

PATHWAY ANALYSIS OF BIOCHEMICAL NETWORKS WITH PETRI NETS

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Dep. of CS

BASIC NOTIONS

-
- minimal T-invariants -> Lautenbach 1973
- elementary modes -> Schuster 1991
- extreme pathways -> Schilling, Schuster, Palson 1999
-

MODULAR COMPUTATION

- approach -> Zaitsev 2005
- (preliminary) results -> Lehrack 2006 (to appear)

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- ❑ elementary T-invariants -> Pascoletti 1986
- ❑ minimal T-invariants -> Lautenbach 1973
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BASIC NOTIONS

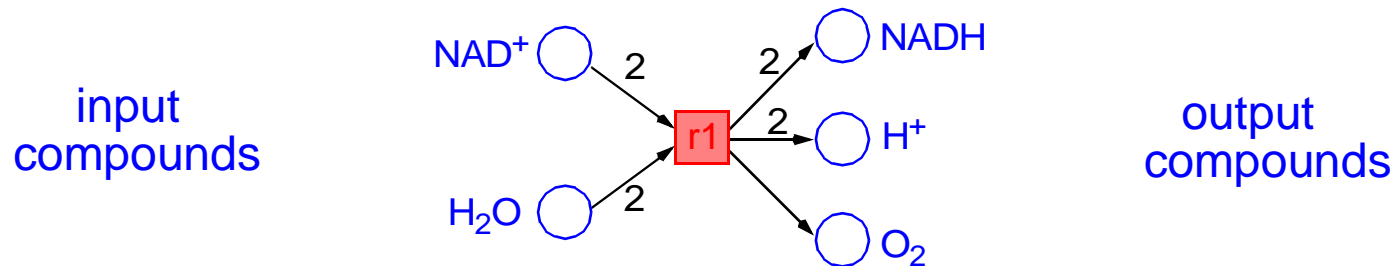
- ❑ **proper T-invariants** -> Pascoletti 1986
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- ❑ **extreme pathways** -> Schilling, Schuster, Palson 1999
- ❑ **generic pathways** -> Bockmayr 2005

MODULAR COMPUTATION

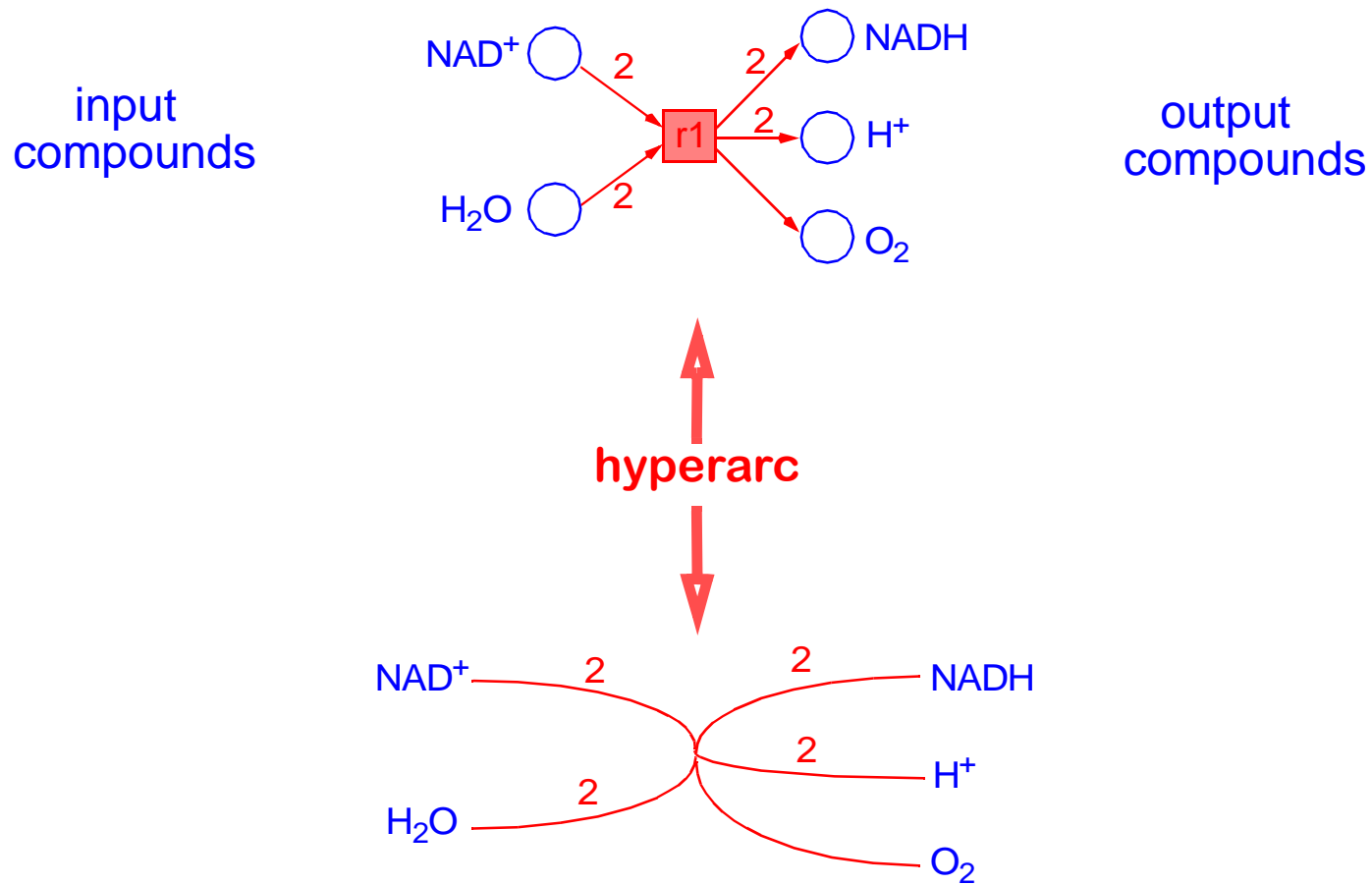
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PETRI NETS - BASICS

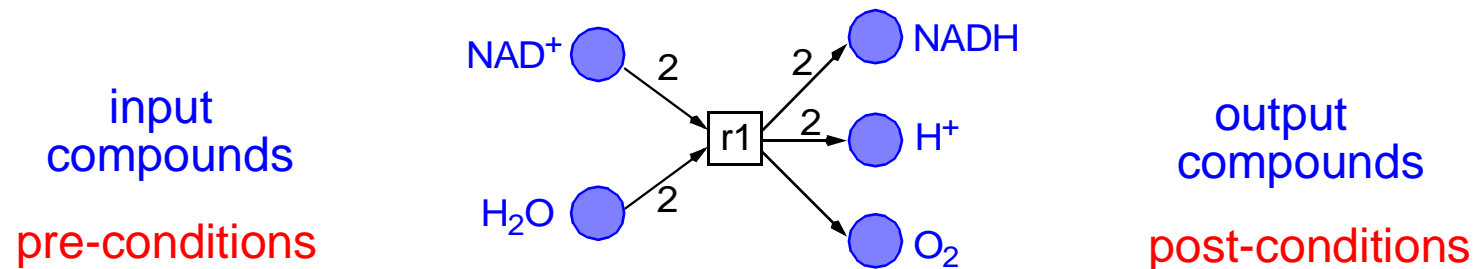
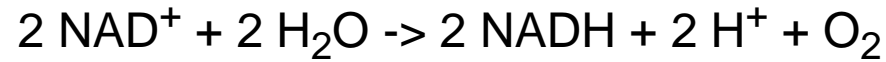
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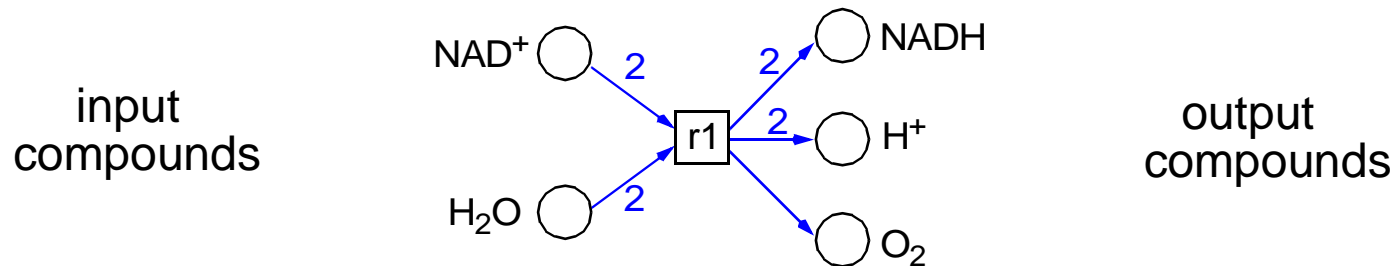
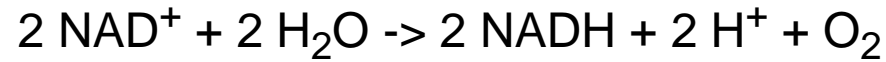


□ atomic actions → Petri net transitions → chemical reactions



□ local conditions → Petri net places → chemical compounds

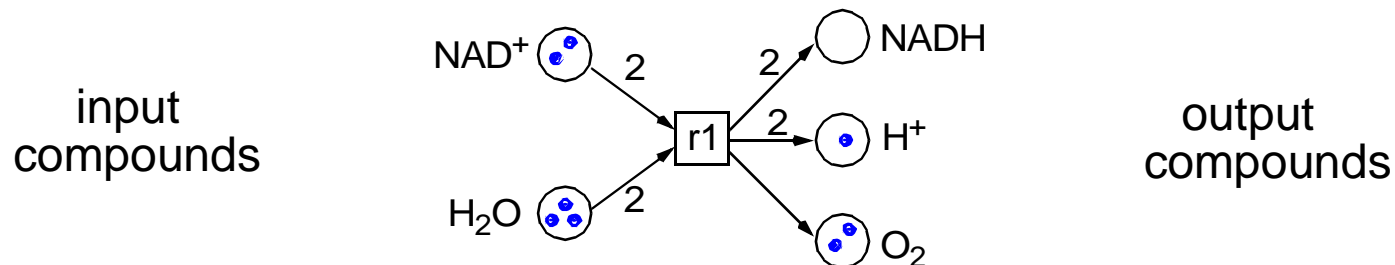
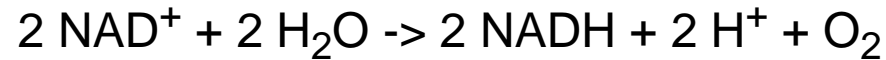
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□ local conditions -> Petri net places -> chemical compounds

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

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



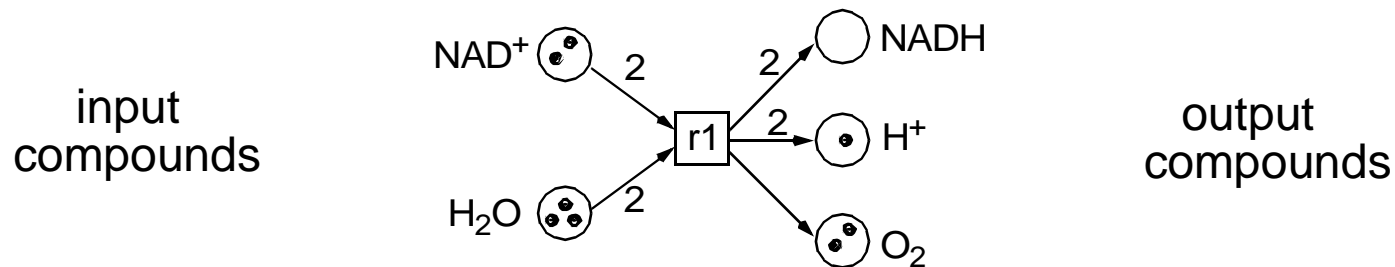
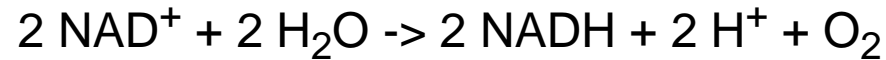
□ local conditions -> Petri net places -> chemical compounds

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□ 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



