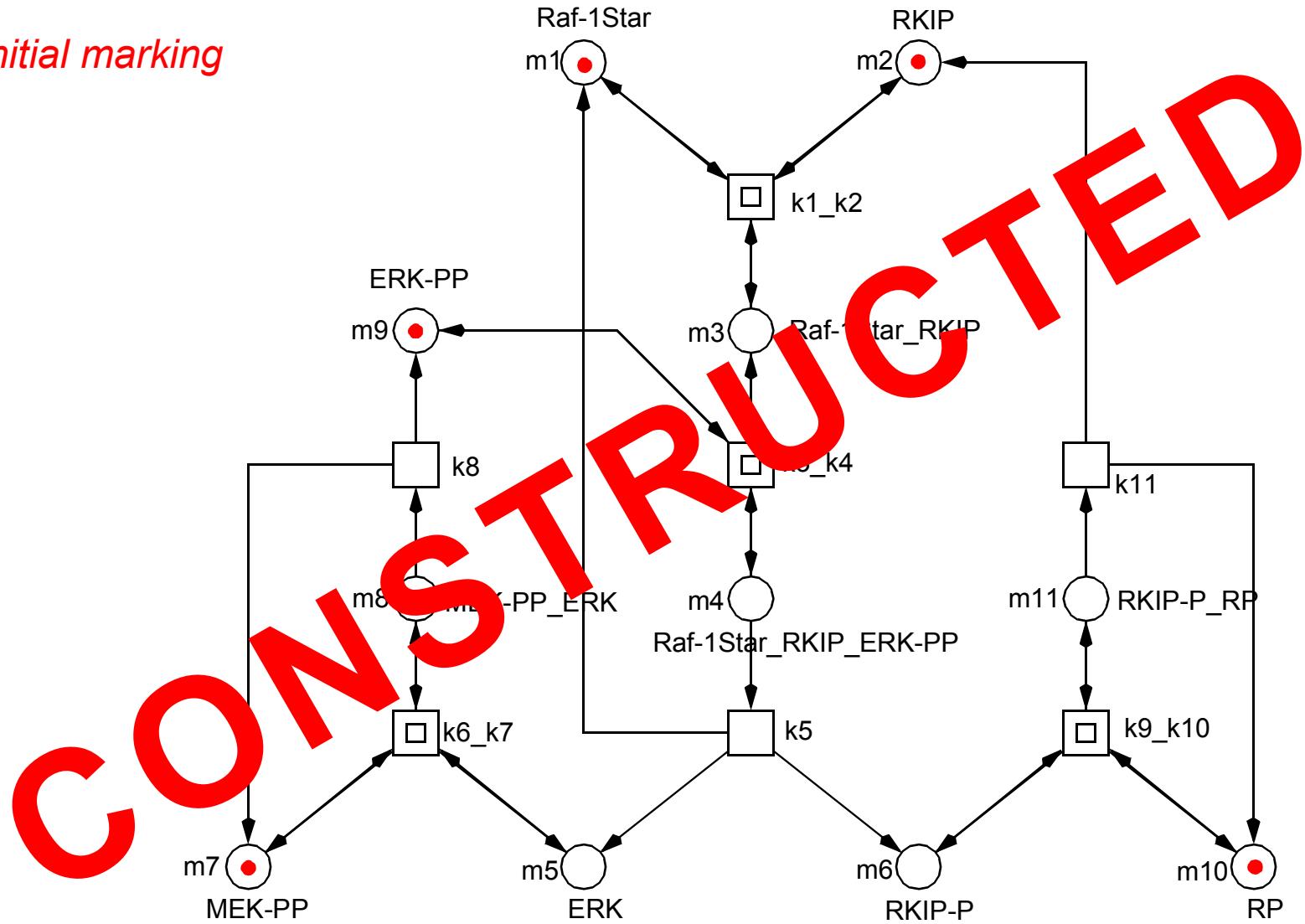
*initial marking*



# RKIP PATHWAY, HIERARCHICAL PETRI NET

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initial marking



# **Petri nets, typical properties**

□ How many tokens can reside at most in a given place ?

->  $(0, 1, k, \infty)$

-> **BOUNDEDNESS**

□ How many tokens can reside at most in a given place ?

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□ How often can a transition fire ?

->  $(0\text{-times}, n\text{-times}, \infty\text{-times})$  -> **LIVENESS**

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□ How often can a system state be reached ?

-> *never* -> **UNREACHABLE** -> **SAFETY PROPERTIES**

-> *n-times* -> **REPRODUCIBLE**

-> *always reachable again* -> **REVERSIBLE (HOME STATE)**

-> *reversible initial state* -> **REVERSIBILITY**

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□ Are there behaviourally invariant subnet structures ?

-> *token conservation* -> **P - INVARIANTS**

-> *token distribution reproduction* -> **T - INVARIANTS**

□ ... and many more -> temporal logics (CTL, LTL)

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# Petri nets, typical analysis techniques

*MODEL ANIMATION (?)*

- static analyses**      -> no state space construction
  
- dynamic analyses**      -> total/ partial state space construction

- ❑ static analyses → no state space construction
    - > structural properties (graph theory)
    - >  $P / T$  - invariants (linear algebra)
  - ❑ dynamic analyses → total/ partial state space construction



# **Dynamic analyses**

## Ex1 - RKİP, REACHABILITY GRAPH (STS)

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- simple algorithm**
- nodes : system states**

# Ex1 - RKİP, REACHABILITY GRAPH (STS)

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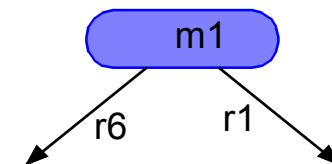
- simple algorithm**
- nodes : system states**

m1

## Ex1 - RKİP, REACHABILITY GRAPH (STS)

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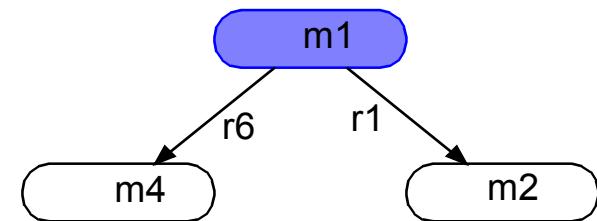
- simple algorithm
- nodes : system states
- single step firing rule
- arcs : the (single) firing transition



## Ex1 - RKİP, REACHABILITY GRAPH (STS)

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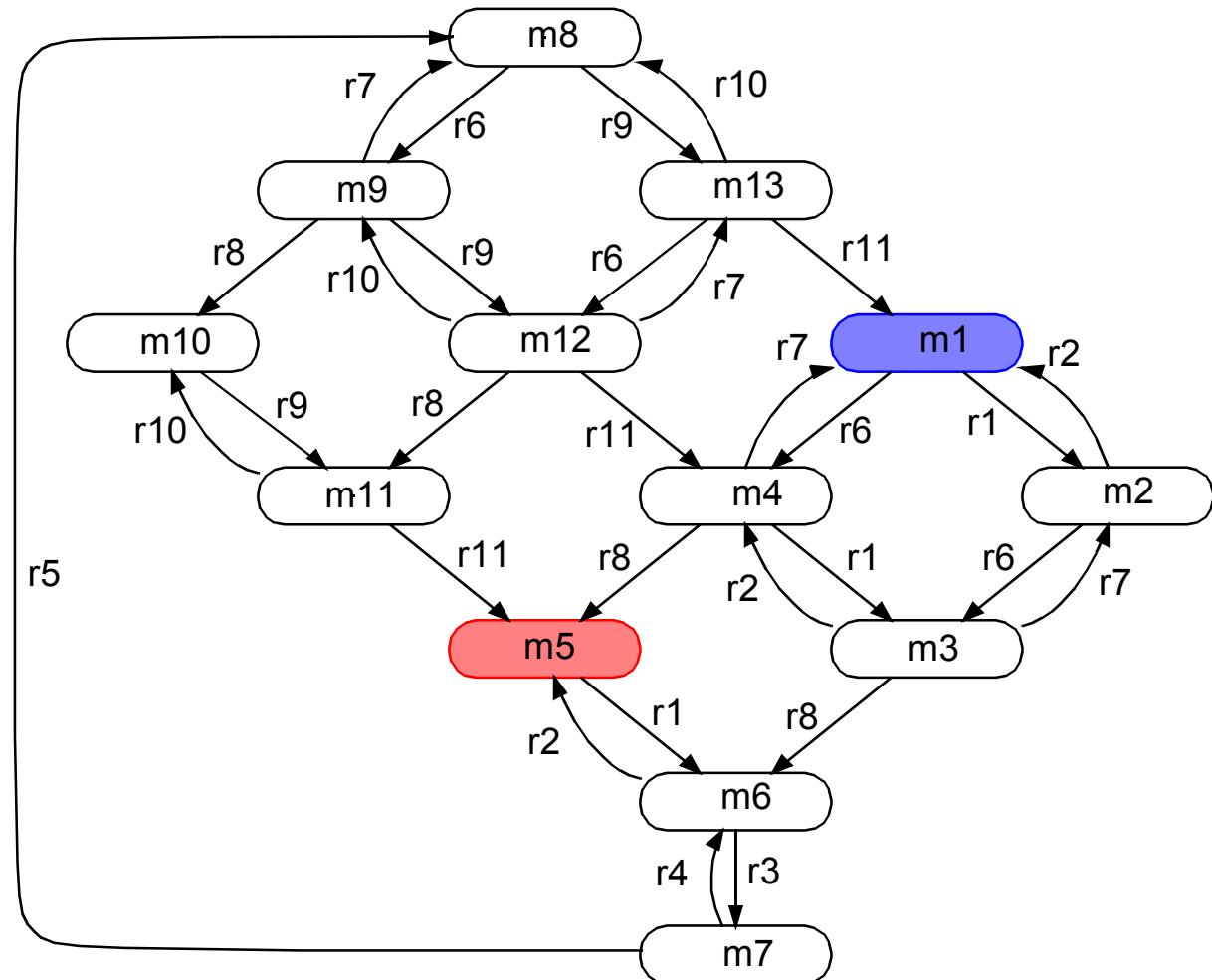
- simple algorithm
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# Ex1 - R<sub>KIP</sub>, REACHABILITY GRAPH (STS)

PN & Systems Biology

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



- **reachability graph (*occurrence / marking graph*)**
  - > *(labelled) state transition system* (-> graph)
  - > *Kripke structure, CTMC, . . .*
- **interleaving semantics**
  - > *(sequential) finite automaton*
  - > *concurrency == enumerating all interleaving sequences*
- **reachability graph construction - simple algorithm**
  - > *depth-first construction*
  - > *breadth-first construction*
- **efficiency depends on efficiency of data structures**

- **boundedness**

-> *finite graph*

- **reversibility**

-> *one Strongly Connected Component (SCC)*

- **dead states (deadlock)**

-> *terminal nodes*

- **liveness**

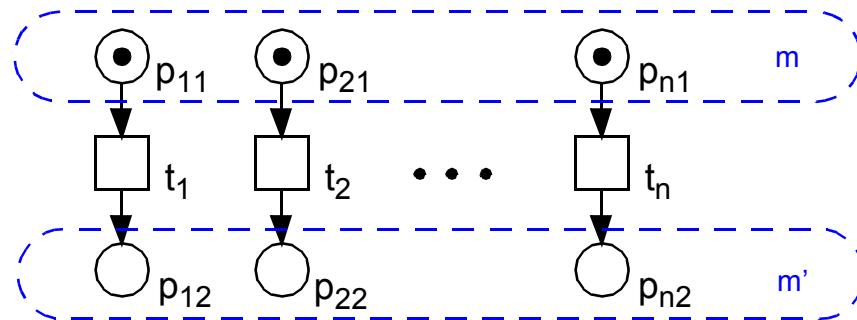
-> *every transition contained in all terminal SCC*

- **boundedness**  
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- **reversibility**  
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**-> reachability graphs tend to be huge <-**

# STATE SPACE COMPLEXITY, CAUSES

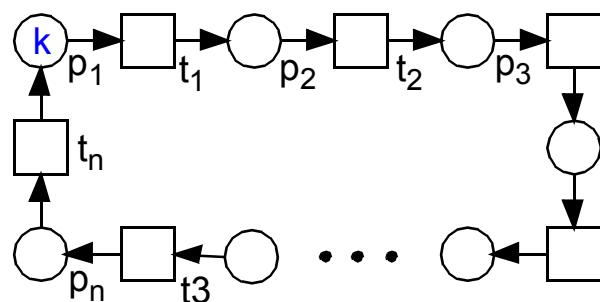
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$n!$  interleaving sequences

$m \rightarrow m'$

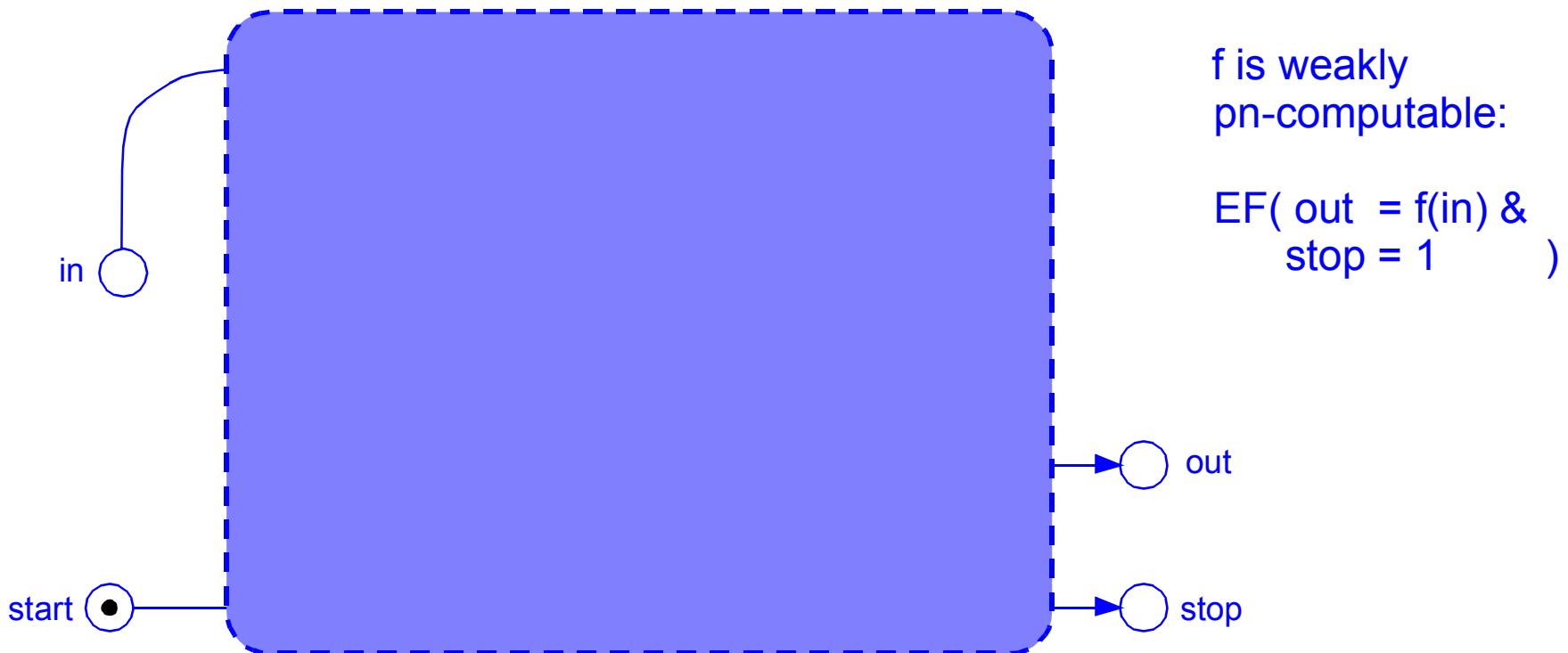
$2^n - 2$  intermediate states



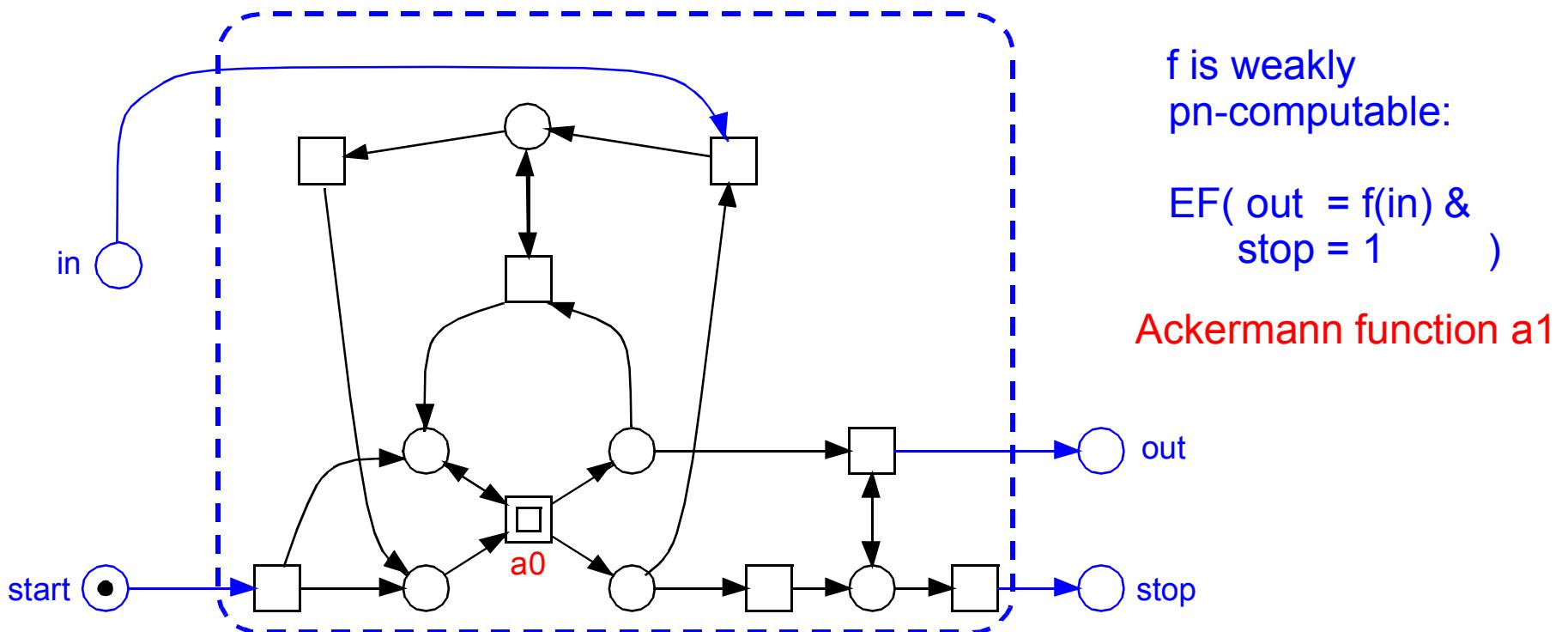
$\frac{(n + k - 1)!}{(n - 1)! k!}$  states

(combination with repetition)

- infinite for unbounded nets
- worst-case for finite state spaces [Priese, Wimmel 2003]  
... *cannot be bounded by a primitive recursive function* ...
- proof -> Petri net computer for a function  $f: \mathbb{N}_0^m \rightarrow \mathbb{N}_0$



- infinite for unbounded nets
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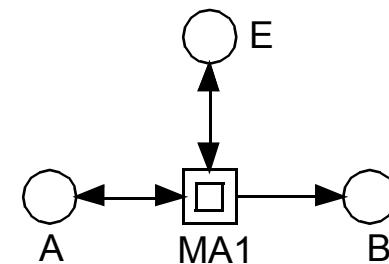
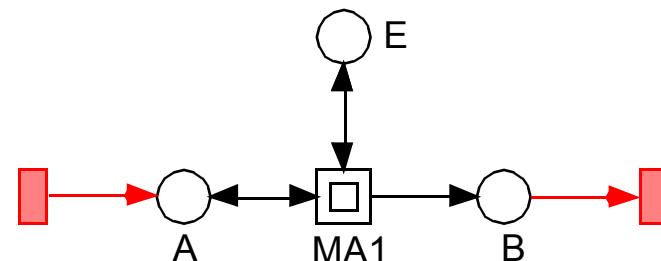
# **STATIC ANALYSES**

- no (full / partial) state space construction
    - > *works also for unbounded models = infinite state spaces*
  - structural analysis
    - > *boundary nodes, . . .*
    - > *Siphon Trap Property*
  - net reduction
- 
- > *to decide liveness  
(hopefully)*

- no (full / partial) state space construction
    - > works also for unbounded models = infinite state spaces
  - structural analysis
    - > boundary nodes, . . .
    - > Siphon Trap Property
  - net reduction
  - invariants
    - > P-invariants - mass-preserving modules
    - > T-invariants - state-repeating modules
  - modularization by T-invariants
    - > abstract dependent transition sets (ADT-sets) define building blocks
  - core network identification
- 
- > to decide liveness  
(hopefully)
- > to decide boundedness

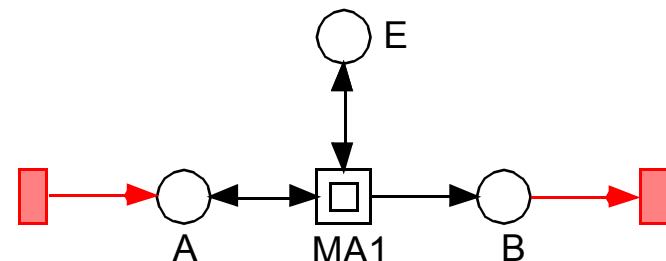
## □ boundary nodes

- > *input transitions* -> not BND
- > *input places* -> not LIVE
- > *LIVE & BND* -> no boundary nodes



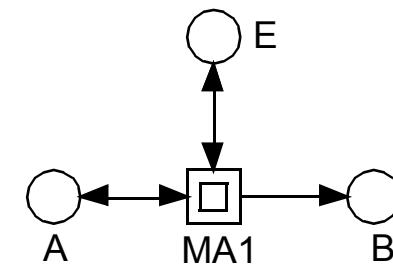
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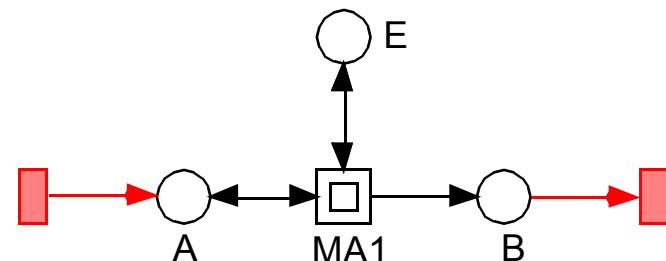
□ **conservative -> BND**

- > *all transitions preserve token number*



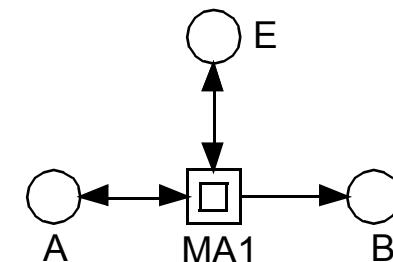
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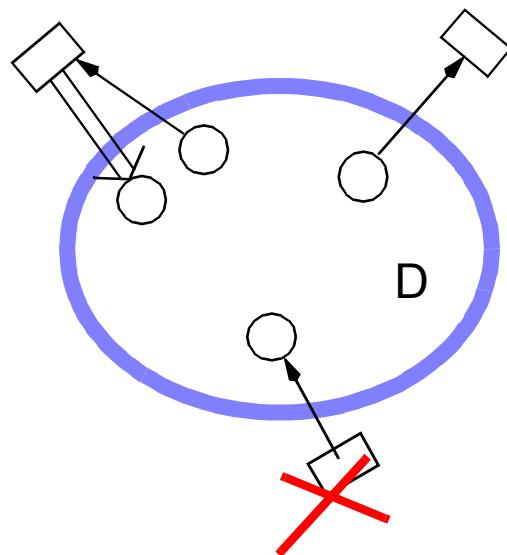


□ **Siphon Trap Property (STP)**

## SIPHON TRAP PROPERTY (STP)

**Siphon D**

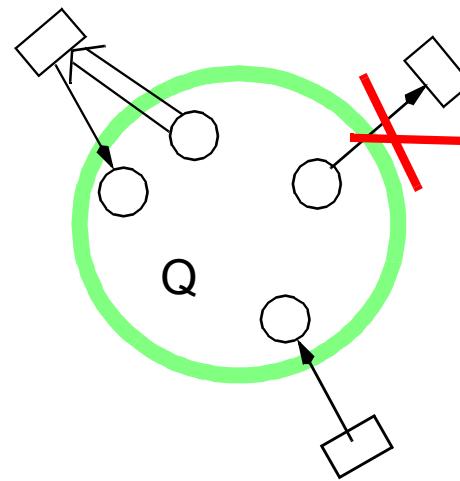
FD DF



any transition putting token into the set  
also takes token from it:  
an empty siphon will never get marked