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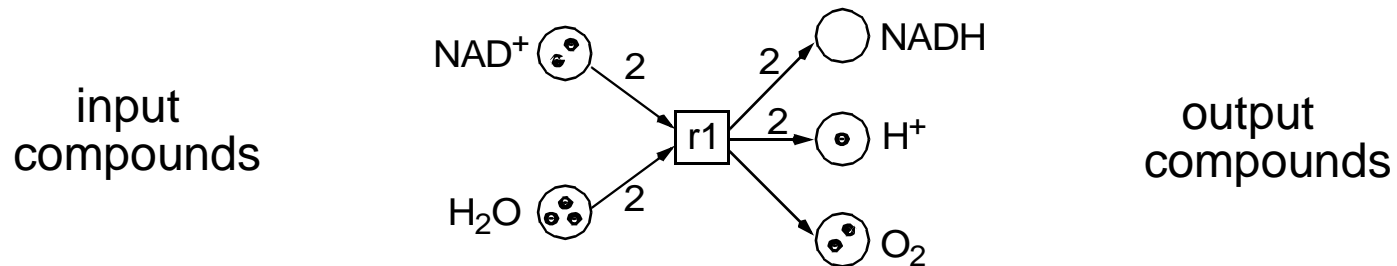
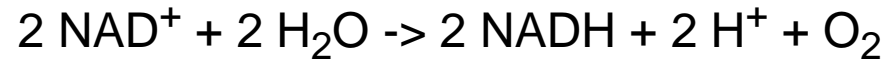
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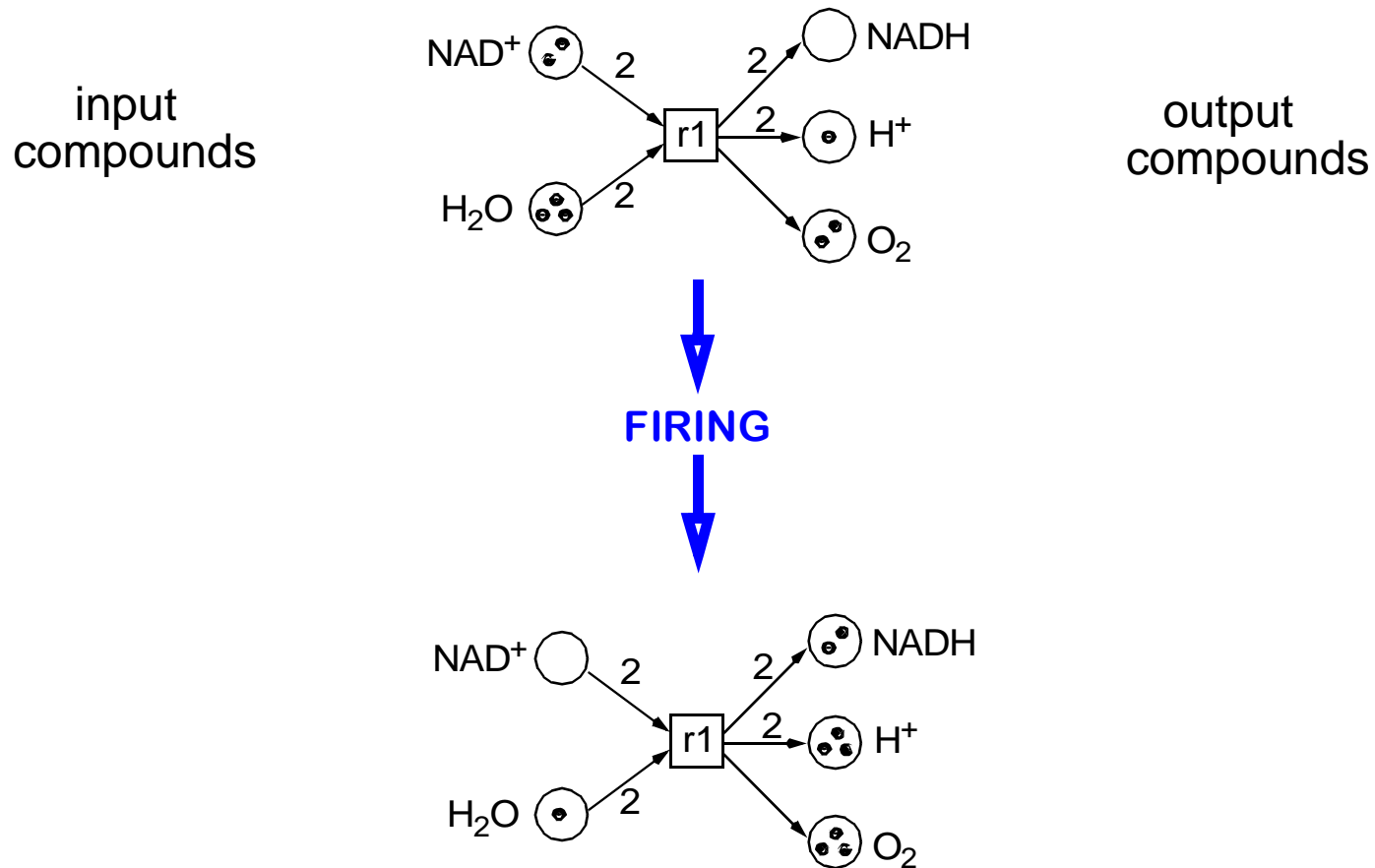
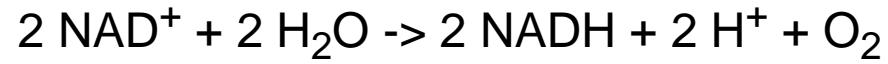
□ system state → marking → compounds distribution

□ $\text{PN} = (\text{P}, \text{T}, \text{F}, m_0)$, $\text{F}: (\text{P} \times \text{T}) \cup (\text{T} \times \text{P}) \rightarrow \mathbb{N}_0$, $m_0: \text{P} \rightarrow \mathbb{N}_0$

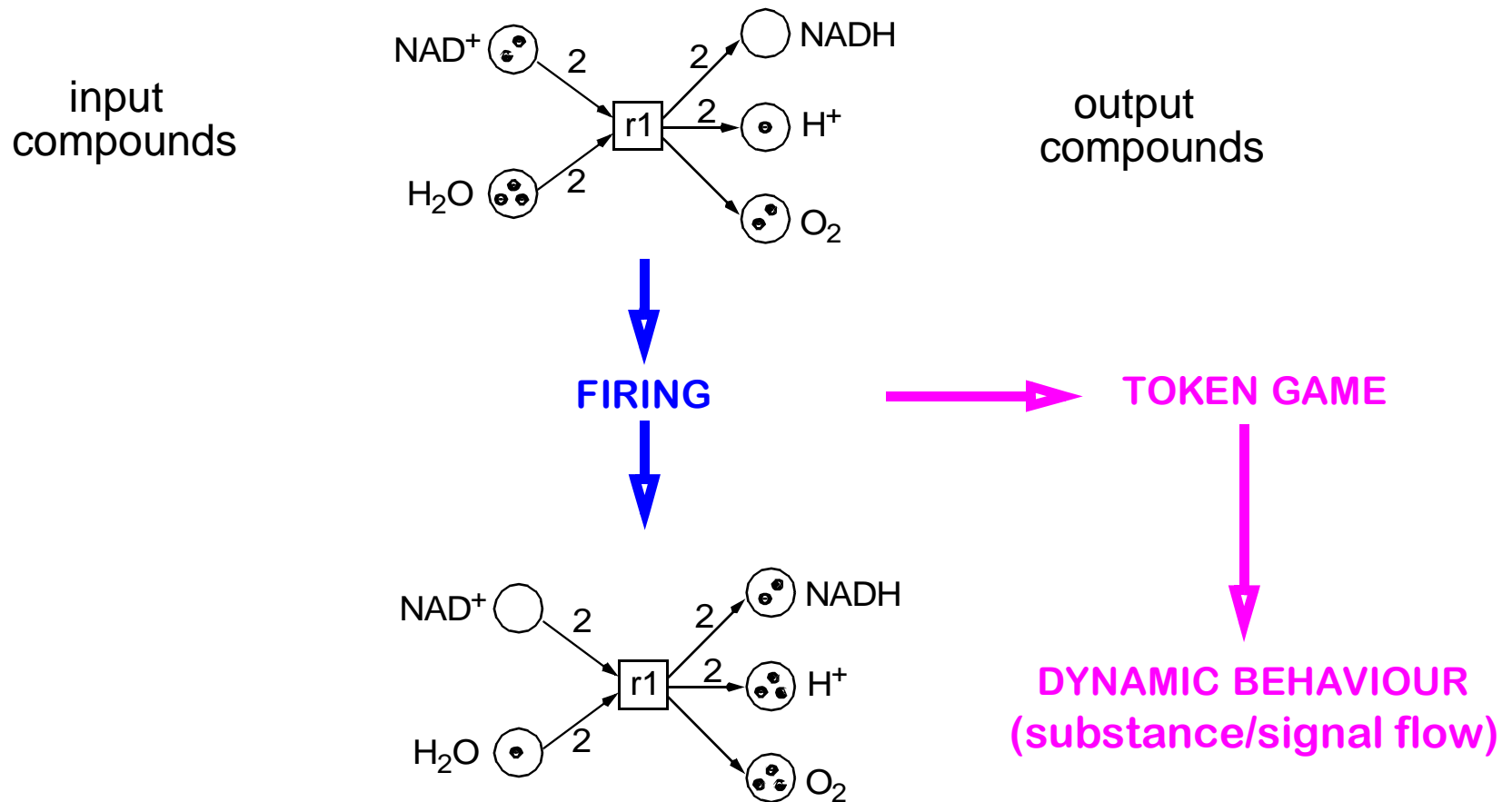
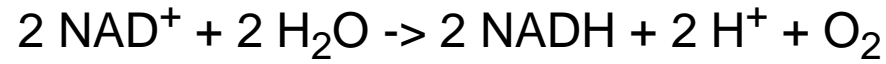
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❑ biochemical networks

-> *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*

INVARIANT ANALYSES

INCIDENCE MATRIX C

- a representation of the net structure

=> stoichiometric matrix

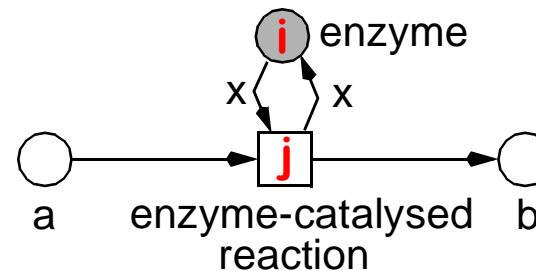
$$C =$$

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*

□ 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*

□ 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

-> ANALOGUE DEFINITIONS FOR P-INVARIANTS

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

- **T-invariants = (multi-) sets of transitions = Parikh vector**
 - > *zero effect on marking*
 - > *reproducing a marking / system state*

- **partially ordered transition sequence** -> **behaviour understanding**
of transitions occurring one after the other
 - > *substance / signal flow*
 - > *signal transduction networks, gene regulatory networks*

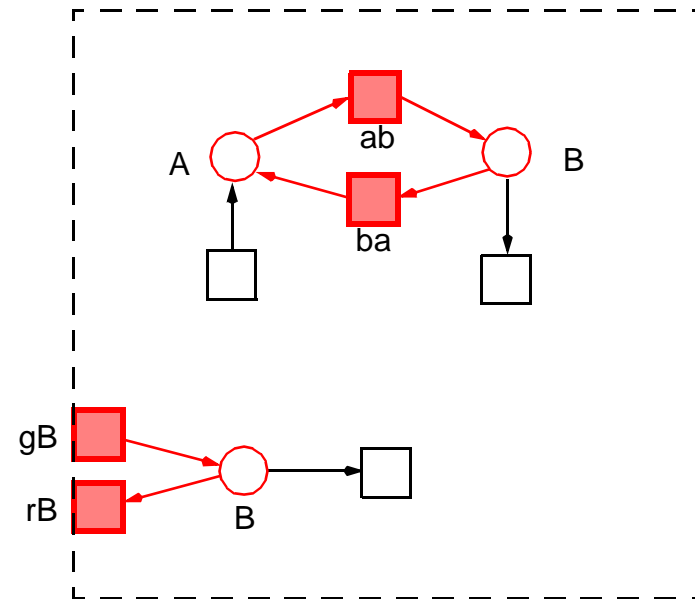
- **relative transition firing rates**
of transitions occurring permanently & concurrently
 - > *steady state behaviour*
 - > *metabolic networks*

□ trivial minimal T-invariants

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

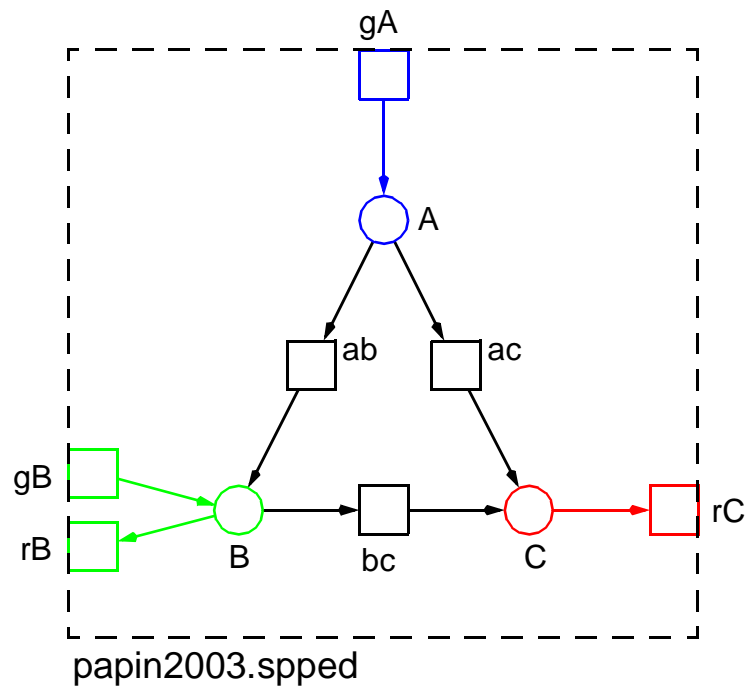
□ non-trivial minimal T-invariants

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



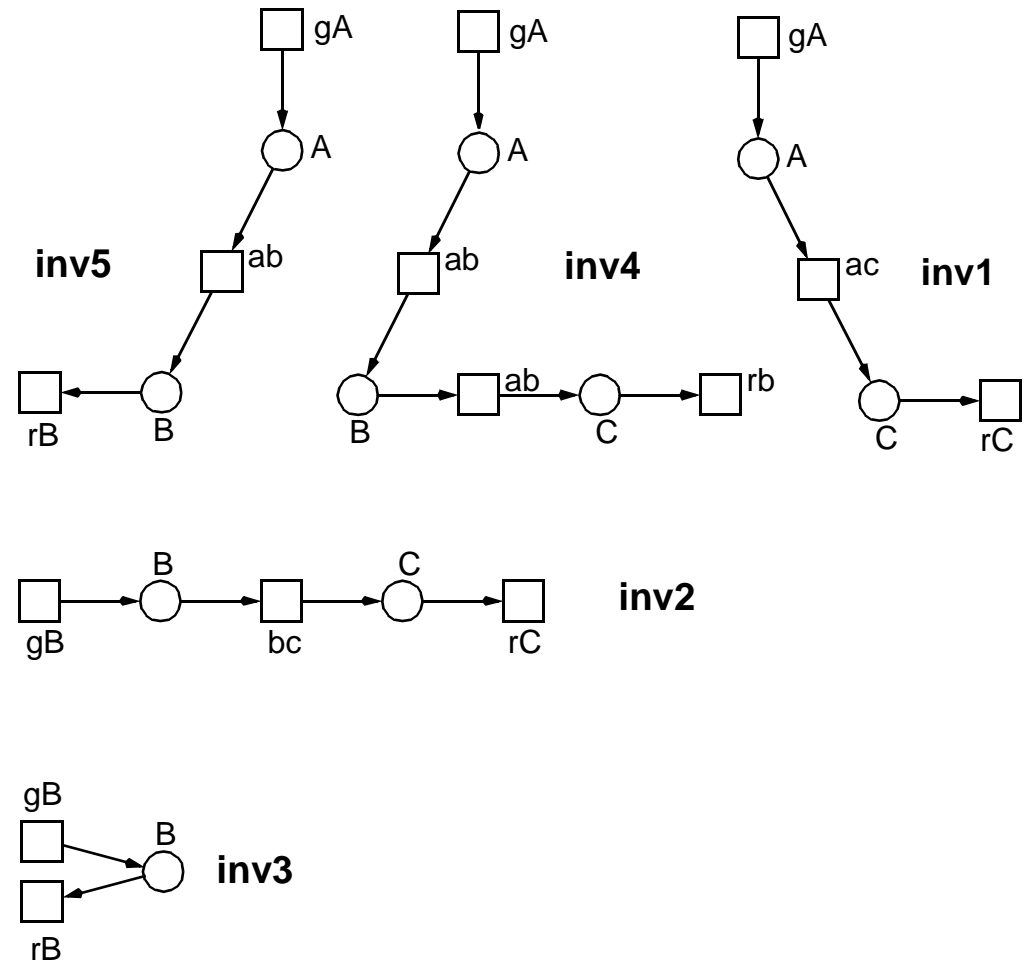
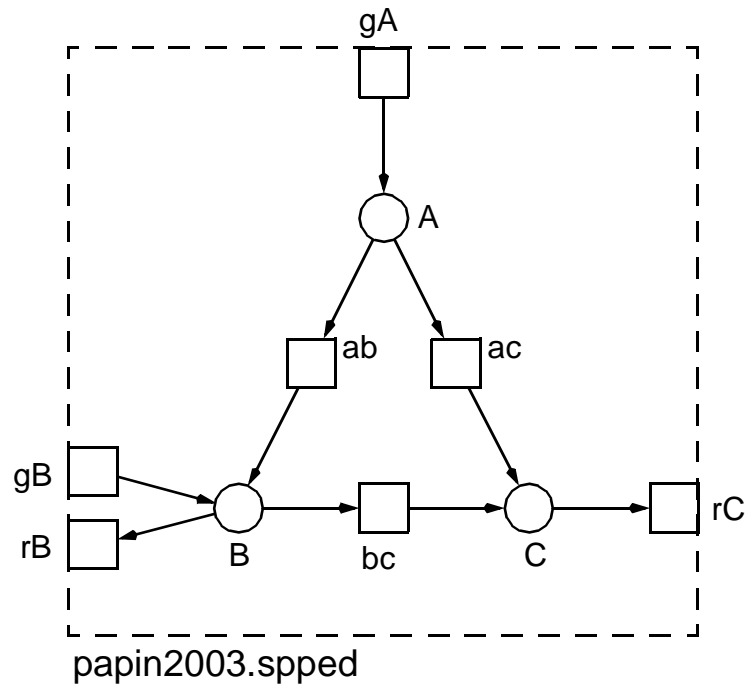
EXAMPLE

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



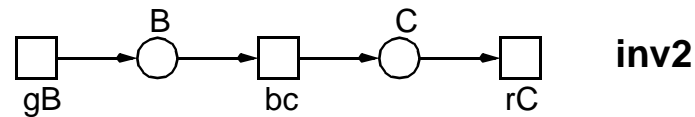
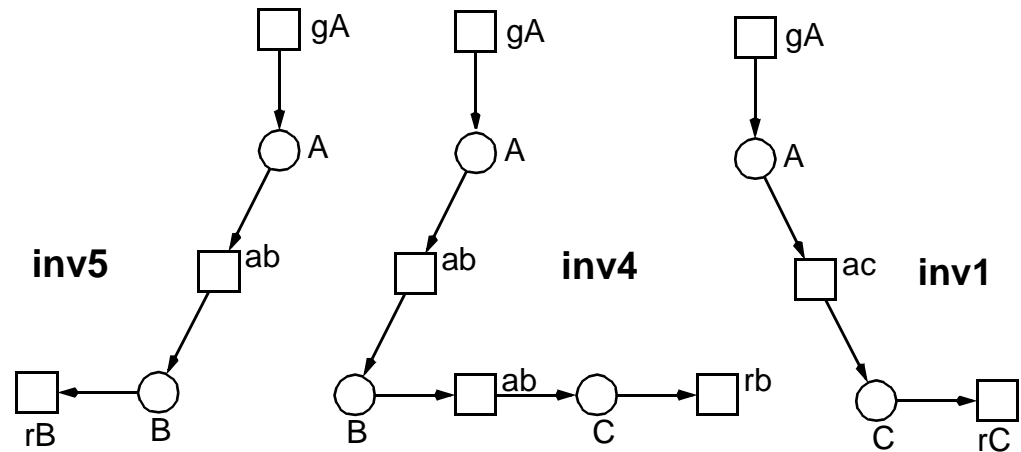
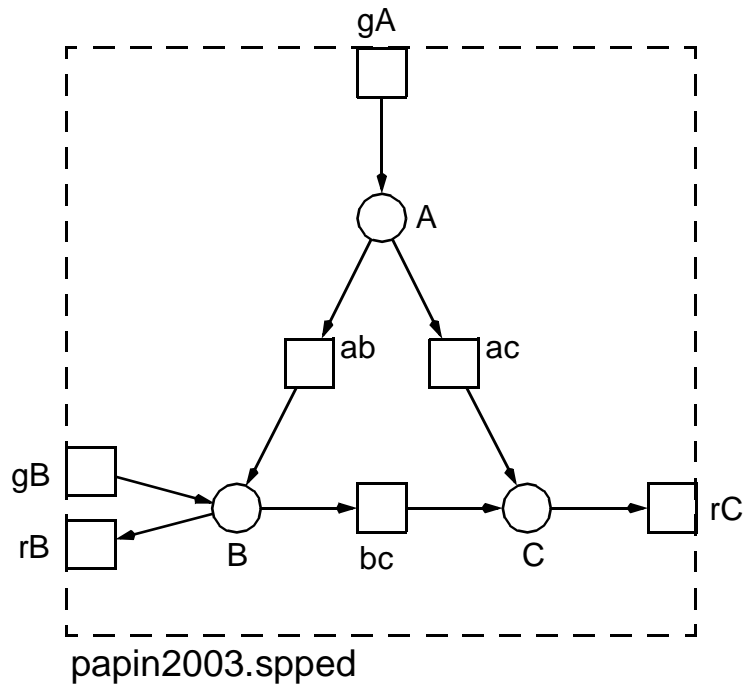
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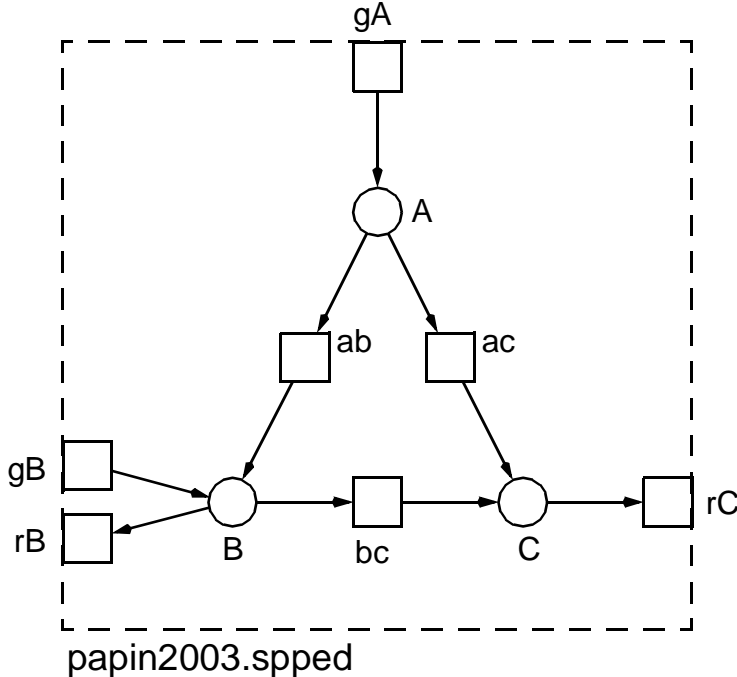
EXAMPLE, ELEMENTARY MODES

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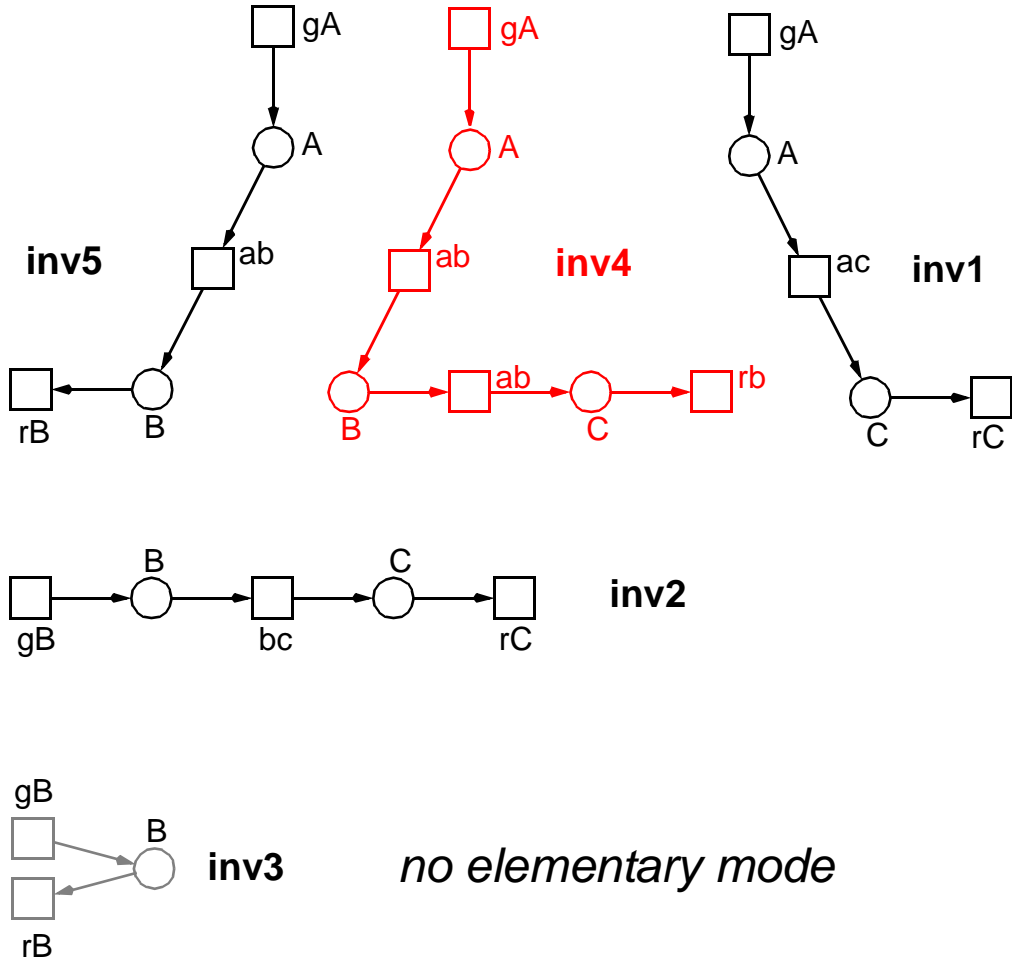