**Trap Q**

QF FQ

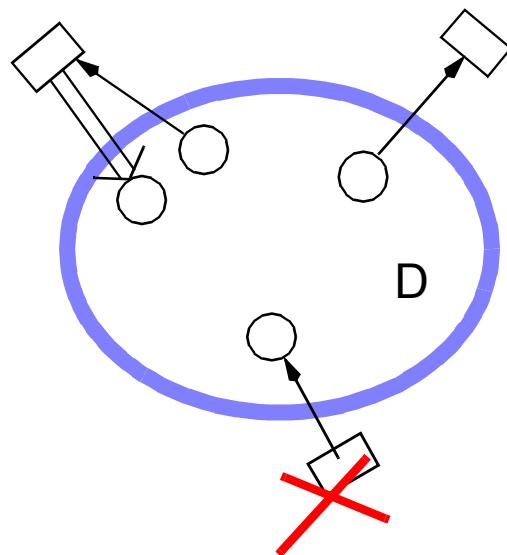


any transition taking tokens from the set  
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## SIPHON TRAP PROPERTY (STP)

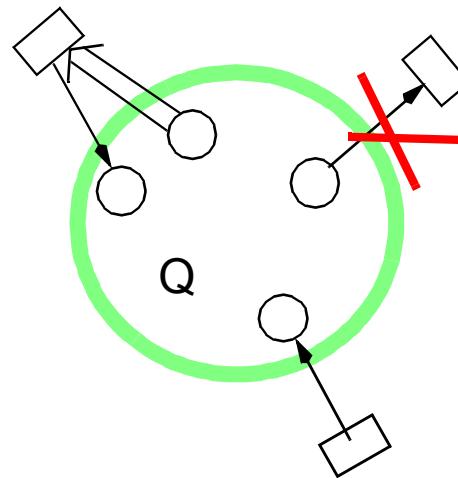
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**Siphon D**  
FD DF



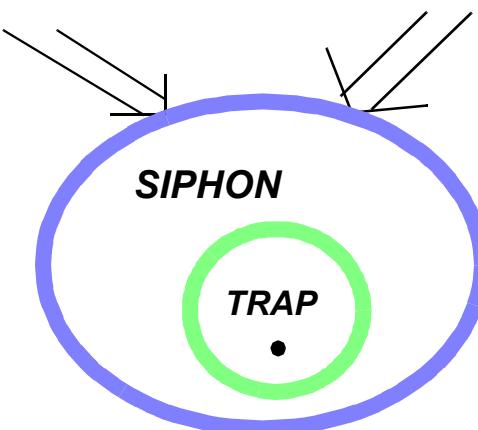
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**Trap Q**  
QF FQ



any transition taking tokens from the set  
also puts token into it:  
a marked trap will never get empty

**STP:** each siphon contains a  
(sufficiently) marked trap (at  $m_0$ )



## SIPHON TRAP PROPERTY (STP)

- allows to decide liveness, sometimes

*EFC -> STP (& HOM & NBM )    <->    live*

*ES & STP ( & HOM & NBM )      ->      live*

*STP ( & HOM & NBM )*      ->      *no DSt*

*no siphon* -> *live*

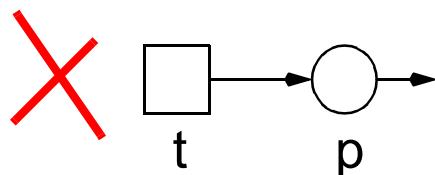
- allow to decide liveness, sometimes

<i>EFC -&gt; STP (&amp; HOM &amp; NBM )</i>	$\leftrightarrow$	<i>live</i>
<i>ES &amp; STP ( &amp; HOM &amp; NBM )</i>	$\rightarrow$	<i>live</i>
<i>STP ( &amp; HOM &amp; NBM )</i>	$\rightarrow$	<i>no DSt</i>
<i>no siphon</i>	$\rightarrow$	<i>live</i>

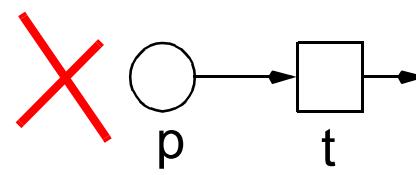
- some examples, where it helps

- > *RKIP pathway (BND): ES & STP -> live*
- > *MAPK cascade (BND): STP & nES -> no DSt*
  
- > *biosensor (not BND): ES & STP -> live*
- > *apoptosis (not BND): no siphon -> live*
  
- > *lac operon (not BND): STP & nES -> no DSt*

- downsizing the net structure while preserving some properties  
-> *liveness, boundedness*
- example of two simple reduction rules



t live  
p unbounded

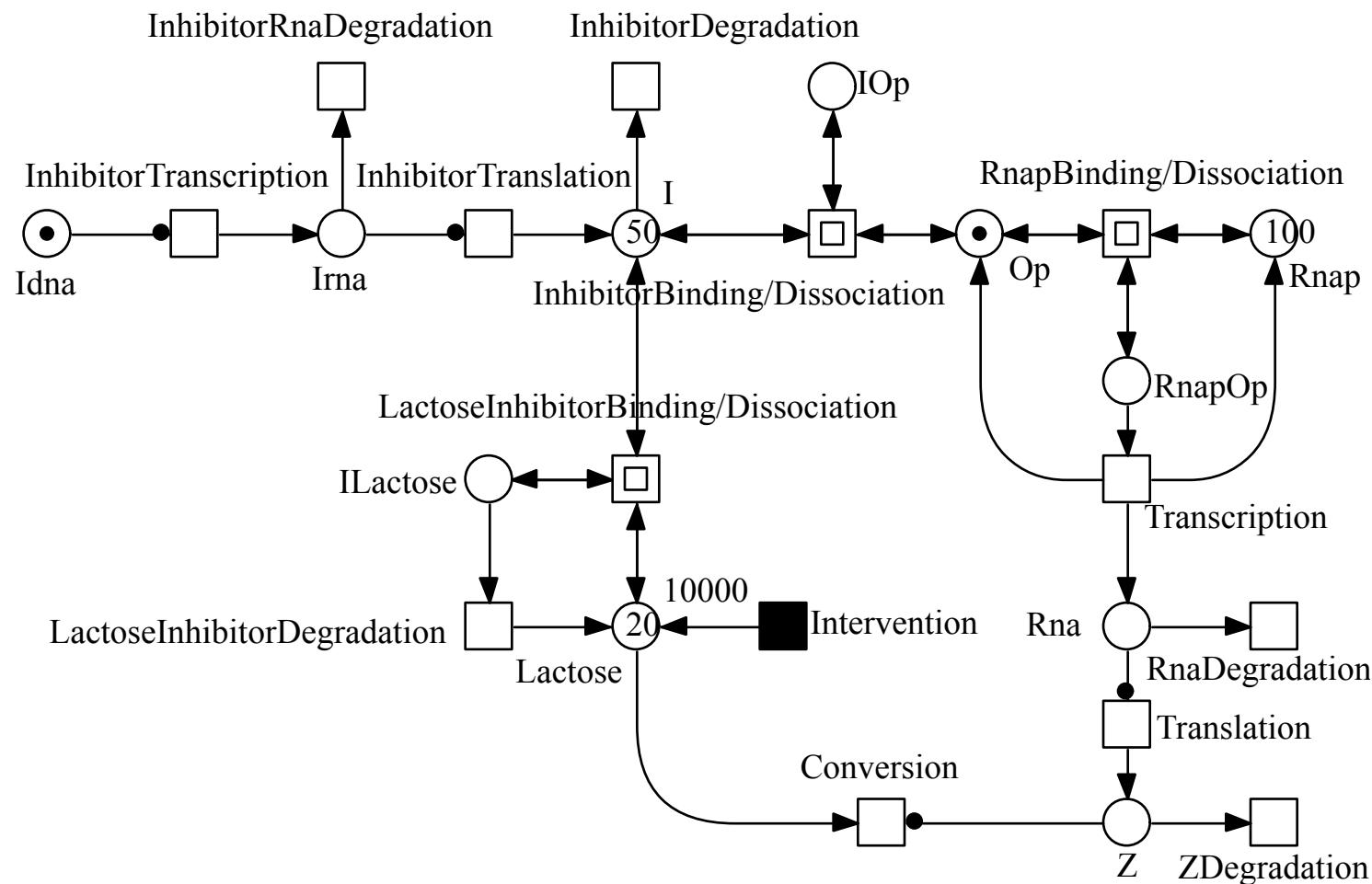


t not live  
p bounded

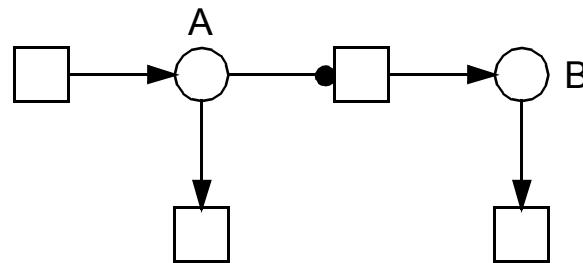
-> *input nodes allow net reduction:*  
*conclude properties -> delete nodes -> conclude properties -> . . .*

- several sets of reduction rules
  - > *relatively weak, in general*
  - > *sensitive to the order they are applied*
  - > *however, sometimes they help*

## □ lac operon (Wilkinson 2006)



- lac operon, reduced by INA/HUB



- liveness becomes obvious

PUR	ORD	HOM	NBM	CSV	SCF	CON	SC	FT0	TF0	FPO	PF0	<b>NC</b>
N	N	Y	Y	N	N	N	N	Y	Y	Y	N	<b>nES</b>
<b>DTP</b>	CPI	CTI	SCTI	SB	<b>k-B</b>	1-B	DCF	<b>DSt</b>	DTr	LIV	RV	
<b>Y</b>	N	Y	-	N	<b>N</b>	N	-	<b>N</b>	-	-	-	

# **NET INVARIANTS, A CRASH COURSE**

$r1: A \rightarrow 2B$

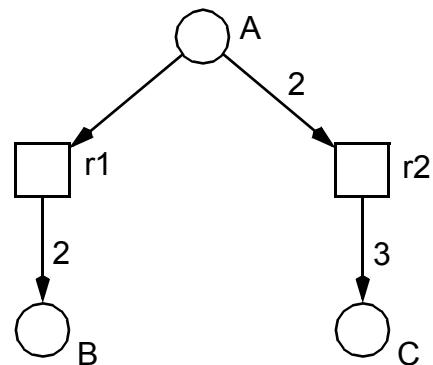
$r2: 2A \rightarrow 3C$

## T-INVARIANTS, Ex1

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$r1: A \rightarrow 2B$

$r2: 2A \rightarrow 3C$

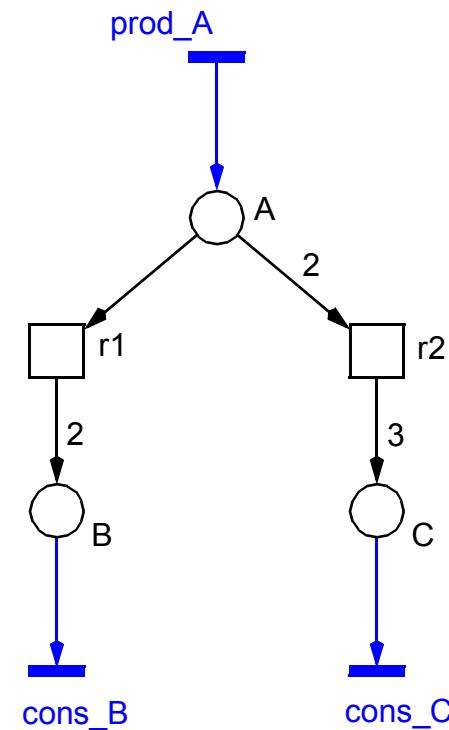
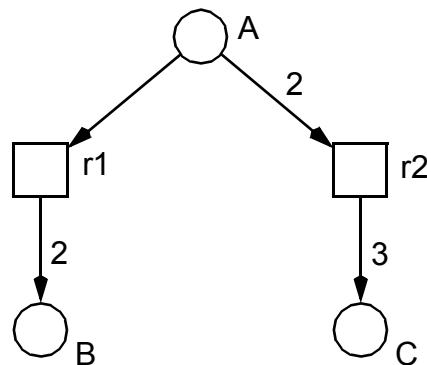


## T-INVARIANTS, Ex1

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$r1: A \rightarrow 2B$

$r2: 2A \rightarrow 3C$



## INCIDENCE MATRIX C

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- a representation of the net structure