EXAMPLE, EXTREME PATHWAYS

- substances involved
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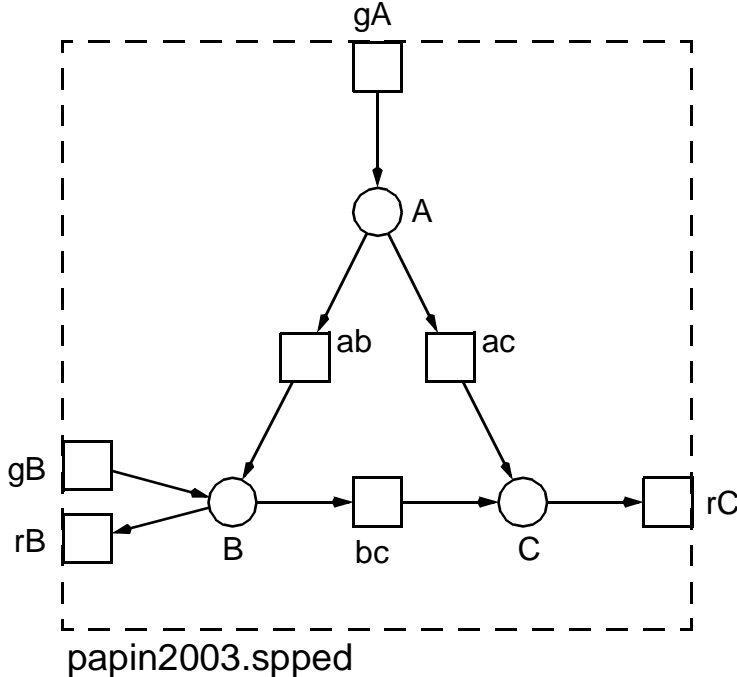


no extreme pathway

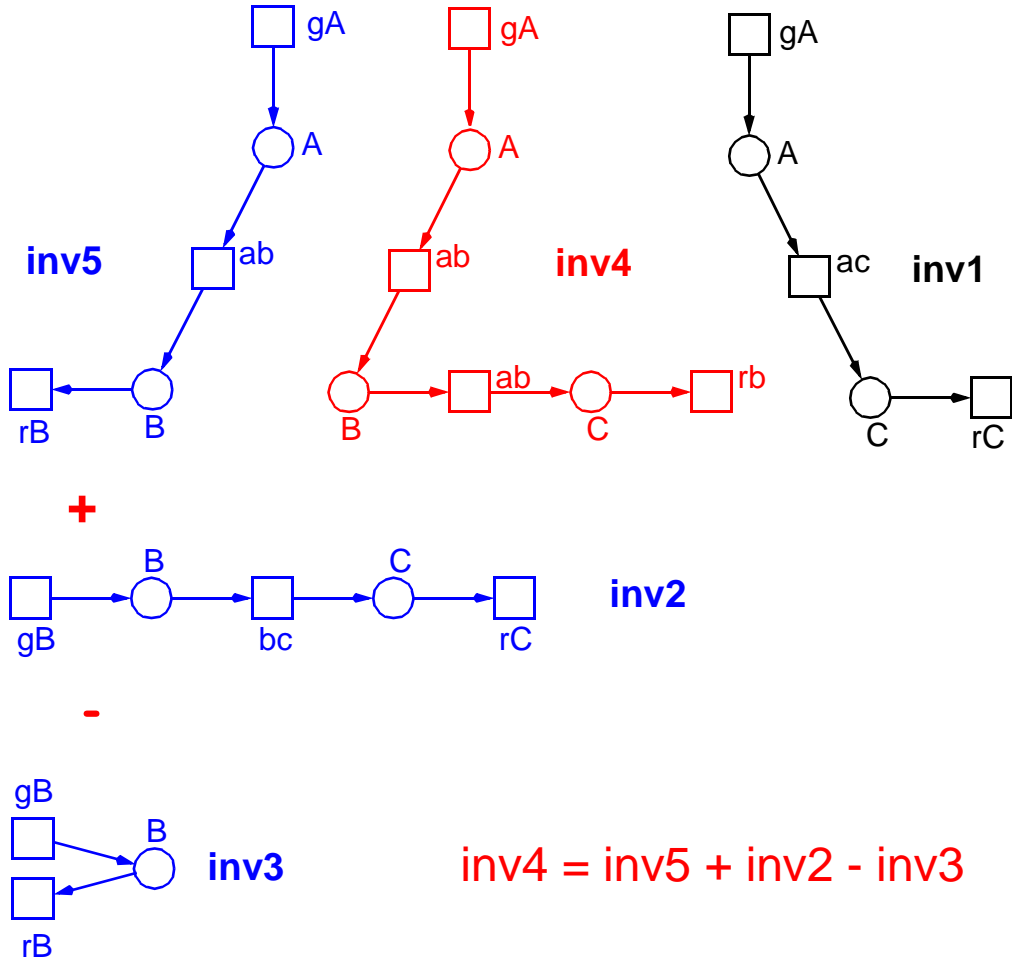


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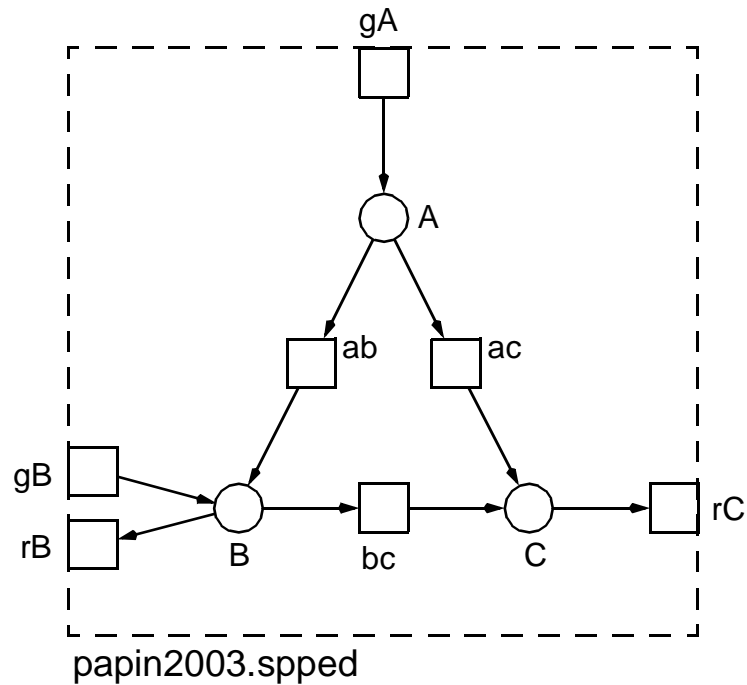


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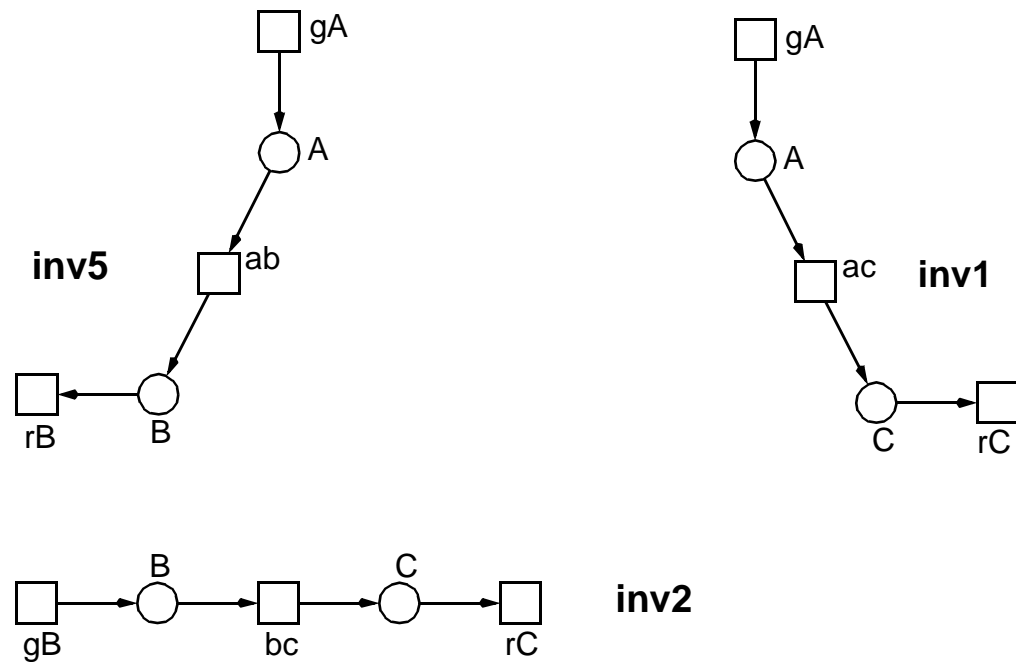