=> stoichiometric matrix

$C =$

P \ T	t1	...	tj	...	tm
p1					
pi			cij		
:					
pn			vtj		

$$c_{ij} = C(pi, tj) = \delta_{tj}(pi)$$

$$vt_j = C(*, tj)$$

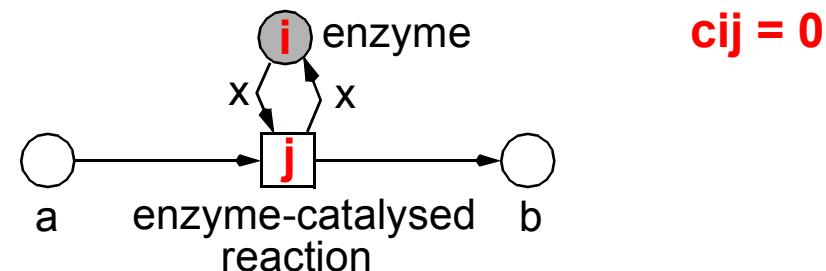
- matrix entry  $c_{ij}$ :

token change in place  $pi$  by firing of transition  $tj$

- matrix column  $vt_j$ :

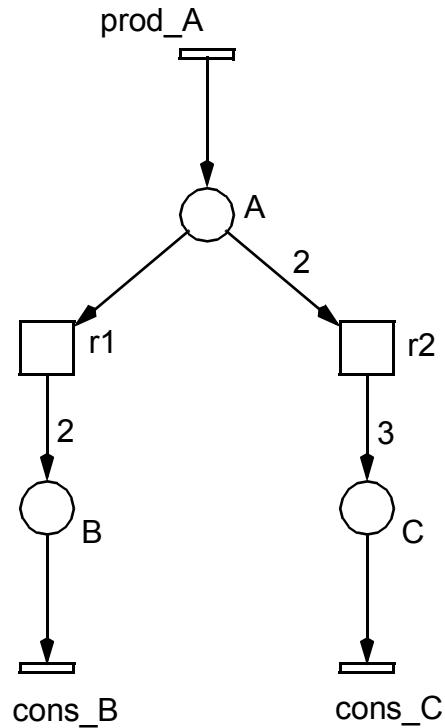
vector describing the change of the whole marking by firing of  $tj$

- side-conditions are neglected



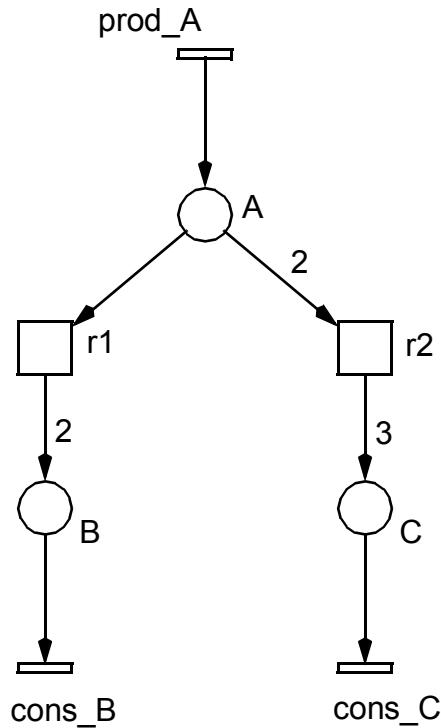
# INCIDENCE MATRIX C, Ex1

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# INCIDENCE MATRIX C, Ex1

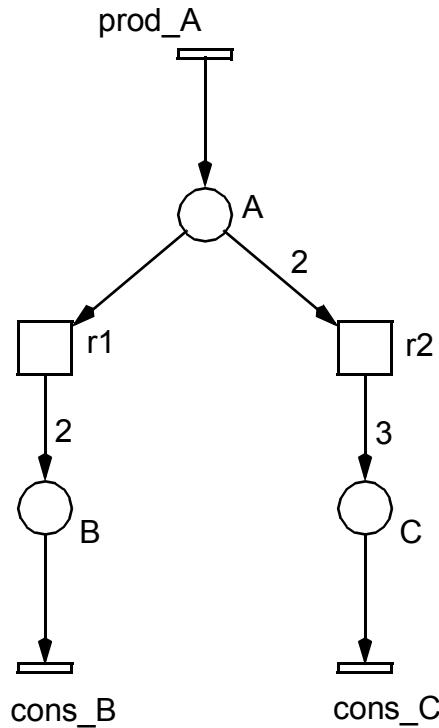
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T P	r1	r2	prod_A	cons_B	cons_C
A					
B					
C					

# INCIDENCE MATRIX C, Ex1

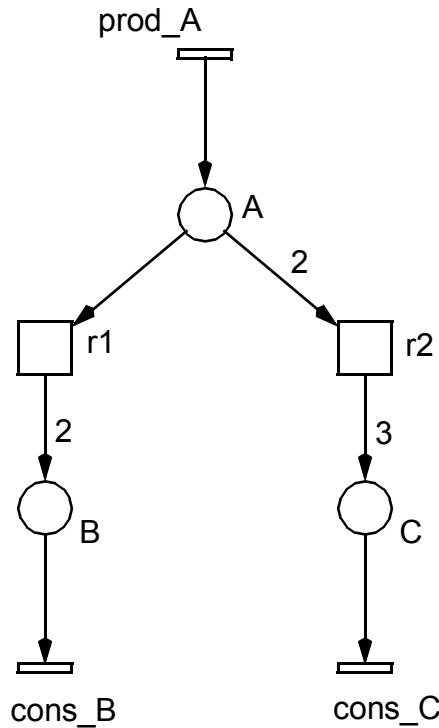
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T P	r1	r2	prod_A	cons_B	cons_C
A	-1	-2	1		
B	2			-1	
C		3			-1

# INCIDENCE MATRIX C, Ex1

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T P	r1	r2	prod_A	cons_B	cons_C
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B	2			-1	
C		3			-1

1                  1                  2

□ Lautenbach, 1973

-> Schuster, 1993

□ T-invariants

-> *integer solutions x*

-> *multisets of transitions*

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

□ Lautenbach, 1973 → Schuster, 1993

□ T-invariants → multisets of transitions

-> integer solutions  $x$

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

□ minimal T-invariants

-> there is no T-invariant with a smaller support → sets of transitions

-> gcd of all non-zero entries is 1

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

-> multiplication with a positive integer

-> addition

-> division by a common divisor

$$kx = \sum_i a_i x_i$$

□ Lautenbach, 1973

-> Schuster, 1993

□ T-invariants

-> integer solutions  $x$

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

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$$kx = \sum_i a_i x_i$$

□ Covered by T-Invariants (CTI)

-> each transition belongs to a T-invariant

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

---

*PN & Systems Biology*

- T-invariants = (multi-) sets of transitions = Parikh vector
    - > zero effect on marking
    - > reproducing a marking / system state
  - two interpretations
    1. *partially ordered transition sequence*  
*of transitions occurring one after the other*
      - > substance / signal flow
    2. *relative transition firing rates*  
*of transitions occurring permanently & concurrently*
      - > steady state behaviour
  - a minimal T-invariant defines a connected subnet
    - > the T-invariant's transitions (the support),
      - + all their pre- and post-places
      - + the arcs in between
    - > pre-set of support = post-set of support
      - > self-contained

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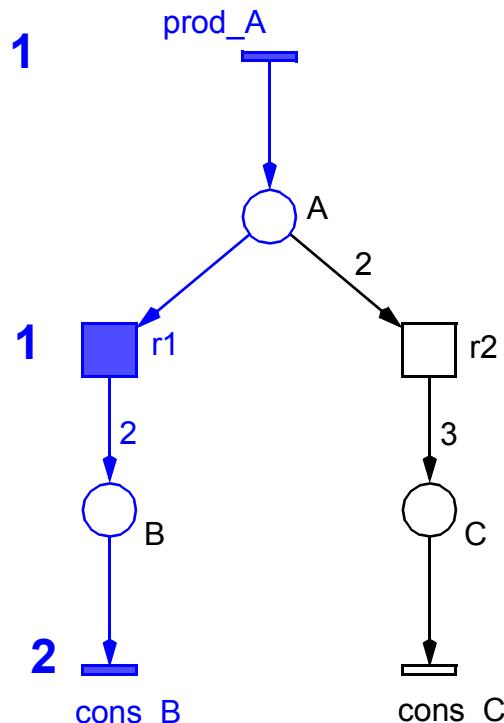
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-> behaviour understanding  
of transitions occurring one after the other
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## T-INVARIANTS, Ex1

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$r2: 2 A \rightarrow 3 C$



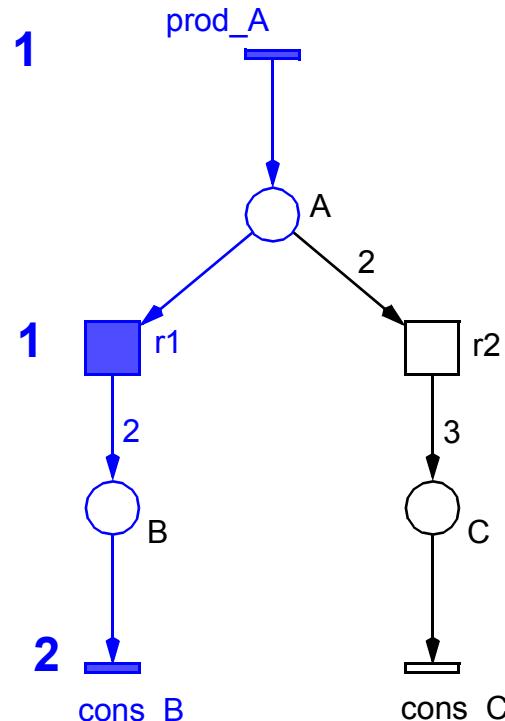
**T-INVARIANT 1**

# T-INVARIANTS, Ex1

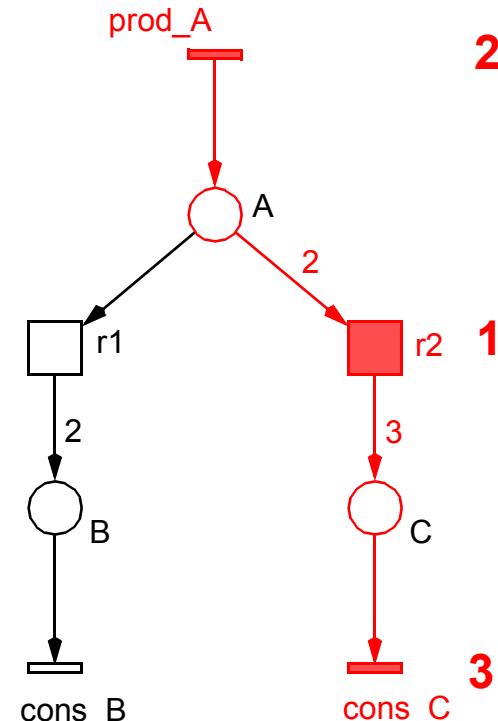
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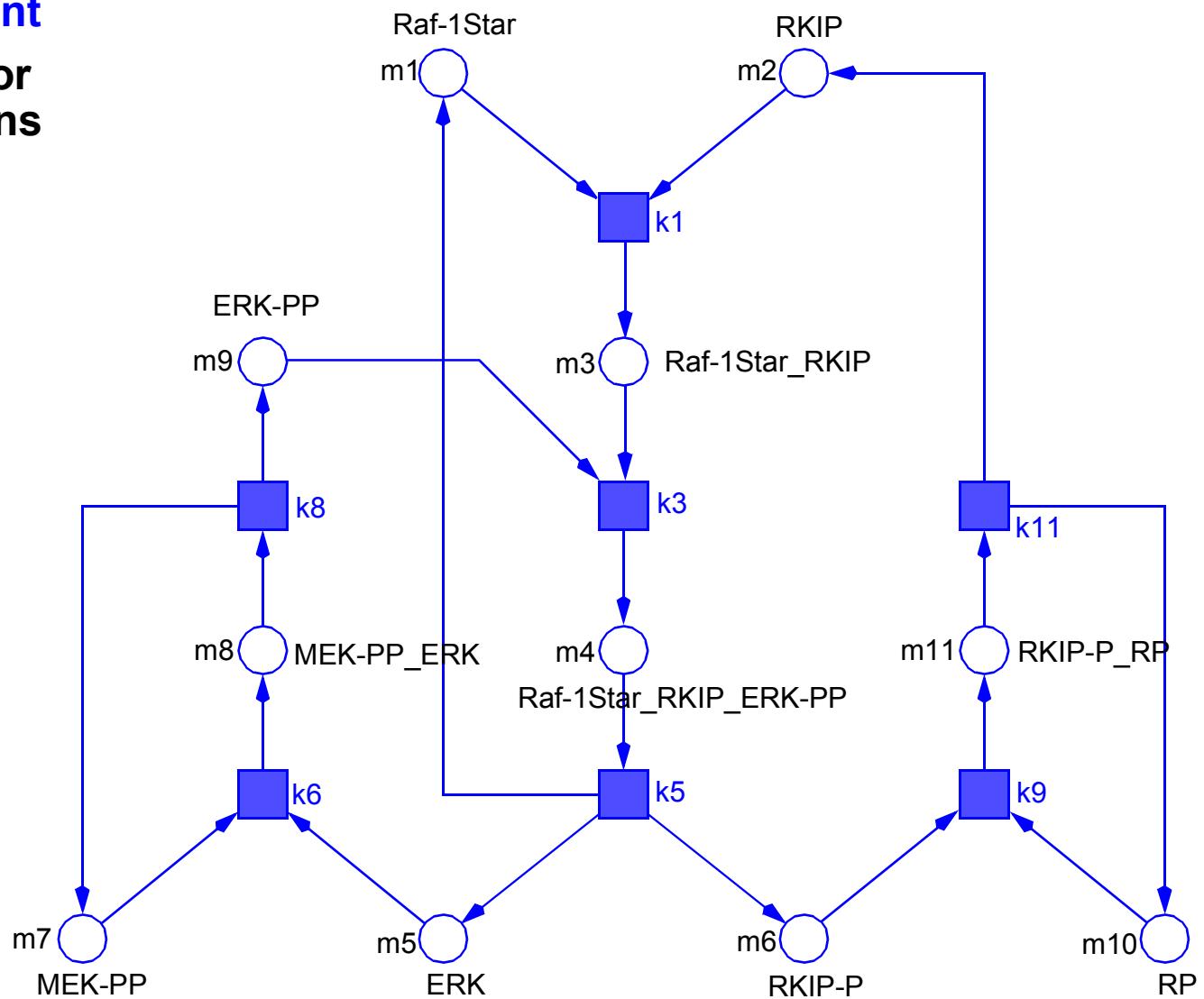
T-INVARIANT 1



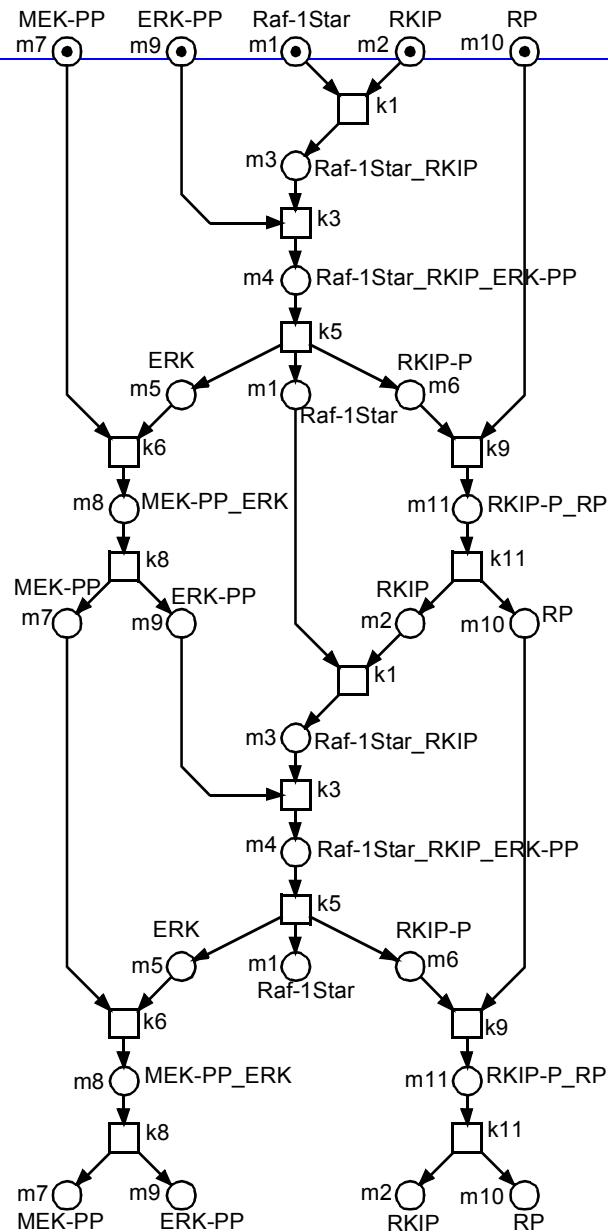
T-INVARIANT 2

-> non-trivial T-invariant

+ four trivial ones for reversible reactions



- feasibility check under the constructed marking**
- T-invariant's unfolding to describe its behaviour**
  - > **partial order structure**
- labelled condition / event net**
  - > *events (boxes)*
    - *transition occurrences*
  - > *conditions (circles)*
    - *involved compounds*
- occurrence net**
  - > *acyclic*
  - > *no backward branching conditions*
  - > **infinite**



- Lautenbach, 1973

- P-invariants

- > integer solutions  $y$

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

- > 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 place belongs to a P-invariant

- > CPI => BND (sufficient condition)

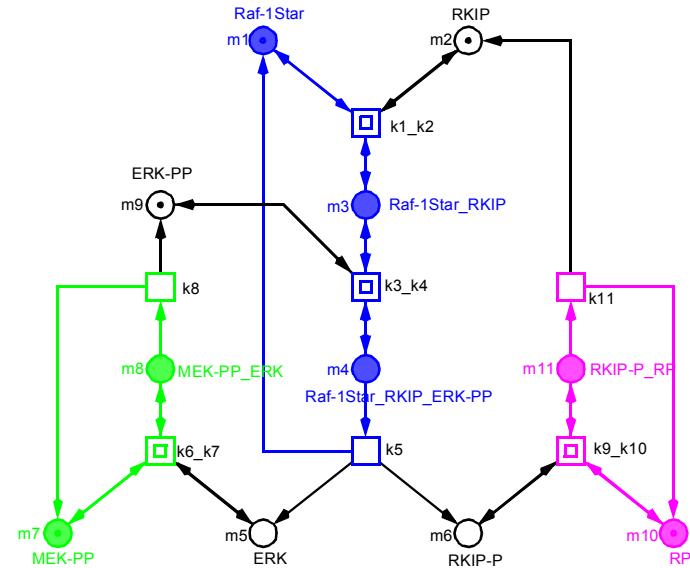
- the firing of any transition has no influence on the weighted sum of tokens on the P-invariant's places
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- a P-invariant defines a subnet
  - > *the P-invariant's places (the support),  
+ all their pre- and post-transitions  
+ the arcs in between*
  - > *pre-sets of supports = post-sets of supports*      -> **self-contained**

# P-INVARIANTS, Ex - RKIP PATHWAY

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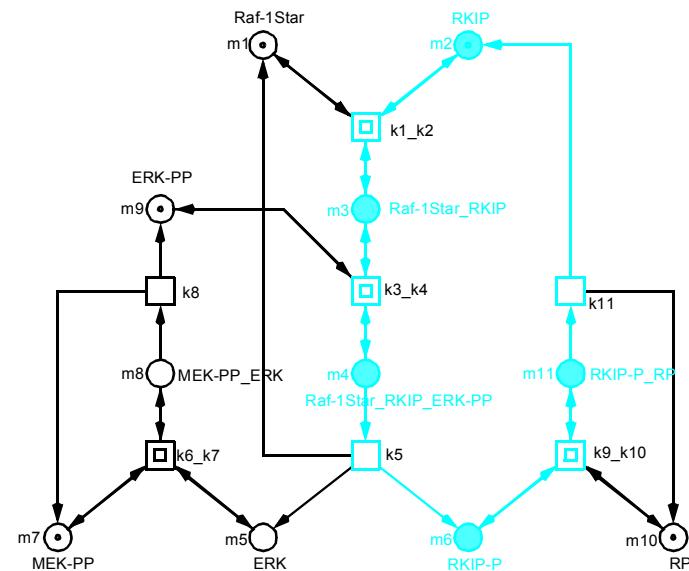
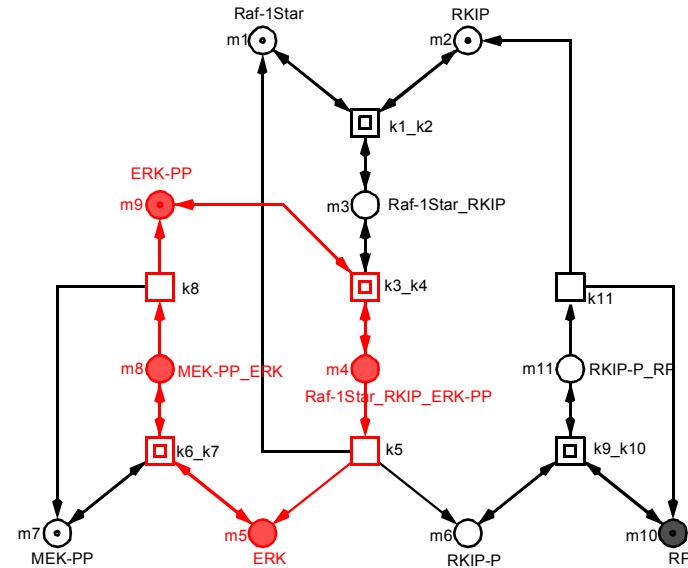
**P-INV1: MEK**

**P-INV2: RAF-1STAR**

**P-INV3: RP**

**P-INV4: ERK**

**P-INV5: RKIP**



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-> *P-invariants are structural deadlocks and traps*

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  - > *exactly one token, corresponding to species conservation*
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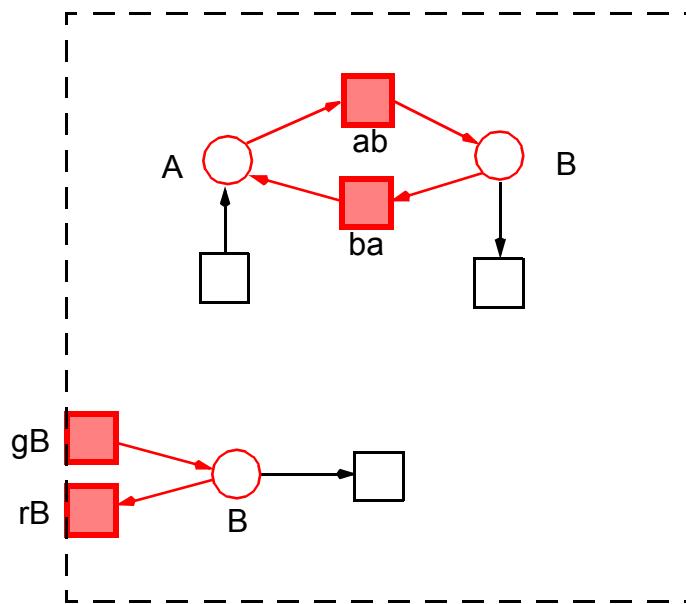
**-> UNIQUE INITIAL MARKING <-**

- trivial minimal T-invariants

- > *reversible reactions*
- > *boundary transitions of auxiliary compounds*

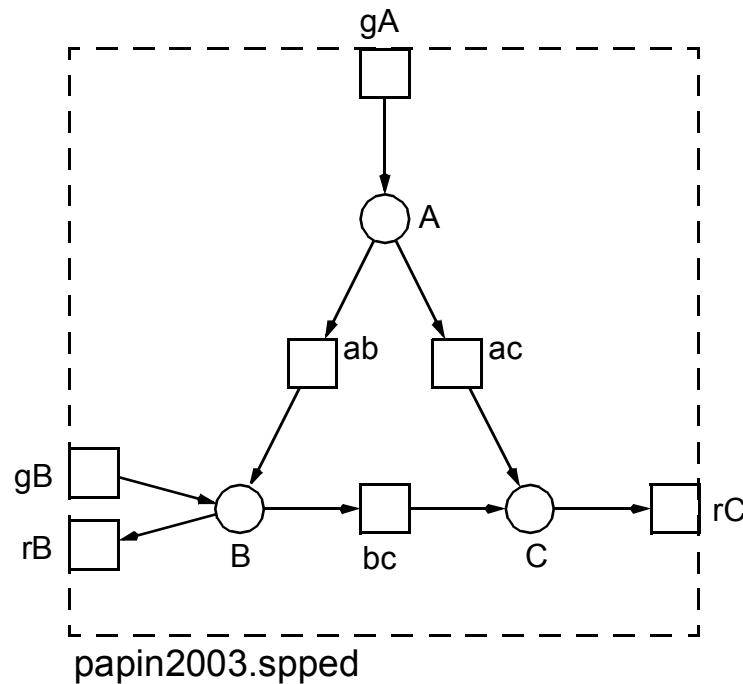
- non-trivial minimal T-invariants

- > *i/o-T-invariants*  
*covering boundary transitions of input / output compounds*
- > *inner cycles*