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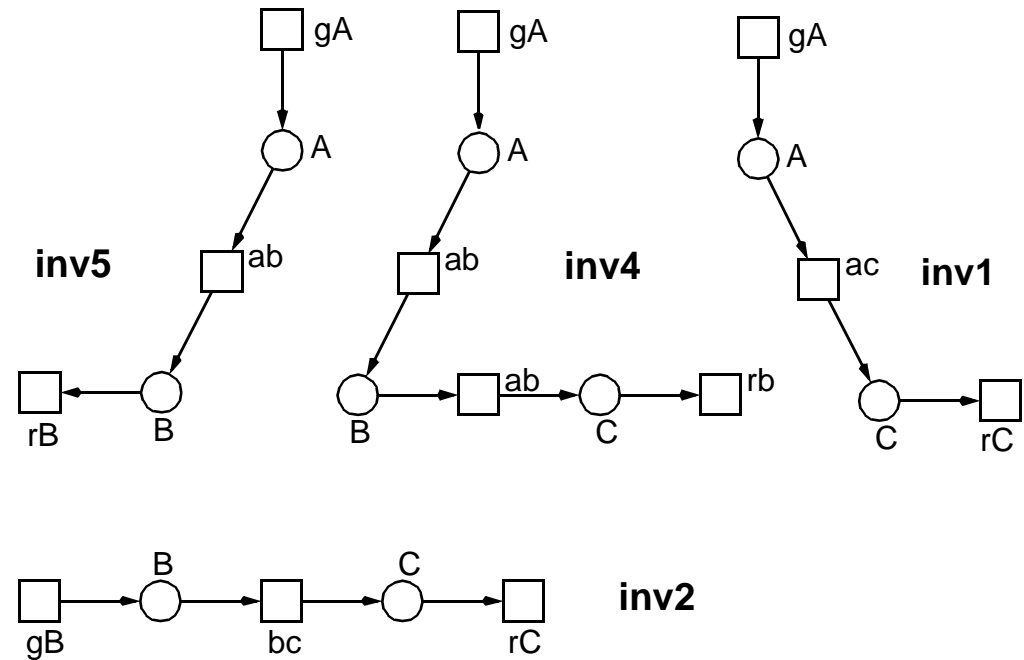
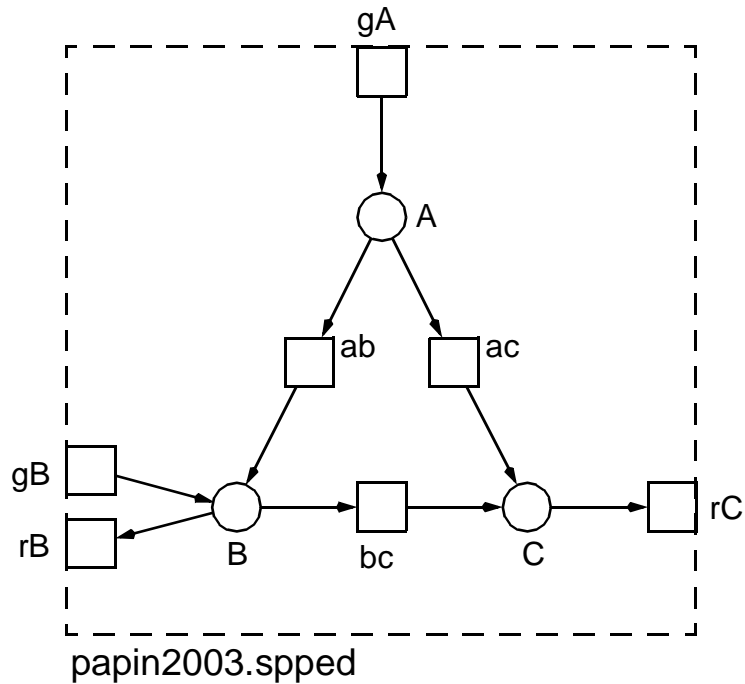
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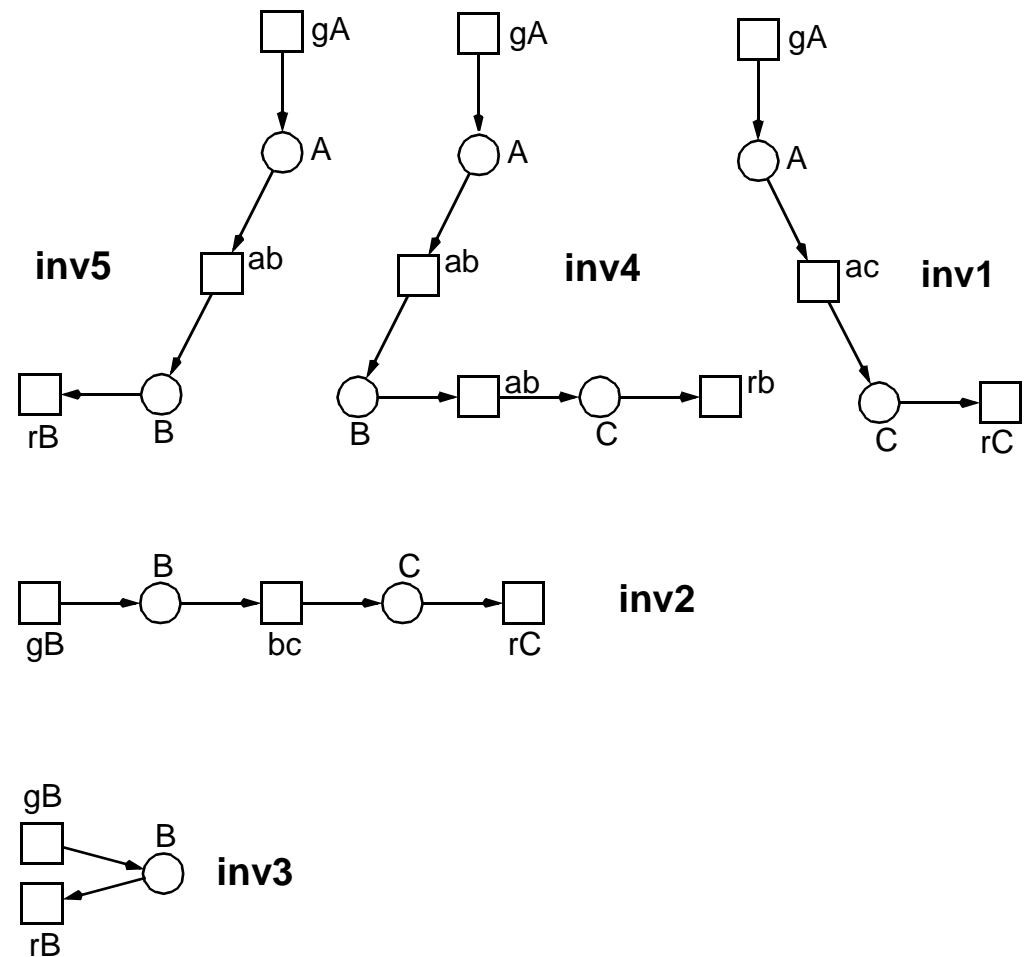
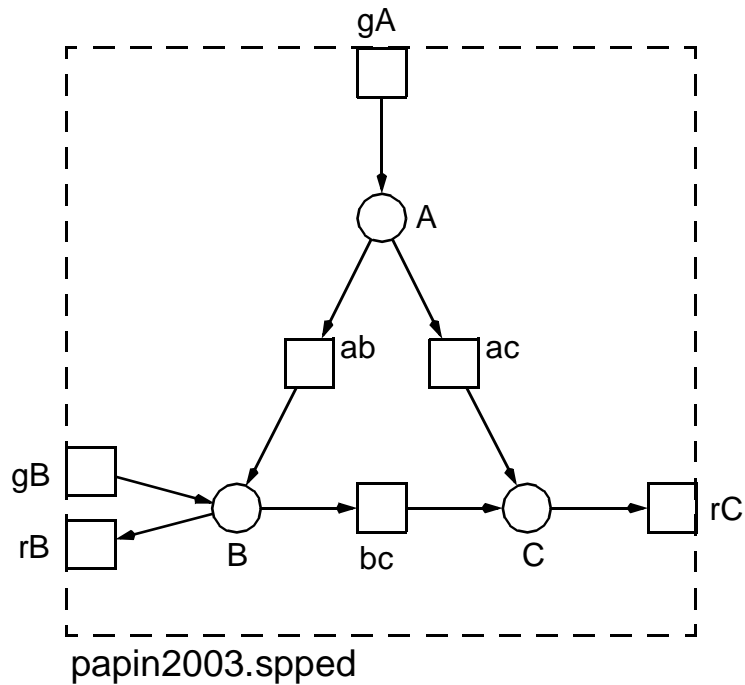
ELEMENTARY MODES

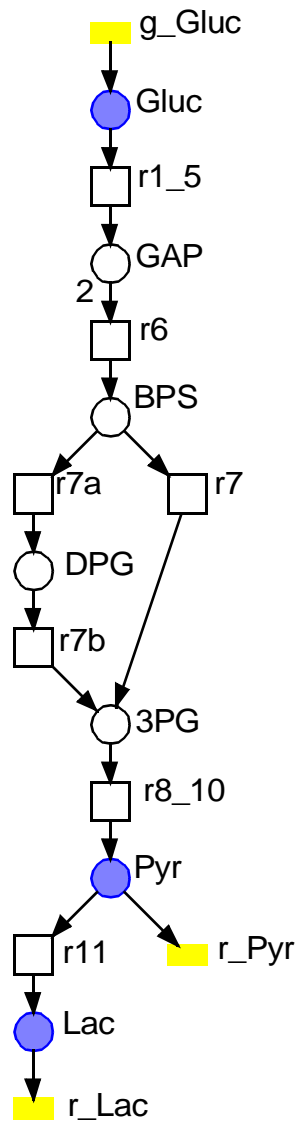


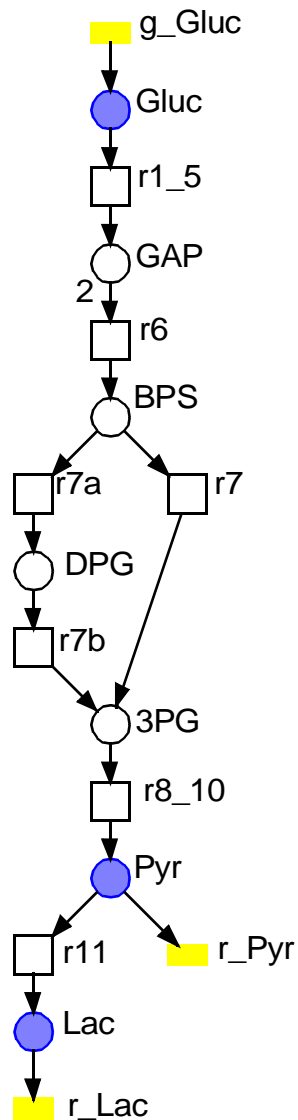
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MINIMAL T-INVARIANTS

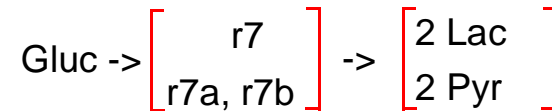




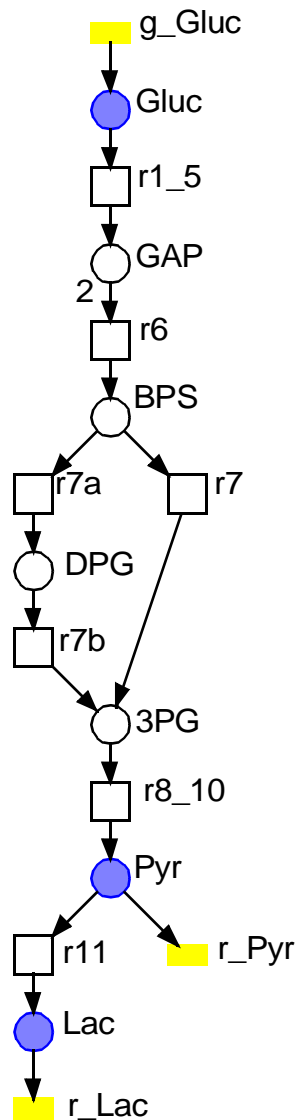


four minimal T-invariants

1. Gluc -> r7 -> 2 Pyr
2. Gluc -> r7 -> 2 Lac
3. Gluc -> r7a, r7b -> 2 Pry
4. Gluc -> r7a, r7b -> 2 Lac

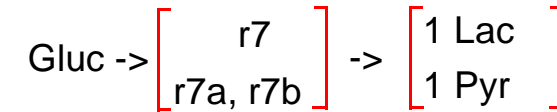
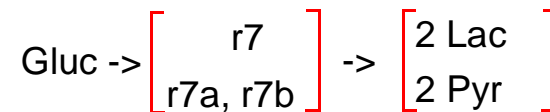


$$kx = \sum_i a_i x_i$$

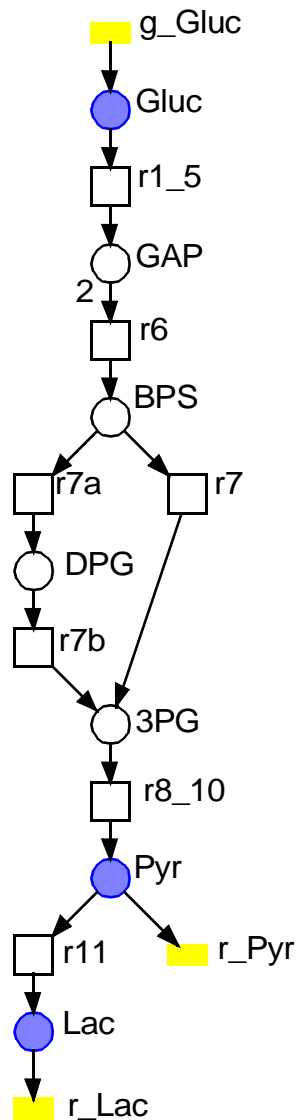


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five additional T-invariants

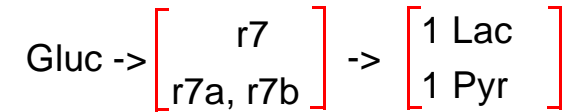
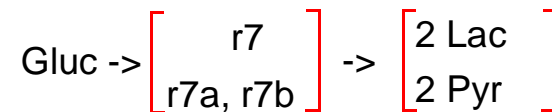
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$$\text{inv6} = (\text{inv2} + \text{inv4}) / 2$$

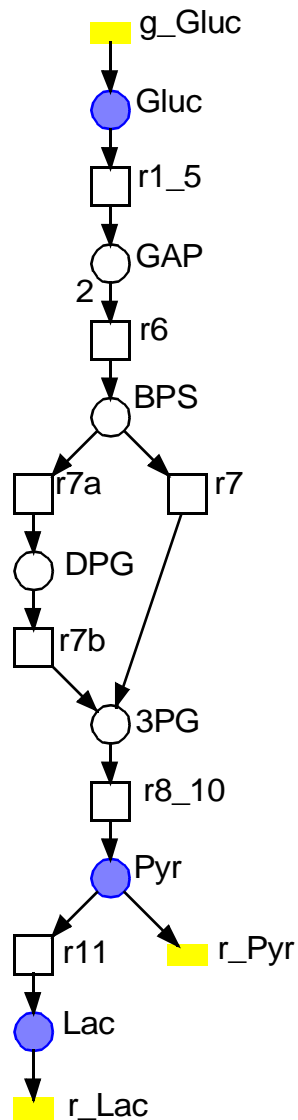
$$\text{inv7} = (\text{inv1} + \text{inv2}) / 2$$

$$\text{inv8} = (\text{inv3} + \text{inv4}) / 2$$

$$\text{inv9} = (\text{inv1} + \text{inv2} + \text{inv3} + \text{inv4}) / 4$$



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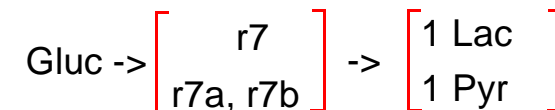
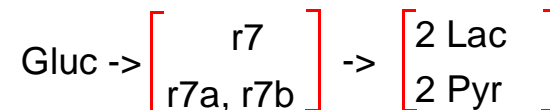
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$$kx = \sum_i a_i \cdot x_i$$

$$x = \sum_i a_i \cdot x_i$$

MODULAR COMPUTATION

- ❑ **decomposition
into subnets**

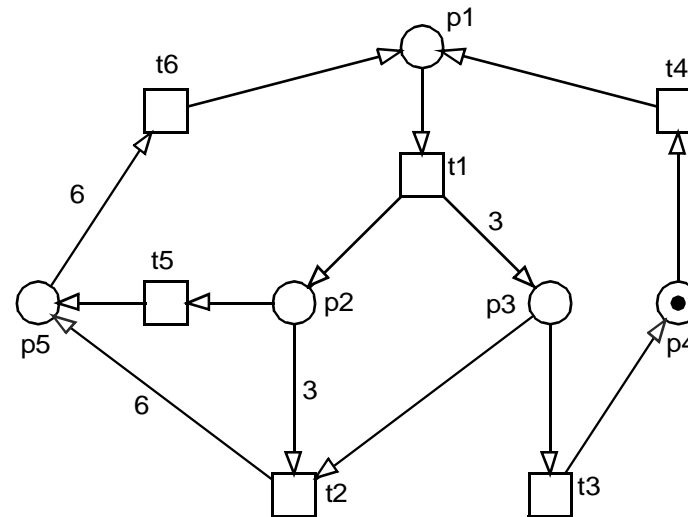
- ❑ **for each subnet:
computation of
(local) invariants**