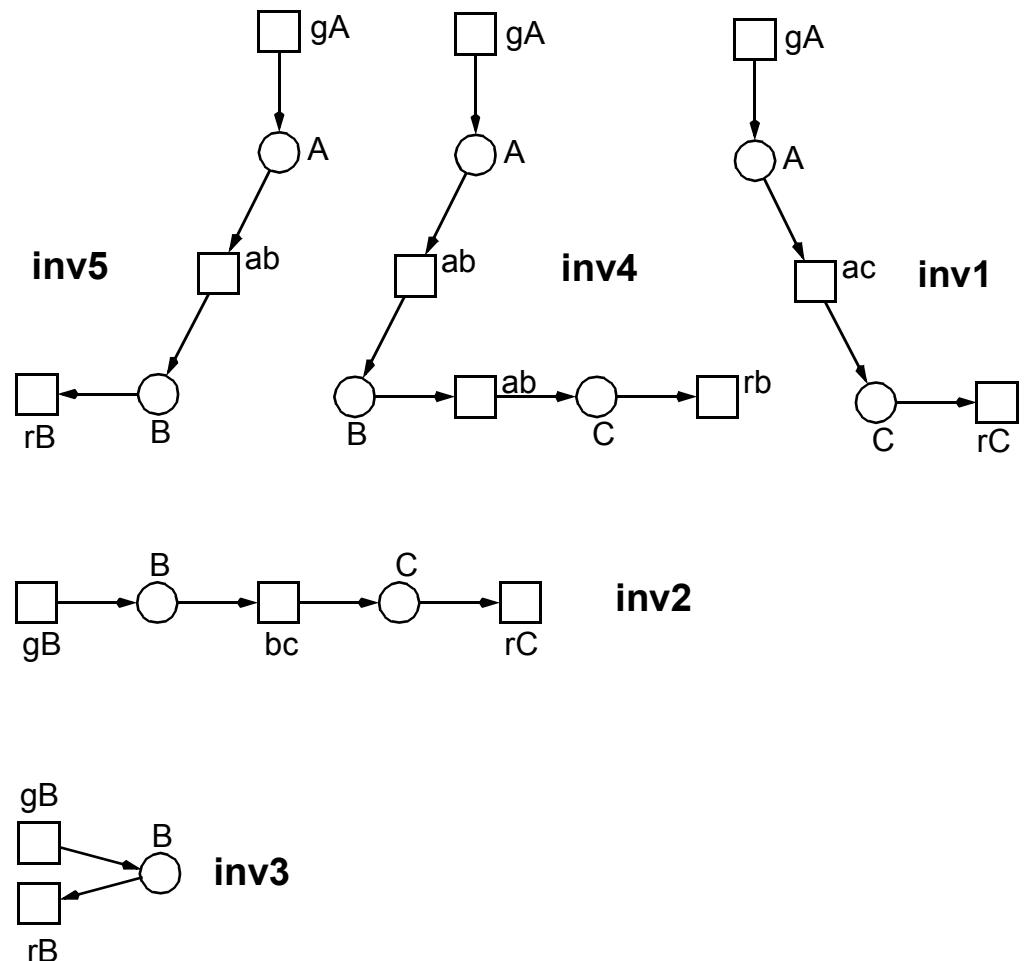
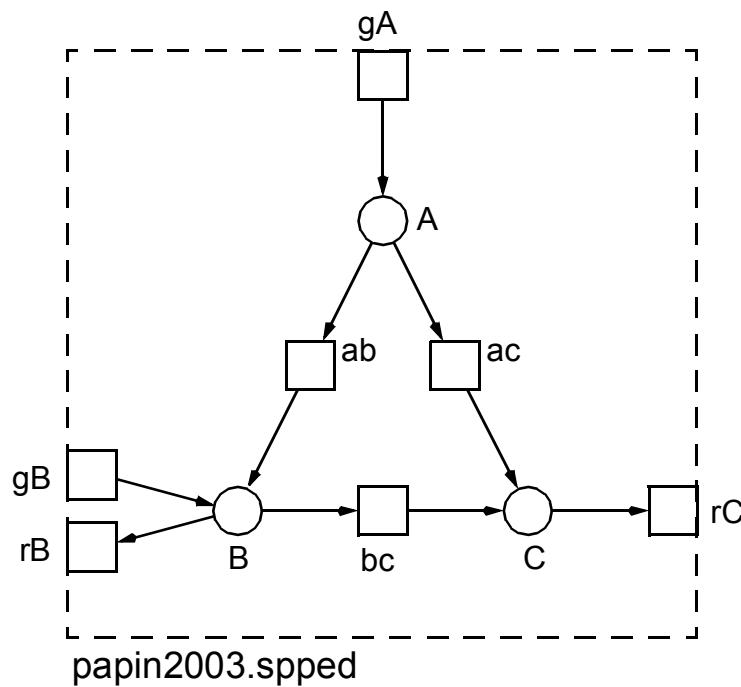
### □ substances involved

- > *input substance A*
- > *output substance C*
- > *auxiliary substance B*



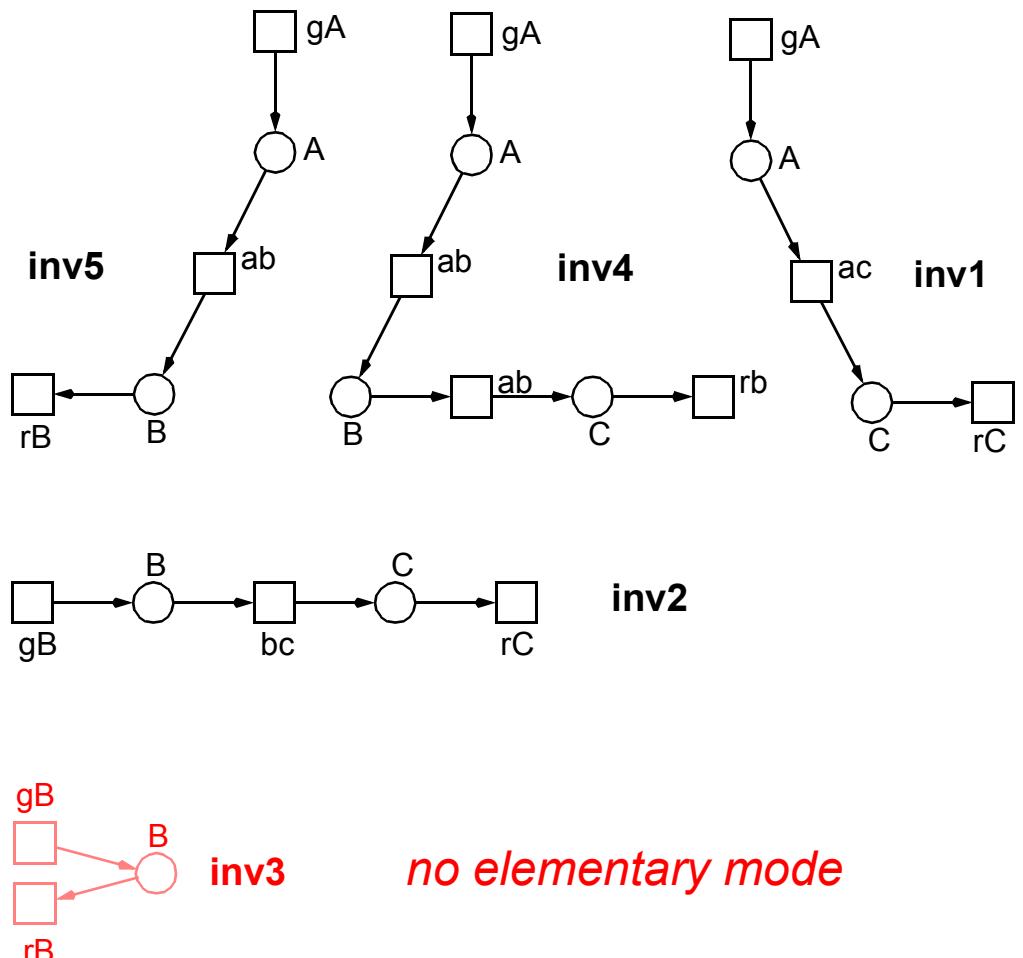
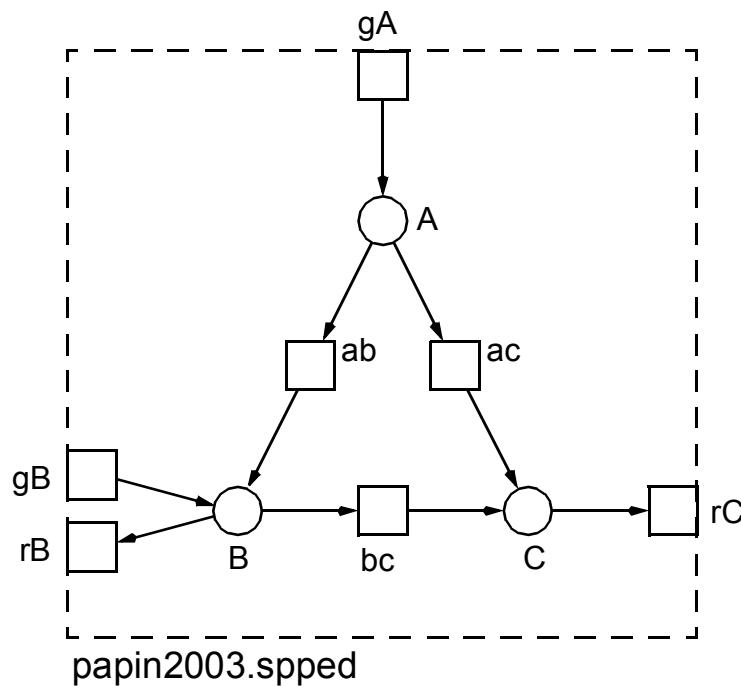
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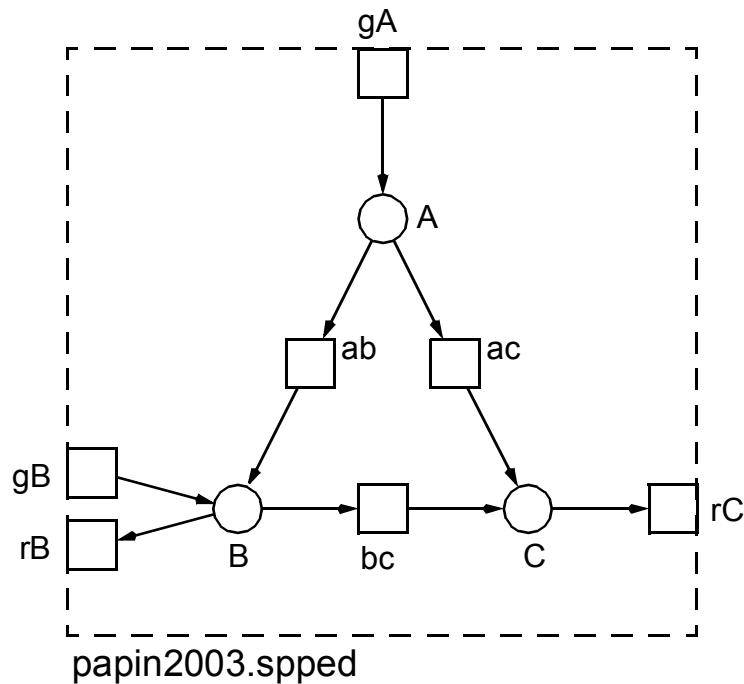
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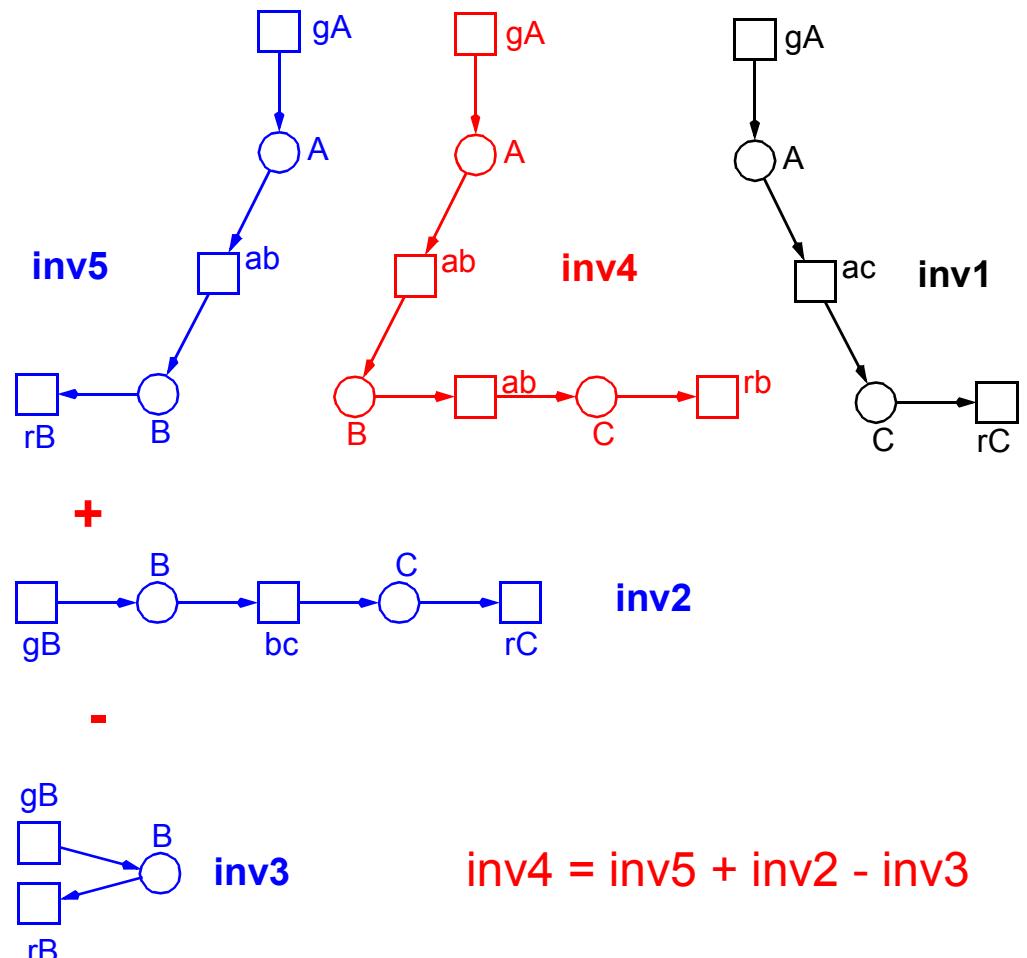