- ❑ **computation of
interface invariants**

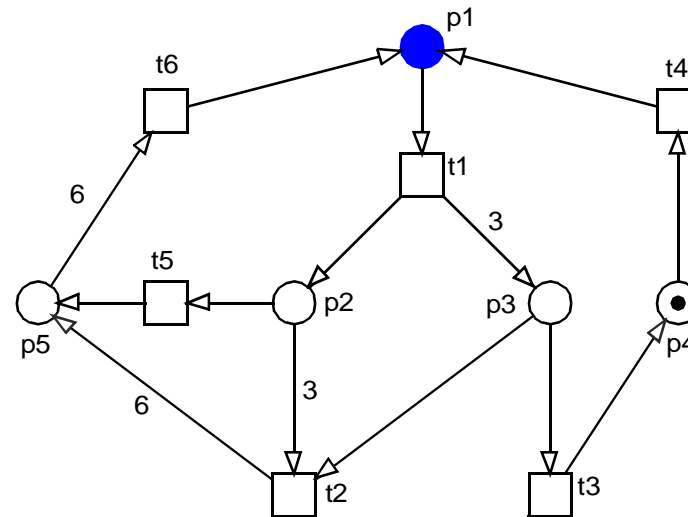
- ❑ **calculation of
system invariants**
 - > *by composition of
subnet invariants*

 - > *guided by
interface invariants*

- ❑ decomposition into subnets
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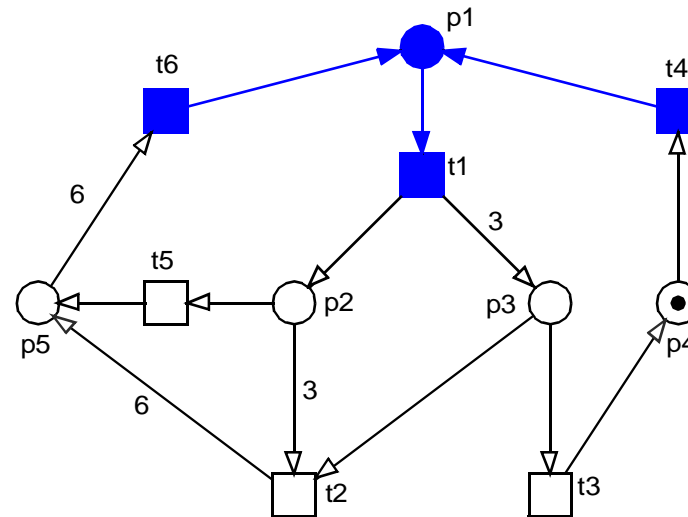


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subnet - transition-bordered conflict cluster C, defined by its places

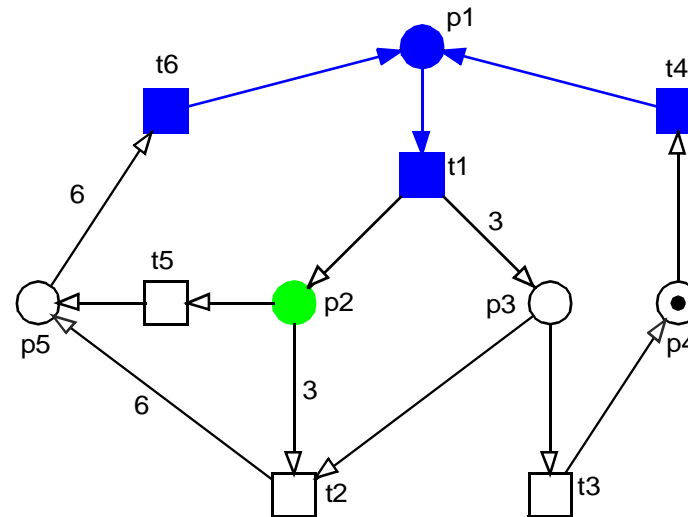
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all preplaces of output transitions belong to C

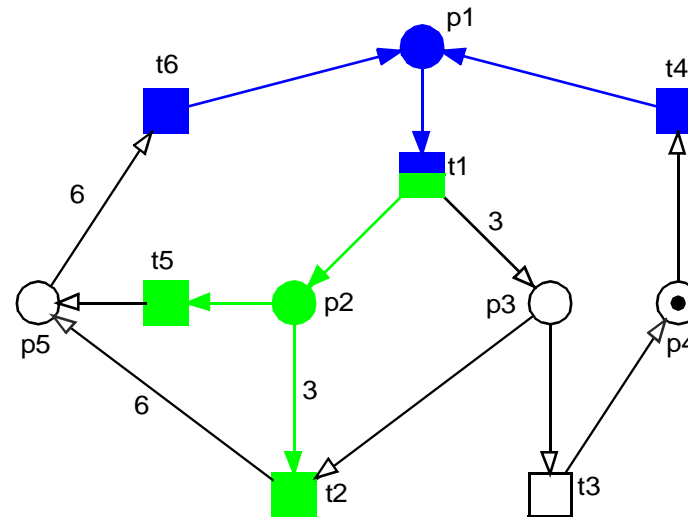
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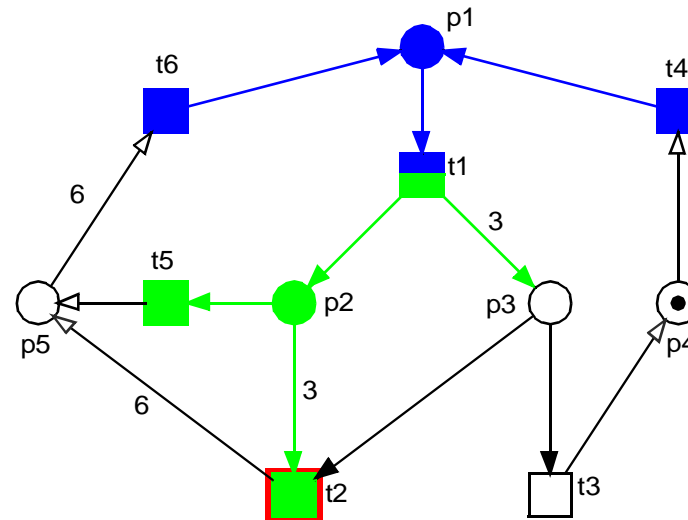
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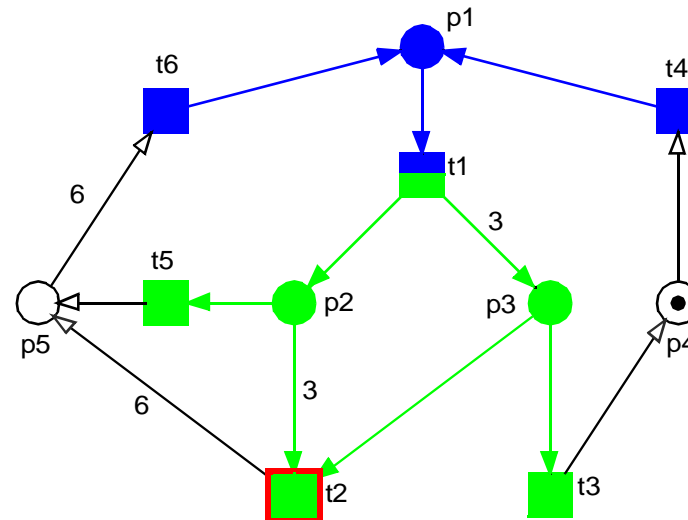


subnets -

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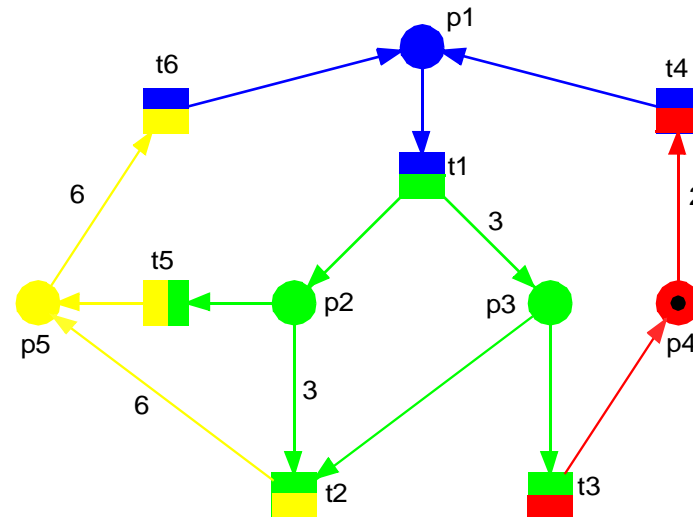
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 - > *guided by interface invariants*



subnet -

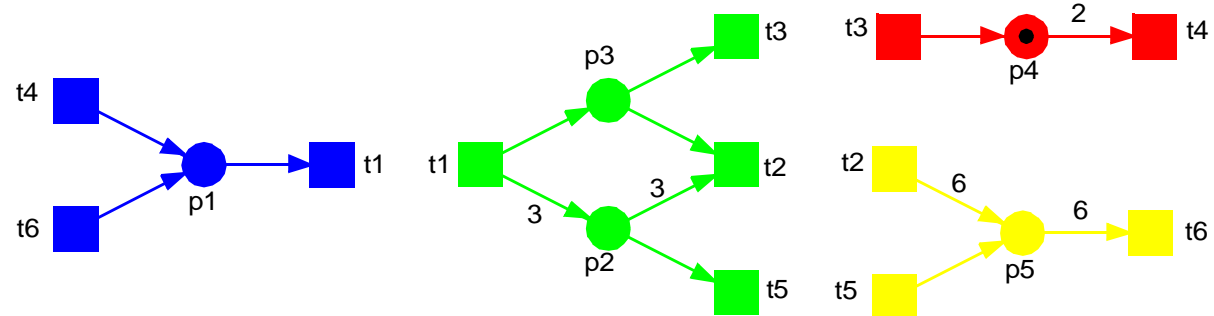
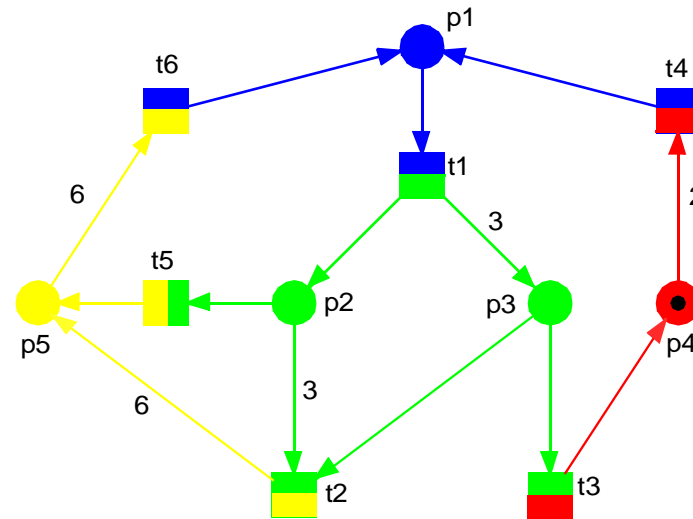
transition-bordered conflict cluster C , defined by its places

*all postplaces of input transitions belong to C
all preplaces of output transitions belong to C*

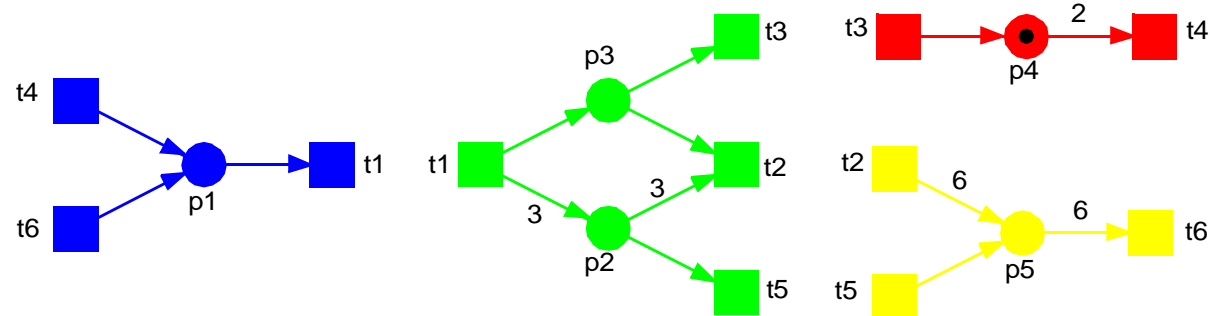
each interface transition has at most

- *one input subnet*
- *one output subnet*

- ❑ decomposition into subnets
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- decomposition into subnets