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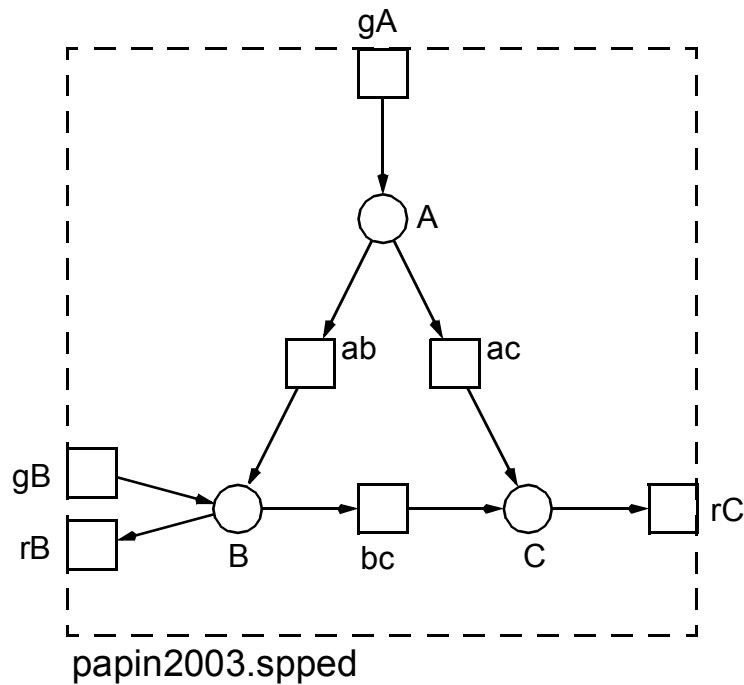


*no extreme pathway*

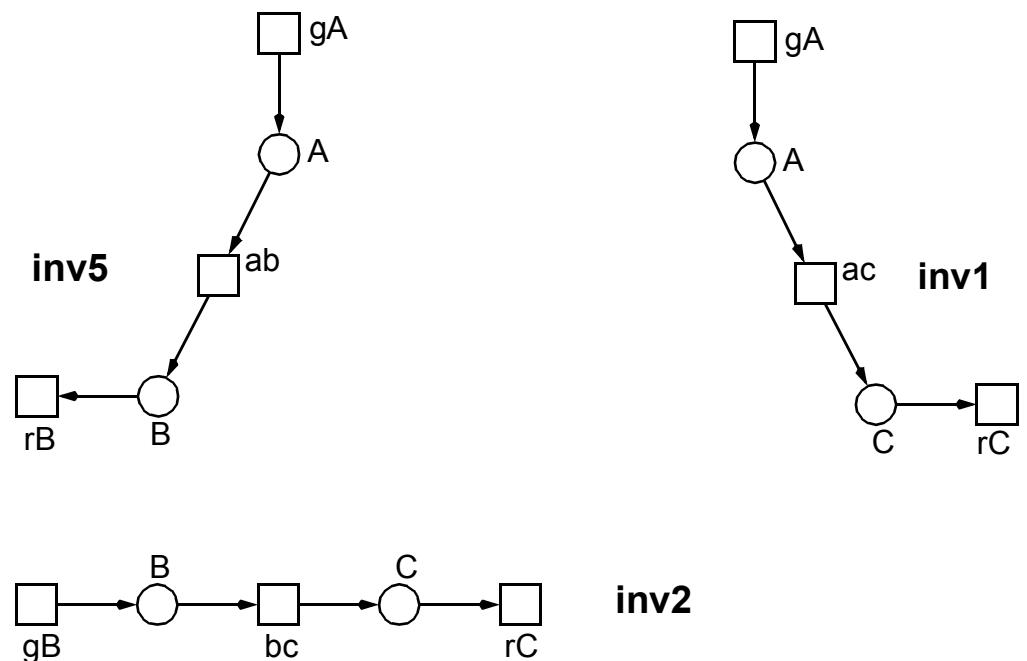


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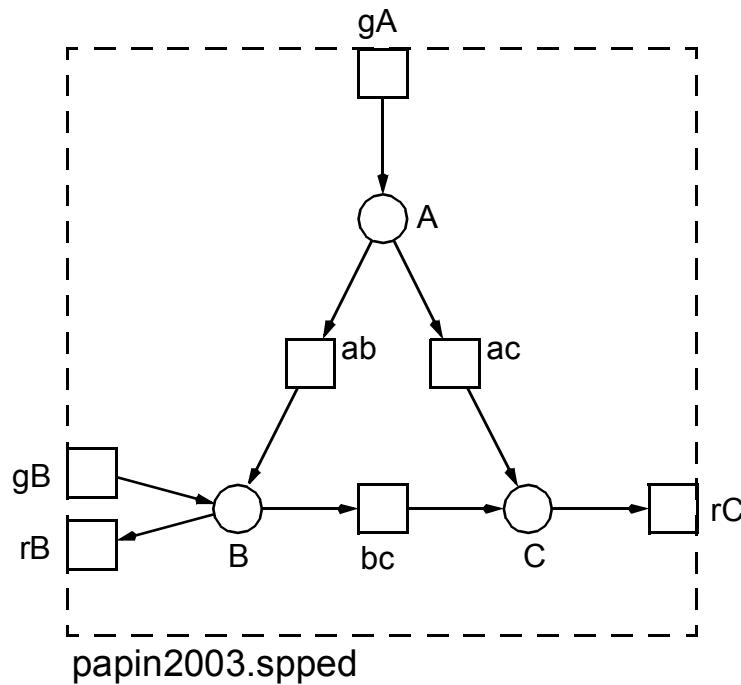


## EXTREME PATHWAYS

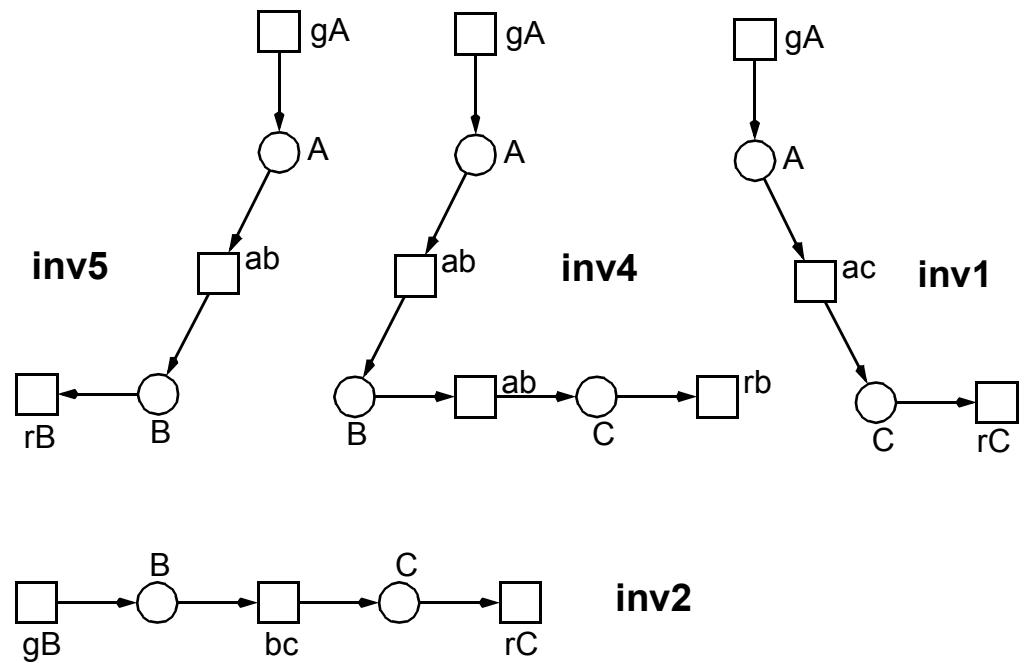


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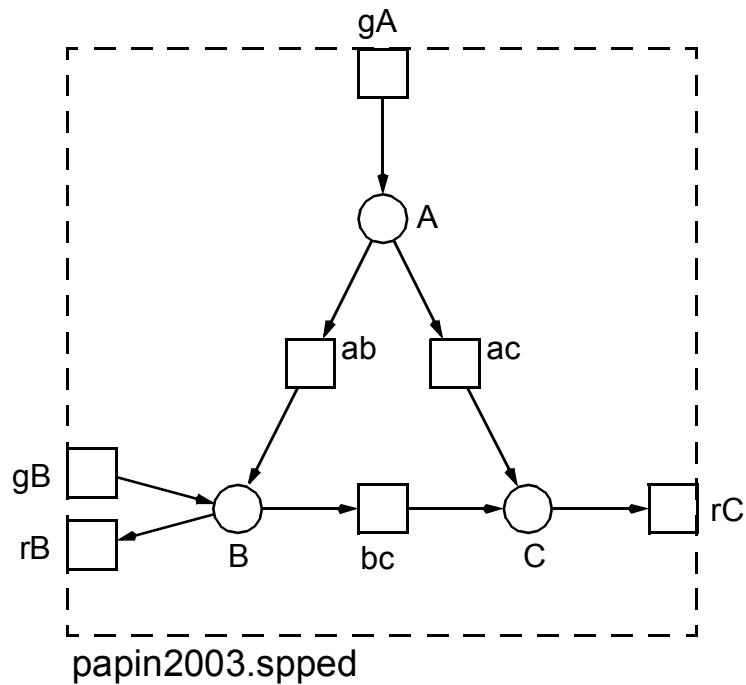


## ELEMENTARY MODES

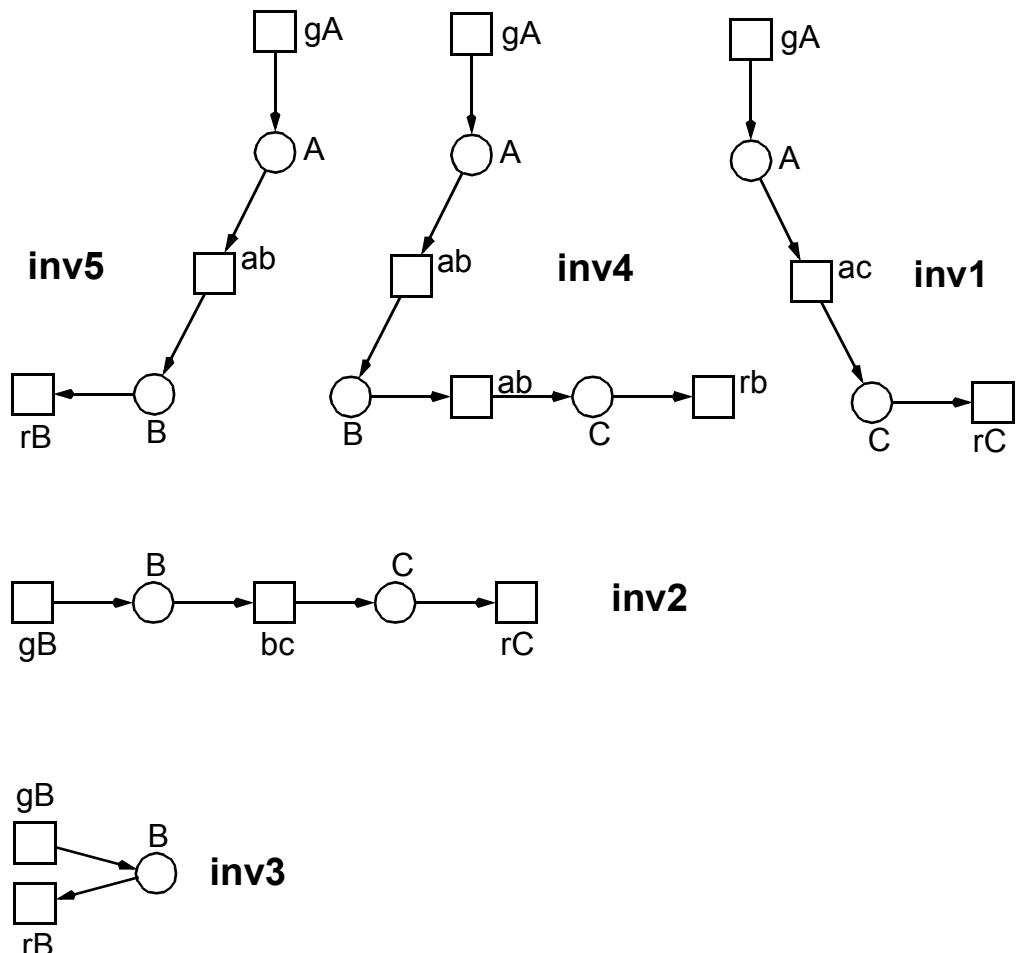


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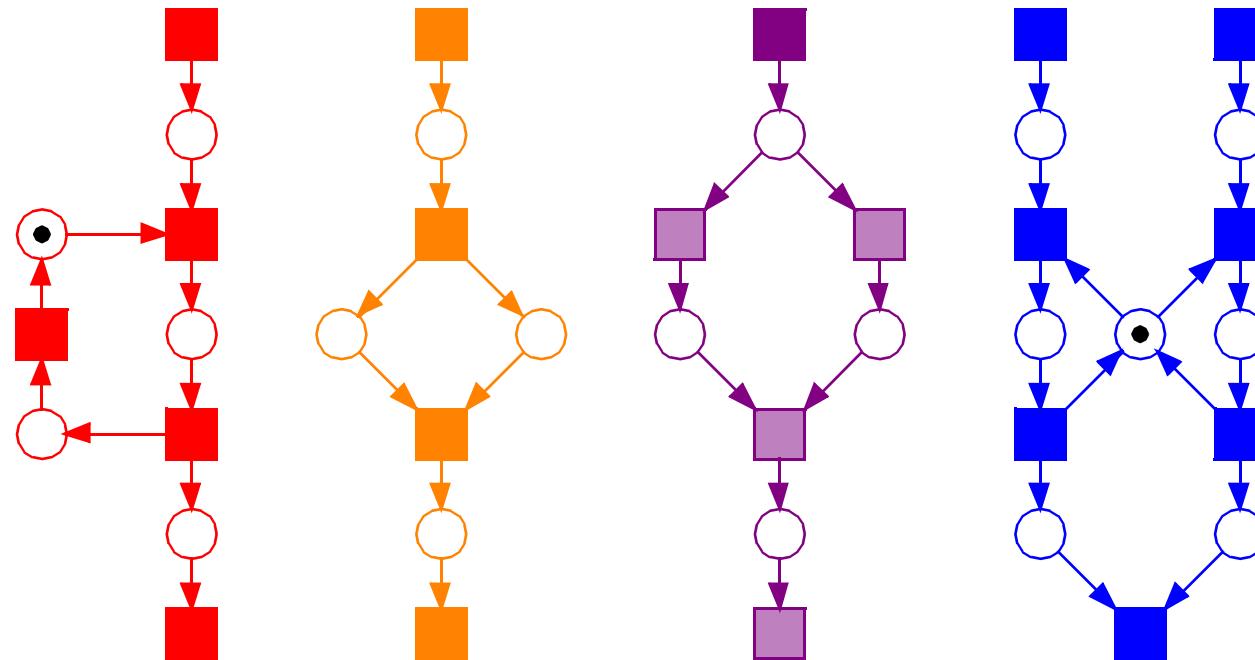
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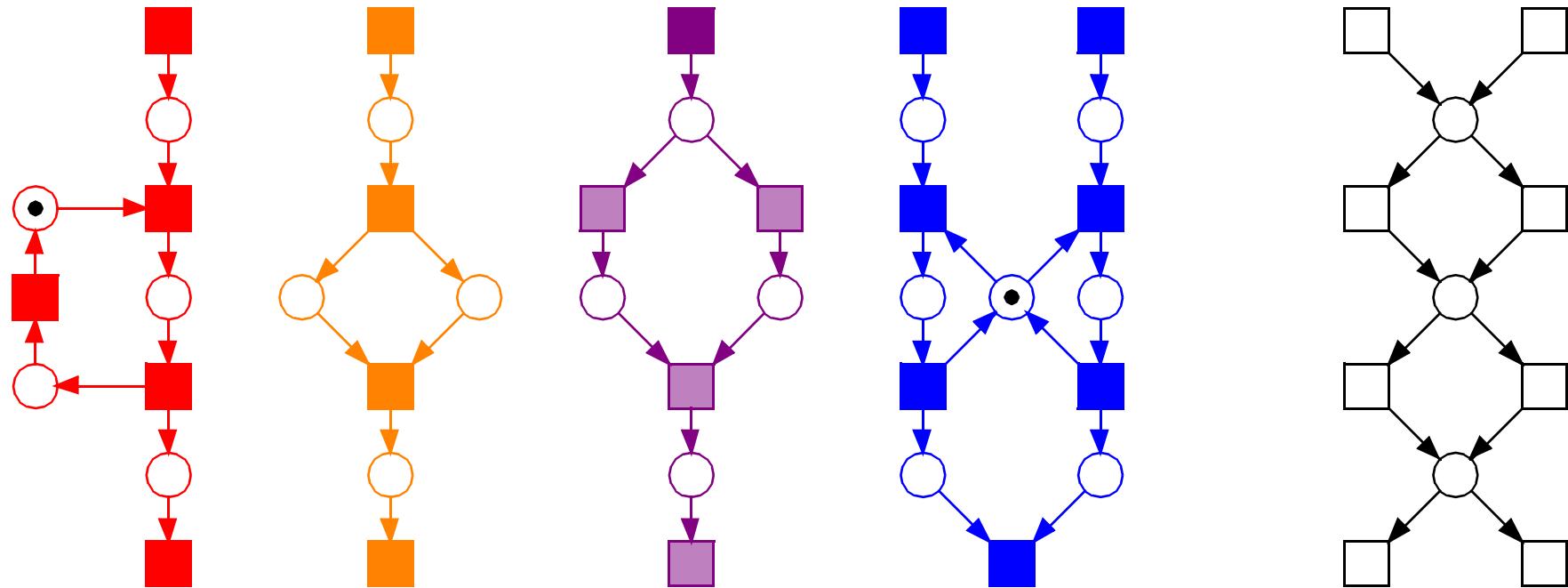
## MINIMAL T-INVARIANTS



- T-invariants may contain any structure



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- T-invariants generally overlap
  - > combinatorial effect brings *explosion* in the number of min. T-invariants ( $2^4$ )
- likewise for P-invariants

## subnetwork identification

- > *P-invariants: token-preserving modules (mass conservation)*
- > *T-invariants: state-repeating modules (elementary modes)*

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- > *CPI (if closed model), CTI*
- > *no minimal P/T-invariant without biological interpretation*
- > *no known mass conservation without corresponding P-invariant*
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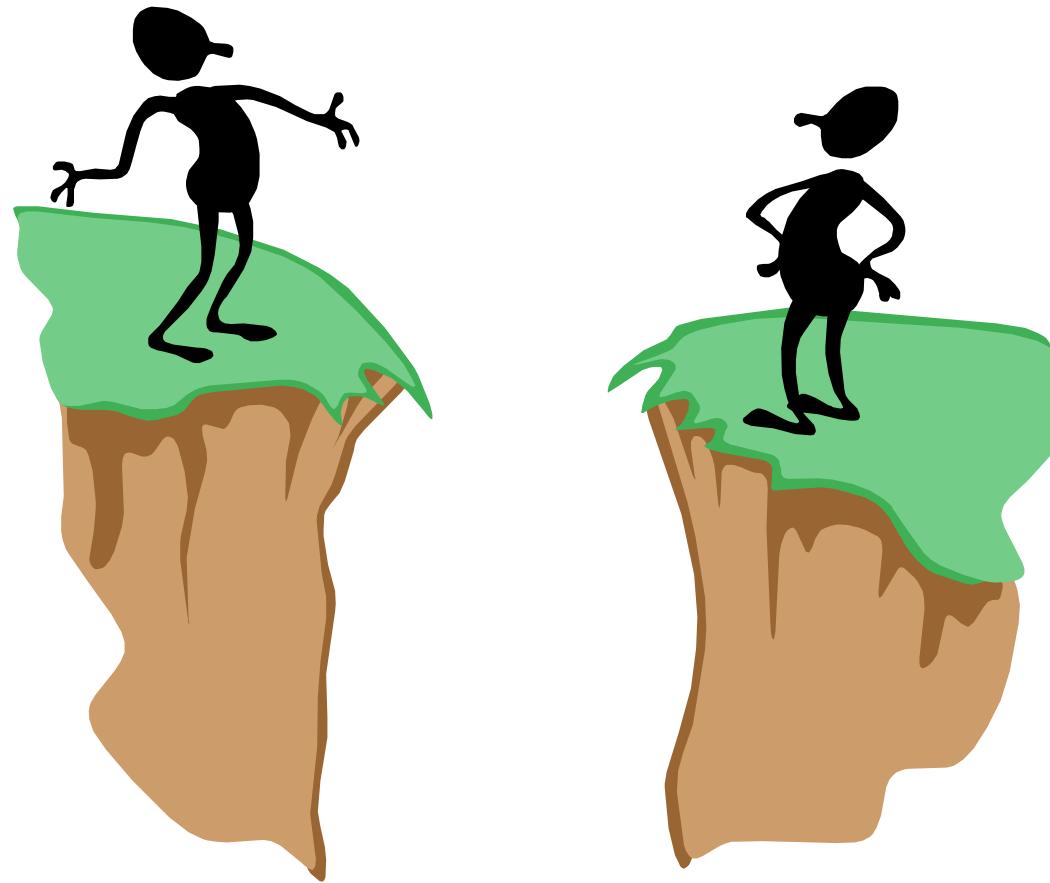
## □ choice of stochastic analysis techniques

- > *bounded*      - *analysis techniques, esp. analytic model checking*
- > *unbounded*    - *simulation techniques, esp. simulative model checking*

- M Heiner, D Gilbert, R Donaldson:  
*Petri Nets for Systems and Synthetic Biology;*  
in SFM 2008, Springer LNCS 5016, pp. 215-264, 2008.
  
- M Heiner:  
*Understanding Network Behaviour by Structured Representations of Transition Invariants - A Petri Net Perspective on Systems and Synthetic Biology;*  
in Algorithmic Bioprocesses; Chapter 19, Springer, July 2009.
  
- M Heiner, K Sriram:  
*Structural Analysis to Determine the Core of Hypoxia Response Network;*  
PLoS ONE 5(1): e8600, doi:10.1371/journal.pone.0008600, January 2010.
  
- M Heiner; R Donaldson; D Gilbert:  
*Petri Nets for Systems Biology;*  
in MS Iyengar (ed.): Symbolic Systems Biology: Theory and Methods,  
Jones & Bartlett Publishers, LLC, 2010.

**THANKS !**

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