- for each subnet: computation of (local) invariants

$$\begin{aligned} x_1 &= (t_4, t_1) \\ x_2 &= (t_6, t_1) \end{aligned}$$

$$\begin{aligned} x_3 &= (t_1, t_2) \\ x_4 &= (t_1, t_3, 3 t_5) \end{aligned}$$

$$x_5 = (2 t_3, t_4)$$

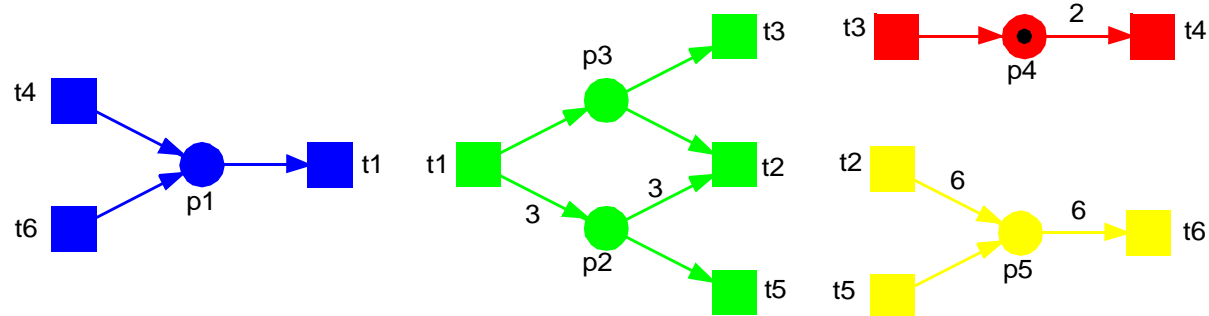
- computation of interface invariants

$$\begin{aligned} x_6 &= (t_2, t_6) \\ x_7 &= (6 t_5, t_6) \end{aligned}$$

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FOR EACH CONTACT TRANSITION

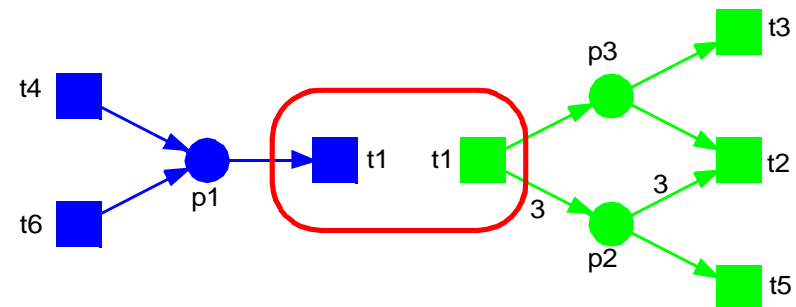
$$\begin{aligned} x_1 &= (t_4, t_1) \\ x_2 &= (t_6, t_1) \end{aligned}$$

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$$x_5 = (2 t_3, t_4)$$

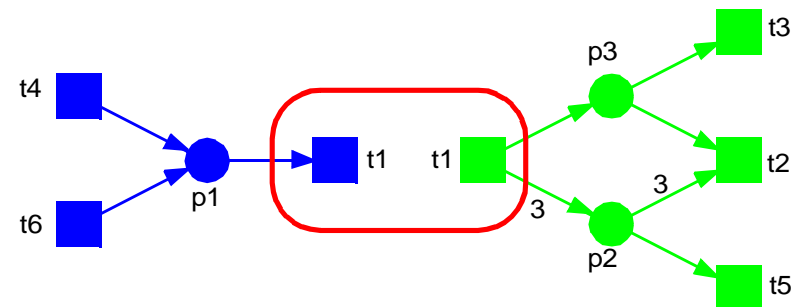
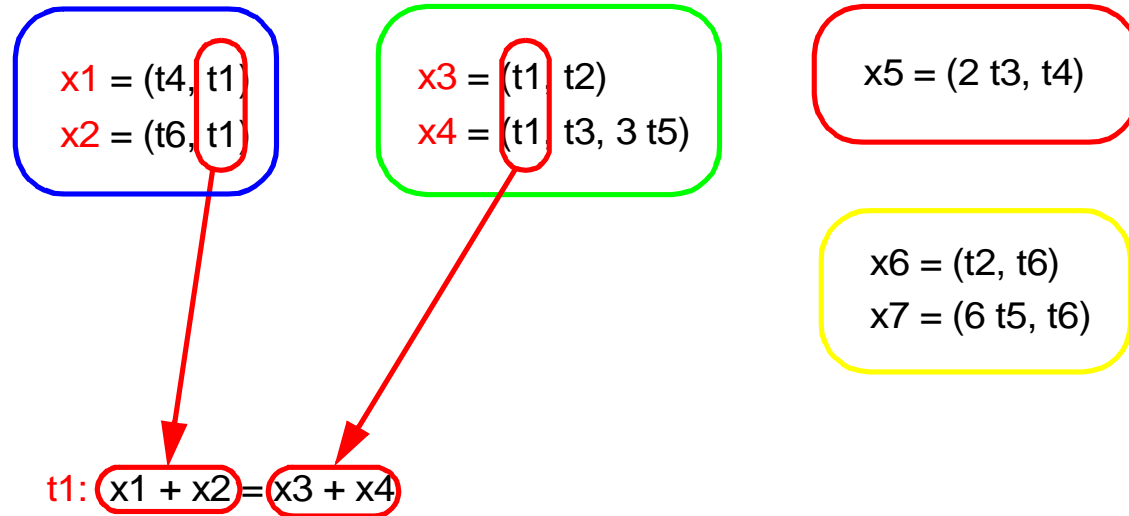
$$\begin{aligned} x_6 &= (t_2, t_6) \\ x_7 &= (6 t_5, t_6) \end{aligned}$$

t1:



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FOR EACH CONTACT TRANSITION

$$\begin{aligned} x_1 &= (t_4, t_1) \\ x_2 &= (t_6, t_1) \end{aligned}$$

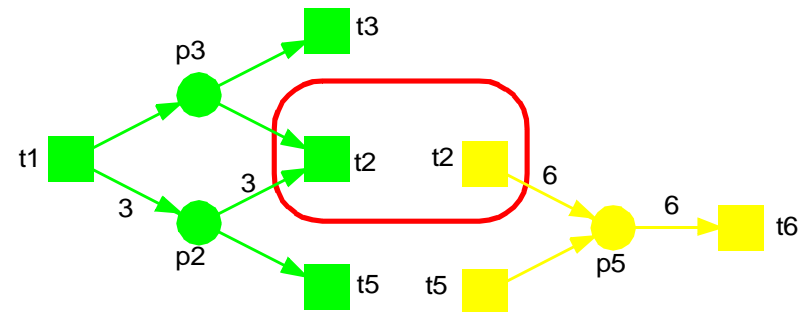
$$\begin{aligned} x_3 &= (t_1, t_2) \\ x_4 &= (t_1, t_3, 3 t_5) \end{aligned}$$

$$x_5 = (2 t_3, t_4)$$

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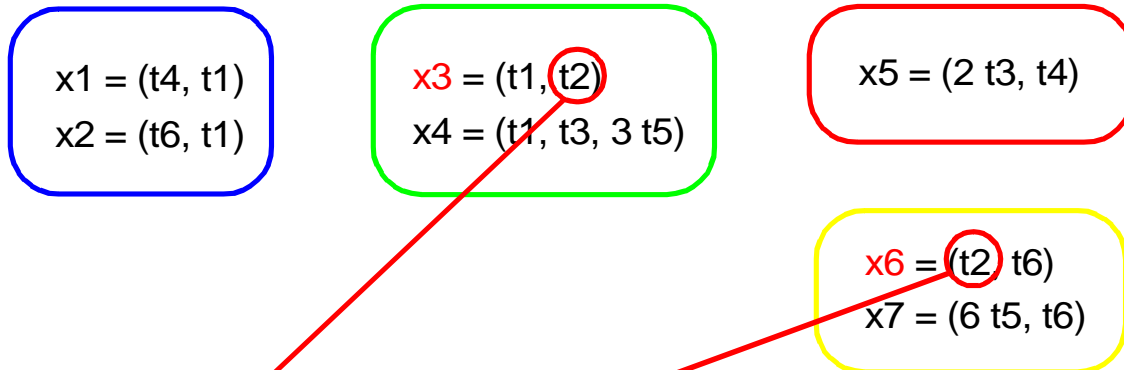
$$t_1: x_1 + x_2 = x_3 + x_4$$

t2:



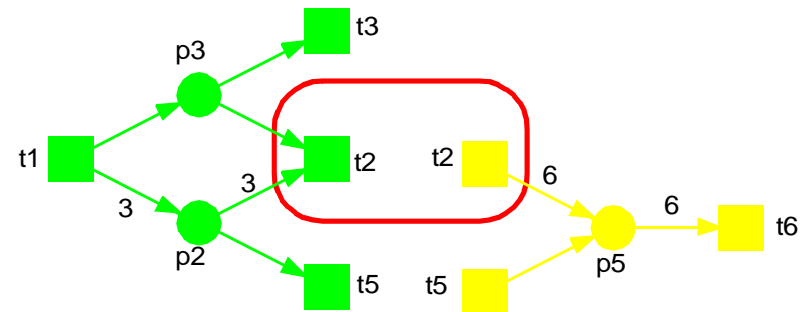
- ❑ decomposition into subnets
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FOR EACH CONTACT TRANSITION



$$t1: x1 + x2 = x3 + x4$$

$$t2: x3 = x6$$



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FOR EACH CONTACT TRANSITION

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$$t_1: x_1 + x_2 = x_3 + x_4$$

$$t_2: x_3 = x_6$$

t3:

t4:

t5:

t6:

- decomposition into subnets

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$$t_1: x_1 + x_2 = x_3 + x_4$$

$$t_2: x_3 = x_6$$

$$t_3: x_4 = 2 x_5$$

$$t_4: x_5 = x_1$$

$$t_5: 3 x_4 = 6 x_7$$

$$t_6: x_6 + x_7 = x_2$$

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$$t_1: x_1 + x_2 = x_3 + x_4$$

$$t_2: x_3 = x_6$$

$$t_3: x_4 = 2 x_5$$

$$t_4: x_5 = x_1$$

$$t_5: 3 x_4 = 6 x_7$$

$$t_6: x_6 + x_7 = x_2$$



$$(x_2, x_3, x_6)$$

$$(x_1, x_2, 2 x_4, x_5, x_7)$$

- ❑ decomposition into subnets

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$$\begin{aligned}(x_2, x_3, x_6) \\(x_1, x_2, 2 x_4, x_5, x_7)\end{aligned}$$



- decomposition into subnets

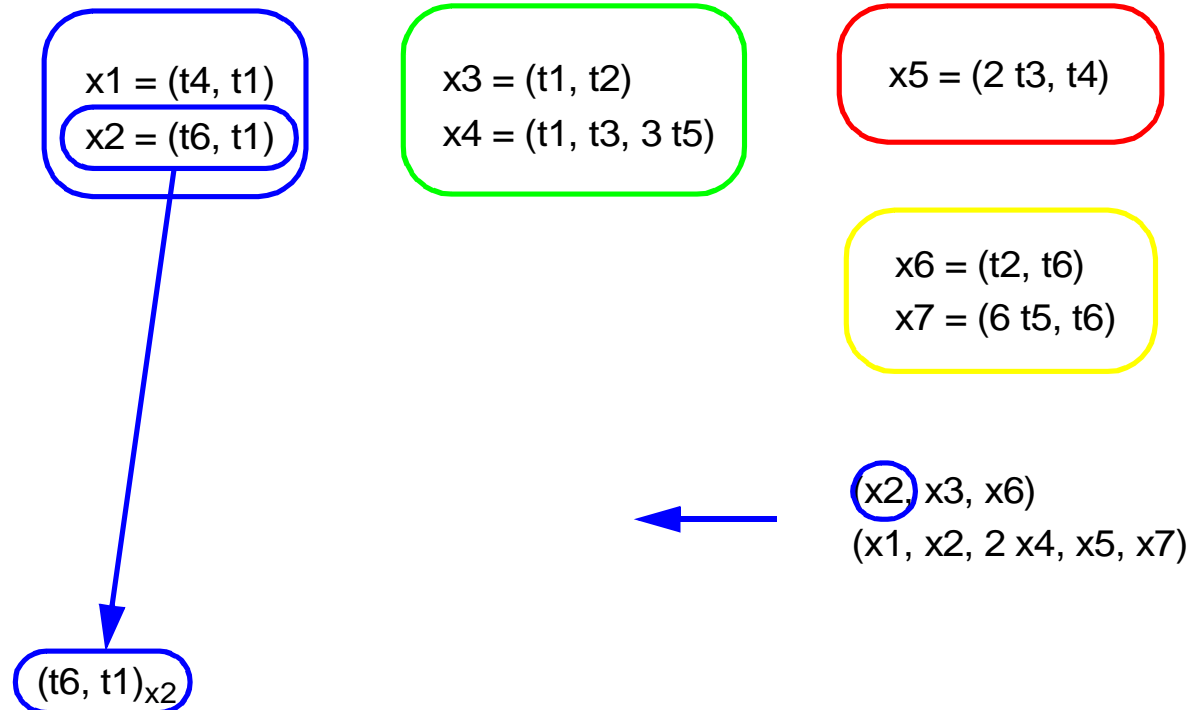
- for each subnet: computation of (local) invariants

- computation of interface invariants

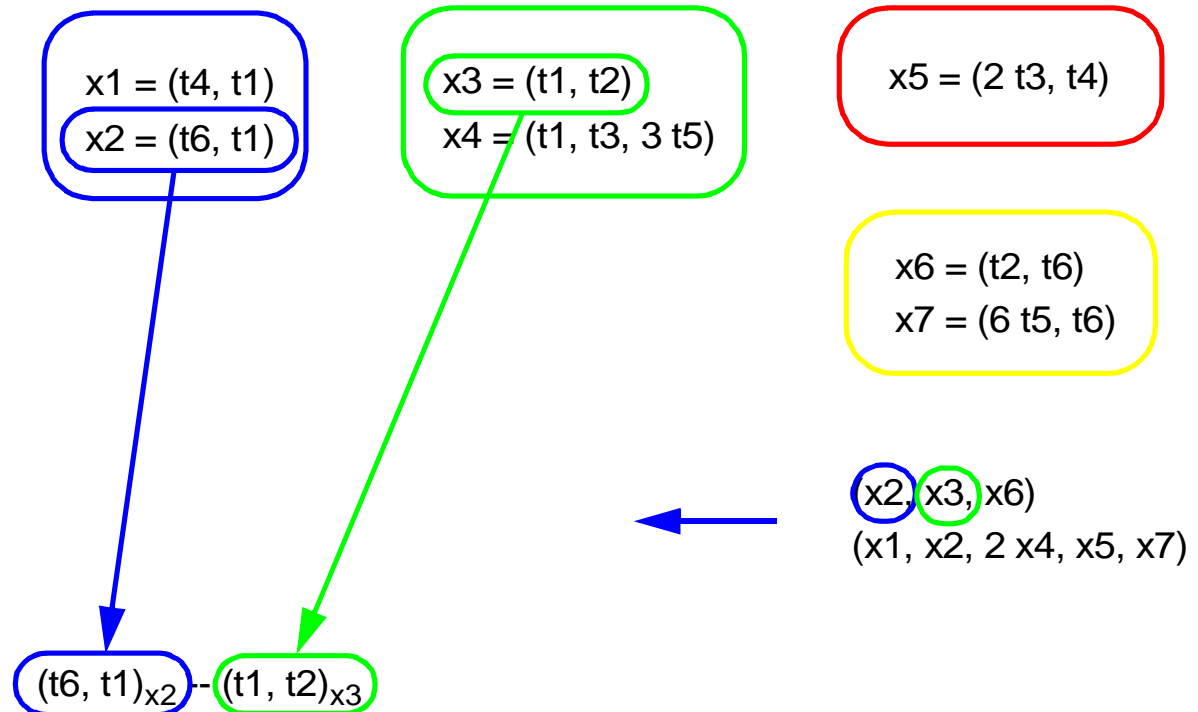
- calculation of system invariants

-> by composition of subnet invariants

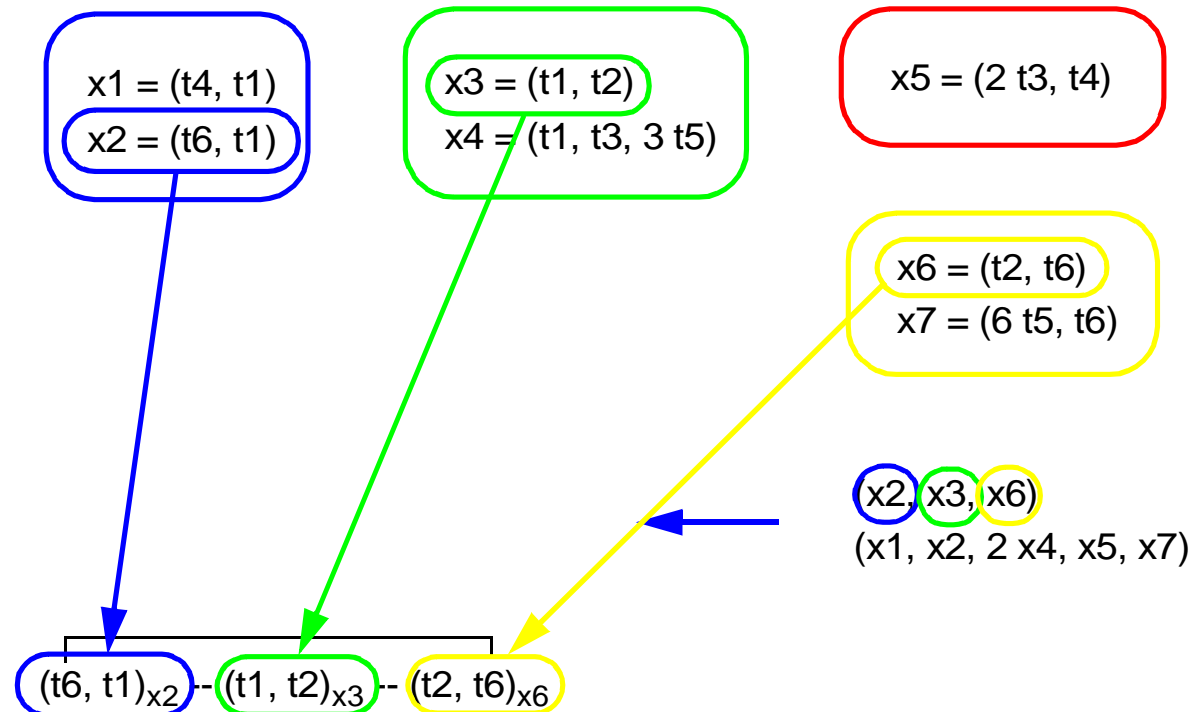
-> guided by interface invariants



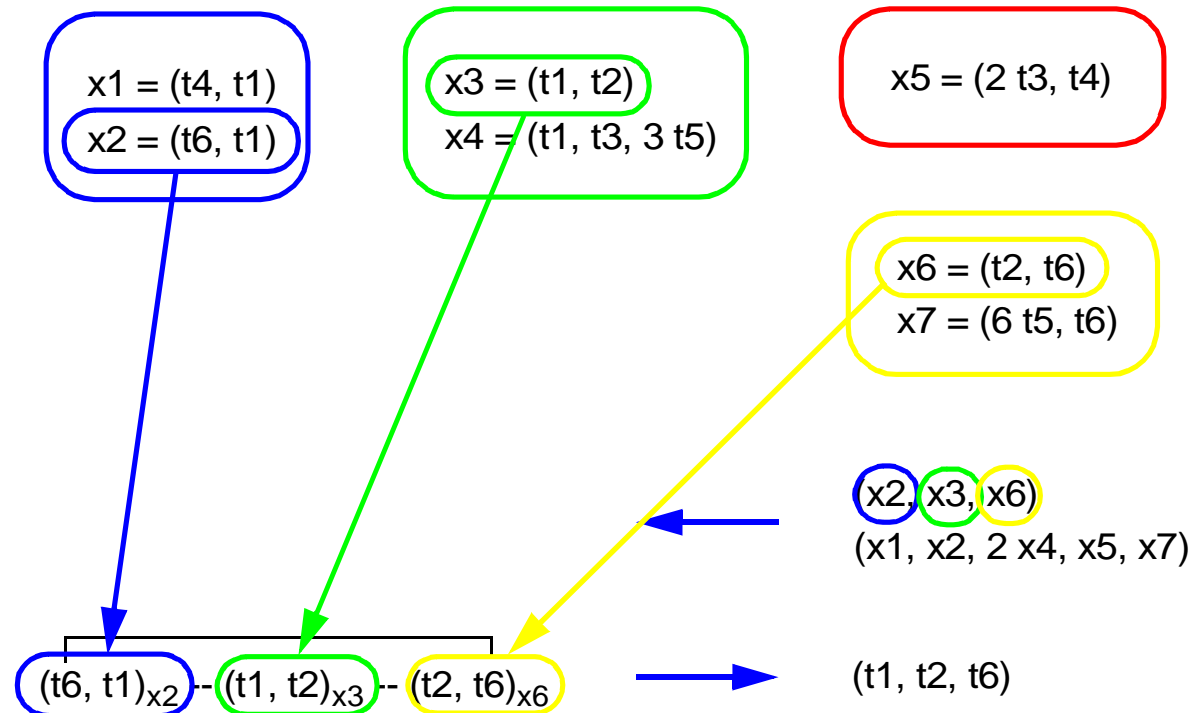
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$$\overbrace{(t_6, t_1)_{x_2} \text{ -- } (t_1, t_2)_{x_3} \text{ -- } (t_2, t_6)_{x_6}}$$



$$\begin{aligned} &(x_2, x_3, x_6) \\ &(x_1, x_2, 2 x_4, x_5, x_7) \end{aligned}$$



$$(t_1, t_2, t_6)$$

$$2 \left(\begin{array}{l} t_1, t_3 \\ 3 t_5 \end{array} \right)_{x_4} \text{ -- } (2 t_3, t_4)_{x_5} \text{ -- } (t_4, t_1)_{x_1} \text{ -- } (6 t_5, t_6)_{x_7} \text{ -- } (t_6, t_1)_{x_2}$$

- decomposition into subnets

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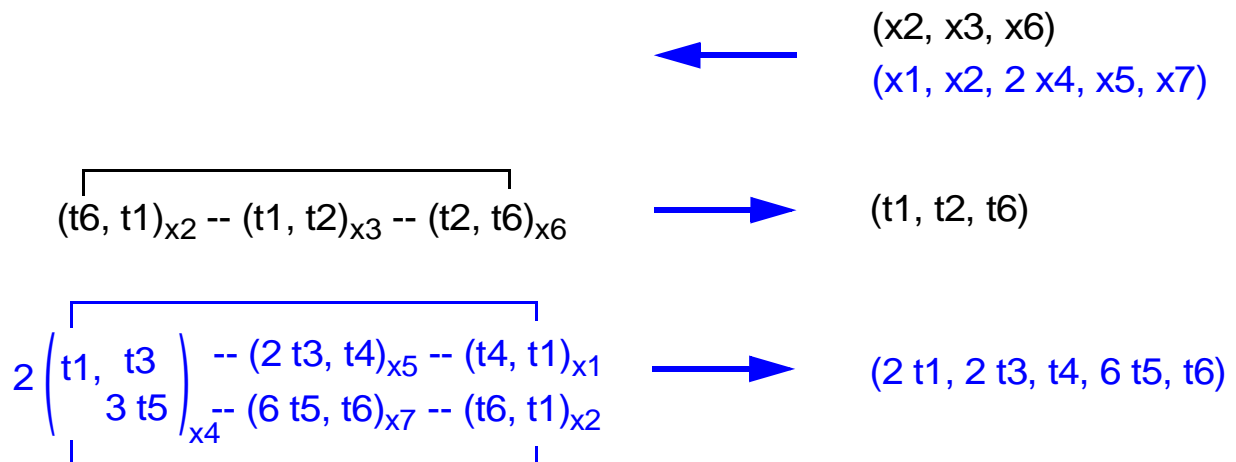
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$$\begin{aligned} & \overbrace{(t_6, t_1)_{x_2} \text{ -- } (t_1, t_2)_{x_3} \text{ -- } (t_2, t_6)_{x_6}} \\ & 2 \left(\begin{array}{l} t_1, t_3 \\ 3 t_5 \end{array} \right)_{x_4} \text{ -- } (2 t_3, t_4)_{x_5} \text{ -- } (t_4, t_1)_{x_1} \\ & \text{-- } (6 t_5, t_6)_{x_7} \text{ -- } (t_6, t_1)_{x_2} \end{aligned}$$

$$\begin{aligned} & \leftarrow (x_2, x_3, x_6) \\ & (x_1, x_2, 2 x_4, x_5, x_7) \end{aligned}$$

$$\begin{aligned} & (t_1, t_2, t_6) \\ & (2 t_1, 2 t_3, t_4, 6 t_5, t_6) \end{aligned}$$

ASSUMPTION

- the solution of many small systems is less time/space consuming than the solution of a single larger one

MAJOR (KNOWN) DRAWBACK

- the computation of system invariants does not only produce minimal invariants

CASE STUDIES

- > excel file

Lautenbach, K.:

Exact Liveness Conditions of a Petri Net Class (in German);
Berichte der GMD 82, Bonn 1973.

Pascoletti, K.-H.:

Diophantische Systeme und Lösungsmethoden zur Bestimmung aller Invarianten in Petri-Netzen;
Berichte der GMD 160, 1986.

Starke, P. H.:

Analyse von Petri-Netz-Modellen;
Teubner 1990.

Zaitsev, D.:

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Univ. Paris-Dauphine, LAMSADE, TR 224, 2005

Schuster, S.; Hilgetag, C.; Schuster, R.:

Determining Elementary Modes of Functioning in Biochemical Reaction Networks at Steady State.
Proc. Second Gauss Symposium (1993) pp. 101-114

Schilling, C. H.; Letscher, D.; Palsson, B. O.:

Theory for the Systemic Definition of Metabolic Pathways and their Use in Interpreting Metabolic Function from a Pathway-Oriented Perspective;
J. Theor. Biol. (2000) 203, pp. 229-248.

THANKS !

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